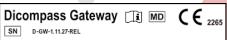
Dicompass Gateway

User guide

Medoro s.r.o.

2025-04-25



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UDI (01)08594212320015(11)250425(8012)1.11.27-REL

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1 Introduction

1.1 Purpose of this document {#intro_Purpose of this document}

This document is a user manual in electronic form, describing the functions and operation of the SW Dicompass Gateway.

The user manual is intended for professional users, i.e. for trained operators of SW Dicompass Gateway, i.e. trained medical and intermediate medical personnel. User training is carried out once after the

implementation of the MD and then with each released version with a change in the MAJOR number, i.e. the first number of the versioning format.

Administration of SW Dicompass Gateway does not apply to the regular user, therefore it will not be described here. It is described in detail in the Administrator's Guide.

Please read this manual carefully before using the Dicompass Gateway medical device.

If necessary, this instruction manual is provided in paper form, at no additional cost to the user, within 7 days of receiving the request. The request can be sent by email to helpdesk@medoro.org.

Name and address of the manufacturer: **MEDORO s.r.o.**, Štrossova 567, 530 03 Pardubice, Czech Republic, ID: 26002612, VAT number: CZ26002612, http://www.medoro.org, http://www.dicompass.cz, e-mail: info@medoro.org

1.2 Reporting serious adverse events {#intro_Reporting serious adverse events}

According to Regulation 2017/745 of the European Parliament and Council (EU) from 5th April 2017 on medical devices, amending Directive 2001/83/ES, Regulation (ES) No 178/2002 and Regulation (ES) No 1223/2009 and repealing Council Directives 90/385/EHS and 93/42/EHS, serious adverse event refers to an event which directly or indirectly leads to, may have or may lead to one of the following consequences:

- 1. death of a patient, user or other person
- 2. temporary or permanent deterioration of patient's, user's or other person's health
- 3. serious threat to public health

Any serious adverse event that has occurred in connection with the device in question should be reported to the manufacturer and the competent authority of the Member State in which the user and/or the patient is established.

In the event of an adverse event, contact us using the HelpDesk, by phone on +420 463 356 445, or by email at helpdesk@medoro.org.

1.3 Description of the intended purpose of use {#intro_Description of the intended purpose of use}

The intended purpose of SW Dicompass Gateway:

The specialized modular software Dicompass Gateway is intended as a tool for working with image data for the purposes of diagnostic and therapeutic actions in the healthcare sector. Dicompass contains not only a DICOM viewer, but also modules for a complete solution for video digitization

from endoscopes, ultrasounds, microscopes, as well as other devices that do not have a direct DICOM output, conversion of recordings from digital cameras, scanners and cameras into DICOM format (DICOMization). Dicompass also offers functions for radiodiagnostics and radiotherapy.

The Dicompass Gateway software is an active medical device that does not come into contact with the patient and its purpose is to display and work with image medical documentation. The Dicompass software is thus intended for all groups of patients undergoing radiological or other (e.g. endoscopic) examinations that generate image documentation.

The Dicompass Gateway software is used for the diagnosis of health conditions requiring radiological or other (e.g. endoscopic) examination generating image documentation.

The indication for the use of MD Dicompass Gateway is connected with the need for a specialized doctor to diagnose the deteriorated condition of the patient using imaging methods. Dicompass enables this image documentation to be archived and subsequently worked with.

Contraindications - Patients in whom it is not possible to obtain evaluable imaging documentation.

Warning: The use of Dicompass is conditioned by the training of users.

Dicompass Gateway is a certified tool for archiving, viewing and postprocessing DICOM data acquired using DICOM modalities, such as X-ray, CT, MR, SONO, ECG and others. This SW uses web technologies and for its purpose it is necessary to use standard HTML browsers (current versions) supporting WebGL (Edge, Chrome, Firefox) across all system platforms (Windows, Apple iOS, Linux, Android). The program's interface aims at user convenience and clarity.

The software is classified and certified as a Class IIb medical device in accordance with applicable legislative requirements.

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Figure 1: img

Dicompass Gateway is a purely web-based DICOM tool that, in addition to a web browser, does not require any additional support resources installed on the workstation. Thanks to traceless technology, it leaves no traces on the user's PC.

Considering the HTML technology used, it can also be operated on any mobile device (a tablet or a smartphone), both inside and outside of your medical facility.

Dicompass Gateway is a very sophisticated tool for remote consultations and all work performed within this SW is fully audited.

Not only does it contain the basic tools for viewing and working with DICOM images, such as distance measurement, area measurement, windows, annotations, rotation options, screen splitting, support

of multiple monitors, including diagnostic ones, etc., but also supports image captions, any image filters, or a wide variety of image displays.

In multiple windows, it is possible to use synchronized image viewing or browsing images marked as important.

Dicompass Gateway supports a number of standardized formats such as JPEG, PNG, SR, MPEG2, MPEG4 and more. Images or recordings can be exported to an external device, burned to a CD/DVD, or sent via metropolitan exchange ePACS / ReDiMed to another medical facility. Dicompass Gateway also supports multiple PACS archives at once. You don't have to search and switch between different functions. The search dialog allows you to set exact parameters. The overall browser settings can be saved to each user account. After logging in, your labels or measurement tools will not be shuffled by your colleagues previous usage.

Thanks to use of web technology, all image documentation is not stored on the end device of the doctor's diagnostic workstation in advance, thus, when it is opened, there may be a delay before loading. This delay is affected by the speed of the internet connection or the response of the line of the internal network of the medical facility.

MD Dicompass Gateway is a separate device and for its intended purpose of use does not use additional accessories and is not part of the system or kits.

MD Dicompass Gateway does not contain a medicinal substance, including derivatives of human blood or plasma, or tissue or cells of human origin or their derivatives, or tissues or cells of animal origin or their derivatives. It is not intended for introduction into the human body.

Dicompass Gateway is not a single-use resource. It is supplied non-sterile and is not intended for sterilization.

Dicompass Gateway is delivered exclusively electronically and thus does not use a storage requirement.

We recommend using the Dicompass Gateway measurement function only on calibrated images by the modality manufacturer. For non-calibrated images, the results of the MD Dicompass Gateway measurement function are only informative, and to clarify the result, we recommend using the calibration tools of the used modality.

The technical description of the device is part of the service manual.

1.4 System requirements {#intro_System requirements}

The chart describes the minimum required configuration per server for SW operation Dicompass Gateway:

Parameter	Requirement
CPU	4 cores
RAM	8 GB
HDD	according to desired volume of data
Network interface	1 Gbps

The chart describes the minimum required configuration of a PC for SW operation Dicompass Gateway:

Parameter	Requirement
OS	Windows 10, 11 (64 bit)
	MacOS
Internet browser	Google Chrome (minimum version 100)
	Microsoft Edge (minimum version 100)
	Mozilla FireFox (minimum version 100)
	Safari (minimum version 15.4)
CPU	2 cores
RAM	4 GB
HDD	0.5 TB
Network interface	100 Mbps

For the correct functionality of MD Dicompass Gateway, it is required that no restrictions be applied to the amount of data transferred or to the number of requests.

The installation and configuration of the MD Dicompass Gateway is carried out according to the service manual, and a handover protocol containing its validation is kept on the correctness of the installation and the functionality of the device at the customer's place. Installation and configuration of the device is performed only by service technicians trained by the manufacturer.

Based on the terms of the customer's service contract, MD Dicompass Gateway is remotely monitored and serviced 5/8, 24/7, or according to other agreed conditions.

Requests can be reported in three ways, by creating a request directly in the Helpdesk system at: -

https://helpdesk.medoro.cz/ - by email at helpdesk@medoro.org - by phone +420775324005, or to a specific service technician, who then enters the request into the Helpdesk system

Regular maintenance is carried out on the basis of released resource updates and their installation.

For the correct and safe functioning of the device, it is not necessary to calibrate MD for the entire specified period of its life.

There are no risks for persons involved in the installation, calibration or servicing of the devices.

The security measures of the Dicompass Gateway resource are determined by the Linux operating system used and its native firewall, which is configured when the system is implemented. Access to the server part of the system is only allowed to trained service technicians by the manufacturer using SSH key encryption. Access to the user part of the system is protected by two-factor user authentication or by connecting to the customer's LDAP/AD.

1.5 List of terms and abbreviations used {#intro_List of terms and abbreviations used}

Term	Explanation	
AD	Microsoft Active Directory	
AE	Application Entity	
DICOM	Digital Imaging and COmmunication System - it is a standard for imaging, distributio storage and printing of medical data	
DPGW	Dicompass Gateway	
DVR	Direct Volume Rendering - in addition to color, it also uses transparency and can display everything at once. All voxels of volume contribute to the final image	
HP	Hanging Protocol - a protocol enabling to change and automate the form of display of the opened study based on many parameters (type of examination, image position)	
MinIP	Minimum Intensity Projection	
MIP	Maximum Intensity Projection	
MPR	Multiplanar reconstruction	
MWL	a service enabling transmission of patient demographic data within the DICOM protocol - Modality worklist	

List of used terms and abbreviations

Term	Explanation
SSD	Shaded Surface Display, 3D projection way, voxels with a density higher than the given limit contribute to the resulting image
SSO	Single Sign On - automatic user login with their identity logged in the operating system
ID	Patient ID number
VRT	Volume Rendering Technique, 3D projection way

2 Login to the system {#arch_Login to the system}

DPGW supports multiple ways of user authentication / authorization when logging in to the web interface:

- with user name and password from local DPGW database
- with a user name and password authenticated against LDAP or Active Directory
- two-factor authentication (username, password and authenticator)
- Single Sign On (SSO) on stations in the domain in the MS Windows environment
- with certificate local user
- with certificate LDAP/AD user
- by passing login information from HIS (information about currently supported HIS which are able to pass user's identity to DPGW can be given to you by the system supplier)
- with temporary one-off token generated directly in DPGW for online consultations

To log in to the web interface, enter the IP address or domain name of the server running DPGW in the address bar of the web browser. If you do not have SSO, login certificate or other advanced authentication method configured, the system will require you to fill in a username and password:

NAME AND PASSWORD
Login name*
This field is required!
Password*
This field is required!
SIGN IN

Figure 2: img

By clicking on the SIGN IN button the user will be logged in. If the login fails (incorrect username or password), a red bar with an error message will be displayed and the login must be repeated with the correct login information. By default, password guessing prevention function is turned on, so that each time you try to log in incorrectly, the time required to authenticate the user to the server increases. If you have forgotten your login details, contact your system administrator or supplier before your account is completely suspended due to random password guessing.

In the case of two-factor authentication, it is also required to enter the numerical code generated in the selected authenticator. If you are logging in to DPGW using two-factor authentication for the first time, you will be redirected to a page with a QR code that you scan with your mobile device in the selected authenticator and you will be assigned an account with a generated unique number for your login to –PRODUCT- SHORT-NAME–. As an authenticator, we recommend, for example, Google Autheticator, which can be downloaded for free from the online distribution service Google Play and the App Store. To work with Google Autheticator, follow the manufacturer's instructions.

After a successful login into the system, the basic screen for searching the system registers will be displayed. The logged-in user may not have all the features described below available. It depends on the type of license and the functional roles that the user has assigned in the system.

3 Basics of user interface {#arch_Basics of user interface}

Web user interface comprises of three main parts:

- Dialog box for managing and searching registers of DPGW, digitizing, dicomization and importing data.
- DICOM box of DICOM viewer with tools for working with image data.
- Administrator interface for system administration and configuration.

You can intuitively switch between sections using interface controls. If you use a multiple monitor setup, it is possible to configure the web environment so that the search and data displaying are shown separately on the preview monitor and the DICOM browser window on other (diagnostic) monitors.

3.1 Searching in system registers {#arch_Searching in system registers}

G ARCHIVES RECENT VIEWED USER STATION WORKING LISTS DICOM IMPORT VIDEOCONFERENCE DIGITALIZATION TEMPORARY DATA

After a successful login into the web interface, the basic dialog box for working with registers of DPGW is displayed. At the top you will find a bookmark bar:

Figure 3: img

- Archives ARCHIVES searching through user-accessible DICOM archives, including the internal database of DPGW (patients, studies, orders). You can edit, delete, or forward the retrieved data to other AEs or exchange networks (ePacs, ReDiMed). The retrieved data can be opened in integrated DICOM viewer by double-clicking.
- **Recent viewed** viewing recently viewed studies. It is possible to select a filter on User/Station/Organizational Unit it is affected by whom the studies were displayed. The User filter displays recently viewed studies only by the logged in user. Station filter displays recent studies displayed on the entire station with respect to access rights. Organizational unit filter shows recently displayed studies of the entire organizational unit, again taking into account access rights.
- User User private DICOM user data box. The user can import data into this box, for example from a CD/DVD, or another user can forward the data there or the user can forward it himself. The data in this box is visible only to the given user and is thus available at any station where the user logs in.
- **Station STATION** private DICOM station data box. The same rules apply to it as to the user's data box. Only the stored data belonging to the station can therefore be seen by any user who logs into the given station. On the other hand, the user will not see this data on any other station.
- Working lists **WORKING LISTS** work (visit) lists created by user or automatically. Each user/group of users can create any number of worklists that they can share with each other. Individual exams can be added to the worklist simply from the search dialog box or directly from the DICOM viewer. DPGW can even create worklists automatically and execute them based on pre-defined rules when saving modality exams, or when receiving an HL7 message from HIS. This setting must be made by the system administrator.
- **DICOM Import** import DICOM data from CD/DVD/folder into user folder, station or archive. The imported data can then be edited in the user/station folder, displayed in a browser or, after checking, forwarded directly to the PACS archive.
- **Videoconference VIDEOCONFERENCE** interface for real-time video and audio sharing via video conferencing call with support for multiple call participants.
- **Digitalization** interface for digitization and dicomization of images and videos from the grab card, or local folder with support of categorization and loading of the Modality Worklist.
- **Temporary data TEMPORARY DATA** management of areas for temporarily stored DICOM data. Above all, these are the so-called "**Unreceived Data**", i.e. data that could not be stored in the archive due to invalid or inconsistent data. You can correct the data in this area and have it saved again, or download the data in dicomdir format. It is also possible to define an area for the so-called "**Recycle Bin**", where the data deleted by the user from the web interface is moved, so that it can

be restored in case of incorrect deletion.

To the right of the tabs, the name of the currently logged in user, the main menu icon and the icon for hiding the entire dialog box are displayed - the DICOM viewer window is displayed . If you have a multiple monitor setup where the search dialog is still open on a separate monitor, this close icon is replaced by the logout icon from the web interface of DPGW.

In the main menu 🧧 you will find the following items:

- Settings settings access to the administrator menu for system administration and configuration
- Logout 🗉 Logout from the web interface
- Change Password
 Change password password change of the currently logged-in user
- User guide User guide displays the user's manual
- **About** About application displays a dialog box with information about the application version, the user and the station to which the user is logged in
- The last item displays information about the instance name of DPGW and the domain to which the user is logged on.

3.2 Searching in Archives {#arch_Searching in Archives}

Figure 4: img

Searching in archives serves the user for searching available DICOM archives or internal databases and thus serves as one of the main pillars of - PRODUCT-SHORT-NAME–. This search engine tool is displayed as soon as you log into DPGW or select the Archives tab. It includes 3 areas for searching and working with visual documentation: - Search toolbar - Search parameters - Search results

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Figure 5: img

3.2.1 Search Toolbar {#arch_Search Toolbar}

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Figure 6: img

The individual functions of the search toolbar are listed in the following table:

lcon	Function	Description
٩	Search	confirmation of patient or study search according to selected parameters in the form
Ð	Clear Tab	clears all added parameters in the form
8	Add to Favourites	selected search parameters can be saved as a favourite search and then brought up from the favourites list instead of time-consuming filling out of the search fields
*	Show Favourites List	displays a list of saved favourite search parameters
	Forms Settings	possibility to define fields that will be offered during the search on individual levels (Patient, Study, Order)
=	Change Layout	possibility to change the layout of search parameters
С	Refresh	reloading of the search box

lcon	Function	Description
φ	Enable Auto Refresh	turns on automatic reload of the search box (if this option is enabled, this icon is green)
A	Select All	selects all search results for bulk processing
	Deselect All	removes the selection from the previous point
•	View Thumbnails	switches on/off series thumbnails preview (preview on - green icon)
-	Hide Manual Filters	switches on/off displaying of manual filters (displaying on - green icon)
Ŧ	Clear manual filters	removes all added parameters in the searched parameters columns
٠	Column Settings	settings for displayed columns in search results
*	New patient	creating a new patient folder
a	Replace and View	displays selected studies with the replacement of the last displayed ones (double-clicking on the line of the given study also works)
	Add to new tab	display of selected studies in a new DICOM browser tab
G	View patients in tabs	display of selected studies in individual DICOM browser tabs
.	Add and View	displays selected studies with adding them to recently viewed
٣	Move series	marks selected study for transfer to another patient
*	Send selected	forwards selected studies to another DICOM node
٥	Export Selected to DICOMDIR	exports selected images to DICOMDIR
C	Burn Selected	prepares selected studies for burning in .img format
e	Export Selected Images	exports selected images to the selected format
B	Add to Worklist	adds selected studies to the worklist - either an existing one or allows you to create a new one
R	Add Tags	adding information to the selected study
•	Delete Selected	deletes selected studies or series

Some of these tools are also accessible in the context menu, after right-clicking on the selected study.

3.2.2 Search Parameters {#arch_Search Parameters}

The search parameters are used to find the required study stored in the selected PACS archive. To get the most accurate search results, we recommend using all fields for data entry.

After entering the required data, it is possible to bring up the search results using the "Search" button , or using the "Enter" key. To clean the form use the function "Clear form" . Both functions are located in the Search Toolbar.

Individual search categories can be expanded up/down using the \square / \square button, located on the right of the category name.

To enter data in each category of search parameters, click in the required field and enter alphanumeric characters. To delete them, it is possible to use the "backspace"/"delete" key or the \times on the right of the text field.

To speed the work up when searching, you can us	se the characters entered in the individual text fields:

Character	Description	Example
*	Wildcard for series of characters	when you enter "Tes*" in the "Name" field, it searches for all results whose name starts with Tes + all characters, e.g. Test, Tester, Tested, etc.
?	Wildcard for one character	when you enter "Tes?" in the "Name" field, it searches for all results whose name starts with Tes + one character, e.g. Test

The search parameters contain 3 main areas that allow you to search for:

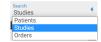


Figure 7: img

- Patients contains categories Patient and Archive Selection
- Studies contains categories Patient, Study, Other, Modality, Tags and Archive Selection
- Orders contains categories Patient, Order, Examination and Archive Selection

The individual categories are listed and described below:

Patient	^
Identification	A
Last name	
First name	
= • Date of birth	—
Gender All gender	•

Figure 8: img

3.2.2.0.1 Patient The "Patient" category contains a field to identify the patient being searched for and contains the following search fields:

- Identification allows entering the patient's birth number
- Last Name allows entering the patient's surname
- Name allows entering the patient's name

	< <	AUG	UST 2	022	> >	
Mon	Tue	Wed	Thu	Fri	Sat	Sun
1	2	3	4	5	6	7
8	9	10	11	12	13	14
15	16	17	18	19	20	21
22	23	24	25	26	27	28
29	30	31				
	CLOS	E		т	ODAY	

Figure 9: img

For the advanced search of the "Date of birth", it is possible to use special characters =, >, < and "between" which can be offered by using the "Expand" icon -:

=	
=	
>	
<	
Between	

Figure 10: img

- 1. The "=" character used to search for patients with the date of birth on a specific selected date
- 2. Character ">" used to search for patients with date of birth after the selected date
- 3. Character "<" used to search for patients with date of birth before the selected date
- 4. "Between" option used to search for patients with date of birth between the selected dates

Betwe 🔻	
Date since	
Date until	

Figure 11: img

• Gender - allows you to enter the patient's gender. The drop-down menu is used to select it:

Gender All gender	•
All gender	
Male	
Female	
Other	

Figure 12: img

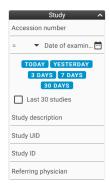


Figure 13: img

3.2.2.0.2 Study The "Study" category contains fields to identify the study you are looking for and contains the following search fields:

- Accession Number allows you to enter the number of order on which the examination was performed
- Date of Examination allows you to enter the examination date in a similar way as described above in the field "Date of birth", or by selecting one of the offered options
- Last 30 studies enables the display of the last 30 examinations received in PACS
- Study Description allows you to enter the name of the study
- Study UID allows you to enter a unique study number
- Study ID allows you to enter a study ID
- Referring Physician allows you to enter the referring physician

Others	^
Series description	
Source AETs	/

Figure 14: img

3.2.2.0.3 Others The "Others" category contains the following search boxes:

- Series Description allows searching by series name
- Source AETs allows searching by DICOM device name (AE Title) from which the data was sent to DPGW

	Modalit	y ^			
ECG	CR	СТ			
DS	DX	ES ES			
01	КО	MG			
MR	NM	OP OP			
ОТ	PR	PT			
PX	RF	sc 🗌			
ST	SR	US			
XA	XC				
	9				
Custom n	Custom modality				

Figure 15: img

3.2.2.0.4 Modality The "Modality" category contains boxes for refining the search for studies based on the selection of the source modality:

Abbr.	Description Abbr.	Description	Abbr.	Description	Abbr.	Description
ECG	Electrocardiog Ca phy	Computed Radiogra- phy	СТ	Computed Tomogra- phy	DS	Digital Sub- traction Angiogra- phy
DX	Digital Ra- ES diography	Endoscopy	Ю	Intra-Oral Radiogra- phy	КО	Key Object
MG	Mammograph y MR	Magnetic Resonance	NM	Nuclear Medicine	OP	Ophthalmic Photogra- phy

Abbr.	Description	Abbr.	Description	Abbr.	Description	Abbr.	Description
ОТ	Other	PR	Presentation State	РТ	Positron emission tomogra- phy (PET)	ΡΧ	Panoramic X-Ray
RF	Radio Fluo- roscopy	SC	Secondary Capture	ST	Single- photon emission computed tomogra- phy (SPECT)	SR	Structured Report
US	Ultrasound	ХА	X-Ray An- giography	XC	External- camera Photogra- phy		

To speed up the search, it is possible to use the functions listed in the table:

Character	Description	Example
	Select All	Marks all modalities
	Deselect All	Deselect all modalities
Ð	Invert	Inverts marked modalities

If you do not find the required modality, you can search for it in the "Custom modality" text box by entering the modality name.

Tags	^
Study contains	-
At least one tag	•
SELECT CLEAR	

Figure 16: img

3.2.2.0.5 Tags The "Tags" category contains the option to select and then search for labeled studies, i.e. studies containing this added information.

• studies can be searched by value for all selected tags or one of the selected tags:



Figure 17: img

Using the "SELECT" button, it is possible to open a table with the option of filtering and marking defined tags:

		Selec	t tags	>
TREE SELECTION FO	RM SELECTION			
^{roup} lategorizace				•
ilter by tag name				
ags to select			Selected tags	
Onemocnění			Komplikace dilatace	8
└→ CN		0	Lokalizace Ostatní	8
L→ UC		0		
→ Neoplázie		0		
ل→ Malignita		0		
→ Divertikulóza		0		
└→ Celiakie		0		
→ Ischemie		•		
L→ Barrett		•		
→ Esofagitida		•		

Figure 18: img

In the "TREE SELECTION" dialog, it is possible to search for and then assign already created labels. In the "Group" bar, you can select the desired set of created labels. In the "Filter by label name" window, you can quickly find this label by entering the desired label name. The left column "Select Label" contains all available labels for the given group and can be selected using the icon • to move it to the right column "Selected Labels". Selected labels can then be removed using the icon •.

In the "FORM SELECTION" dialog, you can similarly select labels for a given study, but here in the "Group" bar you can find e.g. individual departments of the hospital and according to the group selection, labels can be selected by ticking them \blacksquare / \square .

Dicompass Gateway

Select tags						
TREE SELECTION	FORM SELECTION					
Group ORL						
Lokalizace						
Ucho	Nos	🗌 Ústa	Hltan	Hrtan	Jícen	
Krk	🗌 Kůže	Skelet	🗸 Ostatní			
Patologie						
🗌 zánět	nádor	Úraz	Vývojová vada	Exoraezofageáln	CMP/NEURO	
ronchopatie	ostatní					
Terapie						
bez terapie	konzervativní ter	Chirurgická tera				
CANCEL						SELEO

Figure 19: img

After selecting individual labels, select the "SELECT" action to select and then search for the selected labels. With the "CANCEL" action, you cancel the action.

Use the "CLEAR" button to remove labels from the search

Warning: Labels are created and managed only by the system administrator. The user is not allowed to add, modify or delete these labels.

List of flag	^
Select flag	•
CLEAR	

Figure 20: img

3.2.2.0.6 List of flag {#arch_List of flag} The category "List of flag" allows you to search for studies according to the assigned flags. Use the "Select Flag" scroll bar to select the desired flag to search for. Individual flags can be edited, see more in the "Add Flags" chapter. To remove the selected flag from the search, select the "CLEAR" action.

		Order	^
Orde	er ID		
-	•	Date and tim	ie 🛅
	TODA	YESTERD	AY
	3 D/	AYS 7 DAYS	5
		30 DAYS	
	y UID		

Figure 21: img

3.2.2.0.7 Order The "Order" category contains a field to identify the searched orderes and contains the following search boxes:

- Order ID allows you to enter the number of order which the examination was performed on
- Date and time of creation allows you to enter the date of creation of the order in a similar way as described above in the field "Date of birth", or by selecting one of the options offered
- Study UID allows you to enter a unique study number

	Procedure	
Procedur	e ID	
Queue		
	Start	Ċ
TOD	AY YESTE	RDAY
3 [DAYS 7 DA	YS
	30 DAYS	
Dhue ove	mination co	de

Figure 22: img

3.2.2.0.8 Procedure The "Examination" category contains fields to identify the exam you are looking for and includes the following search boxes:

- Procedure ID allows you to enter the examination number
- Queue allows you to enter the name of the workplace where the examination is performed
- Start date allows you to enter the date of the examination in a similar way as described above in the field "Date of birth", or by selecting one of the options offered
- Physician examination code allows you to enter the examination code
- Status allows you to search for active and already completed examinations. The status contains 3 examination options: "All statuses", "Active" and "Completed", which can be marked in the menu opened by the "Expand" icon -:

		Status All statuses All statuses Active Completed	•
Figure 23: img			
		Select archives OPGWTEST8	^

Figure 24: img

3.2.2.0.9 Select Archives {#arch_Select Archives} The "Select Archives" category is used to specify the source PACS repositories from which the results are to be searched and displayed. This category can contain multiple archives depending on the customer's SW infrastructure.

Use the \blacksquare / \square checkboxes to select archives.

	Form items configuration							
Patient	Stu	dy	Or	der	Pre	ocedure		Others
Identification	Accessio	n number	🗸 Order ID		V Proce	dure ID		Series description
Identifier name	🗸 Date of e	xamination	🗾 Date and	time of creation	🔽 Queue			Source AETs
🔀 Last name	🔽 Study des	scription	🛃 Study UID		🔽 Start			
First name	🔽 Study UID				Phys.	examination code		
🔽 Date of birth	🛃 Study ID				🔽 Status			
Gender	Referring	physician						
	Modality	т	ags	List of	flag	Dom	ains	
Mod	dality	✓ Tags✓ Study c	ontains	🖌 List of flag	I	V Domains		
CANCEL								SAVE

Figure 25: img

3.2.2.1 Form Items Configuration {#arch_Form Items Configuration} The form items configuration determines displaying/hiding *■* / □ fields of each search category. fields of each search category. This setting is located in the search toolbar under "Form items configuration" .

3.2.3 Favourite Search Parameters {#arch_Favourite Search Parameters}

3.2.3.1 Saving Filters {#arch_Saving Filters} To make the user's work easier, it is possible to save the most frequently used parameters as filters for searching using the "Add to Favorites" function a. This function is located in the Search Toolbar and saves the complex of currently set search parameters with the option of naming it.

When using the search box "Examination date", DPGW always uses the current date and time.

Save favourite parameters	×
Level User	-
0561	
Name*	
Auto search	
Default after logging	
	SAVE

Figure 26: img

The "Save favorite parameters" feature includes the following saving options:

• Level - possibility to save to a logical drive in the menu opened by the "Expand" icon -:

Level User	•
User	
Station Organizational unit	

Figure 27: img

- User saves the filter to the currently logged in user's account
- Station saves the filters to the currently used station (e.g. PC). The parameters then can be used by all users using this station
- Organizational unit saves the filters to the user's current organizational unit (e.g. doctors, nurses, assistants, etc. according to the current settings by the administrator)
- Name used to name the favorite filter
- Automatic search after selecting a favorite filter, it will automatically search for the desired results and it is not necessary to use the "Search" function
- Default after login after the user logs in, it automatically fills in the search filter according to the specified criteria

3.2.3.2 Load Filters {#arch_Load Filters} The "Favourite Items" function ***** located in the Search Toolbar allows you to select an already saved search filter.

Filer by favourite item name	avourite items				
All today User		0	Q	1	
CT All Organizational unit			Q	1	Î
CT 3 days User		0	Q	1	Î
test2 Organizational unit			۹	1	Ĩ

Figure 28: img

The consequently opened "Favourite Items" table allows you to select or specify the saved filters using the "Filter by favourite item name".

The tools for the saved filters administration are on the right of the saved filters:

lcon	Function	Description
	Default after login	After the user logs in, it automatically fills in the search filter according to the specified criteria
Q	Search	Searches for results according to the parameters of the saved filter
/	Edit	allows you to edit a saved filter. Selecting this function changes the Search Toolbar with the option "Save Changes" 📮 , or "Cancel changes" 💌
	Delete	Removes the saved filter from the favorite parameters list

3.2.4 Search Results {#arch_Search Results}

The "Search Results" are used for displaying the search results corresponding to the entered search parameters.

÷	÷	÷	Ŧ	÷	Ŧ
Full name +	Identification	Date of birth	Modality	Tags	Date of examination
ANONYMNI Anonym (6)	1111	1953-05-16	CT, SR		2015-12-07 07:31
> 🚊 🗖 ANONYM-JZ (6)	ct1236	2018-05-01	CT, SR		2011-03-28 15:16
> 🖹 🗋 ANONYM-OA (18)	petct	2018-05-01	CT, PT, SR		2009-01-28 00:00

Figure 29: img

3.2.4.1 Table Columns Configuration {#arch_Table Columns Configuration} The search results are sorted by value columns which can be set with the "Column Settings" function **2**. After opening it, it is possible to specify displaying of individual columns using the checkboxes. The "Save" button then saves the required columns to the user's account. The "Column settings" parameters may vary depending on the selected search parameter area:

Patient

Table columns configuration						
Full name	🗸 Date of birth	Identification				
		SA	VE			

Figure 30: img

• Study

		e columns configui	ation \
	Full name	Identification	Tation
	Modality	Tags	Date of examination
	Accession number	Archive - AET	Archive - name
	Referring physician	Description	Viewed
	Study ID	Domain	Number of serie
	Finding	🖌 List of flag	
			SAVE
img			
-			
2			
	Tabl	e columns configur	ration
2	Tabl	e columns configur	ration >
	Full name	✓ Identification	Date of birth
Orders	Full name Order ID	IdentificationDepartment	 Date of birth Referring physician

Figure 32: img

After finding the required study parameters, these results can be sorted in descending/ascending order ■/■ by clicking on the value of the required column.

Full name •

Figure 33: img

To specify the searched parameters, it is possible to use text boxes for individual column values:

Figure	34:	img
--------	-----	-----

These text boxes can be hidden by using the "Hide manual filters" function **s**, located in the search toolbar.

When entering parameters in the text boxes of the columns of the searched parameters, these can be removed by the "Clear manual filters" function s, located in the Search toolbar.

If the "Viewed" box is checked in the column settings of the study table, the "Viewed" column will be displayed:

ू Full name +		Tewed
> 🖻 🗖 ANONYM Anonym	1111	
> 🖻 🗖 ANONYM Anonym	1111	

Figure 35: img

The viewing status of the studies is expressed by a colour bar, the colours representing: - green - viewed by logged in user - red - displayed on the station on which the user is currently working (the study could have been viewed by another user) - blue - viewed in the healthcare facility (the study could have been viewed by another user and on another workstation) - grey - study not viewed (this colour replaces one or more of the previous colour statuses)

3.2.4.2 Categories of found data {#arch_Categories of found data} DPGW works with data depending on following sequence:

- PATIENT the main folder containing all performed examinations and stored data, i.e. Studies and Orders
 - Study a component of the examination (X-ray, CT, MRI, etc.), containing individual series
 - * Series a folder with individual instances (e.g. performed protocol for CT examination)
 - Instance specific stored files (e.g. X-ray image)

The search results may vary depending on the selected search parameter area:

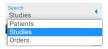


Figure 36: img

• Patients - searches for patients according to the required criteria, i.e. main folders containing individual studies

🛩 🔔 🔲 TEST			1111			<
Level	Accession number	Modality	Date of examination	Study ID	Archive	Description
> 😫 🗋 Study (1)	747489298328849	DX	2007-04-11 11:57		DPGWTEST8	Plice
> 😫 🗋 Study (7)	7474892983288492	DX	2013-10-09 11:32		DPGWTEST8	C, Th
🗸 😫 🔲 Study (15)	7474892983288492	СТ	2017-08-21 11:35		DPGWTEST8	Head^01_Head_Neuro (Adult)
Level	Series preview	Modality	Date & time	Tags	Descriptio	on
III Series 1 (1)		СТ	2017-08-21 11:35		Topogram	n 0.6 T20s
∎∎ 🗖 Series 3 (50)	245	СТ	2017-08-21 11:39		AX Head :	3.0
	1.000					

Figure 37: img

• Studies - searches for individual studies according to the required criteria. The same patient may then appear multiple times in the search results, however, with a different study. When selecting this area, we recommend entering as many search parameters as possible.

✓ 🚊 🔲 TEST (1)		1111			MG	2017-12-19 11 <
Level	Series preview	Modality	Date & time	Tags	Description	
💵 🛄 Series 11 (4)		MG	2017-12-19 11:34		MG FOR PRESENTATION	
✓ 🚊 🔲 TEST (1)		1111			DX	2007-04-11 11:57
Level	Series preview	Modality	Date & time	Tags	Description	
🖷 🗋 Series 1954 (2)		DX	2007-04-11 11:00		Wrist	
	1.100					

Figure 38: img

• Order - searches for input orders according to the required criteria. The same patient may then appear multiple times in the search results, however, with a different order. When selecting this area, we recommend entering as many search parameters as possible.

🖌 📃 🔲 EINSTEIN Albe	ert (1)		7903140003	1879-	03-14	AC8025	cos_a <
Level	Procedure ID	Queue	Start	Status	Phys. examination code	Examination text	
Procedure	AC8020.1	COS_62	2021-04-05 16:00	Active	PROC-001	Gastro	

Figure 39: img

3.2.4.3 Creating a new patient {#arch_Creating a new patient} The "New Patient" tool is used to create a new patient folder if this folder is missing in the PACS system. It is then possible, for example, to move to this folder the documentation of a patient where demographic data were incorrectly selected already on the device modality.

Selecting the "New Patient" action 🖆 brings up the "New Patient" table:

irst name	Last name	Middle name	
refix	Suffix	Date of birth	
ender ither	← Email	Phone	
entifiers			
dentifier name 🔺 ID	Value		Actions
DD IDENTIFIER			

Figure 40: img

In this table, you can enter patient demographic data in the text fields. In the identifiers section, enter the identifier's name (ID) and also its value; if necessary, you can add another identifier using the "ADD IDENTIFIER" action.

The "CREATE" action creates a new patient folder with the selected parameters. The "CANCEL" action cancels the work performed.

If you enter patient information that already exists in the PACS system, you will be notified by a message:

Figure 41: img

Warning: the created new patient folder can only be found in the "Patient" search category, because it does not contain any studies.

3.2.5 Actions with Search Results {#arch_Actions with Search Results}

After finding the required parameters, DPGW allows you to work with this documentation using the functions located in the search toolbar or in the floating panel on the line of the found patient, study, series and order. These features may vary depending on the search category selected.

The individual steps are described in more detail in this chapter:

3.2.5.1 Search Toolbar Actions {#arch_Search Toolbar Actions}

3.2.5.1.1 Selecting Search Results {#arch_Selecting Search Results} The "Select All/Deselect All" function

If the user wants to select/deselect individual search results, he can do so with a checkbox in the search results column on the left of the patient's name /.

Marking multiple consecutive search results is enabled using the "Shift" key on the keyboard.

3.2.5.1.2 Viewing Search Results {#arch_Viewing Search Results} Search results can be opened and displayed in the integrated DICOM viewer in three modes:

- Replace and View <a>
 -replaces the data opened in the DICOM viewer with the selected search results data
- Add to new tab 🖻 will display the selected studies in a new DICOM browser tab

• Add and View 🖻 - adds the selected search results data to the already open DICOM browser data

You can also view the search results in the DICOM viewer by double-clicking on the result itself. This function substitutes the "Replace and View" mode of a specific result.

According to the above-mentioned "Select Search Results" function of DPGW described above, specific search results can be marked and viewed at the same time.

3.2.5.1.3 Moving Series {#arch_Moving Series} After selecting the chosen series in the "Studies" filter category of the "Archives" window, these series can be moved into the selected study using the "Move Series" icon Societ in the Search toolbar.

Selecting this function changes the icon and the function becomes active. Click on this icon to cancel the action.

To insert selected series into another study, perform the action on the floating panel of the study by using "Insert Selected Series" **1**. To confirm this action, it is necessary to agree to the change of DICOM tags of the series contained in the table "Edit tags value" to move the series by clicking "Save":

Edit tags value X								
VR	Name •	Pattern image	Original image	Result				
SH	AccessionNumber	7474892983288492	7474892983288492	7474892983288492				
🗸 LO	AcquisitionDeviceProcessingDescription	UNIQUE: S:200 L:4.0 SCL d:1.2 g:3 sb:3 eq:-0.3 nr:0 dc:3.9 ba	Wrist	UNIQUE: S:200 L:4.0 SCL d:1.2 g:3 sb:3 eq:-0.3 nr:0 dc:3.9 ba				
🖊 LO	AnatomicRegionSequence[1]/CodeMeaning		Wrist					
🖊 AS	PatientAge							
DA	PatientBirthDate							
TM	PatientBirthTime							
LO	PatientID	1111	1111	1111				
PN	PatientName	Test	Test	Test				
CS	PatientSex	M	M	M				
LO	SeriesDescription	BEDERNI PATER	Wrist	BEDERNI PATER				
UI	SeriesInstanceUID	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22940.0	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22908.0	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22940.0				
LO	StudyDescription	BEDERNI PATER	MAMMOGRAFIE DIAGNOSTICKA	BEDERNI PATER				
CLOSE				SA				

If the series are successfully moved, the system notifies you at the bottom of the screen:

Figure 42: img

Series has been successfully moved

3.2.5.1.4 Forwarding Data {#arch_Forwarding Data} In the "Archives" window, the marked data can be forwarded to the selected destination with the "Forward selected" **•**.

Selecting this function displays a table for possible forwarding of the selected data:

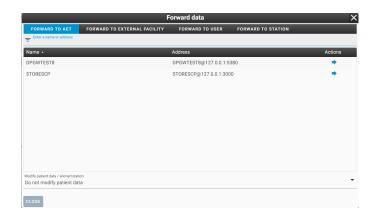
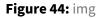


Figure 43: img

This table contains: - Forwarding area: - AET - forwards the selected data to AEtitle (another DICOM modality) - External facility - forwards the selected data to another medical facility

			Forward dat			
FOR	WARD TO AET FORWARD TO EXTERN	AL FACILITY	FORWARD TO USER	FORWARD TO STATION		
- Ent	er a name or address					Show favorit
	Name *	State	City	Address	Information	Actions
*	1. Oční s.r.o.	CZ	Ostrava	Ostrava		ReDiMed 🔻 🔿
☆	3.LF CUNI	CZ	Praha	Praha		ReDiMed 🔻 🔿
*	4D medica sro.	CZ	Brno	Brno		ReDiMed 🔻 🔿
☆	A.H. Media s.r.o.	CZ	Brno	Brno		ReDiMed 🔻 🔿
☆	ABC alergologie s.r.o.	CZ	Brno	Brno		ReDiMed 🔻 븆
☆	Acetabulum	CZ	Uherské Hradiště	Uherské Hradiště		ReDiMed 🔻 븆
☆	Affidea Brno s.r.o.	CZ	Brno	Brno		ReDiMed 👻 🔿
☆	AGEL Clinic	SK	Bratislava	Bratislava		ReDiMed 👻 🔿
☆	AGEL Diagnostic a.s Banská Bystrica	SK	Banská Bystrica	Banská Bystrica		ReDiMed 👻 🔿
	tient data / anonymization modify patient data					



In the list, you can search for the desired medical facility and choose the type of exchange network ePACS, ReDiMed, or mDEX. The selected study can then be sent using the "Forward" action **•**. If you select the mDEX exchange network, you can choose whether to send the data normally or urgently (mDEX statim), data sent in this way will be prioritized for upload.

Individual medical facilities can be marked with a star icon and included as favorites. By checking the "Show favorites" box, you will subsequently filter out only marked medical facilities.

- User saves the selected data to the currently logged in user's account
- Station saves the selected data on the currently used station (e.g. PC) and the selected data can be seen by all users using this station
- Text filter for easy search

- List of possible destinations after selecting a specific destination to send data to, data can be sent with the action "Forward" •
- Modify patient data/anonymization before forwarding, the selected data can be modified using

Modify patient data / anonymization Do not modify patient data	
Do not modify patient data	
Modify patient data	
Automatic anonymization - random values	
Automatic anonymization - Czech names	

the drop-down bar function: Automatic anonymization - Czech names Automatic anonymization - English names

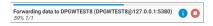
- Do not modify patient data DICOM tags will not be changed during forwarding
- Modify patient data manual editing of DICOM tags of selected data; the table will be

Modify patient data Modify patient data
First name
Last name* Test
Middle name
Prefix
Suffix
Patert ID* 1111

extended by the following lines: [[[]]

- Automatic anonymization random values
- Automatic anonymization Czech names
- Automatic anonymization English names

By selecting the "Forward" action, the process of sending data to the chosen destination is started. The status of the upload process is displayed in the lower right corner of the screen:



```
Figure 45: img
```

After the forwarding process is completed, the system informs about the status of the result in the "Complete data forwarding" window:

Complete data forwarding	×
Forwarded to: DPGWTEST8	
(DPGWTEST8@127.0.0.1:5380))
Successfully forwarded 51/51:	_
CLOSE (5)	ETAIL

Figure 46: img

Warning: if you are sending anonymized data, the patient renaming information will be listed in the table called by the "DETAIL" action in the format "Identity changed from [PATIENT NAME (ID)] to [PATIENT NAME (ID)]".

3.2.5.1.5 Exporting to DICOMDIR {#arch_Exporting to DICOMDIR} In the "Archives" window, the selected patients, studies or series can be saved into the local storage of the station in the set destination selected in the web browser settings. This data is stored in the DICOMdir format, i.e. the folder of DICOM files compressed by the zip method. The action can be performed by using "Export to DICOMDIR" icon .

Clicking the icon opens "Export to DICOMDIR" table:

Export to DICOMDIR	×
Filename DICOMDIR-1111-Testzip	
Modify patient data / anonymization Do not modify patient data	•
CANCEL	EXPORT

Figure 47: img

This table contains the following parameters: - Filename - text box for selecting the name of the saved .zip file - Modify patient data/anonymization - allows anonymization of patient data, similar to the "Forward" function described above

Selecting the "Export" button starts the downloading process. The status of the export process is displayed in the lower right corner of the screen:

Creating DICOMDIR (Test) 24% 1/1	1	0
--------------------------------------	---	---

Figure 48: img

Warning: To enable file downloads, it is necessary to enable pop-ups in your web browser (Chrome, Firefox, etc.).

3.2.5.1.6 Burn selected {#arch_Burn selected} In the "Archives" window, the selected patients, studies or series can be saved in the local storage of the station in the set destination selected in the web browser settings. This data is stored in ISO format, i.e. a CD image. The action can be performed by using "Burn" function **@**.

Clicking the icon opens "Preparing data for burning" table:

Preparing data for burning	
Filename Test-	
Attach embedded viewer	
Modify patient data / anonymization	
Do not modify patient data	•
CANCEL	CREATE

Figure 49: img

This table contains the following parameters: - Filename - text field for selecting the name of the saved .ISO file - Attach embedded viewer - an executable DICOM viewer will be integrated into the ISO file for possible displaying of burned DICOM data - Modify patient data/anonymization - allows anonymization of patient data, similar to the "Forward" function described above

Selecting the "Create" button starts the downloading process. The status of the export process is displayed in the lower right corner of the screen:

Preparing data for burning 57% 1/3	00
5/% 1/3	

Figure 50: img

Warning: To enable file downloads, it is necessary to enable pop-ups in your web browser (Chrome, Firefox, etc.).

3.2.5.1.7 Exporting Images {#arch_Exporting Images} In the "Archives" window, the selected patients, studies or series can be saved in the local storage of the station in the set destination selected in the web browser settings. This data is saved in the selected file format (JPG, PNG, TIFF) compressed by the zip method. The action can be performed by using the "Export images" function **@**.

Clicking the icon opens "Export images" table:

Export images	×
Filename EXPORT-1111-Test.zip	
Print patient data to the images	
Export file format JPG	•
Modify patient data / anonymization	•
Modify patient data / anonymization Do not modify patient data	EXPO

Figure 51: img

This table contains the following parameters: - Filename - text field for selecting the name of the saved .zip file - Print patient data to the images - adds OSD labels directly to the image - Export file format - option to select exporting to JPG, PNG and TIFF - Modify patient data / anonymization - allows anonymization of patient data, similar to the "Forward" function described above

Selecting the "Export" button start the downloading process. The status of the export process is displayed in the lower right corner of the screen:

Exporting images (Test null) 8% 1/1	00

Figure 52: img

To enable file downloads, it is necessary to enable pop-ups in your web browser (Chrome, Firefox, etc.).

3.2.5.1.8 Adding to Worklist {#arch_Adding to Worklist} In the "Archives" window, you can add selected patients or studies to worklist using "Add to worklist" **.**Clicking this icon opens "Add to working list" table:

	Add to working list	
Filtrer by list name		
List name 🔺	Note	Add
gastro		+
test		+
CREATE WORKING LIST		CLO

Figure 53: img

This table contains the following parameters: - Text filter to facilitate searching for created work lists which can be sorted in descending/ascending order $\blacksquare / \blacksquare$ - List of created worklists - List name - Note - the option to insert a note on the study shortcut added to the worklist - Add + - adds a study shortcut to the selected worklist - Create working list - opens a table with the option to create a new worklist and name it:

	Working list	×
List name*		
CANCEL		SAVE

Figure 54: img

• Cancel - cancels performed action

If the data shortcuts have been successfully added to the worklist, the system notifies you at the bottom of the screen:

The study was successfully added to working list

Figure 55: img

For more information on using worklists, refer to the "Worklists" chapter.

3.2.5.1.9 Adding Study Flags {#arch_Adding Study Flags} In the "Archives" window, you can assign flags to individual studies to add additional information. Clicking the "Add flags" icon **S**, which brings up the following table:

Add Study flags	X
= Filter by name	
▲ Urgent	
Událost	
亂 Second read	
🖉 In progress	
For shredding	
CLOSE CREATE STUDY FLAG	SAVE

Figure 56: img

This table contains the following parameters:

- Filter by name text field for finding a specific flag name
- List of created flags with the option to select a specific one
- Close cancels the current action
- Create study flag creates a new flag with the possibility of naming it and assigning an icon to it (icons follow Google fonts and can be found here: https://fonts.google.com/icons, after entering their name, the selected icon will appear):



Figure 57: img

• Save - saves the selected flag for the study. If the flag was successfully added to the study, the system notifies you at the bottom of the screen:

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"Event" has been successfully create

Figure 58: img

Once saved, the individual flag icons will be visible in the search results window in the "Available Flags" column.

The flags assigned to the study can be managed using the "Manage Study Flags" function \Box located in the floating study panel. The opened table allows you to add and remove flags:



Figure 59: img

Individual flags can be managed in "Settings" of DPGW in the "Study Flags" tab. More detailed information on flag management is described in the "Settings" chapter.

3.2.5.1.10 Shredding In the "Archives" window, the selected data can be deleted using "Shred items" function **1**.

Warning: This data will be irreversibly deleted from the archive, so use this action with caution.

Clicking the "Shred Items" icon opens "Shred" table with the following options:



Figure 60: img

- Shredding reason Text field for entry of required data shredding reason
- Information about shredded data
- Cancel cancels the performed action

• Shred - performs the action

Warning: For possible shredding of selected data, the user's authorization based on his roles is required. If you do not see this option and it is needed, contact your administrator.

Figure 61: img

3.2.5.2 Floating Panel Actions {#arch_Floating Panel Actions} In the window of search results, it is possible to open a menu of floating panel actions for individual patients, studies, series and orders using the arrow for showing it < and hiding it <.

Floating panel actions vary depending on selected data areas (patients, studies, series, and orders).

Floating panel actions can also be called using the context menu by a right mouse click on the selected study:



Figure 62: img

If you have multiple studies selected and you call up the context menu by right mouse click on a selected study, you are alerted to this in the context menu by double checking to the right of each tool:

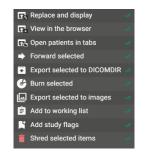


Figure 63: img

The individual functions of the floating panel are listed in the following table:

lcon	Function	Description
ព	View in Viewer	displaying of selected studies with replacing currently displayed (for more information, see chapter "Viewing Search Results")
F.	Add and View in Viewer	displaying of selected studies with adding to already displayed (for more information, see chapter "Viewing Search Results")
R	Add to new tab	display of selected studies in a new DICOM browser tab (more information in the chapter "Displaying search results")
0	Detailed Information	opens a table with detailed information about selected data
ΕK	Select Patient for Digitalization	Selects patient's initials and inserts them in the "Digitalization" window
/	Edit Patient/Procedure	allows editing of patient's information or editing information about procedure on order
ľa	Edit Study/Order	allows editing of information about the selected study/order
	Forwarding data	forwarding selected studies to another DICOM node (for more information, see chapter "Forwarding Selected Data")

Icon	Function	Description
•	Export to DICOMDIR	exports selected images to DICOMDIR (for more information, see chapter "Export Selected to DICOMDIR")
F	Burn	prepares selected studies for burning in CD image format (for more information, see chapter "Data Burning")
	Export Images	exports images to the selected format (for more information, see chapter "Exporting images")
٩	Study Forwarding to External Network	allows sending selected studies using external networks (ePACS, ReDiMed, etc.)
大	Patient Merge	merges two patients into one
۲	Move Study/Series	marks study/series for moving them to different patient/study (for more information, see chapter "Moving Series")
Ĩ	Move Images	marking for moving selected images to another patient/series
Ē	Add to Worklist	adds selected studies to either existing worklist or allows to create a new one (for more information, see the chapter "Add to Worklist")
Ø	Add to Last Worklist	adds selected studies to last selected worklist (for more information, see chapter "Add to worklist")
	Flags Management	adding and removing flags of selected study (for more information, see chapter "Adding Study Flags")
	Add Last Flag	adds last selected flag to selected study (for more information, see chapter "Adding Study Flags")
	Shred	deletes selected patient (for more information, see chapter "Shredding")

lcon	Function	Description
	Shred instances	delete individual images

0

Figure 64: img

3.2.5.2.1 Detailed Information {#arch_Detailed Information} Clicking the "Detailed Information" icon opens a table with overview of detailed information of the selected data:

	Detail	X
Patient info		
Patient name ANONYM Anonym Gender Male	Date of birth 2023-01-01 UUID 3eb7250d-efce-4eef-9ce5-49835470b6ec	Identification
Study info		
Date of examination 2023-01-01 10:00	Modality CT, SR	Tags
Accession number 111112223333	Study ID 1234	Study description Abdomen
Referring physician	Requesting physician	Number of series 6
uid 1.2.840.113619.2.55.3.4271045733.996.1449464144	uuo c995c8b2-a1b0-48e8-b7b3-d3174f7a63e9	
Other info		
Archive - AET DPGWTEST8	Archive - name DPGWTEST8	
CLOSE		

Figure 65: img

It has only informative function without the possibility of data modification.

Figure 66: img

3.2.5.2.2 Edit Patient/Exam {#arch_Edit Patient or Exam} DPGW allows you to edit patient demographics and exams on orders. This paragraph describes the procedure for editing the data of individual areas:

1. Edit Patient

Clicking the "Edit Patient" icon opens a "Pacient" table with following data to be edited:

Last name* Test	Middle name	
Suffix	Date of birth	Ē
•		
Value		Actions
		Actions
	Test Suffix	Test Middle name Suffix Date of birth

Figure 67: img

- Name
- Last name
- Middle name
- Academic title before
- Academic title after
- Date of birth date of birth format in the form of YYYY-MM-DD, or the option to open a calendar (described in more detail in the section "Search criteria->Patient->Date of birth")
- Gender allows you to select the gender of the patient by selecting a parameter in the drop-down bar (described in more detail in the section "Search criteria->Patient->Gender")
- Identifiers a list of added identifiers and the possibility of removing them
- Add identifier creating of new patient identifiers (in this case birth number with value 1111)
- Cancel cancels the performed action
- Save saves the modified patient data and if the modification was successfully saved, the system notifies you at the bottom of the screen:

Figure 68: img

Patient has been successfully edite

2. Edit Procedure

This action is located in the floating panel of the selected procedure.

Clicking the "Edit Procedure" icon opens a "Procedure" table with following data to be edited:

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	Procedure		×
Procedure ID* AC8020.1	Queue COS_62	Status Active	•
Phys. examination code PROC-001	System Medoro	Start 2021-04-05 16:00	
End			
Examination text Gastro	Description Gastro		
CANCEL		-	SAVE

Figure 69: img

- Procedure ID procedure designation
- Queue the name of the workplace performing the procedure
- Status indicates whether the patient has already undergone the procedure or not. This can be changed in the drop-down menu with the values Active/ Completed:

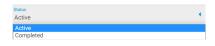


Figure 70: img

- Examination code designation of the examination performed
- System the system in which the order was created
- Start the format of the examination commencement in the form of YYYY-MM-DD, or the option to open a calendar (described in more detail in the section "Search criteria->Patient->Date of birth")
- End the format of the examination completion date in the form of YYYY-MM-DD, or the option to open a calendar (described in more detail in the section "Search criteria->Patient->Date of birth")
- Examination Text the name of the performed procedure
- Description the reasoned description of the required procedure
- Cancel cancels the performed action
- Save saves the modified exams

AB

Figure 71: img

3.2.5.2.3 Edit Study/Order {#arch_Edit Study or Order} DPGW allows you to edit study data and orders. This paragraph describes the procedure for editing the data of individual areas:

1. Edit Study

Clicking the "Edit Study" icon opens a "Study" table containing the data to be edited:

	Study		×
Study ID	Accession number 7474892983288492	Description Coronary*Diagnostic Coronary Catheterization	
Referring physician	Requesting physician	Date of examination 2019-02-10	
Study time 05:08:44	0		
CANCEL			SAVE

Figure 72: img

- Study ID study designation
- Accession number the designation of the order on the basis of which the procedure was performed
- Description name of the performed procedure
- Referring physician the doctor performing the examination
- Requesting physician the doctor creating the request for the examination
- Date of examination examination date format in the form of YYYY-MM-DD, or the option to open a calendar (described in more detail in the section "Search criteria->Patient->Date of birth")
- Study time examination time format in the form HH-MM-SS
- Cancel cancels the performed action
- Save saves the modified study, if the modification was successfully saved, the system notifies you at the bottom of the screen:

Study has been successfully edited



2. Edit Order

Clicking the "Edit Order" icon opens a "Order" table containing the data to be edited:

	Order	×
Order ID* AC8025	External ID AC8025	DAC group
Priority Routine	Status Active	Department
Referring physician	Requesting department	Requesting physician
Weight	Height	Age Unit 👻
Description Gastro		
CANCEL		SAVE

Figure 74: img

- Order ID the designation of the order on the basis of which the examination was performed
- External ID the identification of the order made by the external SW that sent the request (RIS)
- DAC group the item used for data access control purposes
- Priority the option to select the urgency of the examination request by selecting the priority in the drop-down menu:

Priority Routine	•
Statim	
Routine Referring physician	

Figure 75: img

• Status - indicates whether the patient has already undergone the examination or not. This can be changed in the drop-down menu with the values of Active/Completed:

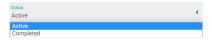


Figure 76: img

- Department the designation of the department in which the patient is hospitalized
- Referring physician the doctor performing the examination
- Requesting department the designation of the department of the doctor creating the order for the examination
- Requesting physician the doctor creating the order for the examination
- Weight
- Height
- Age the ability to specify a unit of time in the case of a patient who has not reached 1 year of age in the drop-down menu:



Figure 77: img

- Description the name of the performed procedure
- Cancel cancels the performed action
- Save saves the modified order

Q

Figure 78: img

3.2.5.2.4 Study Forwarding to External Network {#arch_Study Forwarding to External Network}

This table contains audit records of the patient's referral to another medical facility through the excternal network.

Study forwarding to external network							
Created •	Destination name	Login name	User name	Station name	Network	Domain	Forwarded
2022-07-25 11:30	nemocnice			172.16.23.61	ePACS	Nemocnice	545/545

Figure 79: img

Figure 80: img

3.2.5.2.5 Patient Merge {#arch_Patient Merge} This function is used to merge two patients into one.

大

This action can be performed in the searched data window in the "Search Patients" area.

The implementation of the action process: 1. Mark the patient whose demographic data you want to keep by using the floating panel "Patient Merge" **1**. When the patient is selected, the system notifies you at the bottom of the screen:

Patient selected. Now you can select other patients to merge with this patient.

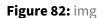
Figure 81: img

After selecting a patient to merge, a new icon will appear in the "Patient Merge" search toolbar with the number of selected patients 1.

2. Mark the patient to be merged into the first patient by clicking "Patient Merge" I on the floating panel. If the patient has been selected, the system notifies you at the bottom of the screen:

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Patient for merge was selected. You can select another patient.



After selecting a patient to merge, a new icon will appear in the "Patient Merge" search toolbar in the number of selected patients 2.

- 3. If you want to merge more patients into the first, proceed as in point 2. The toolbar icon will change depending on the number of patients selected.
- 4. To merge, click the "Patient Merge" in search toolbar to open a table of patient merge options:



Figure 83: img

The Patient Merge table offers the option to swap patients using this function 🗟.

To perform the action, press MERGE or to cancel the action, press CANCEL.

If the patients are successfully merged, the system notifies you at the bottom of the screen:



ኘ

Figure 85: img

3.2.5.2.6 Move instances {#arch_Move instances} The function serving the purpose of selecting specific images in the "Study" filter area in the "Archives" window, and moving these images into the selected series. Clicking the "Move instances" opens "Select images" table.

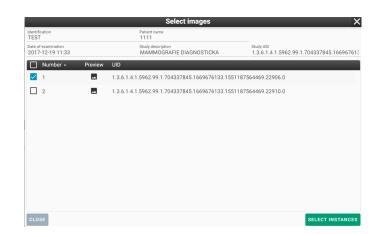


Figure 86: img

In this table, you can use the checkboxes \blacksquare / \square to select specific images and then confirm their selection by clicking "Select Instances".

If the images are successfully selected, the system notifies you at the bottom of the screen:

Study has been successfully selected. Please select target series.

Figure 87: img

Using this function adds the "Delete selected data" it to the toolbar. Click this icon to cancel the action.

To insert selected images into another series, use the "Insert selected images" Σ . on the floating panel. To confirm the action, it is necessary to agree to the change of DICOM series tags by clicking "Save" in the table "Edit tags" to move images:

	Edit tags value X								
	VR	Name •	Pattern image	Original image	Result				
	SH	AccessionNumber	7474892983288492	7474892983288492	7474892983288492				
<u>~</u>	LO	AcquisitionDeviceProcessingDescription	UNIQUE: S:200 L:4.0 SCL d:1.2 g:3 sb:3 eq:-0.3 nr:0 dc:3.9 ba	Wrist	UNIQUE: S:200 L:4.0 SCL d:1.2 g:3 sb:3 eq:-0.3 nr:0 dc:3.9 ba				
\checkmark	LO	AnatomicRegionSequence[1]/CodeMeaning		Wrist					
\checkmark	AS	PatientAge							
	DA	PatientBirthDate							
	TM	PatientBirthTime							
	LO	PatientID	1111	1111	1111				
	PN	PatientName	Test	Test	Test				
	CS	PatientSex	M	M	M				
\checkmark	LO	SeriesDescription	BEDERNI PATER	Wrist	BEDERNI PATER				
	UI	SeriesInstanceUID	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22940.0	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22908.0	1.3.6.1.4.1.5962.99.1.704337845.1669676133.15 51187564469.22940.0				
\checkmark	LO	StudyDescription	BEDERNI PATER	MAMMOGRAFIE DIAGNOSTICKA	BEDERNI PATER				
CLOS	SE				SAVE				

Figure 88: img

If the images are successfully moved, the system notifies you at the bottom of the screen:

Dicompass Gateway User guide

 Images has been successfully moved
 ×

 Figure 89: img
 Images has been successfully moved

 Figure 90: img
 Images has been successfully moved

3.2.5.2.7 Shred instances {#arch_Shred instances} The function for deleting individual images. Clicking the "Shred instances" icon on floating panel opens a table for selecting individual images:

Identification		Select images	
TEST		1111	
Date of examination 2018-04-13 13:23		Study description BEDERNI PATER	Study UID 1.3.6.1.4.1.5962.99.1.704337845.1669676
Number - P	review	UID	
1	-	1.3.6.1.4.1.5962.99.1.704337845.1669676	5133.1551187564469.22938.0
1	-	1.3.6.1.4.1.5962.99.1.704337845.1669676	5133.1551187564469.22906.0
2	-	1.3.6.1.4.1.5962.99.1.704337845.1669676	5133.1551187564469.22941.0
3	-	1.3.6.1.4.1.5962.99.1.704337845.1669676	5133.1551187564469.22912.0
4	-	1.3.6.1.4.1.5962.99.1.704337845.1669676	5133.1551187564469.22914.0
CLOSE			SHRED INSTANCE

Figure 91: img

In this table, you can use checkboxes explore check boxes to select specific instances and then confirm their shredding by clicking "Shred Instances".

To confirm the action, you will be asked to enter the reason for shredding:

Shred	×
Are you sure that you want to shred these items?	
Please enter shredding reason*	
This field is required!	
Patient: TEST (1111)	
Study: DX (2018-04-13 13:23)	
Series: 1 - DX	
Instance: 1 (1.3.6.1.4.1.5962.99.1.704337845.1669676133.1551187564469	9.22938.0)
CANCEL	SHRED

Figure 92: img

Warning: This data will be irreversibly deleted from the archive, so use this action with caution.

Warning: For possible shredding of selected data, the user's authorization is required based on his role. If you do not have this option available and it is desirable, contact your administrator.

3.3 Recent Viewed {#arch_Recent Viewed}

The "Recent Viewed" tab RECENT VIEWED is used for displaying recently viewed studies in a viewer window.

It is possible to select a level filter of User/Station/Organizational Unit that affects whom the studies were displayed by:

Level User		•
User		
Station		
Organiza	ational unit	

Figure 93: img

- User displays recently viewed studies only from the user that is logged in
- Station displays recent studies displayed on the workstation
- Organizational unit displays recently viewed studies of the entire organizational unit

Working with these studies is then similar to working in the "Archives" tab, i.e. using the "Search toolbar" and "Floating panel actions" whose functions are described above in this user manual.

3.4 Local data - station / user {#arch_Local data on station and user}

DPGW allows you to forward data to a specified organizational unit, i.e. to a specific user or station. To forward, use the "Forward selected" function in the "Search toolbar", or "Floating panel actions" . The opened "Forward data" table contains several tabs. Select the bookmark **FORWARD TO USER** to forward data to the user , or select the bookmark **FORWARD TO STATION** to forward the data to the station. When you select the required tab, you can then search and browse existing users and stations. Use the "Forward" button to forward to the found destination . The forwarded data is stored in the organizational unit folder of the user/station located on the server with installed DPGW, so the user does not download the data directly to the used workstation. Access to the forwarded data thus depends on the selected organizational unit:

3.4.1 User

To access the data forwarded to the user, use the "User" tab

The "User" tab is a private DICOM box for user data. The user can import data into this box, for example from a CD/DVD, or another user can forward it to him there or he can forward the data himself. The

data in this box is visible only to the given user and is thus available at any station where the user logs in.

3.4.2 Station

Use the "Station" tab station to access data forwarded to the station.

The "Station" tab is a private box for DICOM data. The same rules apply to it as for the user's box. The stored data belongs only to the station and can therefore be seen by any user who logs on to the given station. Conversely, the user will not see this data on another station.

3.4.3 Work with local data {#arch_Work with local data}

Working with local data, whether by selecting the "User" or "Station" tab is similar to the "Archives" tab. The work is performed on data separated from the PACS archive, so it does not have a direct impact on the data already stored in the production archive. To work with local data, use the functions located in the "Toolbar" or "Floating panel actions".

• Tool bar:

lcon	Function	Description
C	Refresh	reloading of local data window
6 ⁸	Replace and View	displays selected studies with replacing already displayed (alternated by double-clicking on the line of the given study)
B	Add and View	displays selected studies in addition to already displayed
Ĩ	Delete	deletes selected studies or series from the local data tab

• Floating panel action:

lcon	Function	Description
1	Edit patient	allows editing of patient demographic data
	Data forwarding	forwards selected studies to another DICOM node (for
		more information, see the "Forward selected" chapter)

lcon	Function	Description
Î	Delete	deletes selected studies or series from the local data tab

4 Working lists {#wl_Working lists}

The tab "Working lists" working lists allows the user to manually or automatically create work (visitation) lists. Each user / group of users can create any number of working lists that they can share with each other. Individual examinations can be added to the working list simply from the search dialog or directly from the DICOM browser. DPGW can automatically create and fill in working lists based on defined rules when saving examinations from modality, or when receiving an HL7 message from NIS. This setting must be made by the system administrator.

To add a study from the archives tab to the working list, follow the chapter User interface basics-> Search in archives-> Actions with retrieved data-> Actions of the search toolbar->**Add to worklist**

The tab Working lists contains three main areas for working and the displaying archives and studies in them. These areas will be described in the following paragraphs:

- Working list toolbar
- Manage working lists
- Working list studies

D AF	RCHIVES	RECENT VIEWED	USER STATION	WORKING LISTS	DICOM IMPORT	VIDEOCONFERENCE	DIGITALIZATION	TEMPORARY DATA				147
+		C = •	- 🛩 🖬 🔹									
Filtrer by list	t name	Test										
ist name -	Actions				₹	Ŧ	Ŧ	Ŧ	Ŧ	Ŧ	Ŧ	Ŧ
èst	1</td <td>* Full</td> <td>ANONYM Anonym</td> <td></td> <td>Identification</td> <td>Date of birth 2023-01-01</td> <td>Note</td> <td>Accession number 1112223344555</td> <td>Modality CT, PT</td> <td>Date of examination 2023-01-01</td> <td>Source AET DPGW</td> <td>Referring physician</td>	* Full	ANONYM Anonym		Identification	Date of birth 2023-01-01	Note	Accession number 1112223344555	Modality CT, PT	Date of examination 2023-01-01	Source AET DPGW	Referring physician
est1	1</td <td></td> <td>ANONYM Anonym</td> <td></td> <td>1111</td> <td>2023-01-01</td> <td></td> <td>1111122233344333</td> <td>CT</td> <td>2023-01-01</td> <td>DPGW</td> <td></td>		ANONYM Anonym		1111	2023-01-01		1111122233344333	CT	2023-01-01	DPGW	
est2	1</td <td></td> <td>ANONYM Anonym</td> <td></td> <td>1111</td> <td>2023-01-01</td> <td></td> <td>112345</td> <td>CT, PT</td> <td>2023-01-01</td> <td>DPGW</td> <td></td>		ANONYM Anonym		1111	2023-01-01		112345	CT, PT	2023-01-01	DPGW	
			ANONYMOUS		ANON	2023-01-01 2020-11-24		112345	CT, SR	2022-01-01	DPGW	
			TEST Dicom		1234567890	2020-11-24			XC XC	2020-04-15	DPGW	
			TEST test		test				xc	2020-07-21	DPGW	
			TEST TEST		TEST	2022-10-23			ES, SR	2022-12-01	DPGW	
		• 4									_	

Figure 94: img

4.1 Working list toolbar {#wl_Working list toolbar}



Figure 95: img

Individual functions of the working list toolbar are listed in the following table:

lcon	Function	Description
C	Reload	realoads working list
+	Create Working List	creates a new working list
=	Change Working list Order	allows change of order of studies in a working list
V	Open All Items	displays individual studies by expanding patients information in "Studies in working list"
J	Mark All	marks all results in "Studies in working list" for processing more of them at once
0	Deselect All	deselects all fromt the previous option
6	Replace and View	displays selected studies with replacing of already displayed (double-clicking on the line of the given study also works)
0 [*]	Add and View	displays selected studies with adding them to already displayed
	Add all and View	displays all studies in a working list with adding them to already displayed
	View patients in tabs	display of all marked studies in the search dialog in individual DICOM viewer tabs
۵	Replace, View and Delete	displays selected studies with replacing already displayed and removing them from the working list at the same time
•	Delete	deletes selected patients off a working list

lcon	Function	Description
•	Table Columns Configuration	configures displayed columns in "Studies in a working list"

4.2 Management of working lists {#wl_Management of working lists}

Management of working lists is located in the left panel in the tab "Working List". Already created working lists are displayed in this area and DPGW allows working with them.

To simplify work, worklists can be filtered by their name using the "Filter by list name" text field:

Figure 96: img

Working lists can be sorted alphabetically by clicking on the "List Name" line in ascending order List name -, or in descending order Listname -.

After creating worklists using the "Create working list" function 🔹, the names of these lists appear in the "List name" column in this area. The "Actions" column Actions allows you to work with lists and contains the following functions:

4.2.1 Share

Figure 97: img

This feature allows the user to share individual working lists with other users or groups of users created in the system DPGW.

<

"Working list share management" table can be opened by using the "Share" function:

Dicompass Gateway

		Working list share manage	gement					
 Enter the user login, username 	ne or group name							
vailable users		Selecte	d users					
Login name kateg	User name KATEG	Add Login n	ame User name	Share	Add	Comment	Delete	Remove
nonloc	NONLOC	digi	DIGI					•
bac	PAC	0						
test1	TEST1	0						
test2	TEST2	0						
test3	TEST3	•						
vailable groups		Selecte	d groups					
Group name		Add Group r		Share	Add	Comment	Delete	Remove
2FactorAuthCInt		0						
Admin		•						
Archive Query Only		0						
BackOffice		0						
CACHE_errorCache		0						
CACHE_trash		Ð						

Figure 98: img

In the table header, the available users and user groups can be filtered using the text box $\overline{\mathbf{r}}$ Enter the user login, usemame or group name. The search result is applied to both areas at the same time.

To share with a specific available user or group, it is necessary to move the user/group to the area of selected by using action "Add" •. After successful addition, the user or group will be displayed in the "Selected users / Selected groups" area.

Shared users/groups can be given permissions for working with shared worklists using the check boxes / in the columns of "Selected users / Selected groups", containing following functions:

- Share allows selected user / group to view the created working list
- Add allows the selected user / group to add studies to the created working list
- Comment allows the selected user / group to add comments to studies in the created working list
- Delete allows the selected user / group to delete studies in the created working list
- Remove - removes a user / group from the "Selected users / Selected groups" area

After finishing work in the table "Working list share management", "CANCEL" can be used to cancel the performed action, or "SAVE" to save the performed action.

If worklists have been successfully shared with users / groups, the system will notify you with the following bar at the bottom of the screen:

Users with access to working list was successfully updated.

Figure 99: img

4.2.2 Edit

Figure 100: img

The "Edit" function allows the user to change the name of the created working list. Selecting the "Edit" action brings up the "Working list" table:

1

	Working list	×
List name* test		×
CANCEL		SAVE

Figure 101: img

The table contains a text box for editing the name of the worklist, the "CANCEL" button - cancels the performed action or "SAVE" button - saves the performed action.

4.2.3 Remove

Figure 102: img

The function "Remove" allows to delete created working list. Selecting "Remove" action brings up "Question" table:

Î



Figure 103: img

The table contains buttons: "NO" - cancels performed action and "YES" - deletes selected working list.

4.3 Working list studies {#wl_Working list studies}

test								
- Full name		Date of birth				- Modality	Date of examination	Source AET
E TEST Dicom	1234567890	1957-04-15	2021-04-21 08:50			xc	2021- 🖳 😭	
🚊 🔲 TEST Dicom	1234567890	1957-04-15	2021-04-21 08:50			SR, XC	2021-04-19	DPGWTEST8
🚊 🗖 TEST Dicom	1234567890	1957-04-15	2021-04-21 08:51			XC	2021-04-19	DPGWTEST8
🚊 🗖 ANONYMNI Anonym	1111	1953-05-16	2022-07-22 09:40		182-185	CT, SR	2015-12-07	DPGWTEST8
🚊 🗖 TEST	1111		2022-08-04 13:49		7474892983288492	ХА	2019-02-10	DPGWTEST8

Figure 104: img

The "Working list study" area contains individual studies of created working list. To view the studies, it is necessary to select a specific worklist in the left panel "Manage working lists". The name of the worklist (in this case TEST) is displayed in the header of this area and it is possible to work with it further.

4.3.1 Table columns configuration {#wl_Table columns configuration}

Results can be sorted according to value column which can be set by using function "Table columns configuration" . After opening the table it is possible to set displaying of individual columns using the check boxes. The "Save" button then saves required columns.



Figure 105: img

After selecting specific working list, it is possible to sort results in descending/ascending order \square/\square by clicking on the value of the required column.



Figure 106: img

To specify the searched parameters, it is possible to use text fields by the individual column values:



Figure 107: img

4.3.2 Change working list priority {#wl_Change working list priority}

The "Change working list priority" function located in "Worklist toolbar" allows user to change the order of patients added to a specific working list in the "Working list study" area. Selecting the action will bring up the "Change worklist order" table.

Reorder working list		Х
TEST Dicom • 1234567890 63 - 1x dlouhe video, 2x snimek	[XC]	=
TEST Dicom ● 1234567890 62 - 2x video, jedno dlouhe, 3x snimek	[XC,SR]	=
TEST Dicom • 1234567890	[XC]	=
ANONYMNI Anonym ● 1111 Bricho	[CT,SR]	=
TEST • 1111 Coronary*Diagnostic Coronary Catheterization	[XA]	=
CANCEL		ок

Figure 108: img

The change of order of selected patient is then performed by the drag&drop style, i.e. by dragging to the selected position.

The button "CANCEL" - cancels performed action and "OK" - saves order of patients into a working list.

This function is active in case of adding more patients into a working list.

4.3.3 Floating panel action {#wl_Floating panel action}



Figure 109: img

In the "Working list study" area, a menu for individual patients can be opened by the action of floating panel, using the arrow for opening \blacksquare and then hiding ⊇ the menu.

Warning: The floating panel action expands when you add a note.

Individual functions of the floating panel are following:

lcon	Function	Description
厚	Add note	adds note to a selected patient in a working list
民	Edit note	allows to edit added note
睅	Remove note	removes added note
ត	Replace and open	displays selected studies with replacing of already displayed ones (more information in chapter "Displaying of search results")
ត	Replace, open and delete	displays selected studies with replacing of already displayed ones and deleting them from the working list
Î	Delete	removes selected patiens from a working list

4.3.4 Note action {#wl_Note action}

DPGW allows user to add notes to individual patients and work with them in a specific worklist in the range of Add, Edit and Delete:

• Add note 🖪

Choosing the function opens the "Study note" table:

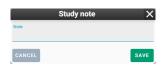


Figure 110: img

The table contains text box for inserting a note and the "CANCEL" button - cancels the performed action or "SAVE" - saves the performed action.

In case that a note is successfully inserted, the system notifies about this with the following bar at the bottom of the screen:

The note was successfully saved

Figure 111: img

• Edit note 🗳

Choosing the function opens the "Study note" table with already filled-in test note for modification:

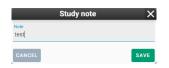


Figure 112: img

The table contains text box for modification of the note, the "CANCEL" button - cancels the performed action or "SAVE" - saves the performed action.

• Remove note 🗳

Choosing the function opens "Question" table:

Question	×
Are you sure that you want delete note?	
NO	YES

Figure 113: img

The table contains buttons "NO" - cancels the performed action and "YES" - deletes the note.

In case that the note is successfully deleted, the system notifies about this with the following bar at the bottom of the screen:

The note was successfully saved

Figure 114: img

5 DICOM data import {#import_DICOM data import}

DICOM data import tab recommender allows the user to upload DICOM files to system logical drives. Folders and .dcm or .zip files located in local storage or on removable media (flash drive, CD / DVD) can aslo be imported.

Warning: Follow your IT administrator's security instructions when importing data from removable media.

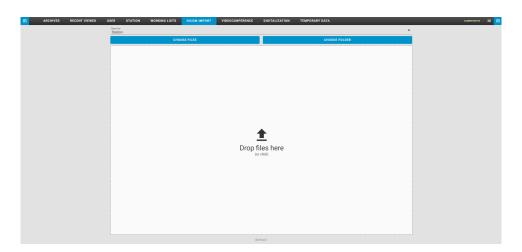


Figure 115: img

5.1 Destination selection {#import_Destination selection}

The destination selection for data entry is located in the header of the "Import DICOM" folder and contains the following system logical units:

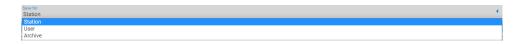


Figure 116: img

- User imports data for currently logged-in used into "User" folder
- Station imports data on currently used station (PC) into "Station" folder. The data can be used by all users using this station
- Archive imports data directly into PACS archive DPGW

5.2 Data input {#import_Data input}

Only folders and files of the .dcm or .zip (containing .dcm files only) type can be imported from the local storage or removable media inserted into the workstation. To enter data, use one of the following options:

5.2.1 Choose files {#import_Choose files}



Figure 117: img

Selecting the "Choose files" options opens a table with local files:

- → ` ↑ 🖡	> This P	C > Downloads	~	O P Search Dov	wnloads
Organize - New	folder			8	•
	^	Name	Date modified	Туре	Size
Quick access		Brain Aneurysm Treatment.mp4	7/25/2022 11:27 AM	MP4 File	20,023 KB
Desktop 🤉	*	1 Test.zip	7/25/2022 11:18 AM	Compressed (zipp	633 KB
🕹 Downloads 🦻	e	videoplayback.mp4	7/25/2022 11:11 AM	MP4 File	626 KB
🛅 Documents 🦻	e	0000.jpg	7/25/2022 10:53 AM	JPG File	9 KB
Notures 🛛 🕫	e 🗸 👘	DICOM	7/25/2022 11:16 AM	File folder	
51		0000.ipg		 All Files (*.*) 	

Figure 118: img

In this table, select the file you want to import. To select multiple files, use the ctrl or shift keyboard shortcuts in this table with the left mouse button or keyboard arrows. Only .dcm and .zip (containing .dcm files only) files can be imported.

After choosing the required files, insert them with the "Open" action, or use "Cancel" to cancel the action.

Files can also be inserted by clicking or using drag & drop style, i.e. by dragging them to the "Drop files here" area:



Figure 119: img

The inserted files appear in the list and other ones can be added as described above:

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Figure 120: img

Use the cross icon on the right of the file name to remove files from the list.

Import of inserted files into chosen destination can be done by clicking the "Import" button.

Figure 121: img

5.2.2 Choose folder {#import_Choose folder}

Figure 122: img

Clicking the "Select folder" button opens a table with local folders:

	• • • • • • • • • • • • • • • • • • • •	PC > Downloads >	~	Ö 🔎 Sear	ch Downloads
Organize 🔻	New folder				E≡ ▼
📌 Quick acces		Name V Last week (1)	Date modified	Туре	Size
♣ Downloads ★ ■ Documents ★	ls ≠	DICOM	7/25/2022 11:16 AM	File folder	

Figure 123: img

In this table, choose the folder you want to import and choose the "Upload" action to insert the data into the list for import. The import list will then contain all the files from the selected folder:



Figure 124: img

Use the cross icon on the right of the file name to remove files from the list.

Import of inserted files into the selected destination can be done by clicking the "Import" button:

Figure 125: img

5.3 Viewing progress and results of import {#import_Viewing progress and results of import}

After clicking the "Import" button, you will be informed about the status of the import process.

1. Information about the status of upload to the browser cache with the display of a progress bar for each file in the list of inserted data:

464/2216

Figure 126: img

And the overall progress of all files:

Figure 127: img

2. Information about the status of upload to selected system destination in DPGW. Information about the start of the import operation is displayed in the middle of the screen:

A job has been started, it may take a while to complete

Figure 128: img

In the lower right hand corner, you will simultaneously be informed about the status of the data import:



Figure 129: img

• action can be used to display more information about the import status. • action can be used to cancel the import.

The data import time varies depending on the volume of the imported data.

- 3. After the data import is completed, you will be informed of the result by the table "Import of DICOM files has been completed". These results may vary depending on the imported data, and one of the following may occur:
- Data imported successfully This option reports that the data import was successful:



Figure 130: img

• Data partially imported - This option reports that the data import was successful only for valid files, the rest ended in error:

DICOM import done	×
DICOMDIR-1111-Testzip - 4 images were successfully imported; 1 file rejected • DICOMDIR - File is unsupported or invalid	
	DISPLAY IN VIEWER

Figure 131: img

• Data not imported - This option reports that the data could not be imported and all data ended in error:



Figure 132: img

When exporting data to the DICOMDIR file folder, a compressed .zip folder is created and it contains a DICOMDIR file and a subfolder with individual images, i.e. .dcm files. In this case, a separate DICOMDIR file was imported but ended in error due to the fact that this file is only a representative of the study and does not contain data for possible import (individual slides).

Successfully imported data can be opened directly from the table of import results using "Display in viewer" button. This function is similar to the "Replace and view" function described in the chapter "Displaying search results".

6 Digitalization and conversion of data into DICOM format {#digi_Digitalization and conversion of data into DICOM format}

6.1 Digitalization tab {#digi_Digitalization tab}

"Digitalization" tab Contains an interface for digitizing and dicomization of images and videos from the grab card, or local folders with support for categorization and loading of the Modality Worklist. This tab contains several main areas for uploading data:

- 1. Select digi station
- 2. Adding the patient and adding information to the procedure
- 3. Dicomization/digitization of data and its modification
- 4. Working record set
- 5. Saving and deleting of uploaded data

These areas are described in the following paragraphs:

6.2 Select digi station {#digi_Select digi station}

If multiple digitizing stations are connected to the DPGW system, the user is allowed to select required digitizing station.

Select digi stat	ion		
C 🖿			
Name •	Host	Connecte Free d	Used by user Actions
Advantech	10.20.0.228	X 🗸	B *
Advantech 4k test	10.20.0.229	X 🗸	
Axiomtek	10.20.0.166	X 🗸	-
COS .65	10.20.0.65	X 🗸	
Digi composite	10.20.0.150	X 🗸	-
Gastroenterology .56	10.20.0.56	× ×	
Havirov Test	10.20.0.155	× 🗸	
Orthopedics .62	10.20.0.62	✓ ✓	-
Urology .63	10.20.0.63	\checkmark \checkmark	-
2vstup	10.20.0.153	✓ ✓	

Figure 133: img

"Reload" action 🖸 updates the list of digitization stations connected to the DPGW system.

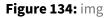
Action "Continue without digitization" 🚬 skips selecting a digitizing station.

The list of digitization stations includes:

- Name the name of the digitizing station
- Host IP address where the digitizing station is located
- Connected information about the availability status of the digitizing station, information indicates the availability of the digitization station, information indicates the unavailability of the digitizing station
- Free information status about using the digitizing station, information station means that the digitization station is not used by another user, information means that the digitizing station is used by another user
- Used by user if the digitizing station is already used by another user, the name of the user is indicated here
- Action by clicking the "Select" action replace the selected digitization station for work (you can replace this action by double-clicking the selected line of the digitization station)

Select the "Digitization" tab to display a dialog for working with records:

ARCHIVES	RECENT VIEWE	D USER	STATION	WORKING LISTS	DICOM IMPORT	VIDEOCONFERENCE	DIGITALIZATION	темро	RARY DATA			administrator 🔳 🖬
Manual	Worklist					Videomatr	rix Dicomization	Edit items	Management	O Quese	Working set	Unprocessed items
A Patient	٩										Files	Select
Identification	▲			Choose files	Choose folder						2024-04-15 (74) 🗸	
Last name												
First name											14.30.48	1430/48
Gender Other	*											0
Date of birth											M-30-50	14 30 52
Procedure											14.050	
Order ID											1	
Study description						1					14:00:54	14:43:11
					Dr	op files here						S
						tmp4, *jpg, *pdf					10.4813	Makia
												1446-02 (00-00-07)
											100	
											144604 (000000)	14.4928
											-	
🔚 Save 🚺 Cancel	«						<u>+</u> 10				»	Selected Delete



6.3 Adding the pacient and adding information to the procedure {#digi_Adding the pacient and adding information to the procedure}

Patient demographics can be entered manually or by selecting from the worklist.

Warning: it is very important to enter patient data with 100% accuracy! If you enter the wrong birth number, last name, first name, it is possible to accidentally duplicate one patient under different details and the images will not be paired.

6.3.1 Manual entry {#digi_Manual entry}

To manually enter patient's demographic data, click the "Manual" tab Manual . Selecting this tab will allow you to enter data using the text boxes:

Realignment	۲
Identification	
Last name	
First name	
Gender Other	~
Date of birth	
En Procedure	
Order ID	
Study description	

Figure 135: img

• Identification - allows entering the patient's birth number - this field is required

- · Last name allows entering the patient's last name this field is required
- First name allows entering the patient's name
- Gender allows entering the patient's gender. The drop-down menu is used to select it
- Order ID allows entering the order number on which the examination was performed
- Study description allows entering the name of the study

Whispering can be used to fill text into text fields, i.e. it automatically completes text when typing in the text field.

To delete text in text fields, you can use the delete function \times located in the field, to the right of the entered text.

For automatic completion of manual entry fields, it is possible to use the actions in the "Archives" dialog, "Floating panel actions" and select the "Select patient for digitization" action for the selected patient \blacksquare , after saving the digitization records, a new study will be created for this patient with the record saved in the PACS archive.

6.3.2 Worklist

The second option is to select the patient from the worklist worklist . It includes patients who are preregistered, booked or waiting for an examination. The image below shows the patients waiting for the examination from the worklist. Click the patient on the list to select them.

	•
scription	
EINSTEIN Albert 7903140003	
HAVEL Vaclav 361005123	
SEDESAT A Dva 6262626262	
SMOLJAK Ladislav 311209123	
	~
	EINSTEIN Albert 7903140003 HAVEL Vaclay 96000123 SEDESAT A Dra 65636525262 SMOLJAK Ladislay 311209123

Figure 136: img

The worklist can be filtered by time period using the scrolling menu in the "Search" section:

Search All days	<
Today	
Yesterday	
Custom date	
All days	

Figure 137: img

If you select "Custom date", you can enter a specific day in the "Date" field or select it using the calendar .

The list of patients in the worklist can be updated using with the "Reload" button $\,{}^{\, \Theta}$.

Select the patient from the worklist, which will move to the "Patient" field, including filling in the "Study description" field:

옥 Patient		
EINSTEIN Albert 7903140003	~	
iption		
	EINSTEIN Albert 7903140003	

Figure 138: img

With this action, a patient will be selected for digitization or dicomization.

Warning: Carefully select the demographic data of the patient from the worklist section, to avoid possible substitution of the patient. If you have selected a patient other than the one you want in the worklist section, you can click on another patient from the worklist to make a change. If you have selected a patient from the worklist and it is necessary to enter the patient manually, click on the "Manual" tab and enter the patient data, the data will be applied according to the active tab.

6.4 Dicomization/digitization of data and its modification {#digi_Dicomization or digitization of data and its modification}

This area allows the import of nonDICOM data and its modification with subsequent conversion to DICOM format. The options for each category is described below:

Warning: The individual categories of dicomization/digitization may differ depending on the selected digitization station, skipping the selection of digitization stations, or assigned user roles and the license used.

6.4.1 Dicomization

The category "DICOMIZATION" <u>to reconstruction</u> is used for conversion of nonDICOM data into DICOM format and subsequent storage in the system of DPGW.

Only files and folders with files such as .jfif, .pjpeg, .jpeg, .jpg, .m4v, .mp4, .pdf can be dicomized from the local storage or removable media inserted into the workstation. To enter data, use one of the following options:

Choose files

Figure 139: img

6.4.1.1 Choose file {#digi_Choose file} Clicking the "Choose files" button opens a table with local files:

⊢ → ~ ↑	• 1	'his P	C > Downloads	~	ບ 🔎 Sea	arch Downloads
Organize 👻 N	ew fol	der				E • 🔳
Quick access	^		Name	Date modified	Туре	Size
Desktop	1		🛋 0001.jpg	7/25/2022 1:43 PM	JPG File	54 KB
Downloads			Brain Aneurysm Treatment.mp4	7/25/2022 11:27 AM	MP4 File	20,023 KB
			videoplayback.mp4	7/25/2022 11:11 AM	MP4 File	626 KB
Documents			0000.jpg	7/25/2022 10:53 AM	JPG File	9 KB
Pictures	* -		DICOM	7/25/2022 11:16 AM	File folder	
	File or		0000.jpg		Custom	Files (*.jfif;*.pjp;*.jpg;*.p

Figure 140: img

In this table, select the file you want to import. To select multiple files, use the ctrl or shift keyboard shortcuts with the left mouse button or keyboard arrows in this table. Only .jfif, .pjpeg, .jpeg, .jpg, .m4v, .m94, .pdf files can be dicomized.

After selecting the required files, insert them using the "Open" button, or use "Cancel" to cancel the action.

Files can also be insterted by clicking or using drag & drop style, i.e. by dragging them to the "Drag files here" area:



Figure 141: img

The imported files then appear in a list and you can add more to them in above-mentioned way:

Filename		
000	2.mp4	创
000	1.jpg	创
000	10.jpg	₫

Figure 142: img

Use the "bin" action to the right of the file name to remove the file from the list.

Import the imported files into the "Working set" by selecting the "Import" action on the right under the list of selected files.

🛃 Import

Choose folder

Figure 143: img

Figure 143: Img

Figure 144: img

6.4.1.2 Choose folder {#digi_Choose folder} Clicking the "Select folder" action opens a table with local folders:

→ ✓ ↑ ↓ > This PC > Downloads	~	U 🔎 Searc	h Downloads
Organize - New folder			• =
This PC Name	Date modified	Туре	Size
3D Objects			
Desktop	7/25/2022 11:16 AM	File folder	
Cocuments			
Downloads			
· · · · · · · · · · · · · · · · · · ·			
Folder: DICOM			

Figure 145: img

In this table, select the folder you want to dicomize and select the "Upload" action to put the data onto the dicomization list. The "Dicomization" category then contains all the files from the selected folder:

Filena	ame	
Ð	0002.mp4	D
Ð	0001.jpg	创
Đ	0000.jpg	₪

Figure 146: img

Use the "bin" action to the right of the file name to remove the file from the list.

Import the imported files into the "Working set" by selecting the "Import" action on the right under the list of selected files.

	Import
–	Import

Figure 147: img

If you select files of any unsupported type, you will be notified by a web browser pop-up window and the files will not be added to the list for dicomization:



Figure 148: img

6.4.1.3 Displaying the process and results of dicomization {#digi_Displaying the process and results of dicomization} After clicking the "Import" button, you will be notified about the status of the data upload process into the "Working set".

1. Information about the status of the data upload progress into the "Working set" with displaying the progress indicator for the individual files in the list of imported data:

	0002.mp4	
B	0001.jpg	ច
	0000.jpg	t

Figure 149: img

The data import time varies depending on the size of the imported files.

2. After the data import is completed, you will be informed about the result by the "Uploaded files info" table:

Importe	d 3/3	
	0002.mp4	~
	0001.jpg	~
	0000.jpg	~

Figure 150: img

Shortcuts to the uploaded files will then appear in the "Working set" toolbar.

6.4.2 Digitization

Digitization is used for conversion of analog data from examination modalities (e.g. endoscope) to DICOM format and subsequent storage in the DPGW system.

Warning: The video preview must not be used for diagnostic purposes. The preview might be delayed.

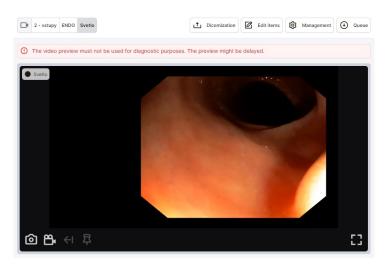


Figure 151: img

If the digitizing station contains multiple outputs, it is possible to display multiple inputs (previews):

Dicompass Gateway

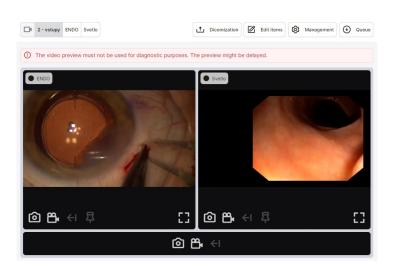


Figure 152: img

To capture a single image from the input of the digitizing station, select the action "Take snapshot" **a**. The image will be captured in the "Working set".

To capture video from the input of the digitizing station, select the "Start video recording" action **B**. Video recording will be marked with an icon, the name of the recording modality and the time of the recorded video **e** recording in the upper left corner of the preview. After starting the video recording, the user is allowed to add **B** marks to the video to mark important parts (see more in the "Edit video" section). The **B** mark action will change color to verify the marking of the video. To stop video recording, choose the action "Stop video recording" **B**. The video will be captured in the "Working Set".

Action "Maximaze video preview" 🖬 allows you to expand the window with a preview of the input from the digitization station.

Action "Timeshift" 🚭 allows the user to go back a period of time in the recorded video, using the displayed table:



Figure 153: img

When reverse recording is active, this information will be displayed next to the video recording icon:

Warning: The "Timeshift" action may vary depending on the configuration of the DPGW product.

Digitization control supports control directly from the camera head of the device or through the pedal control, touch monitor or tablet.

6.4.3 Edit items {#digi_Edit items}

Edit items

Figure 154: img

The "Edit items" category allows the user to edit the uploaded data after selecting the specific record from the "Working set". Data editing options vary depending on the type of record being edited, namely video, image, or pdf:

6.4.3.1 Edit Image {#digi_Edit Image} By selecting an image from the "Working set" opens a preview of the selected image in the "Edit items" category with the following functions:

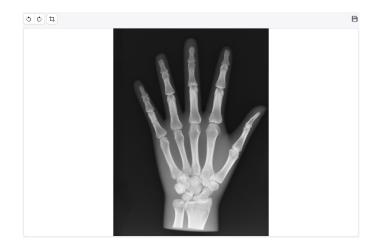


Figure 155: img

• Crop tool 🗅 - this tool is used to crop out a certain area in the image:



Figure 156: img

You can resize the crop area by dragging the red squares located in the upper left and lower right corners of the crop area. When you hover the mouse over the crop area, the numeric values for these squares represent their location in pixels, relative to the cropped image. A numeric value in the center of the crop area indicates the resolution size of the image after cropping. Click in the image to be edited by cropping to display the "Crop area" table for possible manual entry of crop values:

Width	Height	
640	858	
x	Y	
80	107	

Figure 157: img

- Rotate 90° o o the image will be rotated 90° clockwise or counterclockwise, depending on the selected rotation direction
- Download related the image will be downloaded to the local storage of the PC, in case you select "Original" the image will be downloaded without modifications, in case you select "Edited" the image will be downloaded with modifications, in .jpg format
- Save changes 🖻 this is an informative icon indicating the save status of the edit

6.4.3.2 Edit video {#digi_Edit video} Selecting a video from the "Working set" opens the video viewer with the following functions in the "Edit items" category:



Figure 158: img

• "Timeline" - red bar which indicates the location in the selected video. Click or drag right/left to move in the timeline of the video. You can change the timeline scroll speed by dragging up/down:

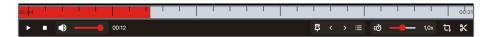


Figure 159: img

- "Play" ► the play function is used to play the video. The beginning of the video can be changed using the timeline.
- "Stop" 💶 the video stop function
- "Volume" 🖸 the function for adjusting the volume of the playing video. You can change the volume by dragging the line
- "Time stamp" for example indicates the current position on the timeline
- "Add mark" 🗳 the function for adding a marker to the selected part on the timeline. Clicking the action adds the mark to the timeline as showed:

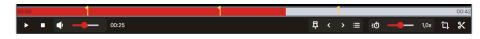


Figure 160: img

You can move between the markers with the left/right arrow <>>.

To delete a selected mark, click it on the timeline. The original "Add mark" action will change to "Remove mark" . Clicking the icon removes the mark.

Action "Marks" 🖬 allows the user to manage added marks. By selecting this action, you will display the "Marks" table, in which it is possible to manually edit individual mark times and add their description:

^{Time} 6.54	s	Description Test
Time 18.88	s	Description
^{Time} 29.95	s	Description

Figure 161: img

Hovering over a mark with a description will display it to the left of the mark insertion action $\blacksquare \blacksquare \bullet \bullet \blacksquare$.

After digitizing the video with the marks into the PACS archive, these marks will be visible when the video is opened in the DPGW DICOM viewer.

- "Speed" C the function for determining the video playback speed. The video can be slowed down to 0.1x playback speed, or accelerated to 4x playback speed using the line 104
- "Crop tool" 📮 for more info, see chapter "Edit image"
- "Editing tool" this tool is used for editing video that will be digitized and then sent to the PACS archive. Selecting this tool will display a bar for defining editing:



Figure 162: img

The bottom bar indicates the location of the video cut. To mark the beginning of video editing, click in the selected part of the bar. You then determine the start and end of the video by moving the squares to the requested area. The marked part of the video can be moved by dragging the yellow part of the section.

The edit tool allows you to add multiple edits to the selected video. To add, click outside the already selected cut and specify the length of the video cut:



Figure 163: img

- "Download" ± rounded the video will be downloaded to the local PC storage in .mp4 format.
- "Save changes" 🖻 this is an informative icon indicating the save status of the edit

6.4.3.3 Edit pdf {#digi_Edit pdf} Selecting a PDF file from the "Working set" will open an external pdf viewer in the "Edit items" category with the option to download it ***** Download it

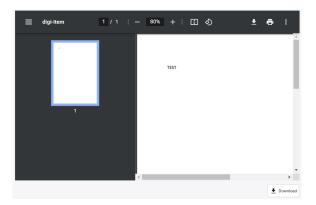


Figure 164: img

6.5 Working set {#digi_Working set}

After uploading the data by dicomization/digitization, this data is moved to the "Working set" area in the right column:

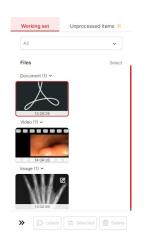


Figure 165: img

A working set allows the user to work with the imported data. When you click on the selected file, it is marked with a red frame and you will be redirected to the "Edit items" category described above.

The pencil icon to the right of the file in the "Working Set" indicates the edited file.

The "Working set" section allows you to perform the following actions:

• "Select" select - allows you to select files for processing in the "Working set" section. Click the left half of the preview to select the image, and click the right half of the preview (with the eye icon) to view the image in the image editing window.

Files	Mark all Cancel
🗌 Video (1) 🗸	
14:48:16 (00:00:01)	
✓ Image (1) ✓	
14:30:44	

Figure 166: img

The "Select" action changes to "Mark all" to mark all files, or "Cancel" to deselect files.

If you select the "Select" action and mark the selected file, the following actions will be available:

- "Delete" 💼 removes selected files
- "Labels" D tabels this action allows the user to assign predefined labels to a file using the displayed table:
- 1. Table for label search search for a label by entering text in the "Filter by tag name" field, click on the searched label in the "Tags to select" field, move it to the "Selected tags" field, and then choose to save:

Tree selection	on	1	Form selection	
Filter by tag name				
Tags to select		Selected tags		
× Cancel				Save 🗸

Figure 167: img

2. Table for label selection - First select the group of labels from the "Group" scrolling menu, then select the required label by clicking in the check box. If you have selected all the required labels, perform the save action:

	Tree selection		Form selection				
Group Gastroenterolog	ie						
Dnemocnění							
CN		Neoplázie	Malignita	Divertikulóza			
Celiakie	Ischemie	 Barrett 	Esofagitida	Infekce			
okalizace							
🗋 jícen	🗌 žaludek	duodenum	🗋 jejunum	🗌 ileum			
Colon	rektum	perineum	pouch	🗌 term. ileum			
Komplikace							
stenóza	absces	🗋 píštěl	Vřed	krvácení			
🗌 skin tags	dilatace	🗋 fisura	perforace				
Morfologie							
🗆 R i0	🗆 R 11	🗌 R i2	🗆 R i3	🗆 R i4			
Mayo 0	Mayo 1	Mayo 2	Mayo 3	sessilní			
stopkatý	C LST	angiodysplázie	poradiační				
Terapie							
EBD	C EPE	EMR	🗆 ST	stent			
FTR	APC						
				_			

Figure 168: img

Warning: Labels may vary depending on DICOM viewer configuration. If you cannot find the label you are looking for, please contact your administrator for this product.

After choosing a label and saving it to a selected file, this file will be marked with a label:



Figure 169: img

• "Selected" - this action moves the selected files from the "Working set" to the "Unprocessed items" tab. The files moved in this way will not be subsequently stored in the PACS archive and it will be possible to work with these files in the future.

6.5.1 Unprocessed items {#digi_Unprocessed items}



Figure 170: img

"Unprocessed items" are those files that have been moved from the "Working set" tab using the "Select" action *reserved* or have been moved automatically, according to a configurable time limit by the product administrator DPGW.

After selecting the selected files, these files can be moved to the "Working set" or deleted. After moving these files to the "Working set", they can be assigned to the required patient and then stored in the PACS.

6.6 Saving, deleting data {#digi_Saving and deleting data}

The last step for saving/deleting acquired data in to the PACS archive is located in the "Save, delete data" area, in the lower left part of the screen.

The following functions are located in the data storage/deletion section:

• "Save" 🖻 🔤 - action for saving acquired data in the PACS archive. By choosing this action, you will display the informative table "Result of saving records":

New items	result	
Imported i	tems to queue: 4/4	
Close (2)	Queue	

Figure 171: img

Select the "Queue" action to display the details of the import of stored data. If all the data has been saved correctly, you will be informed about this with a green check mark to the right of the name of the saved record:

2	TESTR Test 3/3 🔮	()
8	dcm_221213151458_0002.mp4	0
	dcm_221213151508_video_1_20220725_135602_Q8tT11654954272059754281.mp4	0
8	dcm_221213151511_Brain Aneurysm Treatment.mp4	0

Figure 172: img

• "Cancel" 💼 - removal of all records and patient demographic data

Warning: Before uploading data to the PACS system, carefully check the data included in the work set to prevent possible misassignment of data to a patient.

DPGW allows the user, if necessary, to leave a job in the "Digitization" category with the possibility to return to this job later. The entered data is saved and the user can switch between categories or launch the DICOM viewer. Even if the user logs out of the system, the data entered in the "Patient" and "Working Set" areas are kept or moved to the "Unprocessed Records" tab, depending on the configuration and the time since the last logout. After a new login to the system, the user is then notified of this by a table:

Unsynchronized files found	×
New items found when the digi station has been started or the connection has been restored. Please double check the new items if they belong to the current examination or move them to the unprocessed items. The creation date and item for double carbon 223-10-27 13:22-32 Number of found items: 4	⚠
Lunderstand	

Figure 173: img

6.6.1 Videomatrix

The "Videomatrix" • videomatrix category is used for management of video input signals to individual display monitors.

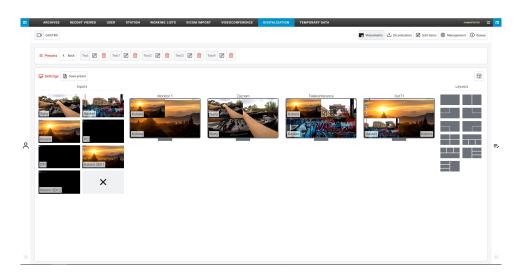


Figure 174: img

The "Videomatrix" category includes: - **Presets** ≔ Presets - selection of saved layouts of individual display monitors

To create a preset, edit the layout of each monitor and select the action "Save Preset" ^{Save preset}, to display the "New Preset" table:

New preset	×
Description Test	
× Cancel	Save 🗸

Figure 175: img

The "Cancel" action cancels the work in progress, the "Save" action saves the preset in the "Presets" field.

The "Edit" rest action allows the user to edit the created preferences. Selecting this tool extends the individual presets you have created to edit them:



Figure 176: img

Use the "Edit" tool [™] to open the "Edit preset" preset name editing table:

Edit preset	×
Description Test	
× Cancel	Save 🗸

Figure 177: img

Use the "Remove" tool 🖆 to remove the created preference.

• **Settings** - **-** Settings - the settings tab contains the current input management, display monitor layout and the option to change the monitor layout.

Use the change view action of the settings tab ^(a) to change the display layout of inputs, display monitors, and layouts:

Settings tab layout 1:



Figure 178: img

Settings tab layout 2:



Figure 179: img

Changing the layout and input of signals:

To change the display monitor layout, select the desired layout from the "Layout" tab and then apply the selected layout to the selected display monitor:

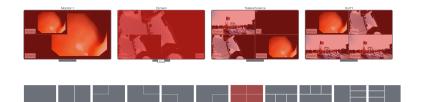


Figure 180: img

To change the display of a video input to a display monitor, click the desired input with the mouse, and then click in the location of the selected display monitor:

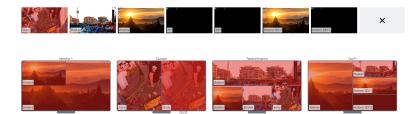


Figure 181: img

This changes the layout on the selected display monitor.

7 Videoconference

The "Videoconference" tab videoconference is used to share video and audio in real time using a videoconference call with support for multiple call participants.

7.1 User videoconference access {#digi_User videoconference access}

As a user of the DPGW system, select the "Videoconference" **VIDEOCONFERENCE** tab to open the videoconference dialog window:

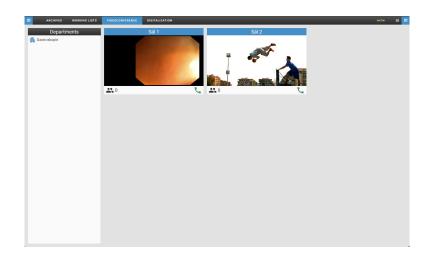


Figure 182: img

In the left column "Departments" you can see the individual departments, by selecting the desired department you can see the available rooms for possible connection via video conference calls. To join a videoconference call, select the desired department and room and then use the "Phone" function \mathbb{V} to request to join a videoconference call. If the call is approved, the videoconference call dialog will be displayed:

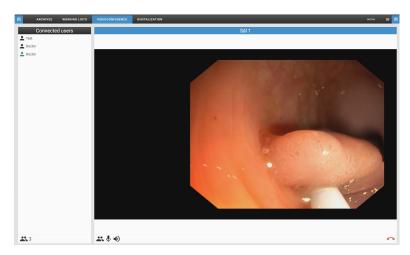


Figure 183: img

Warning: Some users may have privileged access. No confirmation is displayed for their room login and they are automatically logged into the room. For this setting, contact your system administrator DPGW.

In this dialog box, the left column displays the individual connected users. The right column shows the video conference call in session. In this right column, the user is allowed to:

- "Turn off/on microphone" 🎐 / 🗏 this function is used to turn off/on the microphone
- "Mute/unmute" / 🔌 this function is used to mute/unmute the audio of a videoconference call
- "Exit videoconference" 💁 select this function to exit the videoconference call
- "Hide/show users" 🚢 / 🛎 selecting this function hides/shows the left column with the displayed users:

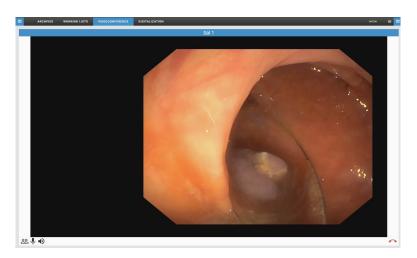


Figure 184: img

7.2 Administrator videoconference access {#digi_Administrator videoconference access}

To create a videoconference call, go to the "Digitization" tab **DEFINITION** and select the desired digitization station (see the Digitization section). The running digitizing station, which is configured for video conferencing, can be accessed remotely by users of this system. If a user wishes to connect to the running digitizing station by videoconference, you will be notified with the option to accept or reject the request:

Connection request	×
The 'Test' user requests access to the video confere	ence.
REJECT	ACCEPT

Figure 185: img

Warning: If privileged users are defined, the call is connected automatically without the need for confirmation.

If you have selected the "Accept" option, the individual connected users will be displayed at the top of the window:

(p) 2	Endoskopie	Matrix
Test Doctor		

Figure 186: img

And in the "Digitization" dialog window, the functions for the management of the videoconference call will be displayed:

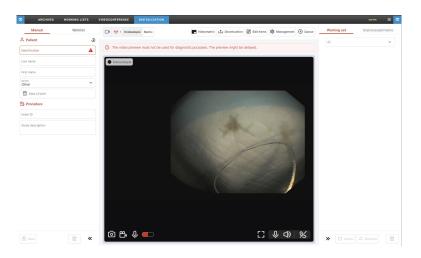


Figure 187: img

Video conferencing management functions: - "Microphone off/on" 🕑 / 🗟 - this function is used to turn off/on the microphone - "Mute/unmute" 🕥 / 🗟 - this function is used to mute/unmute the audio of a videoconference call - "Log out connected users" 🖾 - selecting this function will log out all connected users of the currently running videoconference call

8 Temporary data {#temp_Temporary data}

Tab "Temporary data" THEOREM IS used for managing temporarily stored DICOM data that has not been properly stored in the PACS archive or has been marked for deletion.

Warning: The "Temporary data" tab is dependent on the assigned user role. If you do not have access to the folder, contact your system administrator.

8.1 Types of temporary data {#temp_Types of temporary data}

8.1.1 Cache

Figure 188: img

- Missed data This is data that could not be stored in the archive due to invalid or inconsistent data. You can correct the data in this area and have it saved again, or download the data in dicomdir format.
- Recycle Bin The area where data deleted by the user from the web interface is moved so that it can be restored in case of erroneous deletion. To enable this feature, you must first configure it.

ARCHIVES	RECENT VIEWED USER	STATION	WORKING LIST	S DICOM IMPORT	VIDEOCONFI	ERENCE DIGITALIZATION	TEMPORARY DATA			
_	_	_	_	_	CACHE	EXTERNAL PROCESSING QUEL	IE		_	
Study UID	T Identification		🐨 Patie	nt name	⊤ Sour	rce	〒 Target	= Destination Error cache		
D Patient name	Identification	Date of examination	Modality	Source	Target	Information				Actions
TTESTOVIC mid TTest p	rzd 🚛	1995-07-25	MR	DW	DPGW	org.medoro.dpgw.dicom.C	StoreException: org.med	oro.dpgw.core.CoreException: Mismatch on patient attri	bute [LastName]: DB value=[B	/ 🖬 🗑
		2006-04-12	CR	DPGW	DPGW	org.medoro.dpgw.dicom.C	StoreException: org.med	oro.dpgw.core.CoreException: Mismatch on patient attri	bute [LastName]: DB value=[P	/ 🖬 🗉

Figure 189: img

8.1.2 External processing queue {#temp_External processing queue}

Figure 190: img

This is data that has been sent for processing by third-party applications, e.g. artificial intelligence, a queue of individual processing requests is created here, which can be managed.

EXTERNAL PROCESSING QUEUE

(FI)	ARCHIVES	RECENT VIEWED	USER	STATION	WORKING LISTS	DICOM IMPORT	VIDEOCONFER	RENCE DIGIT	ALIZATION	TEMPORARY DATA			ADMINISTRATOR	= 🛛
							CACHE	EXTERNAL PROC	CESSING QUEUE					
			C	2										
			Ŧ	All statuses								•		
			Na	ame		Status		Sent		Received	Actions			
												*		

Figure 191: img

8.2 Viewing

To facilitate work, lists of data can be sorted and filtered in the tab "Temporary data", using text boxes and drop-down bars which are described below:

⇒ Study UID
 The Identification
 The Patient name
 The Source
 The Target
 The Target

Figure 192: img

- UID study allows user to enter a unique study number
- · Identification allows user to enter patient's birth certificate number
- Patient name allows user to enter patient's name
- Source the original location of the data from which the request came for processing
- Target the location where the data is to be moved
- Destination viewing data lists in individual areas

Ŧ	Destination Nepřijatá data	•
	Nepřijatá data	
	Koš	

Figure 193: img

• Status - displaying of a list of active or inactive data

Ŧ	All statuses	•
	All statuses	
	Active	
	Blocked	
	Running	

Figure 194: img

To update the list, you can use the "Reload" button **c**.

The found list of temporary data is sorted into individual columns which can be sorted by clicking on the name of the selected column. The symbol is then displayed next to this column $\blacksquare / \blacksquare$.

Patient name Identification Date of Modality Source Target Information Actions
examination

Figure 195: img

- Patient name a column containing patients' names
- · Identification a column containing patients' birth certificate numbers
- Date of examination a column containing the date when examination took place
- Modality a column containing modality which was used for examination
- Source the original location of the data from which the request came for processing

- Target the location where the data is to be moved
- Information a column containing information why the study was put in "Temporary data" tab
- Actions a column containing functions for working with temporary data
 - Edit 🗹 data editing
 - Export to DICOMDIR
 saves selected date to local folder on the workstation (PC) in DICOMDIR format
 - Delete 🖲 deletes data from temporary storage

8.3 Editing

Individual data in the list of "Temporary data" tab can be edited using the "Edit" function \checkmark located in the "Actions" column. Select this function to open data editing table.

First name sync1	Last name	Middle name
Prefix	Suffix	
Identification* 1346546546	Date of birth 2022-08-08	Gender Female
Accession number 11013917		
Last known error org.medoro.dpgw.dicom.CStoreException: org.medoro.dpgw.	dicom.lib.DataValidationException: Empty patient Last Name!	
Incoming data from DWTEST8 (DWTEST8@127.0.0.1:5380) DW cache AE	Modality MR	Study description MR L patere
Date of examination 2022-02-11	Study UID 1.3.6.1.4.1.20744.3.1.2.2.12.1328101243577.37892002	Creation date and time 2022-06-24 11:51
Save change as rule		
BACK SAVE		SAVE & SEND

Figure 196: img

The data editing table contains:

- First name option to edit patient's name
- Last name option to edit patient's surname
- Middle name option to edit patient's middle name
- Prefix option to edit academic title before patient's name
- Suffix option to edit academic title after patient's name
- · Identification option to modify patient's birth certificate number
- Date of birth option to edit patient's date of birth by text field or selection
- Gender option to edit the patient's gender (male, female, other)

User guide

- archiveIncoming data from AETitle name of the specific modality from which the data was sent
- Modality types of modality which the data was sent from
- Study description name of the examination
- Date of examination date of performed examination
- Study UID unique study number with the possibility to create new study using the "Generate new UID studies" function <a>o. When you select this function, you will be notified by a pop-up window:

	Attention!	>
	ou sure you want to generate ne ise can break data consistency!	w study
NO		YES

Figure 197: img

- Creation date and time date and time of adding of the study in the tab "Temporary data"
- Save change as rule remembers currently performed correction. If other studies of the same
 patient are stored in the temporary data, the correction will be performed automatically. (Applies
 only to patient demographic data. For example, it is not possible to automate generating of a
 new UID study.)

After finishing editing of selected parameters, made changes can be undone clicking the "Back" button **BACK**, saved and returned to later by clicking the "Save" button **SAVE** or these changes can be saved and sent to PACS archive by clicking "Save&send" **SAVE** & SEND.

8.4 Deleting

If you no longer want to keep the selected data in the temporary data store, you can delete it using the "Remove" action I. If you use this function, you will be notified by a "Question" pop-up window where you need to confirm the deleting:



Figure 198: img

In case the study was successfully deleted from temporary data, the system notifies you at the bottom of the screen:

Record "1.3.6.1.4.1.20744.3.1.2.2.12.1328101243577.371435974 - ^sync1 (1346546546)" has been successfully deleted

Figure 199: img

9 Viewer {#viewintro_DICOM Viewer}

This chapter describes the second main part of DPGW which is a dialog box of DICOM viewer with tools for working with visual data. This dialog box can be opened by opening the patient/study or by action "Back to the exam" , which is located in the upper right hand corner of the basic dialog box for working with registries.

If you need to return to the basic dialog for working with registers from the DICOM viewer dialog, use the action located in the upper left corner of the DICOM viewer.

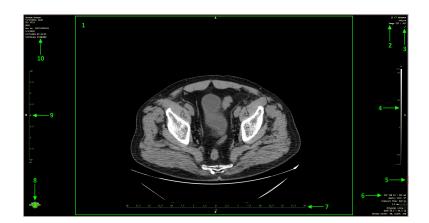
The DICOM viewer dialog with image data tools contains several main areas which will be described below in this user manual. These are: - Image data display window - desktop with open series display - Viewer working set - this set is located in the left column of the DICOM viewer dialog. It contains previews of individual series of examinations and work with image documentation, or tracing of studies of the selected patient - Viewer tools - the tools are displayed in the top bar of the DICOM viewer dialog and they include currently selected tools on individual mouse buttons, fixed tools, a user-configurable toolbar and tabs with all available tools: - Display tools - Measurement tools - Action tools for series - Volume operations, 3D displaying - Online consultation - Viewer settings - Working toolset



Figure 200: img

In case you are working on a diagnostic station equipped with multiple monitors, it is possible to set the DICOM viewer desktop - PRODUCT-SHORT-NAME- to individual monitors. DPGW allows you to

automatically distribute desktops to individual monitors. For this function it is necessary to install the "DICOM viewer extension" add-on in the used internet browser. You can find more about the settings of the multi-monitor station in the chapter "Settings", section "Display settings".



9.1 Image data display window {#viewintro_Image data display window}

Figure 201: img

The main workspace of the DICOM viewer dialog is the actual display of the selected patient series/image. The individual parts of this area are:

- 1. Image this section displays all images, videos, and documents.
- 2. OSD Labels (On Screen Display) information about the currently displayed image/series. In "Viewer settings" information to be displayed can be defined. This setting is applicable to the user/station.
- 3. Key images marking actions for tagging images in the series ☑. The selected image has a double check mark in the button ☑(the opposite indicates unmarked). For description of the marking and its overview, see the section "Action tools for series".
- 4. Brightness display scale the scale shows the range of the selected brightness window. The brightness window settings are affected by two parameters: "center" and "width". Current values of these parameters can be monitored in the lower right corner of the desktop while adjusting the brightness window. (OSD labels, item 6.)
- 5. Scroll bar this bar contains a scroll bar that shows the current scroll position within the series. The behavior of the toolbar is affected by many parameters. These affect several key things:
- initial position of cursor it is possible to parametrically set whether the cursor or the whole

series, open on the first frame or in the middle so that the user can scroll smoothly forward or backward.

- surroundings loading you can parametrically set the number of frames which are automatically loaded around the cursor position. DPGW automatically loads the cursor area because it expects the user to view images while moving around the position of the cursor. If the user jumps to another position within the series, the loading of images from the vicinity of the previous position is automatically interrupted and the loading of the vicinity of the new cursor position begins. This ensures smooth viewing of the series even if it is not fully loaded. However, all images are loaded gradually in the background.
- range of automatically loaded images the user can view images without having to load the
 entire series. If the series has a lot of frames, the cursor area is loaded and then the frames
 are always loaded automatically, omitting the defined number of frames. For example, if all
 10 frames are loaded and the user then starts moving around the series, the preloaded images
 make it easier for him to move around and reduce the time the user would have to wait for all
 the images to load.

Warning: Thus, the behavior of images loading in a series is fully changeable and it depends on the type of modalities and the number of images that the user usually works with.

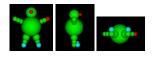
The control of the shift by one image within the series is enabled by using the mouse wheel or the PgDown/PgUp keys. If you require faster scrolling, you can use the scroll bar, or the Shift + PgDown/PgUp keyboard shortcut to scroll 10 frames in a series.

- 6. OSD Labels (On Screen Display) Eposcopic parameters of the modality on which the study was taken. In "Browser settings" you can define what information is to be displayed. This setting is applicable to the user / station.
- 7. Vertical scale indicates the size of the image and its parts horizontally and vertically. If the image is originally DICOM format, it carries information about the actual distance and it is possible to use scales or measuring tools to measure the distance. However, in the case of digitized images, they usually do not carry this information and it is first necessary to calibrate the image = determine the known distance. Based on this data, all measured distances are then recalculated.

Warning: if the image does not show a calibration object and the user does not perform an accurate calibration of the image measurement according to this object, this is only an indicative measurement.

- Pictogram shows the orientation of the image with respect to the position of the patient's body. The pictogram can be displayed in two modes. Schematic figure or cube with lettering on the sides. The displaying can be changed in the "Label layout" settings.
- Schematic figure

Dicompass Gateway



• The cube is marked with letters according to orientation.



The letters can take the following values: - A (anterior) - P (posterior) - R (right) - L (left) - H (head) - F (foot)

The largest letter in the middle indicates the current direction. The lower case letters on each edge indicate the direction the view would be if the patient's cube (orientation) rotated in that direction.

- 9. Horizontal scale indicates the size of the image and its parts horizontally and vertically. If the image is originally DICOM format, it carries information about the actual distance and it is possible to use scales or measuring tools to measure the distance. However, in the case of digitized images, they usually do not carry this information and it is first necessary to calibrate the image = determine the known distance. Based on this data, all measured distances are then recalculated.
- 10. OSD labels (On Screen Display) information about the patient's and study demographic data (name, surname, date and time of creating of study, etc.) In "Browser settings", it is possible to define what information is to be displayed. This setting is applicable to the user/station.

9.1.1 Context menu {#viewintro_Context menu}

In this area, in addition to working on selected data, it is also possible to open the context menu by pressing the right mouse button:

0	Default layout	
~	CT Abdomen 1/1/2023	
r)	1-scout	
r	2-natyve	
þ	3 - Recon 2: natyve	
ß	4-+C	
G	200 - Smart Prep Series	
×	Remove view	

Figure 202: img

The floating panel contains: - Viewer tools - these tools can be set according to user's preference using the settings of individual areas of the viewer tools. For more information on configuring tools in the floating panel, see the section "Configuring tools display" - Series view - allows the user to open

selected series of the open study. These series are listed under the date of the examination - Remove view - closes the open series

The context menu can also be invoked using a keyboard shortcut. This keyboard shortcut must be newly assigned using the "Keyboard Shortcuts" tool in the "Browser Settings" tool tab.

9.2 Viewer tabs {#viewintro_Viewer tabs}

If you display the desired studies using the "Add to new tab" or "Show patients in tabs" action a from the "Archives" dialog, a DICOM viewer dialog opens with individual tabs in its footer:

📄 📰 Anonym Anonym Anonym Anonym Anonym Anonym 👁 🗍 🇊

Figure 203: img

- Use the tool 🖾 to display the currently open image/series of images in full screen mode, hiding any sidebars. To return to the original panel layout, click on the tool again.
- The "List tabs" tool 🔳 allows you to open a menu of open tabs and then display or remove them:

		Anonym Anony	m 🔟					
		Anonym Anony	m 🔟					
		Anonym Anony	m 🔟					
(\bullet)	-	Anonym Anonym	Anonym	n Anonym	Anonym Anonym	۲	٥	î۶

Figure 204: img

• To remove a specific bookmark, move the mouse cursor over the targeted tab and select the trash can icon



Figure 206: img

• Each tab contains an action for editing the name of the tab. This action is called by right-clicking on the selected tab and selecting the "Rename tab" tool



Figure 207: img

to start editing the tab name:



The "CANCEL" action cancels the action being performed. The "SAVE" action renames the tab name according to the text in the text field.

- "Show active tab" action by selecting this action, only the active "List tab" will be displayed. Selecting this action again will show all open tabs of the viewer.
- The "Clone tab" action 🗖 copies the currently open tab and creates a new tab with its contents. When this tool is selected, a new tab will be created with the possibility of editing its name similarly to the "Rename tab" tool.
- The "Remove all tabs" action 🖬 closes all open tabs. Before closing all tabs, you must agree to close them (if you check the option "Do not ask me again" you will not be asked to confirm next time and all tabs will be closed):

Question	
Remove all tabs?	
Do not ask me a	again
NO X	YES 🗸

Figure 208: img

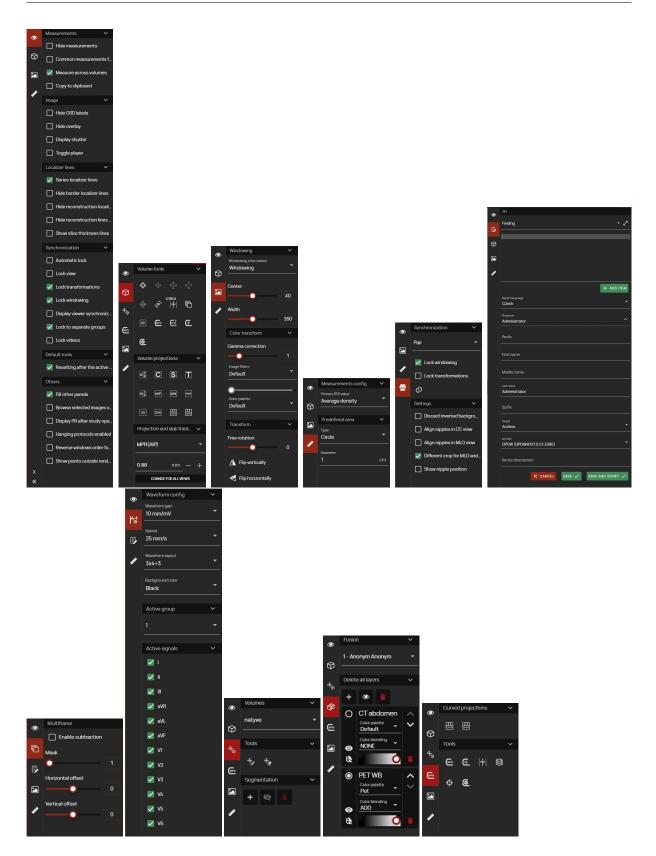
9.3 Working toolset {#viewintro_Working toolset}

Select the "Working tool set" tab , located in the upper right corner of the viewer, to display the sidebar located on the right side of the viewer dialog. This sidebar contains individual tabs corresponding to the modality of the open study. The individual tabs include:

- Display toolset <a>The working display toolset contains items for displaying information over an image, or tools for synchronizing a series of images/videos
- Volume Toolset 💁 The working volume toolset contains items for editing the created volume and changing the side orientation

- Postprocessing toolset <a>The Postprocessing toolset contains items for adjusting the brightness window, colors and filters, or rotating
- Measurement toolset
 The measurement toolset contains items for editing the ROI and its shape
- Mammography toolset 🔤 The mammography toolset contains items for editing mammography images and synchronizing them
- Report toolset 🗹 The report toolset is called up by the "Create SR" tool and is used to create a description of the examination by the user
- DSA toolset 🖸 The DSA toolset contains items for working with the digital subtraction angiography image and allows you to create and move the mask
- ECG toolset 📧 The ECG toolset contains items for setting ECG curves and displaying individual signals
- Segmentation toolset Segmentation toolset contains tools for cutting and editing the image
- Fusion toolset 🚳 The fusion toolset contains tools for displaying and editing fused image series
- Curved Projection Toolset CMPR and SMPR projection views and their associated tools

The tools of the "Working toolset" tabs are described below, according to the individual tab chapters with all available tools. The "Working toolset" contains the following tools:



9.3.1 Display toolset {#viewintro_Display toolset}

Display toolset <a>

 The working display toolset contains items for displaying information over an image, or tools for synchronizing a series of images/videos:



Figure 209: img

The Display tools working set contains the following tools:

Function	Description
Measurements	
Hide measurements	Hides all measurements from the image. Can be used when measured objects interfere with the image documentation, they can be temporarily hidden without removing them.

Function	Description
Common measurements for images in the series	It is used to keep all measured objects in the displayed window for all images in the series. When moving to the next image in the series, the measurement remains unchanged and is displayed on the next images.
Measuring across volumes	It is used to keep all measured objects in the display window for all images in the created volume. When moving to the next image in the reconstructed plane the measurement remains unchanged and is displayed on the next images.
Copy to clipboard	This tool is used to copy the value of the last measurement to the clipboard, then paste it into the text field using the shortcut ctrl+v.
Image	
Hide OSD labels	Function to hide information in the image data display window.
Hide overlay	The tool provides the option to hide/show the saved measurements in the image.
Toggle player	Show or hide the player for automatic play through the series.
Localizer lines	
Series localizer lines	Display of localization lines, between two perpendicular series.
Hide border localizer lines	Hide border localizer lines (border lines shown in green)
Hide reconstruction localizer lines	Hiding localization lines, between reconstructed perpendicular series, in volume operations.
Hide reconstructions localizer lines on the active panel	Hiding localization lines in the active window panel for displaying image data between reconstructed perpendicular series, in volume operations.
Show slice thickness lines	Display of localization lines including display of the selected slice thickness.
Synchronization	
Automatic lock	Synchronous scrolling through series in multiple panels with automatic finding of the corresponding image position in the series
Manual lock	Synchronous scrolling through series in multiple panels from the current image position in the series

Function	Description
Lock transformations	When using the sync function, the image adjustment will be applied to all synchronous series simultaneously
Lock windowing	When using the sync function, brightness and contrast adjustment will be applied to all synchronous series simultaneously
Display viewer synchronizability	Showing the option to lock individual panels in the "Display panel data window"
Lock to separate groups	Grouping of individual panels "Windows for displaying image data" for possible manual or automatic synchronization of series according to the data acquisition plane of the series or by manual entry
Lock videos	Synchronous scrolling through synchronously recorded videos
Default tools	
Resetting after the active viewer change	When switching to another panel in the "Image Data Display Window", the currently assigned tools on the individual mouse buttons will be replaced by the default tools or the tools selected according to the "Tool selection" function
Others	
Fill other panels	Automatic display of series, when changing the number of panels to display in the "Image Data Display Window"
Browse selected images only	Browse only selected keyframes in a series with standard tools (mouse wheel keyboard shortcuts)
Display PR after study opened	Enable/disable automatic display of stored measurements in the image after opening a study
Hanging protocols enabled	Disables/enables the option to display hanging protocols
Reverse windows order for HP	Reversing the order of applied hanging protocols on individual monitors

9.4 Tool search {#viewintro_Tool search}



Figure 210: img

The "Tool Search" function is designed to quickly find the required DICOM viewer tool DPGW.

To find a tool quickly, follow these steps: - display the wanted image/series in the image data display window - press the keyboard shortcut "ctrl + K" - Type part of the name of the tool you are searching for:

Q dista		
▶ ⊷	Distance measurement	
In X:Y	Distance ratio	

Figure 211: img

• hover over the selected tool and click the selected mouse button to assign the tool to that button, if the tool is already assigned to a mouse button, that button will be highlighted:



Figure 212: img

9.5 DICOM viewer status bar and its functions {#viewintro_DICOM viewer status bar and its functions}

🗐 Default 🔒 🤠 📟 Default 👙 🕫

Figure 213: img

The status bar of this DICOM viewer contains the following tools: - Default Automatic layout of OSD labels and the ability to lock them - Allocated memory indicator - Default Placement of keyboard shortcuts and their multilevel assignment - Bug reporting - Global configuration

These tools are described in the following paragraphs.

🗐 Default 🤷

Figure 214: img

The "Automatic layout of OSD labels" function is used for automatic switching between different profiles of OSD labels (On Screen Display), if they are configured and assigned to individual modality types (DX, CR, CT, MR...). These profiles can be configured using the "Profile editor" function.

When the examination is displayed in the "Image data display window", the OSD profile will be automatically selected according to the displayed examination and the modality used to perform the examination when the function is enabled.

It is also possible to switch between the OSD profiles manually by clicking the mouse on this function, which will bring up the menu for selecting the OSD profile e.g.:



Figure 215: img

Click on the desired OSD profile to apply it.

In case you want to apply only the selected profile to all examination modalities without auto-selection, it can be locked with the \square , icon, after locking the icon will be undercoloured \square .

9.5.2 Allocated memory indicator {#viewintro_Allocated memory indicator}

Figure 216: img

The "Allocated memory indicator" function is used to display the used memory of the viewer and to adjust its size.

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Each open study in this DICOM viewer needs a certain amount of RAM memory of the PC station you are working on and this function serves both an informative purpose for displaying the used memory and the possibility of setting and allocating additional memory for working with image documentation.

When the allocated memory is exhausted, the viewer may "freeze" and must be restarted.

The default value of allocated memory is 5GB, in case you are working with larger studies or more open studies, we recommend allocating more memory to work properly with this product.

When you hover the mouse over the icon of this function, an informative table will be displayed about the allocated memory of the DICOM viewer "Allocated memory" and about the total memory usage of the PC station you are working on "System memory" e.g.:

Allocated mem	I	0.07 / 5 GB		
System memo	10.67	/ 15.39 GB		
📕 Default 🔒	٥		Default i	

Figure 217: img

By clicking on this function, you can set the size of the "Allocated memory" (the current size of the viewer's allocated memory is displayed below the text field line):



Figure 218: img

Warning: For proper functionality of this tool it is necessary to have the "DICOM viewer extension" web browser extension installed. If this extension is not installed, the "System Memory" will not be displayed.

9.5.3 Placement of keyboard shortcuts and their multilevel assignment {#viewintro_Placement of keyboard shortcuts and their multilevel assignment}

The "Keyboard shortcut layout" tool received, located in the bottom right corner of the browser, is used to visually display the assigned shortcuts on the virtual keyboard.

To select the level of the keyboard shortcut layout, use the "Keyboard shortcut layout" tool reduct, to call up the levels menu:



Figure 219: img

Multi-level switching of the keyboard shortcut layout can also be done with keyboard shortcuts that you can select using the "Keyboard Shortcuts" tool (see more in "Viewer Tools"). You can return to the original "Default" level layout with the "ESC" key.

To view the virtual keyboard with assigned shortcuts, select the icon to the right of the level description in the menu. To view the assigned shortcuts, hover over the target key to view its assigned shortcuts. Keys highlighted in grey do not have an assigned shortcut:

Hotkey								×
DEFAULT		WINDOWIN	IG		ANN	OTATION		
ESC F1 F	2 F3 F4 F	5 F6 F7	F8	F9 F1	0 F11 F12			
							номе	PG UP
TAB Q W	ERT	u I	0 F			DELETE	END	PG DN
	s shift w Show/Hide wo							
SHIFT Z	alt W Lock view and w						UP	
		SPACE				LEFT	DOWN	RIGHT

Figure 220: img

9.5.4 Bug reporting {#viewintro_Bug reporting}

In case the system behaves non-standardly, it is possible to use the "Bug Report" tool **1** to capture the current system status in a log file with the necessary information for the manufacturer's support department. This tool is located in the bottom right corner of the screen.

Selecting this tool will call up the "Question" table:



Figure 221: img

Selecting "NO" cancels the job, selecting "YES" creates an error message and displays information about the report:



Figure 222: img

9.6 Viewer working set {#viewws_Viewer working set}

Viewer working set is located, by default, in the left column of the DICOM viewer dialog, it contains previews of individual series of examinations and work with image documentation, or finding studies of the selected patient.



Figure 223: img

Viewer working set is sorted according to the sequence of categories ("Patient" and "Study"):

1. Patient - Displays the patient's name, surname and ID number (the functions of the action icons are described in the aricle "Workset display options").



Figure 224: img

2. Study - Displays the date and time of the examination. The anchor icon **B** indicates "Reference Study Flag", i.e. the study determining the primary examination to which the timeline and other operations will relate.



Figure 225: img

By selecting the empty field below the anchor icon, you can select the action that will be displayed in this field. Right-click in the empty field to call up the context menu:



Figure 226: img

- Display findings this action allows you to view the text description of the examination that is part of the study stored in the structured examination description format "SR", or in case of integration is pulled from the external system in which the examination description is stored
- Watch study for changes Display alerts when the number of frames in the displayed study has changed, and reload a series of frames to update the number of frames. If you have an incomplete series displayed in the "Image Data Display Window" and additional frames are added to this series, you will be alerted by a flashing "Watch study for changes" tool icon, clicking on this tool icon will reload the series with the current frame count. After the series of images have been reloaded, the measurements taken and the image postprocessing will be removed. The blinking "Track study changes" tool icon is located at the bottom of the " Viewer working set":



Figure 227: img

- Forward data action for sending the image documentation to another DICOM device, see section "Forward selected"
- Export action to export data to local storage or print it
- Expand/collapse action to expand/collapse all series (thumbnails in the viewer working set) of the selected study
- 3. Series Displays previews of the series in the selected patient study



Figure 228: img

Warning: If the alert icon **I** is displayed in the viewer's working set, patient studies with the same birth number but different name, gender, or date of birth are opened in the viewer. Take extra caution.

The selected series can be displayed by double-clicking on the preview of the series in the viewer working set, or by drag&drop style into the "Image Data Display Window". A preview of an open active series is highlighted with a red border and marked with an eye icon:



Figure 229: img

If the selected study contains more series than can be displayed in the monitor resolution, the user can scroll through the viewer working set using the mouse wheel.

The viewer working set can be configured in "Display settings" to the left or right side of the monitor. If you are working on a multi-monitor workstation, the browser working set can be displayed on each monitor.

9.6.1 Viewer working set display options {#viewws_Viewer working set display options}

The viewer working set can be used with the "Hide/show working set" action \checkmark or pull it to the edge, or expand it to a larger size, using the "Normal/Wide view working set" > < event to display multiple series:



Figure 230: img

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Figure 231: img

9.6.1.1 Viewer working set configuration {#viewws_Viewer working set configuration} Selecting the "Work Set Configuration" action calls up the "Working Set Configuration" table, which allows the user to display the viewer working set, to select the side of the viewer working set display, and to switch between the default and optimal working set display:



Figure 232: img

Warning: turning off viewer working set display can be reverted in the viewer configuration tools tab by turning it on in the "Working set configuration" tool.

"Work Set Configuration" allows you to align the work set in a "Flexible" or "Optimal" way and to display a map of the panels for dragging and dropping. These features are listed below:

9.6.1.1.1 Flexible viewer working set display {#viewws_Flexible viewer working set display} Flexible view shows enlarged previews of series. If the number of series exceeds the views in the displayed working set, the user is allowed to move using the mouse wheel:

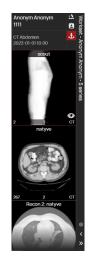


Figure 233: img

The default view follows the sequence of view levels: patient -> study -> series -> subseries -> instances -> volumes -> fusion -> segmentation.

The display of thumbnails in the flexible workset view can be changed using the "Workset Configuration" tool described in the previous paragraph.

9.6.1.1.2 Optimal viewer working set display {#viewws_Optimal viewer working set display} By selecting the "Optimal working set display" action, all series of all studies will be displayed and their size will be adjusted to the monitor resolution:



Figure 234: img

This feature of the viewer serves to better orient the user within the view of one or more studies of a single patient.

Move the mouse over the thumbnail of the series to view information about the series number, number of frames, modality, and series description:



Figure 235: img

9.6.1.1.3 Map of panels for dragging {#viewws_Map of panels for dragging} The "Drag and Drop Panel Map" tool, when turned on, displays a panel map for dragging the image to a specific panel in case you have multiple monitors set up for display.

If this function is not enabled, drag&drop the series to the "Image Data Display Window" of the selected monitor to display the series of images:

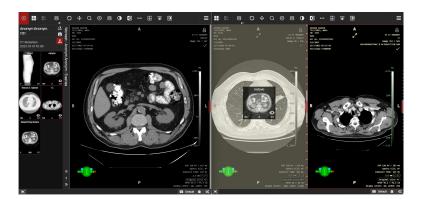


Figure 236: img

If this function is enabled, or if you hold the "CRTL" key while dragging, drag & drop the series to a specific panel using the "Drag and Drop Panel Map":

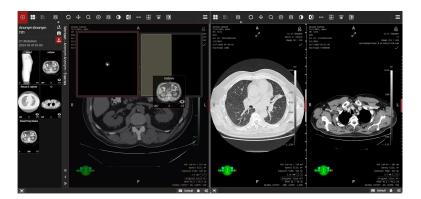


Figure 237: img

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Figure 238: img

9.6.1.2 Function All patient studies {#viewws_Function All patient studies} By selecting the "All patient studies" function, you will call up a table at the bottom of the screen displaying all the performed examinations of the selected patient:

Description						
	Accession number	ality Date of examination	Date of birth M	Study ID	Identification	name +
CT Abdomen	111112223333	2023-01-01 10:00	2023-01-01 C1	13	1111	🖹 🗸 ANONYM Anonym (5)
PET Abdomen	1112223344555	PT 2023-01-01 10:00	2023-01-01 C1	14	1111	🚊 🗖 ANONYM Anonym (5)
PET Brain	112345	PT 2022-01-01 10:00	2023-01-01 C1	15	1111	🖹 🗖 ANONYM Anonym (5)
PET Brain	112345	PT 2022-01-01 10:00	2023-01-01 C1	15	1111	ANONYM Anonym (5)
	112345	PT 2022-01-01 10:00	2023-01-01 C1	15	1111	ANONYM Anonym (5)

Figure 239: img

The "All Patient Studies" functions is listed in this table:

lcon	Function	Description
G	Replace and display	display of selected studies with replacement of the last displayed ones
œ	Add to new tab	display of selected studies in a new DICOM viewer tab
	Add and view	display of selected studies with addition to the last ones displayed (double-clicking on the selected study line also works)
Ð	Back to examination	Closes the "All Patient Studies" table

The "All patient studies" table contains all studies of the selected patient stored in the PACS archive and is divided into columns with individual values. You can search and sort in these columns, see more in the "Search results" chapter.

After hovering the mouse over a specific line of the patient study, a floating panel containing the following actions can be used:

lcon	Function	Description
Ø	View in DICOM viewer	display of selected studies with replacement of currently displayed
F	Add to new tab	display of selected studies in a new DICOM viewer tab
ß	Add and view	display of selected studies with addition to the last ones displayed

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Figure 240: img

9.6.1.3 Timeline A timeline is used for a simple overview of the patient's examination history. After the display, the history of all examinations is clearly shown. Opening the timeline together with the SHIFT key opens the timeline on all set monitors of the multi-monitor diagnostic station. You can open the required historical examination by double-clicking or dragging.



Figure 241: img

In the left part of the timeline, there is information about the name and ID number of the selected patient.

In the middle part is the timeline itself, containing information about the saved studies/examinations of the selected patient. This timeline is divided into individual years showing the type of study/modality on which the study was performed, with the date of the study. The anchor icon II indicates the "Reference Study Flag", the study identifying the primary examination. The eye icon II indicates the study displayed in the "Image Data Display Window". In case the patient has too many studies to display all of them in the monitor resolution timeline, the user is allowed to scroll through the timeline using the mouse wheel. More detailed information about the study in the timeline, can be obtained by hovering the mouse over the desired study:



Figure 242: img

Warning: if there are studies opened in the viewer for a patient with the same ID number but different name, gender, or date of birth, you will be alerted to this in the study description located in the timeline. So take extra care:

Workset - An	Anonym Anonym 1111	2020	11/24 - CT	2022	1/1 PET Brain CT, PT	2023
Date						11/24/2020
Data locat	tion					DPGW
Number o	f series					8
Number o	finstances					512
Please	check patient name	e, Ple	ase check patient :	sex,	Please check patie	nt birth date

Figure 243: img

In the right part of the timeline there are actions "Pin" I for inserting the timeline to the action bar, "Configure filter" , "Filter" and "Hide Timeline" used to close the timeline.

The "Filter" action brings up a table with options for displaying results in a timeline.

Time	line						×
Custom modality							
СТ	MR	PT	US	SC	CR	DX	MG
от							
Кеуж	ords						
Max.s	study a	ge					
F	Sort	ed by a	iscend	ing			
~	Disabl	e filter	s on pa	atient o	:hange		
	Lowp	rofile					
			CAN	icel		SAV	E

Figure 244: img

The timeline filtering table includes the following options:

- 1. Custom modality/modalities text field to indicate the modality shortcut
- 2. Modalities this section contains checkboxes to specify the modalities to be displayed in the timeline, for easy work there is the option of "Mark all", "Unmark all" and "Invert" actions to flip marked/unmarked modalities
- 3. Keywords text field for entering, for example, the name of the study, etc.
- 4. Max. examination age text field for specifying the display of studies in a timeline bordering the age period in months
- 5. Sorting direction by clicking the sorting direction arrow, you determine the order in which the study will be displayed in the timeline according to the examination date
- 6. Turn off filters when changing patient option to turn off/keep the timeline filter when another patient is displayed
- 7. Low profile option to display the timeline in minimized/compact mode:



Figure 245: img

8. Cancel, Save - the "Cancel" action cancels the performed filtering, the "Save" action saves and applies the applied filter to the timeline

9.6.2 Workset Floating Panel {#viewws_Workset Floating Panel}

The viewer's working set allows you to perform actions, above the open patient examination, contained in a floating panel that can be called up by pressing the right mouse button in the patient/study/series area. The actions of the floating panel may be differ depending on the call from a specific area and includes the following functions:

lcon	Function	Description
^	Expand/collapse	Action to expand/collapse all series (thumbnails in the viewer working set) of the selected study
۲	View	Displays the images in the "Image data display window"
89	View connected	Displays the images in all open panes in the "Image data display window" and synchronizes the passage through the series
₽	Create report	Action to call up a table to create a "Structured Report", i.e. a description of the examination
Ŷ	Record voice note	Allows you to record an audio track, e.g. a spoken note for the selected study
Ê	All studies of patient	Actions to call up the table "All patient studies" see article above
0	Reverse item order	Actions for changing the order of images, e.g. for CT axial section, changing the direction of scanning a series of images from craniocaudal to caudocranial.
\mathfrak{O}	Create volume	Actions to create multiplanar planes and 3D volume
Ф	Create fusion	Action to merge two series with overlay
Ę	Forward data	Actions for sending image documentation to another DICOM device, see article "Forward selected"
B	Export	Action to export data to local storage, or print

lcon	Function	Description
<	Share the study	Action for sharing a study to an external user
	Add to worksheet	Selecting the action adds the study to the worklist, see paragraph "Add to worklist", if this action is selected at the "Patient" level, all open studies in the " Viewer work set" of the selected patient will be added to the worklist
₹ <u>₽</u>	Split the series	Devide series into subseries, for example, in dual scans of MR examination
	Edit tags	Allows you to assign a label in the selected series, see article "Labels"
ũ	Watch study for changes	Display a notification of the changed number of images in the displayed study and the ability to reload a series of images with the updated number of images
×	Remove	Removes a patient/study/series from the DICOM viewer workset

5

Figure 246: img

9.6.2.1 Create report {#viewws_Create report} Selecting the "Create report" action from the floating toolbar will call up a side toolbar with the Findings toolset and the option to create a description of the examination.

۲	SR
0	Finding
₽	
Ø	
Ψ	
1	
	+ ADD ITEM
	Report language
	Czech
	Physician Administrator
	Prefix
	First name
	Middle name
	Last name Administrator
	Suffix
	Target Tarchive
	Archive DPGW (DPGW@127.0.0.1:5380)
	Series description
	X CANCEL SAVE V SAVE AND VERIFY V

Figure 247: img

The finding toolset contains the following text boxes and drop-down menus:

• Information Disclosure Area - when the Finding toolset is opened, the name of the area "Finding" is selected, but the user has the option of selecting the area from the drop-down menu that it contains:

Finding	•
Addendum	
Conclusion	
Finding	
History	
Impression	
Patient Presentation	
Previous Finding	
Procedure Description	
Recommendation	
Request	
Summary	

Figure 248: img

It is also possible to add additional message areas with the "Add" action ***** Add additional message areas with the close action *****:

۲	SR
_	Finding – $\mathbf{k}^{\mathbf{X}} \times$
₽	Description of the find.
\heartsuit	
*	
	Summary
	Finding described.
	+ ADD ITEM

Figure 249: img

When describing the examination, the user can enter keywords using predefined shortcuts and then call them up using the shortcut key "ctrl+spacebar":

۲	SR
Ð	Finding - 2 ^N
Ø	bb B buffer base
	+ ADD ITEM

Figure 250: img

Adding and editing keyboard shortcuts can be done in the "Settings" dialog, using the "SR Macro Editor" tool.

Warning: access to the "SR Macro Editor" is controlled by user roles. If you do not have access to this editor, please contact your system administrator DPGW.

The text field of the examination description can be enlarged to full screen with the 2 tool, for shrinking to the original layout select the 2 tool.

Full-screen mode of SR finding tab:

Dicompass Gateway



Figure 251: img

• Language of the report - determination of the language in which the description of the examination is written, Czech is selected by default for Czech users, but the following languages can be selected from the drop-down menu:

Report language	4
Czech	
English	
Czech	
Slovak	
Russian	
German	
French	
Spanish	
Portuguese	
Polish	
Italian	
Greek	

Figure 252: img

- Physician by clicking on the "Physician" area, you can specify the doctor who creates the description of the examination, by default the doctor is determined according to the logged-in user, but when changing these values can be filled in the text fields
- Target In this section you can specify whether the created description should be saved to the PACS archive with its selection, or to the user or station.

To save the concept of the created examination description, use the "Save" action, this saves the SR examination description, which can still be edited.

Patient	2 ⁷¹
Anonym Anonym *1/1/2023 , male ID: 1111	
Study	Co
Accession number : 11 Date of examination : Report date : 4/17/20 Study description : A	1/1/2023 10:00:00
EDIT CONCEPT	1
	tem and Descendants: English
Observer Type: Person Person Observer Name:	
Findings :	pac
Finding: Description	C (1) 1

Figure 253: img

The saved SR concept will be displayed in the viewer working set, as another patient study.



Figure 254: img

Edit the concept by displaying SR in the image data display window and then selecting the "EDIT CONCEPT" action to open the sidebar with the finding toolset.

Select the "Save and Close" action to complete the SR examination description. This action will save the SR finding according to the target destination specified for saving and you will no longer be able to edit the SR:



Figure 255: img

If you do not want to save the created examination description, select the "Cancel" action and the "Finding toolset" will be cancelled.

Ŷ

Figure 256: img

9.6.2.2 Record voice note {#viewws_Record voice note} The "Record voice note" function is used to record and archive an audio track, e.g. a spoken note for a selected study and its speech-to-text conversion options.

Opening the "Record voice note" function will bring up the "Voice Note" toolset sidebar:



Figure 257: img

- This sidebar contains the voice note tools and its options for recording speech to text:
 - Start recording a tool to start/stop recording a voice note, after starting the recording the time of recording will be displayed



Figure 258: img

- OReset recording select this tool to overwrite the current voice note
- D Stop recording and save a tool to save a voice note that you have created
- Cancel recording tool to delete a created voice note
- Settings:
 - Voice note language A drop-down bar to select the language in which the voice note will be recorded
 - Allow speech-to-text conversion selecting this tool will bring up the "SR" toolbox sidebar for writing findings after the voice note has been completed and saved, where the text of the recorded voice note will be inserted
 - Push to talk selecting this tool allows you to record a voice note by holding down the "spacebar" key, if you do not select this tool you can also start/stop the voice note by pressing the "spacebar" key
 - Auto-Recording on start selecting this tool will start automatic recording of the voice note immediately after selecting the "Record voice note" function from the viewer working set on the next start

To record an Voice note, proceed as follows:

• select the "Record voice note" function to bring up the "Voice Note" toolset sidebar

- select whether the voice note is to be recorded as text by using the "Allow speech-to-text conversion" tool in the "Settings" section of the toolset sidebar
- to record a voice note, select the """ "Start recording" or press the "spacebar" key (if "Push to talk" is selected, hold the spacebar key while recording)
- to end the voice note, select the **a** "Stop Recording" tool, or press the "spacebar" key (if "Push to talk" is selected, release the spacebar key to end the recording)
- when you have finished recording your voice note, select one of the following actions:
 - • Reset recording select this tool to overwrite the current voice note
 - Stop recording and save a tool to save a voice note that you have created
 - Cancel recording tool to delete a created voice note
- if you select the action **•** "Stop recording and save" you will bring up the "Voice Note" table:

		^
Target		
Archive		-
Archive		
CLOUDDEMO@medoro (CLOUDDEMO@127.0.0.1:5380)		
Series description		
Voice note		

Figure 259: img

- select the "Target" (archive, station, user), if the archive destination is selected, choose the "Archive" and name the saved voice note in the "Series description"
- by selecting "CANCEL" you cancel the action performed, by selecting "SAVE" a new series will be created in the selected study containing the voice note of the AU modality.
- after saving, the voice note can be played directly from the "Viewer working set" by clicking on the play button:



• playback is also possible after moving the voice note from the "Viewer working set" to the "Display window for image data":

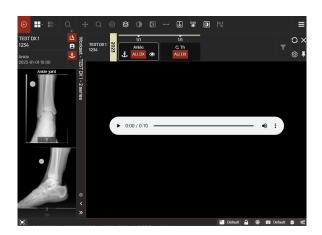


Figure 260: img

After saving the voice note, if you have selected the option "**Allow speech-to-text conversion**" you will be informed about the success of generating text from the voice note:



Figure 261: img

After confirming the success of the text generation, will be opened a "SR" sidebar toolset for typing findings, where the text of the recorded voice note will be inserted:

۲	SR					
	Finding 👻 🖍					
Þ	This is a test of the speech to text function					
	This is a test of the speech to text function					
	+ ADD ITEM					
	Report language					
	English					
	Physician					
	pac					
	Target					
	Archive					
	(CLOUDPACS@127.0.0.1:15380)					
	Series description					
	Voice note					
	X CANCEL SAVE 🗸 SAVE AND VERIFY 🗸					

Figure 262: img

Warning: Always check the generated text from a voice note and make corrections.

Figure 263: img

9.6.2.3 Create volume {#viewws_Create volume} The function "Create volume" is used for the spatial reconstruction of series data with multiple images, e.g. CT, MR with the creation of MPR planes and 3D volume. By selecting this function and if both license and hardware resources are available, volume = new series will be created. The volume loading speed depends on the size of the given series of images, you will be informed about the loading progress by the "Loading volume..." progress window:

 \bigcirc



Figure 264: img

After the volume has been successfully loaded, a new volume series will be created with a cube label in the corner of the preview, which is intended for spatial processing:



Figure 265: img

It can be opened in two ways, by double-clicking or by dragging it into the viewing window. Each of these methods will cause a different response from the DICOM viewer:

1. Double-click to open

After opening the volume series marked with a cube by double-clicking, the display selection tab will be called up:

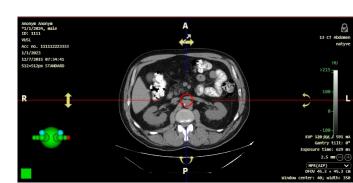


Figure 266: img

- Series 🖻 displays the selected series of images without applying volume operations
- MPR 🜃 displays the selected image series in multiplanar reconstruction with 3 planes display
- MIP 🔤 displays the selected image series in multiplanar reconstruction with 3 planes displayed in the MIP maximum intensity projection
- CMPR 🕮 displays the selected image series in a curved multiplanar reconstruction with 3 planes and curved reconstruction
- SMPR 🕮 displays the selected series of images in a straightened multiplanar reconstruction with 3 planes and a straightened reconstruction

When selecting MPR or MIP from the menu using a mouse double-click, each displayed window corresponds to a different viewpoint. Each window displays the axes, which can be moved by 1, or rotated by holding the rotated arrows 2. Changing the position of any cross of the axes automatically changes the image in the other two windows. Scrolling is also made possible by a wheel at the imaginary intersections of the localization lines: 1. If the mouse cursor is outside the area, this wheel is hidden 2.

If the mouse cursor is in the area, this circle is shown as a minimized **and and a** 3. If the mouse cursor is positioned at the intersection of the localization lines, this wheel is zoomed in and if you press the left



mouse button and drag, you will scroll in the MPR view

Figure 267: img

2. Drag and drop to open

When opened by dragging a volume series into the "Image data display window", you call up the "Display" table:

0 -



Figure 268: img

By choosing a specific action in this menu, it is possible to display only the required volume data reconstruction.

The "Display" table contains the following functions:

Series - 🖸 Series - displays the selected series of images without applying volume operations

Volumes - Acquisition plane - the volume will be displayed in the same plane as the original collection. - Top reconstruction view - the volume will be displayed in the plane that is the top view of the patient. It varies according to the original orientation of the patient (if he was lying, standing, ...) - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction view - the volume will be displayed in the plane that is the side view - Side reconstruction - Side re

Change layout - MPR 🖬 - displays the selected image series in multiplanar reconstruction with 3 planes display - MIP 📓 - displays the selected image series in multiplanar reconstruction with 3 planes displayed in the MIP maximum intensity projection - CMPR 🔛 - displays the selected image series in a curved multiplanar reconstruction with 3 planes and curved reconstruction - SMPR 🕮 - displays the selected series of images in a straightened multiplanar reconstruction with 3 planes and a straightened reconstruction

You can find more about displaying and working with volumes in the paragraph "Volume operations, 3D display".

6

Figure 269: img

9.6.2.4 Create fusion {#viewws_Create fusion} Example of a created fusion including the "Fusion toolset":

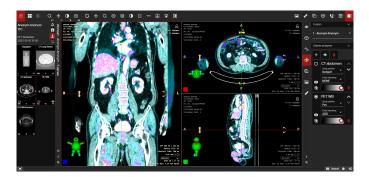


Figure 270: img

The function "Create fusion" is used to merge two series of the patient with an overlap on top of each other. This function can be used, for example, in PET examinations in nuclear medicine, when it is necessary to display accumulated radiopharmaceuticals on a CT examination. By selecting the "Create fusion" action, you will see a table for selecting series to merge:

Fusion		×
Fusion name 1 - Anonym Anonym		
Show all patier	nts	
⊠□ø		0/4
🗸 🚊 Anonyma	Anonym (0/1)	
V 🖻 PET Abd	omen (0/4)	
🔳 5-CTLung	#Bones [CT] (385)	
💼 6-CTabde	omen [CT] (385)	
💼 10 - PET W	3 [PT] (154)	
11 - PET WE	Uncorrected [PT] (154)	
		I
		I
		I
CANCEL	CREATE FUSION	CREATE AND VIEW FUSION

Figure 271: img

This table includes: - Name - name of the created fusion - Show all patients - if you have several patients open, this will show their series as well, if this box is not checked, only the series of the patient whose series you chose the "Create fusion" action will be displayed - Patient's name and the option to select

series for fusion - this area allows the selection of selected series to be merged, scrolling in this area is enabled with the mouse wheel or the scroll bar located to the right of the series area

If you have selected specific series to merge, additional actions will appear at the bottom of the table:



Figure 272: img

- Cancel cancels the performed action
- Create fusion by selecting this action, the fusion of the selected series will be created and displayed only by the illustrated volume cube in the "Viewer working set" without any change in the "Image data display window". By dragging the thumbnail into the image data display window, the fusion in the selected projection will be displayed in the called-up tab:

Displ	ay				Х
Series					
G					
Volume	es				
Ø	ହ	Ø			
MIP	MIN	FMIP	3D	DVR	
Ħ					
Fusion	1-Anc	inym Ai	nonym		
Q	Ø	Ø			

Figure 273: img

To view the fused layers in MPR mode, display the desired fused series by double-clicking the thumbnail located in the viewer working set and selecting the fusion action:



Figure 274: img

• Create and display fusion - select this action to create a fusion of the selected series with display in the "Image data display window" with a three-window display for each of the MPR planes, the "Fusion toolset" will also be displayed



Figure 275: img

The "Fusion toolset" contains:

• Fusion - this area contains the selection of the created fusion using the drop-down menu:



Figure 276: img

- Layers this area contains tools for editing the selected fusion
- Add layers 🛨 select this action to call the tab to add additional series for fusion:



Figure 277: img

- View 💁 tool for redisplaying the created fusion panel
- Delete fusion 🛄 selecting this action removes the created series fusion
- Fused Layer Tab this tab is used to work with the fused layer/series



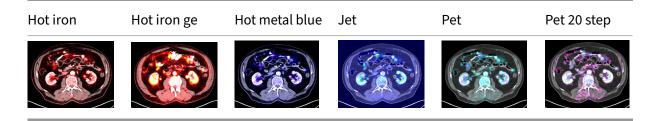
Figure 278: img

The fused layers tab contains: 1. Active layer marking - by marking the active layer, the user is allowed to work with a specific layer in the fused panel in the "Panel data display window". The active layer marking is shown by a filled circle to the right of the series name **O CTACORD**, the inactive layer marking is shown by an empty circle **O CTACORD**.

- 2. Show layer option to show and hide the selected layer/series using the "Show layer" action <a>[/ <a>
- 3. Invert option to invert the view of the selected layer/series using the "Invert" action 💁 .
- 4. Palette the color palette allows the user to change the color display of the fused layer by selecting it from the drop-down menu:



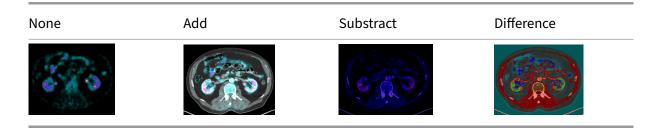
Figure 279: img



5. Color blending - choosing how the final fusion is calculated, whether the individual series are simply laid over each other, added, subtracted or projected where there is a difference:



Figure 280: img



6. Alpha - selection of the display strength of the layer/series, determining its transparency

This option can be changed by dragging the scroll wheel on the % bar or by using the mouse wheel:

Figure 281: img

7. Move down/up - option to change the layer/series order in the "Fusion Toolset" using the "Move down/up" action ∠ / △.

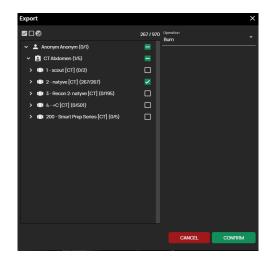
0

8. Remove Layer - option to remove the selected layer/series from the "Fusion Toolset" using the "Remove" action

B

Figure 282: img

9.6.2.5 Export The "Export" function allows the user to download selected image data to the workstation's local storage. If you have edited the image in the "Image data display window", these changes will be reflected in the export.



By selecting the "Export" action, you will call up the "Export" tab:

Figure 283: img

The "Export" table header action allows the user to "Tag All", "Untag All" and "Invert tags" to flip tagged/untagged exam series 2000. To the right of these actions is the number of currently selected images from the total amount marked for export. In this case, 267 images are tagged out of a total of 970 images in the entire study 267/970.

The middle part of this table allows the user to select data for export by checking the \square / \square to the right of the selected area (patient, study, series, image). Individual areas can be downloaded or expanded using the arrow to the left of the area name, up to individual images. By hovering the mouse over the selected area for export, a preview will be displayed to the user, to simplify the work:

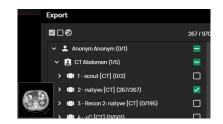


Figure 284: img

If the study contains more items than can be displayed simultaneously in the "Export" table, the user is allowed to scroll in this table using the scroll bar located to the right of the middle part, or by moving the mouse wheel.

The right part of the tab allows the user to select the method for exporting the data called "Operations" using the drop-down menu:



Figure 285: img

By selecting the desired operation and confirming the "Confirm" action, you will invoke one of the selected actions described below. By choosing the "Cancel" action, you cancel the export action.

If you do not select any data, the export operation will display "This field is required!".

9.6.2.5.1 Create a video from a series of images {#viewws_Create a video from a series of images} Selecting this option from the "Operations" drop-down menu will display a table of options for creating a video from the selected series of images:

Operation Create video from series of images
File name: Anonym_Anonym_1111_CT_Abdomen_na
Image format: MPEC
Number of frames per second: 25
Reverse order of frames

Figure 286: img

This tab includes: - File name - allows the user to choose the name of the saved file - Image format - allows you to choose the format of the saved file, namely MPEG or AVI - Number of frames per second - option to choose video playback speed - Reverse order of frames - checking this box will invert the position of the frames in the video

The "Confirm" action will start the export process, where an informative window will then appear in the bottom right corner about the progress of uploading the export for download:

Exporting images to video ()

Figure 287: img

By choosing the action "Stop" • you cancel the action being performed.

By choosing the "Info" action • you can call up a detailed tab to preview the export progress:

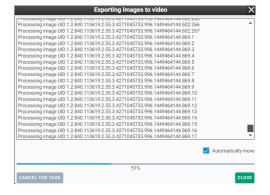


Figure 288: img

After a successful export for download, the exported data is automatically downloaded to the workstation's local storage.

9.6.2.5.2 Create archive {#viewws_Create archive} See more in "Export selected images".

9.6.2.5.3 Print The "Print" function allows the user to print selected images on the printer. You only need to select one series or specific images from one series to be able to print. Selecting this action calls up a tab with print options:



Figure 289: img

This tab contains: - Printer - option to choose one of the available, configured printers - Image layout - option to choose the layout of images on the page 1x1, up to 10x10 - Page Orientation - the option to select the orientation of the image on the page when printing, either Automatic, Portrait, or Landscape - Print multiframe - Format - option to choose the page size print format: A3, A4, A5, A6, PHOTO 10x15, LETTER - Print OSD labels - option to choose to print information about the image - OSD Color - option to select a light/dark color variant of the image information to be printed. This option is only active when "Print OSD labels" is selected - Print measurements

By choosing "OK" you go to the printer settings for printing and then start printing, or by choosing "Cancel" you cancel the action.

9.6.2.5.4 Export to DICOMDIR {#viewws_Export to DICOMDIR} See more in the article "Export selected to DICOMDIR".

9.6.2.5.5 Burn See more in the "Burn selected" article.

Figure 290: img

9.6.2.6 Share study {#viewws_Share study} The "Share study" function is used to create a onetime URL web link for sharing the study to an external user. Right-click on the name of the study, choose the "Share study" function, which brings up the "Create an external link to share data" tab:

<

Create external link to share data	×
Remaining external share links: 100	
Name of the recipient or description of the shared link* Anonym Anonym - Abdomen	
PIN* 4230	
Number of days the link is valid* 30	*
CANCEL	CREATE

Figure 291: img

The called tab contains a pre-filled text field that can be edited:

- Name of the recipient or description of the shared link enter the name of the designated user or a short description of the study being sent
- PIN enter the password for possible opening of the external link (alphanumeric characters can be entered)

Warning: Do not send the shared URL link together with the PIN in one communication due to the security of the data being sent. Always use two types of communication, e.g. email and SMS.

- Number of days of validity of the link determines the period for which the link will be active (the default maximum validity period is 30 days, this value can be configured by the administrator), by selecting the arrows to the right of the set validity period, you select the maximum possible validity period
- Use the "CANCEL" button to cancel the action, use the "CREATE" button to create a URL link to share the selected study, you will be informed about this by the "Share link" table:

Share link information			
Name of the recipient or de	scription of the share Anonym Anonym - Abdomen		
Share link	https://test.dicompass.ck 🖘 🖶 >		
PIN	4230		
Number of days the link is v	valid 1		
CLOSE			

Figure 292: img

- "Save to Clipboard" icon 🖙 copies the created URL link for possible pasting using keyboard shortcut ctrl+v
- The "Print link" icon 👼 will create a document with the appropriate data to access the shared study with the option to print it:



Figure 293: img

Figure 294: img

9.6.2.7 Split series {#viewws_Split series} The "Split series" function is used to separate a series into two or more subseries. This function can be used, for example, with MR dual protocol, when the series contains two series with different sequence parameters:

🗖 Split series

Figure 295: img

By choosing this action, you will create a new series in the "Viewer working set":



Figure 296: img

The selected split series can be displayed by double-clicking on the series thumbnail in the viewer

working set, or by drag&drop into the "Image Data Display Window".

9.7 DICOM viewer tools {#viewtools_DICOM viewer tools}

Figure 297: img

The DICOM viewer tools are used to work with the examination displayed in the "Window for displaying image data" and is divided into several areas:

- 1. Icon to **return to main dialog** of for searching and managing registries.
- 2. **Configure the layout of the panels** in the current window/monitor **E**. Selecting this action allows the user to layout the "Window for displaying panel data" into individual panels according to this configuration:



Figure 298: img

If you work at a workstation with multiple monitors, the selected layout can be applied to all DICOM viewer windows simultaneously using the "Change layout on all monitors" action **I**, if this feature is active, it will appear on a red background **I**.

To save the currently selected panel layout in the "window for displaying panel data" select the "Save current layout" action **I**. When the study is reopened, the panel layout will correspond to the newly set value selected during saving, in case the hanging protocol is not applied.

3. Selecting the Hanging protocol E - action for selecting the Hanging protocol, i.e. displaying the study, according to the parameters set in the "Window for displaying image data", such as the number of panels, selected windowing, display size, etc.

By selecting this action, you will call up a window for selecting a specific Hanging protocol:

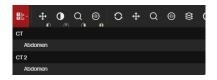


Figure 299: img

This action is made available on the condition that the user has multiple Hanging protocols set for a specific type of study. The names and number of Hanging protocols may differ, depending on the set values.

In the case of opening a mammography study (MG), the Hanging protocols are already prepared. The "Hanging protocol selection" thus contains the following window layouts (the currently selected hanging protocol is then highlighted in red):

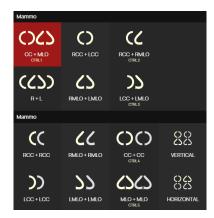


Figure 300: img

4. Currently **selected tools on individual mouse buttons**. They change according to the browser configuration and user saved configuration.



Figure 301: img

Tools, can be set to left, middle, right and left+right mouse buttons.

If a tool is selected from the "User configurable toolbar" or from the "Tab with tools", the active tool on the mouse button will change, according to the mouse button used when selecting the tool.

The configuration of the tools on the mouse buttons after opening the study can be found in the "DICOM viewer settings" chapter.

Use the "ESC" key to return to the default tools for each mouse button.

5. User configurable toolbar

C ⊕ Q ® ≋ O O ⊷ 높 ∞ A ⇔ B O

Figure 302: img

Available tools depend on DICOM viewer configuration and user settings. If the user changes and saves the tools configuration, the tools will be loaded at the next login. They may vary between users and stations.

6. Tabs with all the tools for working with image documentation and their settings



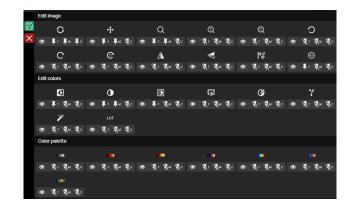
Figure 303: img

Individual tabs with tools will be detailed in the following chapters: - Display tools - Measurement tools - Action tools for series - Volume operations, 3D visualization - Online consultation - DICOM viewer settings - Working toolset

9.7.0.1 Configuring the tool display {#viewtools_Configuring the tool display} The display of tools for working with image documentation can be configured by the "Settings" action located to the left of the open tab:



Figure 304: img



Selecting this action displays the tools for configuring the tool display:

Figure 305: img

The following actions will appear under each tool in the tool group: - Hide/show action 💁 / 🔤 - choosing this action hides / restores the tool of the tool group in the selected tool tab



Figure 306: img

- Unpin/pin from top bar II / II choosing this action will hide / restore the tool in the "User Configurable Toolbar"
- Unpin/pin to top bar for mobile view 🔤 / 🔤 selecting this action will hide/restore the tool in the "User Configurable Toolbar" in minimized/mobile mode
- Unpin/pin from context menu I / N choosing this action hides / restores the tool in the "Context menu" called by the right mouse button in the "Window for displaying image data"

You can save the changes made to the tool tabs using the 🖹 function, or cancel the changes using the 🔀 function located to the left of the open tool tab.

~

9.7.1 Display tools {#viewtools_Display tools}

Figure 307: img

The "Display tools" tab is mainly used for modifying the display of image data, i.e. as postprocessing of the image. In case you want to reset the image editing to the original state, you do it by selecting the "Default layout" tool.

This tab contains the following tool groups: - Edit image - Edit colors - Color palette



Figure 308: img

If a keyboard shortcut is displayed under the tool icon, it can be used to perform the tool action.

9.7.1.1 Edit image {#viewtools_Edit image}	The "Edit image" tool group contains the following
tools:	

lcon	Function	Description
CTRLR	Default Layout	Resets the display - window size, center and width, offset, pseudo-color palette
↔	Move	Enables moving the image within the displayed window
Q	Zoom	After pressing and holding the selected mouse button, scrolling up and down increases and decreases the image
	Zoom 1:1	Selects the image size so that the image is displayed pixel by pixel on the monitor. There is no recalculation

lcon	Function	Description
Q	Optimal zoom	Magnifies and shifts the image so that the maximum of the work area designated for displaying the image is used
∠ ^z	Rotate 90° CCW	Rotates the image 90° counterclockwise
$\mathbf{C}_{_{\mathrm{M}}}$	Rotate 90° CW	Rotates the image 90° clockwise
¢	Free rotation	Allows you to rotate the image by a free angle, smoothly scrolling with
	Flip vertically	Flips the image along the vertical axis
\mathbf{V}	Flip horizontally	Flips the image along the horizontal axis
M3	Waveform config	Allows changing the layout of the ECG waveforms in the display window
6	Magnify glass	After pressing and holding the selected mouse button a circle will appear with an enlarged area under the mouse cursor. This tool is also available without the need to select a tool. After pressing and holding the SHIFT keyboard key, the same tool will appear. In this case, it is possible to gradually and cyclically increase the magnification in 5 steps by clicking the left mouse button.
У	Artificial Intelligence	Submitting a Study to Evaluate Abnormalities Using Artificial Intelligence (the icon is only displayed if AI is integrated into the product interface)

9.7.1.1.1 Free rotation {#viewtools_Free rotation} The "Free Rotation" tool allows you to rotate the image by dragging it to a position selected by the user, to rotate the image, follow these steps: - Select the "Free Rotation" tool with the desired mouse button in the " Display Tools" tool group. - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to select a new rotation of the image to the desired position and then release the mouse button - the image will be rotated according to the position selected by the user

For free rotation, you can also use the side "Postprocessing toolset" allowing you to adjust the brightness window, colors and filters, or rotate:



Figure 309: img

As part of the transformation, the "Working toolset" includes: - Free rotation - by using the scroll bar or by typing a value into the text field, the user can rotate the image by -180° to 180°, the default value is 0 - Flip by vertical axis - flips the image according to the vertical axis - Flip by horizontal axis - flips the image by the horizontal axis

₩3

Figure 310: img

9.7.1.1.2 Waveform config {#viewtools_Waveform config} Selecting this tool calls up the "ECG toolset" for changing and displaying individual ECG curves:



Figure 311: img

This table contains: - Waveform gain - allows you to vary the waveform signal strength from 5mm/mV to 40mm/mV - Speed - allows you to vary the speed of the waveform between 25mm/s and 50mm/s - Waveform layout - allows you to change the layout of individual curves, using the drop down menu:

Waveform layout 3x4+3	•
3x4	
3x4+1	
3x4+3	
6x2	
6x2+1	
12x1	
1x1	
3x1	
3x2	

Figure 312: img

- Background colour allows the user to choose between a dark and light background for the ECG study
- Active Group if the modality contains a multiplexer, this drop-down menu can be used to select the desired active group of ECG signals
- Active group display and hide individual waveform signals

У

Figure 313: img

9.7.1.1.3 Artificial Intelligence {#viewtools_Artificial Intelligence} The "Artificial Intelligence" tool is used to submit a study to evaluate abnormalities using artificial intelligence (AI).

Warning: This tool is only available if you have purchased and installed an AI license and this AI is integrated in the DPGW system.

To submit a study for AI evaluation, follow these steps:

- Open the desired study in a DICOM viewer window
- Select the "Artificial Intelligence" tool **Y** from the "Edit image" tool group
- You will be informed about the initiation of the AI submission and evaluation process:

A job has been started, it may take a while to complete

Figure 314: img

• To update the "Viewer Worksets" sidebar items, we recommend turning on the "Watch study for changes" function, see the "Workset Floating Panel" chapter

Example of X-ray study evaluation

Evaluation of the study using AI, this study will be supplemented with PR (Presentation state) and SR (Structured report) items:



abnormality finding, PR



In the case of a positive abnormality finding, PR will be marked, in the case of a negative



will be marked.

When the image is displayed in the DICOM viewer window, abnormalities will be marked, including text describing the abnormality:



Figure 315: img

To hide the AI evaluation, select the function , located under the upper right OSD label. Selecting this action brings up a drop-down menu for selecting a view:



Figure 316: img

Keyboard shortcuts can be used to toggle AI evaluation or other measurements. These are not set in the default interface and must be assigned using the "Keyboard Shortcuts" tool located in the "Viewer Settings" tab under "Load next/previous saved measurements".

The created SR contains the text of the conclusion evaluated by artificial intelligence.

Patient Anonym Anonym *1/1/2023 , male ID: 1111 Study Accession number : 1122233444555 Date of examination : 1/1/2023 10:00:00 Study description : CR HAND Language of Content Item and Descendants: English Observer Type: Device Device Observer Manufacturer: Milvue Device Observer Model Name: v1.20-8170f5c9

Findings : Finding: hand radiography

No evidence of acute fracture or dislocation. Bone density is normal.

Figure 317: img

Examples of study evaluation using AI by Rayscape and QureAI:



Example of CT study evaluation

Evaluation of the study using QureAI's AI, this study will be added with OT (Other) items:



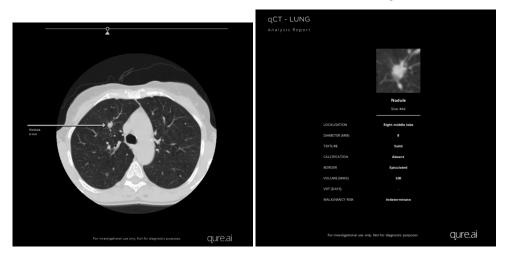
Figure 318: img

The first OT item contains a series of CT scans, including the display of the abnormality marked in the series progress bar:



Figure 319: img

The second OT item contains the displayed abnormality finding and the detected finding:



Warning: The artificial intelligence function serves as a supporting tool for evaluating data from RDG modalities, so it does not replace the expert work of a radiologist. The performing physician is responsible for the final report of the examination evaluation.

9.7.1.2 Edit colors {#viewtools_Edit colors} The "Edit colors" tool group contains the following tools:

lcon	Function	Description
CTRLI	Invert	displays the image in reverse. If the image is black and white, the black/white display will be inverted to white/black. In the case of choosing a color palette, the image is inverted within the given color representation
0	Windowing	after pressing and holding the selected mouse button, it is possible to change the width and center of the brightness window by moving the mouse up/down and left/right to improve the readability of the image or highlight the given structures. If the OSD labels are not changed, the values of the window can be read in the lower right corner, or on the brightness axis (standardly located at the right edge of the image). If you use this tool in a 3D projection, you can change the display of volume structures
	Reset contrast according image values	this tool will select a brightness window calculated from the entire image
ត្	Area windowing	after selection, it is possible to mark the area with a rectangle. From this, the optimal window for displaying all structures will be calculated
CTRLE	Set contrast values	allows manually enter the brightness window through its values
γ	Gamma correction	allows you to change the gamma correction using a called-up sidebar with the option to select gamma values from 0.1 to 5 with a sliding bar
1	Image Filters	for more detailed image processing, smoothing, edge detection and sharpening filters are available
LUT OTRE	Toggle VOI LUT	for some images it is possible to switch the VOI LUT (The Value Of Interest lookup table). It is basically a curve that in a way transforms the display of the given image in a different presentation of contrast, shades of gray, Very often this function is used to switch the display between the display set on the modality and the display using a standard DICOM curve

CTRLE

Figure 320: img

9.7.1.2.1 Set contrast values {#viewtools_Set contrast values} Selecting this tool calls up the "Postprocessing toolset" sidebar:



Figure 321: img

Within the setting of contrast values, the "Working toolset" in the "Windowing" section contains:

• Windowing - allows the user to select predefined values of the brightness window from the dropdown menu (the values of the Center and Width of the sidebar will be subsequently adjusted according to the selected window of values):



Figure 322: img

- Center option to select the center of the brightness window using the text box or the scroll bar
- Width option to select the width of the brightness window using the text box or the scroll bar

1

Figure 323: img

9.7.1.2.2 Image filters {#viewtools_Image filters} Selecting this tool calls up the "Postprocessing toolset" sidebar.

Within the image filter settings, the "Working toolset" in the "Color transform" section contains:

- Gamma correction allows you to change the gamma correction using a text field or sliding bar gamma values from 0.1 to 5
- Image filters allows the user to select individual filters from the scroll bar and specify their intensity using the slider bar:

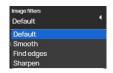
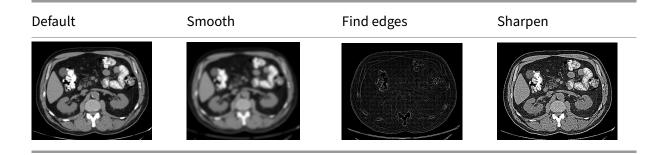


Figure 324: img



• Color palette - contains palettes according to the standard and allows the user to select a palette from the drop-down menu:

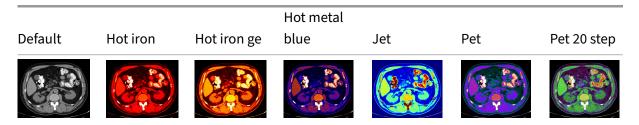
Color palette Default	•
Default	
Hot iron	
Hot iron ge	
Hot metal blue	
Jet	
Pet	
Pet 20 step	

Figure 325: img



Figure 326: img

9.7.1.3 Color palette {#viewtools_Color palette} The group of tools "Color palette" contains palettes according to the standard, converting gray levels to a fixed color palette. The tool group contains the following palettes:



9.8 Measurement tools {#viewmeas_Measurement tools}

The "Measurement tools" tab is used for measuring, labeling and editing measurements, including saving them in an image data frame. In case you want to use the measurement tool, assign the measurement functions to the selected mouse button. This function will then be displayed in the currently selected tools on individual mouse buttons (paragraph 2 of the "DICOM viewer Tools" chapter) and become active.

This tab contains the following groups of tools:

- Distance measurement
- Area measurement
- Measurement other actions



Figure 327: img

9.8.1 Distance measurement {#viewmeas_Distance measurement tool group}

lcon	Function	Description
••	Distance measurement	Measuring the distance between two points in the image
N	Freehand	Distance measurement using multiple points, or to measure the perimeter of an object in the image
۲.	Angle measurement	Angle measurement in the image
COBB A	Cobb angle	Measuring the angle between two line segments in the image
ВОНК	Böhler angle	Measurement of the angle of intersection of line segments in the image
NORB	Norberg angle	Measurement of the angle of the femur heads and their distance
ŧ	Lower limb angles	Measurement of individual angles of the lower limb in the image
	Distance ratio	Measuring the size difference of two line segments in the image
Ð	Note	Inserting text with a directional arrow into the image

The "Distance measurement" tool group contains the following tools:

lcon	Function	Description
¢	Calibration	Calibration of the measured values in the image, by changing the values of the measurement range
Q	Optical density	Informative function to display the blackening value of the photographic layer in units of HU/US
хүг	Coordinates	Informative function for determining the position of the point of interest, relative to the image
XY	XY value	Informative function for determining the position of the point of interest in the XY axis, relative to the image
B	Vertebra note	Marking of individual vertebrae in the image
至	Waveform measurement	Measurement of two values relevant to ECG imaging

Figure 328: img

9.8.1.1 Distance measurement {#viewmeas_Distance measurement} The "Distance measurement" tool measures the distance between two points in the image: - Select the "Distance measurement" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - in the image, the distance will be drawn with a line, with the measured value in mm

•--•



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Figure 329: img

Figure 330: img

9.8.1.2 Freehand The "Freehand" tool measures distance using multiple points, or to measure the perimeter of an object in the image: - Select the "Freehand" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - move the mouse to the next point of interest and click on the selected mouse button with the tool - if you want to complete the measurement, move the mouse to the end point of interest and double-click the selected mouse button with the tool to complete the measurement - the measured distance will be drawn in the picture, with the value in mm

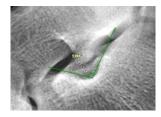


Figure 331: img

<u>ل</u>

Figure 332: img

9.8.1.3 Angle measurement {#viewmeas_Angle measurement} The "Angle measurement" tool measures an angle in an image: - Select the "Angle measurement" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - move the mouse to the next point of interest (vertex) and click on the selected mouse button with the tool - move the mouse to the end point of interest and click on the selected mouse button with the tool - an angle will be drawn in the image, with the specified value in degrees



СОВВ

Figure 333: img

Figure 334: img

9.8.1.4 Cobb angle {#viewmeas_Cobb angle} The "Cobb angle" tool measures the angle between two line segments in the image: - Select the "Cobb angle" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - this will create the first segment, proceed in the same way for the second segment - the angle between two line segments will be drawn in the image, with the specified value in degrees



BÖHL

Figure 335: img

Figure 336: img

9.8.1.5 Böhler angle {#viewmeas_Böhler angle} The "Böhler angle" tool measures the angle of intersection of line segments in the image: - Select the "Böhler angle" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - hover the mouse over the next point of interest (intersection of line segments) and click on the selected mouse button with the tool - move the mouse to the end point of interest and click on the selected mouse button with the tool - the angle at the intersection of the line segments will be drawn in the image, with the specified value in degrees



Figure 337: img

Figure 338: img

9.8.1.6 Norberg angle {#viewmeas_Norberg angle} The "Norberg angle" tool measures the angle of the femur heads and their distance: - Select the "Norberg angle" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest and click and hold the selected mouse button with the tool to create a circle corresponding to the left femoral head - this circle will be copied for possible placement on the right femoral head, click the mouse to confirm - the measurement will be drawn in the image



Figure 339: img

Warning: Use the "Norberg angle" only for veterinary purposes.

+ T

Figure 340: img

9.8.1.7 Lower limb angles {#viewmeas_Lower limb angles} The "Lower limb angles" tool measures the individual angles of the lower limb in the image: - Select the "Lower limb angles" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - this will create the first segment, proceed in the same way for the second segment - the angles of the lower limb will be drawn in the image, with the values in degrees: - HKA - hip-knee-ankle angle - FMA - femoral mechanical angle - TMA - tibial mechanical angle - JLCA - joint line convergence angle



Figure 341: img

₽ X:Y

Figure 342: img

9.8.1.8 Distance ratio {#viewmeas_Distance ratio} The "Distance ratio" tool measures the size difference between two line segments in the image: - Select the "Distance ratio" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - this will create the first segment, proceed in the same way for the second segment - the measurement of the ratio of lengths will be drawn in the image, with the specified value of the difference



3

Figure 343: img

Figure 344: img

9.8.1.9 Note The "Note" tool is used to insert text with a directional arrow into the image: - Select the "Note" tool with the selected mouse button in the "Distance measurement" tool group - move the mouse to the point of interest where the text of the note will be placed, press and hold the selected mouse button with the tool - hover the mouse over the point to which the note will point and release the selected mouse button with the tool - a table will be displayed with the possibility to insert the text note



Figure 345: img

- use the "Esc" key to cancel the action, or type text and use the "Enter" key to insert the selected text into the image
- an arrow with a text note will be drawn in the image

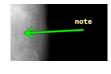


Figure 346: img

Figure 347: img

9.8.1.10 Calibration The "Calibration" tool is used to calibrate the measured values in the image by changing the values of the measurement range: - Select the "Calibration" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - a table will be displayed with the possibility of entering a numerical value in cm for calibration



Figure 348: img

using the measurement tool icon, located on the right side of the displayed table, select the tool
that will be applied after the calibration is completed, on the mouse button that you used to
perform the calibration



Figure 349: img

• use the "Esc" key to cancel the action, or type the value and the "Enter" key will calibrate the measurement according to the specified value

Q

Figure 350: img

9.8.1.11 Optical density {#viewmeas_Optical density} The "Optical density" tool serves as an informative function to determine the measure for the blackening value of the photographic layer in HU "Houndsfield unit" value. It expresses the radiation absorption of a given voxel relative to the absorption of water radiation (for water HU = 0) from different radiation directions. Examples of values are shown in the following table:

Tissue	CT number, HU density
air	-1000
fat	-50100
water	0
liquor	5
cerebral white matter	30
cerebral gray matter	34
blood	47
liver	40-60
muscles	35-75
fibrous tissue	60-90
cartilage	80-130
bone	1000-3000

To view optical density in a CT series: - Select the "Optical density" tool with the selected mouse button

in the "Distance measurement" tool group - this makes the tool active and you can dynamically display the HU in the image by moving the mouse over the area of interest

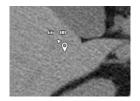


Figure 351: img

- if you want to record a static point of optical density in the image, hover the mouse over the area of interest and click on the selected mouse button with the tool
- a point with a displayed optical density value will be drawn in the image

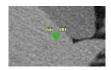


Figure 352: img

• to end the optical density display, select another tool from the "Distance measurement" tab

Warning: the value of the optical density measurement on the X-ray image, i.e. from the CR, DX modality, is expressed using the US unit "Unspecified" or HU "Hounsfield unit", depending on the modality of the examination on which the measurement is performed.

xyz

Figure 353: img

9.8.1.12 Coordinates The "Coordinates" tool serves as an informative function for determining the position of the point of interest, relative to the image, these coordinates of the patient are recalculated according to the DICOM tag "PatientImagePosition": - Select the "Coordinates" tool with the selected mouse button in the "Distance measurement" tool group - this makes the tool active and you can dynamically display the position in the image by moving the mouse over the area of interest



Figure 354: img

- if you want to record a static point of coordinate in the image, hover the mouse over the area of interest and click on the selected mouse button with the tool
- a point with the displayed coordinate value will be drawn in the image

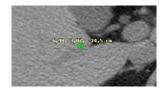


Figure 355: img

• to end the display of coordinates, select another tool from the "Distance measurement" tab

XY

Figure 356: img

9.8.1.13 XY value {#viewmeas_XY value} The "XY value" tool serves as an informative function to determine the position of the point of interest in 2 planes, relative to the image, these patient coordinates are recalculated, according to the DICOM tag PatientImagePosition: - Select the "XY values" tool with the desired mouse button in the "Distance measurement" tool group. - This makes the tool active and you can dynamically display the position in the image by moving the mouse over the region of interest



Figure 357: img

• if you want to record the position in the image, hover over the area of interest and click the selected mouse button with the tool

• a point with the displayed XY value will be entered in the image



Figure 358: img

• to end the display of coordinates, select another tool from the "Distance measurement" tab

:

Figure 359: img

9.8.1.14 Vertebra note {#viewmeas_Vertebra note} The "Vertebra note" tool is used to marking individual vertebrae in the image: - Select the "Vertebra note" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the vertebra to mark and click on the selected mouse button with the tool - a table will be displayed with the option to select the marked vertebra:

Pick fi	rst ver	tebra f	or labe	ling	×
		≣ļc	audal		
Cervical					
C1	C2	C3	C4	C5	C6
C7					
Thoracic					
	T2	Т3	Т4	T5	Т6
Τ7	Т8	T9	T10	T11	T12
Lumbar					
	L2	L3	L4	L5	
Sacrum					
S1	S2	S3	S4	S5	
Соссух					
Co1	Co2	Co3	Co4	Co5	

Figure 360: img

- select the direction of marking the vertebrae Caudal/Cranial
- select the name of the vertebrae to label
- the vertebra will be marked in the image:



Figure 361: img

• to mark more vertebrae, just click on the other vertebrae in the image, they will be marked according to the sequence:



Figure 362: img

• to cancel the marking of vertebrae, select another tool from the "Distance measurement" tab, the marking will also be inactive if you mark a whole series of vertebrae in a sequence

Ħ

Figure 363: img

9.8.1.15 Waveform measurement {#viewmeas_Waveform measurement} By selecting the "Waveform Measurement" tool, it is possible to measure two parameters relevant to the ECG display: - Select the "Waveform measurement" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest, press and hold the selected mouse button with the tool - move the mouse to the end point of interest and release the selected mouse button with the tool - in case of dragging the mouse horizontally between two points on the curve, the time will be measured in ms:



Figure 364: img

• in case of dragging the mouse vertically between two points on the curve, the magnitude of the signal will be measured in μV:



Figure 365: img

9.8.2 Area measurement {#viewmeas_Area measurement}

The "Area measurement" tool group contains the following tools:

lcon	Function	Description
	Rectangle	Measurement of the value contained in the rectangle
0	Ellipse	Measuring the value contained in the ellipse
O	Circle defined by 3 points	Measuring the value contained in a circle
Д	Freehand area	Measurement of the value contained in the polygon
0	Shape	Measurement of the value contained in the object selected by the user
S	Integral	Measurement of area integration values in Doppler ultrasonography
9	Velocity Time Integral	Measurement of blood flow values corresponding to the area under the time-velocity curve in Doppler ultrasonography

lcon	Function	Description
PHT	Preassure Half Time	Measurement of the time required for the pressure gradient to fall to half of its maximum value in Doppler ultrasonography
9	Cardiothoracic index	Measurement of the ratio of the maximum width of the cardiac shadow to the maximum internal chest width

Figure 366: img

9.8.2.1 Rectangle The "Rectangle" tool measures the values contained in the user-created rectangle drawn in the image: - Select the "Rectangle" tool with the selected mouse button in the "Distance measurement" tool group - move the mouse to the point of interest, press and hold the selected mouse button with the tool, this will start recording the rectangle from the corner and expand it by dragging - move the mouse to the end point of interest and release the selected mouse button with the tool - the rectangle area and the measured value of the HU diameter will be drawn in the image

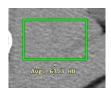


Figure 367: img



Figure 368: img

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Figure 369: img

9.8.2.2 Ellipse The "Ellipse" tool measures the values contained in a user-created ellipse drawn in the image. To create an ellipse, follow the same procedure as when creating a rectangle. This creates an ellipse with the measured values in the image:

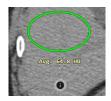
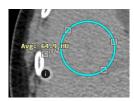


Figure 370: img

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Figure 371: img

9.8.2.3 Circle defined by 3 points {#viewmeas_Circle defined by 3 points} The "Circle defined by 3 points" tool measures the values contained in a user-created circle placed in the image. To create a circle, follow the same steps as for creating a rectangle. This will create a circle with the measured values in the image, which can be adjusted using 3 points:



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Figure 372: img

Figure 373: img

9.8.2.4 Freehand area {#viewmeas_Freehand area} The "Freehand area" tool measures the values contained in a user-created polygon drawn in the image: - Select the "Freehand area" tool with the selected mouse button in the "Distance measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - move the mouse to the next point of interest and click on the selected mouse button with the tool - if you want to complete the measurement, move the mouse to the first point of the free contour, highlighted by a circle, and click the selected mouse button with the tool to complete the measurement



Figure 374: img

• the area of the polygon and the measured value of the HU diameter will be drawn in the image

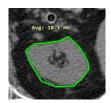


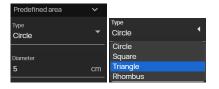
Figure 375: img

• for more information about the measured values, use the icon to display the "Frequency histogram" table, see section "Rectangle"

Figure 376: img

9.8.2.5 Shape The "Shape" tool is used to define the shape and size of the area, using a selection and a text field by the user: - Select the "Shape" tool with the selected mouse button in the "Distance measurement" tool group - use the "Measurement toolset" to select the desired shape type:

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- enter a numeric value for the area size in cm in the "Diameter" field
- after selecting the desired values, click the mouse in the selected part of the image to plot the selected area shape and the measured HU diameter value

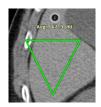


Figure 377: img

• if you have closed the "Measurement toolset" for more information about the measured values, use the icon to call up the "Frequency histogram", see paragraph "Rectangle"

Figure 378: img

9.8.2.6 Integral The "Integral" tool serves as a general function for measuring area integration values in Doppler ultrasonography:

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• Select the "Integral" tool with the desired mouse button in the "Distance measurement" tool group

- hover the mouse over the point of interest and click on the selected mouse button with the tool
- This is how you trace the curve in the measured sonogram
- double-click to end the measurement, this will display the measurement in the image with the measured values in mm

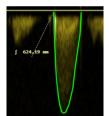


Figure 379: img

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Figure 380: img

9.8.2.7 Velocity Time Integral {#viewmeas_Velocity Time Integral} The "Velocity Time Integral" tool is used to measure blood flow values that correspond to the area under the velocity time curve in Doppler ultrasonography: - Select the "Velocity Time Integral" tool with the desired mouse button in the "Distance Measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - in this way draw a curve in the measured sonogram - double-click to end the measurement, this will display the measurement in the image with the measured values: - Vmax - Velocity maximum in units of m/s - Vmean - Velocity average in units of m/s - Pmax - Peak maximum in units of mmHg - Pmean - Peak average in units of mmHg - VTI - Velocity Time Integral in units of cm

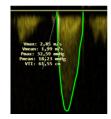


Figure 381: img

Figure 382: img

9.8.2.8 Preassure Half Time {#viewmeas_Preassure Half Time} The "Preassure Half Time" tool is used to measure the time required for the pressure gradient to drop to half of its maximum value in Doppler ultrasonography: - Select the "Preassure Half Time" tool with the desired mouse button in the "Distance Measurement" tool group - hover the mouse over the point of interest and click on the selected mouse button with the tool - draw a curve in the measured sonogram in this way - double-click to end the measurement, this will display the measurement in the image with the measured values in ms:

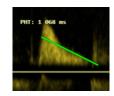


Figure 383: img

Figure 384: img

9.8.2.9 Cardiothoracic Index {#viewmeas_CTR index} The "Cardiothoracic Index" tool is used to measure the ratio of the maximum width of the cardiac shadow to the maximum internal chest width. To use it, proceed as follows:

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- Select the "Cardiothoracic Index" tool with the desired mouse button in the "Distance Measurement" tool group.
- to mark the inner chest width distance, hover the mouse over the point of interest, click on the selected mouse button with the tool and drag the mouse to the second point of interest to measure the maximum inner chest width
- to mark the maximum width of the heart shadow, hover the mouse over the point of interest, click on the selected mouse button with the tool and drag the mouse to the second point of interest of the heart to measure the maximum width of the heart shadow
- the result will be the measured value of the ratio of the maximum heart shadow width to the maximum inner chest width in percentage:

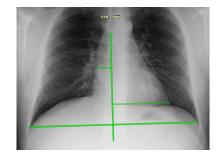


Figure 385: img

9.8.3 Measurement other actions {#viewmeas_Measurement other actions}

The group of tools "Measurement other actions" contains the following tools:

lcon	Function	Description
1	Save all measurements	Create a presentation state with embedded measurements taken over the entire patient study
~o	Remove measurement	Remove the selected measurement from the image
ìo	Remove all measurements	Remove all measurements from the displayed image
ĩó	Remove all measurements from current session	Remove all measurements from all series and all open studies
10	Hide measurement	Hides all measurements from the image
Ø	Common measurement for images in series	Keeping of all measured objects in the displayed window for all images in the series
Ø	Volume of measurements in series	Measurement of the volume in cm ³ between the areas marked "Area measurement"
Ø	ROI measurements config	Specifying the display of the primary value "Area measurement" in the image
	Frequency histogram	Displaying information about the measured values of the measured and selected area
**	Display PR after study opened	Enable/disable automatic display of saved measurements in the image after opening the study

lcon	Function	Description
⇔	Export measurements	Automatic sending of measured values to a third party system

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Figure 386: img

9.8.3.1 Save all measurements {#viewmeas_Save all measurements} The "Save all measurements" tool is used to create a presentation state, or "Secondary scan", saving all measurements taken throughout the patient's study. Thus, this save does not modify the original study stored in the PACS archive, but only adds information about the editable measurements in DICOM PR format, or creates a new snapshot without editing the measurements. To save a measurement to a snapshot, proceed as follows: - Select the "Save all measurements" tool by left mouse button in the "Distance Measurements" tool group to call up the "Save Measurement" table.

Save all measurements	×
Saved measurement format Presentation State (GSPS / CSPS)	
Target Archive	
Archive DPGW (DPGW@127.0.0.1:5380)	•
Series description annotated	
CANCEL SAVE	

Figure 387: img

- select the format of the saved measurement:
 - Presentation State saves only the measurements in PR format over the measured images and the measurements can be edited afterwards.
 - Secondary Capture saves the measurement including the image/series over which the measurement was made in a non-editable format



Figure 388: img

• select the save destination from the drop-down menu



Figure 389: img

• in case you have selected the save target "Archive" and have configured multiple PACS targets, select the desired PACS archive to save (If the user target is selected, the station will make this window inactive)



Figure 390: img

• name the saved series with measurements in the image



Figure 391: img

- "Cancel" to cancel the action, "Save" to create a presentation state or "Secondary capture" with measurement in frame/series
- the newly created presentation state or new series with saved measurements will be added to the "Viewer working set":

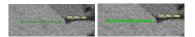


Keyboard shortcuts can be used to view or toggle stored measurements. These are not set in the default interface and must be assigned using the "Keyboard Shortcuts" tool located in the "Viewer Settings" tab under "Load next/previous saved measurements".

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Figure 392: img

9.8.3.2 Remove measurement {#viewmeas_Remove measurement} The "Remove measurement" tool is used to remove the selected measurement from the image: - Mark the selected measurement for deletion by clicking on it with the left mouse button, it will be highlighted



- select the "Remove measurement" tool with the left mouse button in the "Distance measurement" tool group, or press the DELETE keyboard key
- the selected measurement has been removed

Figure 393: img

9.8.3.3 Remove all measurements {#viewmeas_Remove all measurements} The "Remove all measurements" tool is used to remove all measurements from the displayed image: - Select the "Remove all measurements" tool with the left mouse button in the "Distance measurement" tool group, or press the keyboard shortcut SHIFT+DELETE - this removed all measurements from a specific image in the series

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Figure 394: img

9.8.3.4 Remove all measurements from current session {#viewmeas_Remove all measurements from current session" tool is used to remove all measurements from all series and all open studies: - Select the "Remove all measurements from current session" tool with the left mouse button in the "Distance measurement" tool group, which will display the action confirmation table (if you check the option "Do not ask me again" you will not be asked to confirm next time and all measurements will be removed):

Question		
Are you sure to remove all measurements from current sess	sion? This action car	n not be taken back.
	NO X	YES 🗸

Figure 395: img

• press the "NO" button to cancel the action, press the "YES" button to delete all the measurements

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Figure 396: img

9.8.3.5 Hide measurement {#viewmeas_Hide measurement} In the event that the measured objects prevent work with image documentation, they can be temporarily hidden, without the need to remove them with the "Hide measurement" tool: - Select the "Hide Measurement" tool with the left mouse button in the "Distance Measurement" tool group, or in the "Display toolset" - This will hide all measurements, the "Hide Measurements" icon in the "Distance Measurements" tool group will be marked with a red background and you will be alerted with information:

Show measurements: No

Figure 397: img

 in case you want to display the measurement again, select the "Hide measurement" tool again with the left mouse button in the "Distance measurement" tool group or in the "Display toolset", by selecting this action you will be notified with information:

Figure 398: img

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Show measurements: Yes

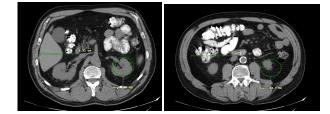
Figure 399: img

9.8.3.6 Common measurement for images in series {#viewmeas_Common measurement for images in series} The tool "Common measurement for images in series" is used to keeping all measured objects in the displayed image for all images in a series. When moving to the next images in the series, the measurement remains unchanged and will be displayed on the next images as well: - In case all the required measurements have been created in the image, select the "Common measurements for images in a series" tool by left-clicking in the "Distance measurement" tool group or in the "Display toolset" - This will anchor the measurements from a specific image for the entire series, the "Common measurements for images in a series" icon in the "Distance measurement" tool group will be marked with a red background image and you will be informed about this:

Common measurements for images in the series: Yes

Figure 400: img

• measurement objects remain unchanged when moving to the next images in the series



• in case you no longer want to use the common measurement action, re-select the "Common measurement for images in series" tool by left-clicking in the "Distance measurement" tool group or in the "Display toolset", by selecting this action you will be notified with information:

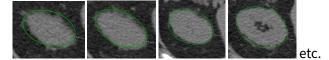
Common measurements for images in the series: No

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Figure 401: img

Figure 402: img

9.8.3.7 Volume of measurements in series {#viewmeas_Volume of measurements in series} The "Volume measurement in series" tool measures the volume in cm³ between the areas marked with "Area measurement": - Mark the area of interest with "Area measurement" in consecutive images in a series, e.g:



• if you have marked all the required areas in the series, select the tool "Measurement volume in series" with the left mouse button in the tool group "Distance measurement" to call up an informative table with the volume calculation:



Figure 403: img

- this table contains information about the measured volume value in cm³ and the number of frames from which this value was created
- click on the "OK" button to close the table

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Figure 404: img

9.8.3.8 ROI measurements config {#viewmeas_ROI measurements config} The "ROI measurements config" tool is used to determine the display of the primary "Area measurement" value in the image: - Select the "ROI Measurement Settings" tool with the left mouse button in the "Distance Measurement" tool group to call up the "Measurement toolset" sidebar enhanced with a histogram and other options

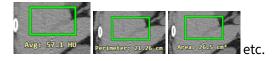


Figure 405: img

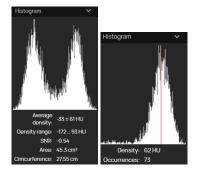
In this toolset, the following settings are possible: - Primary ROI value - a drop-down menu allows the user to select the displayed ROI value in the image:



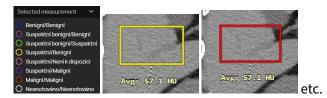
Figure 406: img



• Histogram - used to get more information about the measured values of the measured and selected area, if the mouse is moved to the histogram graph, the specific value is displayed:



- Predefined area using the type selection dropdown menu and scale text box, allows the user to define the shape using the "Type" dropdown menu (circle, square, triangle, rhombus) and the area size of the "Shape" tool using the "Diameter" text box to enter a value in cm
- Selected measurement by selecting from the predefined labels, the colour of the measured area in the image can be changed to indicate, for example, the severity of the condition:



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Figure 407: img

9.8.3.9 Frequency histogram {#viewmeas_Frequency histogram} The "Frequency Histogram" tool is used to get more information about the measured values of the measured and selected area. Select this action to call up the "Measurement toolset" with the display of the Frequency Histogram. This toolset can also be called up with the icon **1** displayed next to the measured area. For more information see the paragraph "Rectangle".

Figure 408: img

9.8.3.10 Display PR after study opened {#viewmeas_Display PR after study opened} The "Display PR after study opened" tool is used to enable/disable automatic display of stored measurements in the image, AI results, etc., in PR (Presentation state) format after opening a study.

By selecting this tool, the icon will be highlighted in red 쭽 and you will be notified when the feature is enabled:

Display PR after study opened: Yes

Figure 409: img

To disable this function, select this tool again, the icon will no longer be highlighted 🛱 and you will be informed about the disabling:

Display PR after study opened: No

Figure 410: img

This tool can also be found in the "Display toolset".

Figure 411: img

9.8.3.11 Export measurements {#viewmeas_Export measurements} The "Export Measurements" tool is used to automatically send the measured values to a third-party system. This tool is used e.g. for integration with NIS (hospital information system), where the measurements in this DICOM viewer are sent directly to the written findings by the radiologist.

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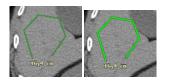
Warning: integration and configuration of this viewer with a third-party application is required for the possible use of this tool. If necessary, please contact your IT administrator of the PACS system.

9.8.4 Editing measurement {#viewmeas_Editing measurement}

Individual measured objects can be edited or the measured value can be moved by editing them.

To edit a measurement: - mark the selected measurement object by clicking the left button on it, which will highlight its outline

Dicompass Gateway



 hover the mouse over the selected measurement object, the individual measurement points will be displayed



Figure 412: img

• by clicking and holding the selected point, you can move the point to the place of interest



Figure 413: img

• the measured value can also be moved, you can do so by moving the point with the value to the place of interest

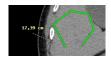


Figure 414: img

• by clicking outside the measured object, or by creating a new measurement object, the modified measurement object will be unmarked with a display of the modifications made

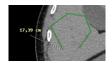


Figure 415: img

9.9 Series action tools {#viewser_Series action tools}

The tab "Series action tools" is used for working with the displayed series, locating the position in the series and synchronizing series.

This tab contains the following groups of tools: - Series actions - Lock view

Serie	s actio	าร			
k	CTRLH	÷.	ß	\$	
0	o	50 æ	٢		
Lock	view				
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8	A	Ф)		

Figure 416: img

9.9.1 Series actions {#viewser_Series actions}

The "Series actions" tool group contains the following tools:

lcon	Function	Description
न्त	Localizer lines	Display of locating lines, between two mutually perpendicular series
1 1	Hide reconstruction localizer lines	Hiding localization lines, between reconstructed mutually perpendicular series, during volume operations
*	Show slice thickness lines	Display of localization lines including display of selected slice thickness
£	Browse selected images only	Browsing only tagged key images in the series
\$	Stacking	Browse through the series with the selected mouse button
() MRI	Dynamic MRI stacking	Dynamic walk-through of the MRI series with the selected mouse button
C	Toggle Player	Display or hide the player for automatic browse through the series

lcon	Function	Description
Ô	Play/Pause in all windows	Shows the video player in all panels "Windows for displaying image data"
	Save key images	Save the marked key images to the new created series
۲	Fill other panels	Automatic series display, when changing the number of panels to display in the "Image data display window"
\odot	Video player tool	Tool for playing and working with video in DICOM format

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Figure 417: img

9.9.1.1 Localizer lines {#viewser_Localizer lines} The "Localizer lines" tool is used to display localization lines between two mutually perpendicular series and displays the current position of the image in the series.

To display "Localizer lines" proceed as follows: - Split the "Windows for displaying image data" into two panels by "Configure the layout of the panels" 🖬 - in these panels display two series whose reconstructions are perpendicular to each other - select the "Localizer lines" tool in the "Series Actions" tool group, or in the "Display toolset" - in the series that is perpendicular to the currently selected series, the localization lines are displayed



Figure 418: img

• to hide the localizer lines, re-select the "Localizer lines" tool in the "Series actions" tool group, or in the "Display toolset"

To hide the border localizer lines (border lines shown in green), use the "Hide border localizer lines" action located in the "Working toolset" tab of the "Display toolset".

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Figure 419: img

9.9.1.2 Hide reconstruction localizer lines {#viewser_Hide reconstruction localizer lines} The "Hide reconstruction localizer lines" tool is used to hide localization lines, between reconstructed mutually perpendicular series, during volume operations.

To hide the "Localizer lines for reconstruction" proceed as follows: - Split the "Image data display window" into the selected panel layout "Configure the layout of the panels" and display the individual series of the reconstructed images in the panels. They will be displayed with localization lines for the reconstructed series



Figure 420: img

- to hide the localizer lines for the reconstructed planes, select the tool "Hide Reconstruction Localizer Lines" in the tool group "Series Actions" or in the "Display toolset"
- localization lines for reconstructed series were hidden



Figure 421: img

• to redisplay the localizer lines for the reconstructed planes, select the "Hide Reconstruction Localizer Lines" tool in the "Series Actions" tool group or in the "Display toolset"

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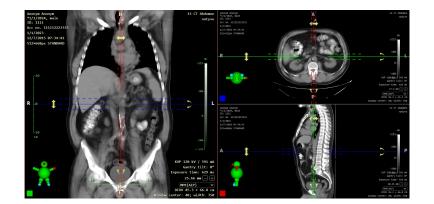
Figure 422: img

9.9.1.3 Show slice thickness lines {#viewser_Show slice thickness lines} The "Show slice thickness lines" tool is used to display the section thickness along localizer lines, between reconstructed perpendicular series, in volume operations. Activate this tool to display the thickness of the slice using dashed lines along the locator lines.

To show "Show slice thickness lines", proceed as follows: - Expand the "Window for displaying panel data" into the desired panel layout by "Configuring the panel layout in the current window/monitor" and display the individual planes of the reconstructed panel in the panels. - Display the localization lines for the reconstructed planes - To display the slice thickness of the localizer lines, select the "Show slice thickness lines" tool in the "Series Actions" tool group or in the "Display toolset". You will be informed about the display of the localizer lines including the slice thickness:

Hide reconstruction localizer lines: Yes

Figure 423: img



• This shows the localization lines including the thickness of the slice

Figure 424: img

• To hide the localizer lines including the slice thickness, select the "Show slice thickness lines" tool again in the "Series actions" tool group or in the "Display toolset". You will be informed about hiding the localizer lines including the slice thickness:

Show slice thickness lines: No

Figure 425: img

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Figure 426: img

9.9.1.4 Browse selected images only {#viewser_Browse selected images only} The tool "Browse selected images only" is used for browsing only marked key images in a series with standard tools (mouse wheel, keyboard shortcuts).

To tag and browse key images, follow these steps: - In the "Windows for displaying image data", display the selected series - when going through a series, you can mark individual images with the area of interest as a so-called key image with the icon of the check mark is located in the upper right corner of the "Image data display window". By marking it, the check mark icon is doubled is. The individual key images will also be displayed in the scroll bar for series passage on the right side of the screen.

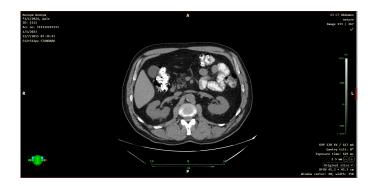


Figure 427: img

• to browse only selected key images, select the "Browse selected images only" tool in the "Series actions" tool group, or in the "Display toolset", selecting the action will notify the system of this information:

Browse selected images only: Yes

Figure 428: img

- this way, when going through the series, you will only skip over marked key images
- to go through the whole series again, select the "Browse selected images only" tool in the " Series actions" tool group, or in the "Display toolset", selecting the action will notify the system of this information:

Browse selected images only: No

Figure 429: img

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Figure 430: img

9.9.1.5 Stacking The "Stacking" tool is used to scroll through the series with the selected mouse button and thus complements the possibility of scrolling through the series with the mouse wheel or arrow keys.

To browse the series with the "Stacking" tool, proceed as follows: - Select the "Stacking" tool with the selected mouse button in the "Series actions" tool group - hover the mouse over the selected display panel with the series - by clicking the selected mouse button and dragging at the same time, you will go through the series - to cancel "Stacking", choose another tool on the selected mouse button

Figure 431: img

9.9.1.6 Dynamic MRI stacking {#viewser_Dynamic MRI stacking} The "Dynamic MRI stacking" tool is used for scrolling and switching MRI series subseries with the selected mouse button and therefore complements the series scrolling possible with the mouse wheel or arrow keys and the "Split series" tool.

To navigate through the series with the "Dynamic MRI stacking" tool, proceed as follows: - Select the "Dynamic MRI stacking" tool with the desired mouse button in the "Series actions" tool group. - hover the mouse over the desired image series display panel in the "Image Data Display Window" - click the selected mouse button and simultaneously drag sideways (left, right) to scroll through the series, drag the mouse up and down to switch between subseries - to exit the "Dynamic MRI stacking" select another tool on the selected mouse button

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Figure 432: img

9.9.1.7 Toggle Player {#viewser_Toggle Player} The "Toggle Player" tool is used to display or hide the player for automatic browse through the series as with a classic video player.



Figure 433: img

The player includes the following tools:

lcon	Function	Description
<	Previous frame	Moves back one frame in the series
	Play	Starts an automatic browse through the series
	Pause	Stops the automatic browse through the series
>	Next frame	Advances one frame in the series
liiil	Selecting playback section	Allows you to select a section of playback in a series
x1	Playback speed	Automatic playback speed selection
15 FPS	FPS playback speed	Selectable autoplay speed by selecting frame per second (FPS)
Ć	Loop	Repetitive automatic browse through series
<u>ت</u>	Loop back and forth	Autoplay through series will continuously play from beginning to end and backwards
×	Close player	Turning off the player

Playback speed

To change the playback speed of a series, select the "Playback Speed" tool and expand the dropdown menu:



Figure 434: img

Select a playback speed from the selected values, or select the "Custom" value to open the FPS speed change tab:



Figure 435: img

Type the desired value in the text field and choose to apply the selected speed to all series in the study. The "Cancel" option cancels the work performed, the "Save" option applies the entered values.

FPS playback speed

To change the playback speed of a series using frames per second values, select the "FPS Playback Speed" tool 📓 and expand the drop-down menu:



Figure 436: img

Select a playback speed from the selected values, or select the "Custom" value to open the FPS speed change tab:



Figure 437: img

Enter the desired value in the text field and choose to apply the selected speed to the displayed series in all panels of the "Image data display windows". The "Cancel" option cancels the work performed, the "Save" option applies the entered values.

Warning: when you enter a custom FPS value, the preset values in the drop-down menu will be affected according to the selected value.

Selecting playback section

The player allows automatic passage of only parts of the series. To select a playback segment, select the "Select playback section" tool . This displays the scroll bar located in the current position bar of the frame in the series:

		(
< > > 🔛	x1 15 🗂 🗁 134		×

Figure 438: img

Drag the end points of the triggered bar to control the size of the playback segment in the frame series:

							(2	
<	►	>		15 FPS	٥	134 267			×

Figure 439: img

DSA - Digital Subtraction Angiography

Figure 440: img

If you have a digital subtraction angiography study open, the "DSA" tool will also be displayed in the playback bar:

DSA



Figure 441: img

Selecting this tool will open the "DSA toolset"

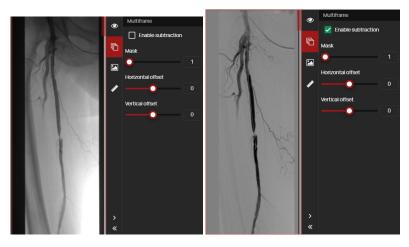
۲	Multiframe	
	Enable subtraction	
Ð	Mask	
P	0(1
	Horizontal offset	
	— •	0
	Vertical offset	
	— •	0

Figure 442: img

This toolset includes: - Enable subtraction - functions to show/hide the DSA mask - Mask - sliding bar

to adjust the DSA mask strength - Horizontal Offset - slider bar to move the DSA mask horizontally -Vertical Offset - sliding bar to move the DSA mask vertically

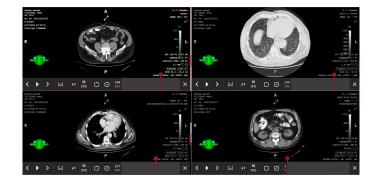
Examples of unapplied/applied DSA masks:



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Figure 443: img

9.9.1.8 Play/Pause in all windows {#viewser_Play or Pause in all windows} The "Play/Pause in all windows" tool will display the player in all "Windows for displaying image data" panels in which a series of images is displayed and automatically start browsing them.



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Figure 444: img

Figure 445: img

9.9.1.9 Save key images {#viewser_Save key images} The "Save key images" tool is used to save marked key images into the new created KO series.

• Select the "Save key images" tool in the "Series actions" tool group to display the "Save key images" table

Save key images	×
Target Archive	
Archive	_
DPGW (DPGW@127.0.0.1:5380)	
Series description	
CT Abdomen - important	
CANCEL	AVE

Figure 446: img

• select the save destination from the drop-down menu

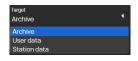


Figure 447: img

• in case you have chosen the storage destination "Archive" and you have configured multiple PACS destinations, select the selected PACS archive for saving (If the user destination is selected, this window will be inactive)



Figure 448: img

• name the new saved key images series



Figure 449: img

- the cancel button cancels the action, the save button creates a new series with key images
- the new created series will be added to the "Viewer Working Set"



Figure 450: img

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Figure 451: img

9.9.1.10 Fill other panels {#viewser_Fill other panels} The tool "Fill other panels" is used for automatic display of series, when changing the number of panels for display in the "Window for displaying image data". The series are added to the panels according to the order in the study, skiping already displayed series. If this action is not turned on, when other panels are displayed, they remain empty.

By selecting the "Fill other panels" tool in the "Series actions" tool group or in the "Display toolset", you will be notified of this by the system:

Fill other panels: Yes

Figure 452: img

To turn it off, select the "Fill other panels" tool again in the "Series actions" tool group or in the "Display toolset", you will be notified by the system:

Fill other panels: No

Figure 453: img

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Figure 454: img

9.9.1.11 Video player tool {#viewser_Video player tool} "Video player tool" is used for playing and working with video in DICOM format.



Figure 455: img

The "Video Player Tool" contains the following tools:

lcon	Function	Description
Þ	Play	To start video playback
	Stop	Stop video playback
< 5s >	Skip a time period	Moving the video loop according to the defined value
•	Volume	Change the video audio volume
x1,0	Playback speed	Change the video playback speed
 ‡	Crop video	Resize a video cutout for export or saving
liiiil	Cut video	Selecting the time period of the video for export or saving
≣	Marks	Calls up a tab for managing assigned marks
Ŧ	Add mark	Marking a frame of interest in a video
< >	Previous/next mark	Browse the associated marks in the video
	Download video frame	Export the video frame to the user's local storage in .jpg format
I	Download video	Export video to the user's local storage in .mp4 format
6	Save	Archiving edited video

1. Video playback

To play the video, use the "Play" icon **D** in the player toolbar.

To stop the video, use the "Stop" 🗖 icon located in the player toolbar.

To scroll through the video loop, click in the video location bar located in the player toolbar. You can also click and drag the mouse at the same time to scroll through the video.



Figure 456: img

For skipping by a period of time in the video, use the arrows ✓ ⁵⁶ >, where the value between the arrows indicates the time by which the video will be advanced. Clicking on the time value will call up a drop-down menu with predefined values that can be selected:



Figure 457: img

To change the video audio volume, select the volume icon <a>Image to display a drop-down menu with predefined volume values:



Figure 458: img



Figure 459: img

2. Video Editing

Marking images in video

To mark a frame of interest in a video, use the player's scroll bar to display the desired frame and then mark it using the "Marks" icon . When clicked, the icon will turn red and a vertical white line will appear on the player scroll bar:



Figure 460: img

You can navigate between the selected frames by clicking on the flag icons placed in the scroll bar of the player, or by using the

If you click on the **s** icon again, the marked image will be unmarked.

You can manage the selected images in a collective way using the table displayed by the 🗏 icon:

		×
	Description	
Seconds		×
Seconds	Description	×
Seconds	Description	×
	CANCEL	SAVE
		Seconds Description Seconds Description

Figure 461: img

In this tab, the individual parameters of the mark can be edited: - The "Time" field allows you to define the positioned marker of the marked frame in the video using a value in seconds -The "Description" field - allows you to define the name of the placed marker, in the preview of the marked frame of the video this description will be shown:



Figure 462: img

- "Delete" icon 🗵 selecting this action removes the placed marker
- "Add tag" button ADD MARK select this action to add a new tag row to the table with the option to define it:



Figure 463: img

- Use the "Cancel" option to cancel the action, the "Save" button to confirm the changes
- If you do not have any marks in the video, start with the "Add mark" button:



Figure 464: img

Video cutting

In case you want to cut only part of the video, select the "Cut video" icon 🔲 from the player toolbar.

This displays a border to mark a section of the video, located in the player toolbar:



Figure 465: img

By moving the ends of the border to mark a section of the video, you determine the time period of the video that will be cut and then saved to PACS:

I⊷I	00:00		

Figure 466: img

If you want to add another cut of video to a single saved loop, hover your mouse cursor over the player's toolbar to display the grey border of the next cut, and left-click to add that section:

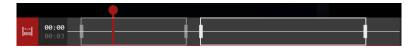


Figure 467: img

To remove a section of a video cutout, hover over the player scroll bar of the selected section and right-click to remove the section.

Video area cropping

By selecting the "Crop Video" tool 🛄 in the player toolbar, you can resize the video crop by dragging the red squares located in the upper left and lower right corners of the crop:



Figure 468: img

3. Archiving edited video and images

Export image

The "Download video frame" tool located in the player toolbar, is used to export a video frame to the user's local storage in .jpg format. This exported image does not contain OSD captions. In case you have defined a "Video area crop", only the selected crop will be saved.

Selecting the "Download video frame" tool will call up the "Download current frame" tab:

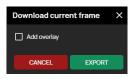


Figure 469: img

- Add overlay if a logo is configured in DPGW, selecting this field will insert a watermark into the saved image located in the upper left corner
- Cancel selecting "Cancel" will cancel the operation being performed
- Export selecting "Export" will start downloading the image to the user's local storage in .jpg format

Export video

The "Download Video" storage in .mp4 format. In case you have defined a "Video crop area" or "Video cut", only the selected crop/section of the video will be saved.

Select the "Download Video" tool to call up the "Download Video" tab:

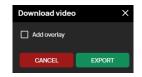


Figure 470: img

- Add overlay if a logo is configured in DPGW, selecting this field will insert a watermark into the saved video located in the upper left corner
- Cancel selecting "Cancel" will cancel the operation being performed
- Export selecting "Export" will start downloading the video to the user's local storage in .mp4 format

Save edited video

The "Save" tool 🖻 located in the player toolbar, is used to archive the edited video.

In case you want to save the selected cut and crop of the video, including the markers, select the "Save" tool, the "Forward data" tab will be called up:



Figure 471: img

• select the save destination from the drop-down menu



Figure 472: img

• in case you have selected the save target "Archive" and have configured multiple PACS targets, select the desired PACS archive to save (If the user target is selected, the station will make this window inactive)



Figure 473: img

• name the edited video for saving



Figure 474: img

• the "Cancel" button cancels the action performed, the "Save" button creates a new video that will be added as another patient study to the "Viewer Work Set"



Figure 475: img

• if the video is not edited, the save icon remains inactive 🗐, edit the video to activate it.

9.9.2 Lock view {#viewser_Lock view tool group}

If you want to browse or work with two or more series synchronously, you can use one of the offered "Lock view" tools:

lcon	Function	Description
	Lock view	Synchronous multi-panels series scrolling from the current image position in the series
í ک	Lock to separate groups	Section into groups of individual panels "Windows for displaying image data" for possible manual or automatic synchronization of series according to the level of data acquisition of the series, or by manual entry
۵	Automatic lock	Synchronous browsing of series in multiple panels with automatic finding of the corresponding image position in the series
	Lock windowing	When using the sync feature, the windowing adjustment will be applied to all sync series simultaneously

lcon	Function	Description
4	Lock transformations	When the sync function is used, the image adjustment will be applied to all the sync series simultaneously
Ð	Display panel synchronizability	Displaying the option to lock individual panels in the "Window for displaying image data"
	Synchronize viewer size between windows	Synchronize the size of displayed images in individual panels of "Windows for displaying image data" on a multi-monitor workstation
₿	Lock videos	Synchronous browsing of synchronously recorded videos
Ф	Mammography Image Alignment	Automatic zooming and placement of mammographic images, according to the ideal parameters of the unfolded panels in the "Image data display window"
)	Set nipple position	Manual adjustment of nipple position for automatic alignment of mammography images

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Figure 476: img

9.9.2.1 Lock view {#viewser_Lock view} The "Lock view" tool is used to synchronously scroll through series in multiple panels. By selecting this tool in the "Series actions" tool group, or in the "Display toolset", all series will be scrolled from the current frame position in the series. Synchronization will be applied to all series and reconstructed views in the same data acquisition plane.

Figure 477: img

9.9.2.2 Lock to separate groups {#viewser_Lock to separate groups} The "Lock to separate groups" tool located in the "Series actions" tool group or in the "Display toolset" is used to divide into groups the individual panels of the "Image data display windows" for possible manual or automatic synchronization of series according to the data acquisition plane of the series or by manual input.

In the event that you have placed and displayed series of studies in several "Windows for displaying image data" panels, by selecting the "Lock to separate groups" tool and then selecting automatic or

B

manual synchronization, they will be marked into individual groups according to the level of data collection:



Figure 478: img

If the "Display panel synchronizability" tool is selected (see below), it is possible to manually mark and unmark individual "Windows for displaying image data" panels for their synchronization in groups:



Figure 479: img

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Figure 480: img

9.9.2.3 Automatic lock {#viewser_Automatic lock} The "Automatic lock" tool located in the "Series action" tool group or in the "Display toolset" is used for synchronous scrolling through series in multiple panels. Automatic synchronization automatically finds the same patient positions according to the data in the DICOM file, aligns the series to the same location, and then locks their simultaneous scrolling. The synchronization will be applied to all series in the same data acquisition plane.

Figure 481: img

9.9.2.4 Lock windowing {#viewser_Lock windowing} The "Lock windowing" tool located in the "Series actions" tool group, or in the "Display toolset" can be applied to manual or automatic synchronization and allows the user to synchronously edit the image, in all synchronized series, using the "Windowing" tool, see paragraph "Display tools".

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Figure 482: img

9.9.2.5 Lock transformations {#viewser_Lock transformations} The "Lock transformations" tool located in the "Series actions" tool group, or in the "Display toolset" can be applied to manual or automatic synchronizations and allows the user to synchronously edit the image, in all synchronized series, using the tools in the "Display tools" and "Measurement tools" group, except for the "Windowing" tool, see paragraph "Edit image".

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Figure 483: img

9.9.2.6 Display panel synchronizability {#viewser_Display panel synchronizability} The "Display panel synchronizability" tool located in the "Series actions" tool group, or in the "Display toolset" is used for the possible locking of individual "Windows for displaying image data" panels and their subsequent manual or automatic synchronization. Selecting this tool will display a lock icon in the upper right corner of the "Windows for displaying image data" panel. Left click on the lock icon to lock this panel for sync

Figure 484: img

9.9.2.7 Synchronize viewer size between windows {#viewser_Synchronize viewer size between windows} The "Synchronize viewer size between windows" tool is used to automatically synchronize the size of the display of images in each panel of the "Image data display windows" on a multi-monitor

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workstation. This utility preserves the size of the display of images in the panels in case only one monitor is displaying the "Viewer Working Set" on a multi-monitor workstation and the panel sizes differ.

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Figure 485: img

9.9.2.8 Lock videos {#viewser_Lock videos} The "Lock videos" tool located in the "Series actions" tool group, or in the "Display toolset" is used for synchronous playback of synchronously recorded videos. This tool is thus applicable to videos obtained using "Capture" (the digitization tool of this product), which enables the so-called synchronous recording of multiple video inputs, when the videos are of the same length. If the videos were not taken by synchronous recording, you will be warned about this with the information "Not enough synchronizable videos from the same group".

9.9.2.9 Synchronisation of mammography images {#viewser_Synchronisation of mammography images} Advanced options for synchronizing and setting up the display of mammography images can be displayed by selecting the "Mammography toolset" tab:

۲	Synchronization V
	Pair •
1	Lock windowing Lock transformations
•	Ф
	Settings 🗸 🗸
	Discard inverted backgro
	Align nipples in CC view
	Align nipples in MLO view
	Different crop for MLO and
	Show nipple position

Figure 486: img

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Figure 487: img

9.9.2.10 Mammography Image Alignment {#viewser_Mammography Image Alignment} The tool "Aligning mammographic images" is used for automatic zooming and placement of mammo-graphic images, according to the ideal parameters of the expanded panels in the "Windows for displaying image data".

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Figure 488: img

9.9.2.11 Set nipple position {#viewser_Set nipple position} The "Set nipple position" tool is used to adjust the position of the nipple, which is the starting point for automatic alignment of mammography images in a pair.

To adjust the nipple position, perform the following steps: - Split the "Image data display window" into two panels by the action "Configure the panel layout in the current window/monitor" - in these panels, display the mammographic images in the same plane to be compared - Recommended: show the current nipple position - open the "Mammography toolset" - check the box "Show nipple positions" - select the "Set nipple position" tool with the desired mouse button in the "Lock view" tool group - click the selected mouse button with the tool on the real position of the nipple to move the yellow marking circle and if the "Align nipples CC/MLO" box in the "Mammography toolset" is checked, the position of the images will be automatically moved

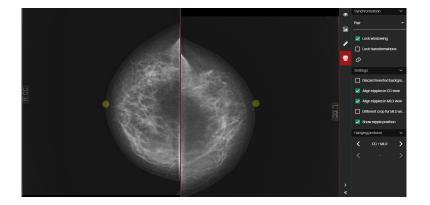


Figure 489: img

9.10 Volume operations, 3D visualization {#view3d_Volume operations and 3D visualization}

The "Volume operations, 3D visualization" tab is used for working with created volumes or fusions from a series of images. To create a volume, proceed according to the paragraph "Create volume", to create merged series, proceed according to the paragraph "Create fusion".

This tab contains the following groups of tools: - Volume tools - Volume projections



Figure 490: img

9.10.1 Volume tools {#view3d_Volume tools}

The "Volume Tools" tool group contains the following tools:

lcon	Function	Description
¢	3D Cursor	Localization of the point of interest in mutually perpendicular planes of the created series volume
(Ô)	Fusion active layer move	Manual adjustment of the overlay of layers in the created fused series of volume
Ç	Fusion active layer rotation	Manual rotation of the active layer in the created fused volume series
؈ٛ	Reset active layer transformation	Restoring the original position and rotation of the active layer in the created fused volume series
¢	Save active layer transformation	Store the active layer movements and rotations in the created fused volume series in REG format
20	Rotate volume	Free rotate of planes in the created series volume
	Interactive slab thickness change	Change thickness of the volume slice
Ē	Create new series with the actual orientation	Creation of a new reconstructed series, according to the selected direction of the MPR projection of the patient
<u>fo</u>	Select transfer function	Changing the display of the DVR volume, according to the default options

lcon	Function	Description
	Video from volume	Creating a video from the DVR volume by rotating it by 180° or 360°
MRK	Set curve of Curved MPR	Tool for marking individual points on the axis of the curved MPR
	Curved MPR Height	Specify the display size of the curved MPR volume
Œ	CMPR curve translation tool	Curved MPR curve shift tool
6	Delete all selected CMPR curve points	Curved MPR curve removal tool
*	Segmentation drawing	Draw a cutting area in the created segmentation
et.	Segmentation cutting	Undrawing the cutting area in the created segmentation

\Leftrightarrow

Figure 491: img

9.10.1.1 3D Cursor {#view3d_3D cursor} The "3D cursor" tool is used to locate a specific point of interest in mutually perpendicular planes of the created volume of the series. To locate a point, select this tool from the "Volume Tools" group of tools with the selected mouse button and then apply to the point of interest in the plane of the displayed series.

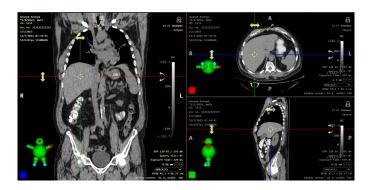


Figure 492: img

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Figure 493: img

9.10.1.2 Fusion active layer move {#view3d_Fusion active layer move} The "Fusion active layer move" tool is used for manual adjustment of layer overlap in created fused volume series. To adjust the overlay of layers, follow these steps: - Create and display the merged layers according to the paragraph "Create fusion" - select the "Fusion active layer move" tool in the "Volume Tools" tool group with the selected mouse button - mark the active layer, see paragraph "Create fusion" - hover the mouse over the view of the fusion plane - press and hold the selected mouse button with the tool, then move the active layer to the position of interest by moving the mouse

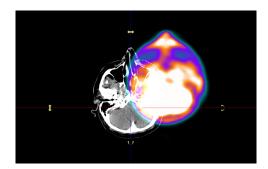


Figure 494: img

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Figure 495: img

9.10.1.3 Fusion active layer rotation {#view3d_Fusion active layer rotation} The "Fusion Active Layer Rotation" tool is used to manually rotate the active layer in the created fused volume series. To adjust the layer overlay, follow these steps: - Create and display the fused layers according to the paragraph "Create fusion". - Select the "Fusion Active Layer Rotation" tool in the "Volume Tools" tool group with the desired mouse button - select the active layer, see paragraph "Create fusion" - hover the mouse over the desired view of the fusion plane - press and hold the selected mouse button with the tool, then move the mouse to rotate the active layer to the desired position

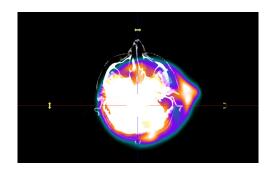


Figure 496: img

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Figure 497: img

9.10.1.4 Reset active layer transformation {#view3d_Reset active layer transformation} The "Reset active layer transformations" tool is used to restore the original position and rotation of the active layer in the created fused volume series. Select the "Reset Active Layer Transformations" action to reset to the original state of the overlay layers, the rotated/offset fusion layers will be returned to their original state before the transformations.

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Figure 498: img

9.10.1.5 Save active layer transformation {#view3d_Save active layer transformation} The "Save Active Layer Transformations" tool is used to save the active layer movement and rotation in the created fused volume series to a newly created series in REG format. To save the performed active layer fusion transformation, proceed as follows: - Create and display the fused layers according to the paragraph "Create fusion". - Perform the required image transformations (move, rotate) - Select the "Save active layer transformations" tool to call up the table:



Figure 499: img

- Select "Target" (Archive, User Data, or Station Data)
- If you have selected the archive save target, select the desired "Archive"
- If necessary, change the name of the active layer transformation to be saved using the "Series description" text box
- Select "SAVE" to create a new series in REG format with the saved position of the active fusion layer, or select "CANCEL" to cancel the performed work
- If the "SAVE" action is selected in the table described above, a new series in the "REG" format will be created and added to the "Viewer working set":



Figure 500: img

Display the saved transformation of the active layer

In case you want to view the saved fusion layer transformation, do the following: - Create and display the merged layers according to the paragraph "Create fusion". - Right-click on the thumbnail preview of the saved transformation state in "REG" format to call up the context menu in which you select the "Activate Spatial Registration" tool:

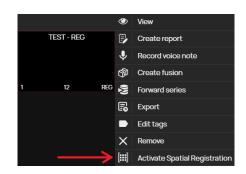


Figure 501: img

• Select this action to recall the saved transformation state of the fusion layer, e.g.:

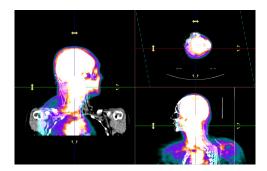


Figure 502: img

• In case you want to return to the original state, the saved fusion layer transformation can be deactivated with the "Deactivate spatial registration" action in a similar way to the activation from the context menu. The original state will then look like this:

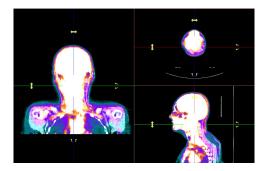


Figure 503: img

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Figure 504: img

9.10.1.6 Rotate volume {#view3d_Rotate volume} The "Rotate volume" tool is used to rotate planes in the created volume. To rotate a plane, proceed as follows:

- Create and view series volume, see "Create volume"
- select the "Rotate volume" tool in the "Volume Tools" tool group with the selected mouse button
- hover the mouse over the view of the volume plane
- press and hold the selected mouse button with the tool, move the mouse to rotate the created volume

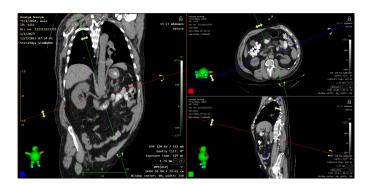


Figure 505: img

Figure 506: img

9.10.1.7 Interactive slab thickness change {#view3d_Interactive slab thickness change} The

"Interactive slab thickness change" tool is used to change the thickness of the displayed slice. To change the slice thickness, proceed as follows:

- Create and view series volume, see paragraph "Create volume"
- select the tool "Interactive slab thickness change" in the tools group "Volume Tools" with the selected mouse button
- hover the mouse over the view of the volume plane
- press and hold the selected mouse button with the tool, move the mouse up and down to change the slice thickness to the required size
- the current size of the slice thickness is displayed in the OSD label of the displayed image, the location of the OSD label is configured by the user, or is set by default in the lower right corner of the display window, e.g.: 2,75 m
- if you have enabled the "Show slice thickness lines" function, the current slice width size will also be visually displayed using the dashed lines of the volume axes:

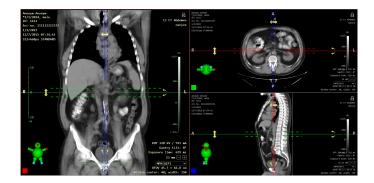


Figure 507: img

9.10.1.7.1 Change slice thickness and volume reconstruction {#view3d_Change slab thickness and volume reconstruction} Changing the slice thickness and volume reconstruction can be changed in the "Volume toolset" if you have created and displayed a volume series:



Figure 508: img

The "Volume toolset" contains the following functions:

Volume tools

The "Volume Tools" tab contains the selected tools of the "Volume Operations, 3D visualization" tab: - 3D cursor - Rotate volume - Interactive slab thickness change - Create new series with the actual orientation - Video from volume

Volume projections

The "Volume projections" tab contains the individual planes and the 3D volume for reconstructing the actively selected window for display.

Projections and slab thickness editor

To change the slice thickness and reconstruct the volume, proceed as follows: - Create and display the volume of the series, see "Create volume". - Open the "Volume toolset" and go to the "Projections and slab thickness editor" tab - select the desired reconstruction using the drop-down menu:



Figure 509: img

• select the desired width of the slice by typing the values into the text box or using the - / + tool



Figure 510: img

Figure 511: img

9.10.1.7.2 Quick switching of slab thickness {#view3d_Quick switching of slab thickness} To quickly change the slice thickness of the displayed series volume, you can use the actions directly from the OSD label location:

2,75 mm (--) (+)

To quickly change the slice thickness, proceed as follows: - Create and view the volume of a series, see "Create volume" - The current size of the slice thickness is displayed in the OSD label of the displayed image, the location of the OSD label is user-configured or set by default in the lower right corner of the display window:



Figure 512: img

using the - / + function you can decrease / increase the slice thickness of the currently selected
volume plane of the image series in the "Image data display window", this function can also be
used by clicking the mouse on a numeric value and moving it sideways to adjust the thickness

Warning: this function can be hidden, to display it you need to set this function using the "Profile editor" tool located in the "Viewer settings" tab.

9.10.1.7.3 Quick volume reconstruction switching {#view3d_Quick volume reconstruction switching} To quickly change the volume reconstruction of the displayed series volume, you can use the actions directly from the OSD label location under the "Change slice thickness" tool:



Figure 513: img

To change the volume reconstruction, click the drop-down menu and select the requested reconstruction. This reconstruction will be applied.

Warning: this function can be hidden, to display it you need to set this function using the "Profile editor" tool located in the "Viewer settings" tab.

C

Figure 514: img

9.10.1.8 Create new series with the actual orientation {#view3d_Create new series with actual orientation} The "Create new series with the actual orientation" tool is used to create a new reconstructed series, according to the selected direction of the patient's MPR projection and the selected slice thickness. To create a new reconstructed series, proceed as follows: - Create and view series volume, see paragraph "Create volume" - Drag the created volume to the selected "Image data display windows" panel and select the desired reconstruction plane from the "Display" tab - Select the "Create new series with current orientation" tool in the "Volume tools" tool group to call up the "Create new series with current orientation" tab:

Create new series with actual orientation			
Slice thickness Custom slice thickness	Target Archive		
Custom slice thickness 2.5 mm	Archive DPGW (DPGW@127.0.0.1:5380)		
Slice count estimation 267	Series description natyve - orientationChanged		
	CANCEL		

Figure 515: img

• In this tab, use the drop-down menu to select the desired slice thickness:



Figure 516: img

• If "Custom slice thickness" is selected, enter the desired thickness in mm in the "Custom slice thickness" text field:



Figure 517: img

• The "Slice count estimate" field in the table serves as an informative message about the number of images in the newly created series.

- Select the save destination of the reconstructed series from the "Target" drop-down menu, if you have set up multiple archives, select the desired archive from the "Archive" drop-down menu and enter the series name in the "Series description" text box
- Selecting the action "Cancel" cancels the action performed, in case of selecting "Save" you create a new reconstructed series, the system will notify you about the start of the series creation
- the newly created series will be added to the "Viewer working set"

*f*_{3D}

Figure 518: img

9.10.1.9 Select transfer function {#view3d_Select transfer function} The "Select transfer function" tool is used to change the DVR volume display, according to the default options. To change the DVR view, proceed as follows: - Create and view series volume, see paragraph "Create volume" - Drag the created volume to the selected view panel and select "Direct volume rendering (DVR)" from the "Display" tab Select the "Select transfer function" tool in the "Volume tools" tools group to display the "Volume toolset":



Figure 519: img

- go to the "Transfer function" tab
- use the "Selected transfer function" drop-down menu to select the display area:



Figure 520: img

• use the drop-down menu "Transfer Function Windowing" to select a change of view:



Figure 521: img

Changing the DVR display will be immediately visible after selecting the display change from the drop-down menu.

Figure 522: img

9.10.1.10 Video from volume {#view3d_Video from volume} The "Video from volume" tool is used to create and then export video from the created DVR volume. To create a video, follow these steps:

- Create and view series volume, see paragraph "Create volume"
- Drag the created volume to the selected view panel and select "Direct volume rendering (DVR)" from the "Display" table
- select the "Video from volume" tool in the "Volume tools" tools group to display the "Video from volume" table:

Video from volume X						
File name: Anonym_Anonym_	_1111_C	Number of frames per second: 10				
Rotation 180°		Video length (s) 5				
Rotate around Vertical axis		Image format: MPEG				
Keep slab thickness						
	CAN	CEL EXPORT				

Figure 523: img

• fill in the required data in the displayed table:

Field	Description	Field	Description
Filenan	New Yorking the exported video	Number of frames per sec- ond	r Selection of playback speed
Rotatio	n Option to choose rotation by 180° or 360°	Video length	Choice of video duration in seconds
	Option to choose rotation horizontally / vertically	Image For- mat	Choice of video export format in .MPEG and .AVI
Keep slab thick- ness	Keeps the currently selected slice thickness width		

• the action "CANCEL" cancels the performed action, the action "CONFIRM" starts the process of exporting the video to the local PC user, you will be informed about the start of the export:



Figure 524: img

• then the table "Video from volume" will be displayed with a queue of export requests, and if the video is created successfully, the download to the user's local PC will start.

9.10.1.11 Curved MPR toolset {#view3d_Curved MPR toolset} The Curved Projection toolset contains the CMPR and SMPR projection views and their associated tools.



Figure 525: img

The "Curved Projections" tab contains: - Curved MPR 🕮 - display of the created curve in volume in four panels of the "Image data display windows" - Straightened MPR 🕮 - display of the straightened projection according to the created curve in the volume in four panels of "Image data display windows"

The "Tools" tab contains:

lcon	Function	Description
(MPR	Set curve of Curved MPR	Tool for marking individual points on the axis of the curved MPR
Œ	CMPR curve translation tool	Curved MPR curve shift tool
	Interactive slab thickness change	Change thickness of the volume slice
	Stacking	Browse through the series with the selected mouse button
¢	3D Cursor	Localization of the point of interest in mutually perpendicular planes of the created series volume

lcon	Function	Description
6	Delete all selected CMPR curve points	Curved MPR curve removal tool
	Create new series with the actual orientation	Creation of a new reconstructed series of curved projection (to create a new series with the created curved projection, follow the "Create a new series with the actual orientation" described above)

The "Curved MPR" tab contains a tool for changing the size of the display area around a curved MPR curve by typing values into the text field, moving the values with the mouse, or using the - and + symbols.



Figure 526: img

The "Slab Thickness" tab contains a tool for changing the width of the displayed slice of the created curved MPR projection by typing values into the text field, by moving the values with the mouse, or by using the - and + symbols.



Figure 527: img

The "Toggles" tab contains the "Show points outside rendered plane" tool, which allows you to display the plotted points of a curved projection created by the user in all planes throughout the entire view volume.



Figure 528: img

If this function is active, the lines between points outside the current view area will be dashed, e.g.:

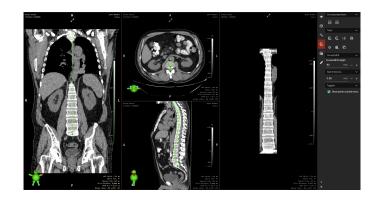


Figure 529: img

Figure 530: img

9.10.1.12 Set curve of Curved MPR {#view3d_Set curve of Curved MPR} The "Set curve of Curved MPR" tool is used to create and display a curved surface created from the volume of the selected series, allowing the user to display, for example, the stretched path of a blood vessel, spine, etc. To determine the curved MPR curve, proceed as follows:

- Create and view the volume of a series, see "Create volume"
- select the "Curved MPR" view in the "Volume projections" tool group (can also be displayed by double-clicking on the series thumbnail with the created volume in the viewer working set)
- to make your work easier, maximize the panel by displaying the desired plane across the entire window with the concated at the top of the panel
- select the "Set curve of Curved MPR" tool in the "Volume tools" tool group with the selected mouse button or from the "Curved MPR toolset"
- scroll through the series to locate the location of the first point and then click the selected tool button to place the point
- as you go through the series, click to place the other points
- individual points can be moved by dragging with the mouse, or by hovering the mouse over the selected point and pressing the DELETE key to delete the point, points can be added to the created curve by hovering the mouse between two created points, when the mouse pointer changes to the + symbol and clicking the left mouse button to add a new point
- to edit the curve in other planes, always find the approximate slice to the point you want to move in the desired plane and drag the mouse to move the point
- if you have placed all the desired points, a curved MPR curve will be displayed, e.g.

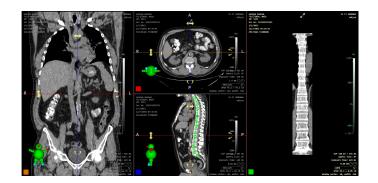


Figure 531: img

To create an "Straightened MPR" follow a similar process, the result in this case is a transversal projection that is straightened compared to the curve of the curved MPR, i.e. it is always perpendicular to the curve, e.g:

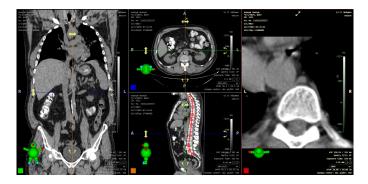


Figure 532: img

Figure 533: img

9.10.1.13 Curved MPR Height {#view3d_Curved MPR height} The "Curved MPR Height" tool is used to specify the size of the curved MPR curve display. To determine the size, proceed as follows: - create a curved MPR curve according to the procedure of the previous paragraph - select the "Curved MPR Height" tool in the "Volume Tools" tool group, this calls up the "Curved MPR Toolset" sidebar with the "Curved MPR" tab:



Figure 534: img

 enter a size value in mm in the text box to display the size range of surrounding structures from specific points on the curved MPR curve, the size will be adjusted automatically after entering the value. This value can also be changed by moving the values with the mouse, or by using the and + signs.

Figure 535: img

9.10.1.14 CMPR curve translation tool {#view3d_CMPR curve translation tool} The "CMPR curve translation tool" tool is used to offset all points of the created curved MPR curve simultaneously. To move the curve, select this tool with the selected mouse button and then the curve can be moved by dragging.

0

Œ

Figure 536: img

9.10.1.15 Delete all selected CMPR curve points {#view3d_Delete all selected CMPR curve points} The "Delete all selected CMPR curve points" tool is used to remove a curved MPR curve. Selecting this tool removes the created curve including all its points.

9.10.1.16 Segmentation toolset {#view3d_Segmentation toolset} The "Segmentation toolset" contains tools for cropping objects in the created volume of a series of frames.



Figure 537: img

The "Tools" tab of the "Segmentation toolset" contains: - 🖄 Segmentation Drawing - tool used to draw an area in the created segmentation - 🔯 Segmentation cutting - tool used to delete part or all of the created segmentation

The "Segmentation" tab of the "Segmentation toolset" contains: -
Create Segmentation - a tool to create a new segmentation layer -
Hide/Show All - tool to hide/show all applied segmentation layers -
Unload Segmentations - tool to remove all created segmentations

To create a new segmentation, do the following:

- 1. Create and display a volume from the selected series of images (see "Create volume" for more information).
- 2. Open the "Segmentation toolset"
- 3. In the "Segmentation" tab, select the "Create Segmentation" button + to call up the "Segmentation" table



Figure 538: img

Name the new segmentation by inserting it into the "Name" text field. If the "Use threshold" field is checked, this table will be expanded to select a threshold value (all points of the created volume containing optical density according to the selected threshold value will be deleted):

Segmentation		×
Name		
This field is required!		
Vse threshold		
Threshold		
		180
Deviation		
-0		50
CANCEL	CREA	TE

Figure 539: img

The "Cancel" action cancels the performed task, the "CREATE" action will add a new segmentation in the "Segmentation toolset" on the "Segmentation" tab:



Figure 540: img

The created segmentation contains: - "Selected" button 🖸 is used to select a segmentation, in case of creating multiple segmentations the other icons are empty 🖸 - The "Hide/Show" button 🔮 is used to hide or show the segmentation - Name of the created segmentation - The "Delete segmentation" button 🖬 is used to remove the created segmentation, after selecting this action you will call the table for confirmation:

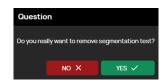


Figure 541: img

- Repeat step 3 of this paragraph to add another segmentation
- 4. Draw the segmentation contours in the selected volume area using the "Draw segmentation" or "Segmentation cutting" tools by selecting from the "Volume tools" tab, or by selecting from the "Segmentation toolset" of the "Tools" tab.

۴,

The cut out areas will then not be displayed in other planes or 3D views.

Figure 542: img

9.10.1.17 Segmentation drawing {#view3d_Segmentation drawing} The "Segmentation drawing" tool is used to draw the area in the created segmentation. To draw an area, proceed as follows: - Create an area segmentation from the selected volume of the image series, see more in "Segmentation toolset" - select the "Segmentation drawing" tool in the "Volume tools" tool group, or by selecting it from the "Segmentation toolset" of the "Tools" tab with the selected mouse button - move the mouse to the starting point in the selected plane of the volume image - by clicking, or by clicking and holding the selected mouse button with the tool, draw the surface of the area of interest - by completing the drawing of the segmentation area, the volume will be recalculated:



Figure 543: img

• the result is cutting out part of the volume:

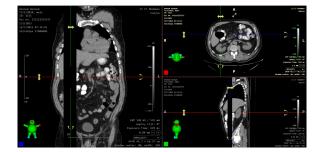


Figure 544: img

°te

Figure 545: img

9.10.1.18 Segmentation cutting {#view3d_Segmentation cutting} The "Segmentation cutting" tool is used to delete part or all of the created segmentation. To remove segmentation, proceed as follows:

- Display the volume with the segmentation created that you want to edit
- select the "Segmentation cutting" tool in the "Volume Tools" tool group, or by selecting it from the "Segmentation toolset" of the "Tools" tab with the selected mouse button
- move the mouse to the starting point in the selected plane of the volume image
- by clicking, or by clicking and holding the selected mouse button with the tool, draw the area you want to cut out
- The result is the cut out part or all of the created segmentation.

9.11 Volume projections {#view3d_Volume projections}

The group of tools "Volume projections" is used to display the projections and planes of the created volume. For more on the possibility of creating volumes, see the paragraph "Create volume". Measurement and post-processing tools of the DICOM viewer can be used in the created volumes, e.g. the "Windowing" tool, 3D volume structures, or MIP, etc. can be added or removed.

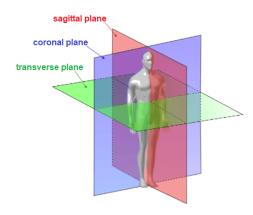


Figure 546: img

DPGW allows these projections/planes to be displayed in volume:

lcon	Projection/plane
MR	Multiplanar reconstruction (MPR)
C	MPR Frontal view
S	MPR Sagittal view
Τ	MPR Transverse view
MP	Max. intensity projection (3 views)
MIP	Max. intensity projection (MIP)
MIN	Min. Intensity projection (MinIP)
FMIP	Fading max. Intensity Projection (Fade MIP)
3D	3D projection
DVR	Direct volume rendering (DVR)
CMPR	Curved MPR
SMPR	Straightened MPR

9.11.1 Multiplanar reconstruction (MPR) {#view3d_Multiplanar reconstruction MPR}

MPR

Figure 547: img

The "Multiplanar reconstruction (MPR)" tool is used to display the reconstructed planes in three panels "Window for displaying image data":

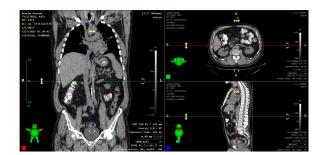


Figure 548: img

9.11.2 MPR Frontal view {#view3d_MPR Frontal view}

Figure 549: img

The "MPR Frontal view" tool is used to display the reconstructed coronal plane in one "Window for displaying image data" panel:

C



Figure 550: img

Figure 551: img

9.11.2.1 MPR Sagittal view {#view3d_MPR Sagittal view} The "MPR Sagittal view" tool is used to display the reconstructed sagittal plane in one "Window for displaying image data" panel:

S



Figure 552: img



Figure 553: img

9.11.2.2 MPR Transverse view {#view3d_MPR Transverse view} The "MPR Transverse view" tool is used to display the reconstructed transverse plane in one "Window for displaying image data" panel:



Figure 554: img



Figure 555: img

9.11.2.3 Max. intensity projection (3 views) {#view3d_Max intensity projection 3 views} The "Max. Intensity Projection (3 Views)" tool serves as a 3D rendering technique that examines each voxel and selects the one with the highest intensity. This is subsequently used in the displayed volume. The others are ignored. (90% of the acquired information will be lost). With this method, calcifications and surgical staples can be better differentiated. Another advantage can be considered greater visibility of bright objects. Disadvantages include a lack of depth information, high-intensity materials (calcifications) can obscure the desired organs. By choosing this tool, the reconstructed MIP planes will be displayed in three panels "Window for displaying image data":

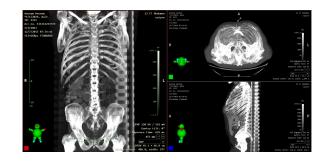


Figure 556: img

Figure 557: img

9.11.2.4 Max. intensity projection (MIP) {#view3d_Max intensity projection MIP} The "Max. Intensity Projection (MIP)" tool is similar to the previous tool, but displays the reconstructed MIP in the currently selected plane in one "Window for displaying image data" panel:

MIP



MIN

Figure 558: img

Figure 559: img

9.11.2.5 Min. Intensity projection (MinIP) {#view3d_Min Intensity projection MinIP} The "Min. Intensity projection (MinIP)" tool is used to reconstruct the projection with the display of only the area with the lowest attenuation coefficient in the given area. It is used, for example, in the description of the bile ducts and pancreatic duct, diagnosis of lung diseases. Generally for very weak signal areas. The tool displays the reconstructed MinIP in the currently selected plane in one panel "Window for displaying image data":



FMIP

Figure 560: img

Figure 561: img

9.11.2.6 Fading max. Intensity Projection (Fade MIP) {#view3d_Fading max Intensity Projection FMIP} The "Fading max. Intensity Projection (Fade MIP)" tool differs from the MIP tool, which displays the highest density from the volume in each projection beam, in that a linear attenuation is also applied to the displayed density according to the distance from the beginning of the displayed section of the volume (slab). The tool displays the reconstructed FMIP in the currently selected plane in one panel "Window for displaying image data":



Figure 562: img

Figure 563: img

9.11.2.7 3D projection {#view3d_3D projection} The "3D projection" tool is used to reconstruct a three-dimensional image of the created volume. The tool displays the 3D reconstruction in the currently selected plane in one panel "Window for displaying image data":

3D



Figure 564: img

Figure 565: img

9.11.2.8 Direct volume rendering (DVR) {#view3d_Direct volume rendering DVR} The "Direct volume rendering (DVR)" tool is used for the complete reconstruction of a three-dimensional image of the created volume, where color intensity = object density. The tool will display the DVR in the currently selected plane in one panel "Window for displaying image data":

DVR



Figure 566: img

Figure 567: img

9.11.2.9 Curved MPR {#view3d_Curved MPR CMPR} The "Curved MPR" tool is used to display the created curve in volume in the four panels of the "Image data display windows", to create a curve, see more in the paragraph "Curved MPR toolset":

	And	O 17 Million whyter whyter 10 Million 10 Million	1	E-3 Advant Miya
- GT		2000	200	
				-
			A DECK	·
			A DESCRIPTION OF	
	?	and and a second	an =40	A DECAMPANE

Figure 568: img

Figure 569: img

9.11.2.10 Straightened MPR {#view3d_Straightened MPR SMPR} The "Straightened MPR" tool is used to display the straightened projection according to the created curve in the volume in four panels of the "Image data display windows". The created transversal projection is aligned against the created curve of the curved MPR, i.e. it is always perpendicular to this curve. To create a curve, see more in "Curved MPR toolset":

SMPR

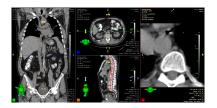


Figure 570: img

9.12 Online consultation {#dwshare_Online consultation}

Action DWShare enables online consultations. This is the online transfer of the image and mouse cursor between connected users.



Figure 571: img

The DWshare tab contains the following tools:

lcon	Function	Description
*	Share cursor	switch to share / unshare cursor position during remote session
(X)	Share actions	switch to share / unshare actions performed during a remote session
Ľ	Create session	creating a new session, defining the name and users who can connect

lcon	Function	Description
•)	Connect	if the user wants to join an existing session to which he is invited, he uses this option
***	Manage session users	option to manage users with session access. Users can be added, removed or external users can be added
Ċ	Hang up	if connected to a session, it is possible to end this session with this tool

9.12.1 Setting up a new online consultation {#dwshare_Setting up a new online consultation}

6

The master user creates a session using the "Create session" tool.

Figure 572: img

This will display the "Create new session" table, in the "Description" text field, choose the name of the online consultation and choose the "Create" action. The "Cancel" action cancels the action being performed.

Create new session	×
Description* Test DWshare	
CANCEL	CREATE

Figure 573: img

By selecting the "Create" action, you will display the following table "Manage users in session" for adding users and managing them (after the session is established, this table can be displayed with the "Manage session users" tool 🖾 located in the "DWshare" tool tab):

User guide

Dicompass Gateway

		Manage us	ers in session		X
INTERNAL EXTERN					
= Enter the user login or username test					
Available users			Selected users		
Login name	User name	Add	Login name	User name	Remove
test1	TEST1	•	Test4	TEST4	•
test2	TEST2	Ð			
test3	TEST3	Ð			
CANCEL				AD	DD EXTERNAL CLIENT SAVE

Figure 574: img

This table contains two tabs "Internal" and "External". Internal users are users who are connected to the same server within the internal network. External users are users who are connecting from an external environment, outside of the medical facility.

Internal users can be searched using the "Enter the user login or username" text field in the header of the "Internal" tab. You can find the selected user in the "Available users" field and add it with the icon • in the "Add" column. The user added to the session will appear in the "Selected users" field and you can remove it if necessary using the • icon.

External users can be added using the "Add external user" action ADD EXTERNAL CLEART , which displays the "Add external client" table:

User name*	
PIN*	

Figure 575: img

In this table, enter the user name "User name" and password "PIN" in the text fields, then choose the "Save" action, the "Cancel" action cancels the action being performed.

After creating a new external user, it will be displayed in the "External" tab, in the "Manage users in session" table:

Dicompass Gateway

	Manage users in session		X
INTERNAL EXTERNAL			
Selected external clients			
User name	Share link	PIN	Remove
Test	https://10.20.0.12:8444/viewer/dwshare?token=DSQN4TFXYM	➡ 1234	•
CANCEL		ADD EXTERNAL	CLIENT SAVE

Figure 576: img

Then send the "Share Link" to the external user (for copying you can use the icon ^(w)) and the selected password. To maintain security, we recommend choosing a different way of communicating link and password information. This will create one-time access for an external user, without access to search in the PACS archive. To remove a user, use the icon •.

After finishing adding users, choose the "Save" action, this will start a new online consultation. "Cancel" action cancels the action being performed.

In case of user login in the created online consultation, you will be informed about the number of logined users by the number displayed next to the icon of the "DWshare" tab 🗟.

To connected users, the shared examination study will be displayed by the master of the consultation, including the mouse cursor and work on it:

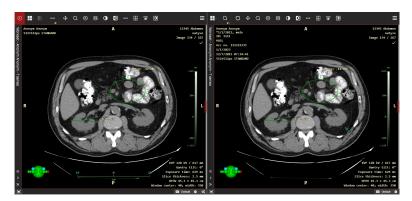


Figure 577: img

9.12.2 Login to the online consultation as an internal user {#dwshare_Login to the online consultation as an internal user}

You have two options for logging in to the online consultation as an internal user, if you are working in DICOM viewer and you are added to the online consultation, the "Incoming request for remote consultation" table will be diaplayed:

Figure 578: img

If you choose "Accept remote consultation", the examination study of the master remote consultation and the shared work on it will be displayed.

If you choose "Reject remote consultation" or you are not working on image documentation in DICOM viewer, you can connect to the created online consultation to which you are added using the "Connect" tool **D**. Select the tool to display the "Join the session" table:

∓ Enter the user login or username			
_ogin name ▲	User name	Description	Connect
admin	ADMINISTRATOR	Test DWshare	7

Figure 579: img

Select online consultation and choose the "Connect" action, the examination study of the master remote consultation and the shared work on it will be displayed.

9.12.3 Login to the online consultation as an external user {#dwshare_Login to the online consultation as an external user}

To log in to the online consultation as an external user, enter the obtained url of the "Share link" into the Internet browser, this will display the login page of the Dicompass Gateway DICOM viewer:



Figure 580: img

Enter the "PIN" and select the "Sign in" action. You will be connected to the online consultation:



Figure 581: img

As an external user, you can use the following tools for DWshare: - Share cursor - Share actions - Hang up

9.13 DICOM viewer settings {#viewsettings_DICOM viewer settings}

The "DICOM viewer settings" tab contains general tools for managing DICOM viewer settings and preferences.

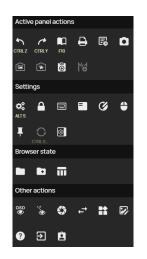


Figure 582: img

This tab contains the following groups of tools:

- Active panel actions
- Settings
- Browser state
- Other actions

9.13.1 Active panel actions {#viewsettings_Active panel actions}

The group of tools "Active panel actions" contains the following tools:

lcon	Function	Description
5	Previous Action	allows you to undo the action taken
<i>(</i> *	Next action	allows you to perform the returned action
	DICOM tags viewer	viewer of DICOM tags that are stored in the displayed image
Ð	Print	enables you to print the displayed study, see more chapter "Viewer Working set" paragraph "Print"

lcon	Function	Description
ß	Export	allows you to export the displayed study, see more chapter "Viewer working set" paragraph "Export"
	Secondary capture	allows the user to archive the currently displayed image without editing
	Capture active viewer	allows the user to archive the currently displayed image of the active panel, including its measurements and OSD labels
	Capture desktop	allows the user to archive the currently displayed screenshots of the desktop, including its layout
6	Capture active viewer to clipboard	select this tool to export a screensho of the selected panel in the "Image data display window" to the local storage in .png format
16	Export ECG	allows the user to export ECG examination results

Figure 583: img

9.13.1.1 DICOM tags viewer {#viewsettings_DICOM tags viewer} The "DICOM tag viewer" tool is used to display DICOM tag information about the selected image/series of the selected study. By selecting this tool, you will display the "DICOM tags viewer" table:

Dicompass Gateway

litter						
	Group	Group, Tag •	Name	Value		
	All tags	(0002,0010)	Transfer Syntax UID	ImplicitVRLittleEndian (1.2.8	0.10008.1.2)	
0002	Meta data	(0008,0005)	Specific Character Set	ISO_IR 100		
0008	Study	(0008,0008)	Image Type	ORIGINAL, PRIMARY, LOCALI	ER	
0009	Private group	(0008,0012)	Instance Creation Date	12/7/2015		
0010	Patient	(0008,0013)	Instance Creation Time	07:32:26		
0018	Acquisition	(0008,0016)	SOPClass UID	CTImageStorage (1.2.840.10	008.5.1.4.11.2)	
0019	Private group	(0008,0018)	SOPInstance UID	1.2.840.113619.2.55.3.427104	5733.996.1449464144.5	99.1
0020	Relations	(0008,0020)	Study Date	1/1/2023		
0021	Private group	(0008,0021)	Series Date	1/1/2023		
0023	Private group	(0008,0022)	Acquisition Date	12/7/2015		
0027	Private group	(0008,0023)	Content Date	12/7/2015		
0028	Image represent	(0008,0030)	Study Time	10:00:00		
0032	Group	(0008,0031)	Series Time	10:00:00		
0040	Group	(0008,0032)	Acquisition Time	07:32:17		
0043	Private aroup	(0008 0033)	Content Time	07:32:26		
					TEXT DUMP	CLOSE X

Figure 584: img

To speed up your search, you can enter the desired parameter in the "Filter" field, e.g:

DICOM	DICOM tags viewer						
•							
Filter Patient Na							
ID •	Group	Group, Tag •	Name	Value			
	All tags	(0010,0010)	Patient Name	Anonym Anonym			
0002	Meta data						
0008	Study						
0009	Private group						
0010	Patient						
0018	Acquisition						
0019	Private group						
0020	Relations						
0021	Private group						
0023	Private group						
0027	Private group						
0028	Image represent						
0032	Group						
0040	Group						
0043	Private aroun						
						CLOSE X	

Figure 585: img

The left column shows groups of DICOM tags, and the right column shows individual DICOM tags of the selected group.

The "Text dump" action is used to display all DICOM tags in text form:

CLOSE X
1

Figure 586: img

The "Close" action allows you to close the "DICOM tags viewer" table.

If you open a sequence with information it is possible to go back to all data using the "*" • , located above the search sequence.

۵

Figure 587: img

9.13.1.2 Secondary capture {#viewsettings_Secondary capture} The "Secondary capture" tool is used to save the currently displayed image in the marked "Image data display window" panel without editing and measuring in the image. By selecting this tool, the "Secondary capture" table is called up:

Secondary capture	×		
Target Archive			
Archive			
Archive			
DPGW (DPGW@127.0.0.1:5380)			
Series description			
scout - secondary			
CANCEL	SAVE		

Figure 588: img

This table contains the following fields:

- Target the option to choose the destination of archiving the image to "Archive", "User data" or "Station data"
- Archive in the case of configured multiple PACS archives, you can choose selected archive for saving, the field is active if you have selected "Target" "Archive"
- Series description allows you to name the saved image
- The "Save" action archives the selected image, the "Cancel" action cancels the performed action

If you have saved the selected image, it will be added to the "Viewer Working Set" and saved to the selected "Target":



Figure 589: img

Figure 590: img

9.13.1.3 Capture active viewer {#viewsettings_Capture active viewer} The "Capture active viewer" tool is used to save the currently displayed image in the marked window "Image data display window", including its measurements and OSD labels. To save, proceed similarly according to the paragraph "Secondary capture".

*

Figure 591: img

9.13.1.4 Capture desktop {#viewsettings_Capture desktop} The "Capture desktop" tool is used to save the currently displayed images in all "Image data display window" panels, including measurements and OSD labels. To save, proceed similarly according to the paragraph "Secondary capture".

ĕ

Figure 592: img

9.13.1.5 Capture active viewer to clipboard {#viewsettings_Capture active viewer to clipboard} The "Copy active viewer to clipboard" tool exports a screenshot of the selected panel in the "Image data display window" to the local storage in the ".png" format. This serves as a quick way to save the active image to the user's PC.

6

Figure 593: img

9.13.1.6 Export ECG {#viewsettings_Epxort ECG} The "Export ECG" tool is used to export the currently displayed ECG series in the active panel "Image data display window" to a text file with ECG examination parameters. Select this tool to start automatic file download.

9.13.2 Settings

lcon	Function	Description
00	Global configuration	Open system configuration Dicompass Gateway
	Change password	Changing the password of the currently logged in user
	Shortcuts	Displaying or defining new keyboard shortcuts
1	Profile editor	Editor for display and location of OSD labels in panels "Image data display window"
Ø	Preset windows management	Defining windowing profiles
0	Tool Selection	Adding and editing saved DICOM viewer configurations
IJ	Position of pinned actions	Determining the display order of tools in "User configurable toolbar" and "Context menu"
0	Restart windows layout	Reloading the displayed images/series in the

The "Settings" tools group includes the following tools:

Working set configuration Allows configuration of the "Viewer Working Set" view containing thumbnails of series

"Image data display window"

\mathbf{Q}_{0}^{0}

Figure 594: img

1

9.13.2.1 Global configuration {#viewsettings_Global configuration} The "Global configuration" tool is used to display the configuration of the Dicompass Gateway system. See more chapter "Settings"

Figure 595: img

9.13.2.2 Change password {#viewsettings_Change password} The "Change password" tool is used to change the password of the currently logged-in user. To change the password, select the "Change password" tool, which display the "Change password" table:

Change password	×
Old password*	
New password*	
Check for a new password*	
CANCEL	SAVE

Figure 596: img

To change your password, follow these steps:

- enter the currently selected password in the "Old password" text field
- enter new password in the "New password" text field
- in the "Check for a new password" text field, re-enter the choosed new password to check it
- use the "Save" action to change the password, the "Cancel" action cancels the action

You will be informed about a successful password change with information in the DICOM viewer footer:



.....

Figure 598: img

9.13.2.3 Shortcuts The "Shortcuts" tool is used for displaying or defining new Dicompass Gateway viewer shortcuts. Selecting this tool will call up the "Shortcuts config" table:

Dicompass Gateway

		Shortcuts config		>
Filter by module/gr	oup/action name	Filter by hotkey		
Module •	Group	Name	Hotkey	
Active panel	Image	3D cursor		×
Active panel	Image	Coordinates		×
Active panel	Image	Default	L	×
Active panel	Image	Flip horizontally		×
Active panel	Image	Flip vertically		×
Active panel	Image	Free rotation		×
Active panel	Image	Hide 3D cursor		×
Active panel	Image	Image zoom in	PERIOD	×
Active panel	Image	Image zoom out	COMMA	×
CLOSE		RESET SHORTCUTS TO DEFAULT		SAVE

Figure 599: img

To find a specific tool for assigning or modifying a new keyboard shortcut, use the "Filter by module/group/action name" or "Filter by hotkey" text fields.

The individual tools in the table are arranged in columns with the sequence Module -> Group -> Name, in the column "Hotkey", the currently selected keyboard shortcut is displayed. To remove a shortcut, click the × icon in the selected line of the tool.

To change or edit a keyboard shortcut, click on the line of the selected tool, which will display the table for entering a new keyboard shortcut:

Figure 600: img

Use the "Save" action to insert a new keyboard shortcut for the selected tool, the "Cancel" action cancels the performed action.

If the keyboard shortcut is already in use, when you select the "Save" button, you will see warning information in the footer of the viewer:

Shortcut is already in use in 'Frontal view'

Figure 601: img

The "Overwrite" action will be added to the table for defining the keyboard shortcut, which allows you to transfer the keyboard shortcut to a new tool:

Figure 602: img

Example - Filtered "Frontal view" tool with new assigned keyboard shortcut ALT+F:

		Shortcuts config		>
∓ Filter by modul	le/group/action name	Filter by hotkey ALT F		>
Module •	Group	Name	Hotkey	
Active panel	Volume	Frontal view	ALT F	×
CLOSE		RESET SHORTCUTS TO DEFAULT		SAVE

Figure 603: img

With the "Save" action in the "Shortcuts config" table, you save new selected keyboard shortcuts to logged-in user. You will be informed of this fact by the information in the footer of the page:

Figure 604: img

Use the "Cancel" action to cancel the action.

The "Reset shortcuts to default" action restores the original layout of the keyboard shortcuts. To restore the original settings, select the "Reset shortcuts to default" action and then the "Save" action. You will be informed about the return to the original settings by the information in the footer of the page:

Shortcuts successfully resets to default	×
Figure 605: img	
Figure 606: img	

9.13.2.4 Profile editor {#viewsettings_Profile editor} The "Profile editor" tool is used to display and position OSD labels in the individual panels of the "Windows for displaying image data". OSD labels

(On Screen Display) represent information about the given images (their acquisition, date, names, etc.) and the content of the information displayed in the image (size, measurements, etc.).

To change the layout of the OSD labels, select the "Profile editor" tool, which display the "Profile editor" table:

Profile editor					Х
+ ADD 🗸	Auto select OSD profile				
Name	Condition	Priority		Ac	tions
Default			0 /	•	Ð
MG	List of modalities: MG		0 /	• •	×
ECG	List of modalities: ECG		0 /	•	×

Figure 607: img

This table contains: - Add - actions to create a new OSD label layout - Autoselect OSD profile Auto select OSD profile - option to choose automatic selection of OSD labels layout according to the condition, i.e. priority and modality of examination, in case of disabling this function only the default OSD labels layout will be displayed - List of saved OSD label layouts - contains Name, Condition, Priority and specific Actions: - Edit Condition - tool for specifying the condition for automatic OSD profile selection - Edit - tool for editing the created OSD profile - Duplicate - create a new OSD profile with the same parameters according to the selected profile - Set as default - selecting this action will set the selected profile as default - Restore - by selecting this action the selected profile will be set to the default values - Remove - selecting this action will delete the selected profile

Edit Condition

\$

Figure 608: img

Tool for specifying the condition for automatic selection of the OSD profile, selecting this action brings up the table:



Figure 609: img

This table contains:

- Manual Configuration a tool to enable/disable the selection of a modality determining condition, selecting this action will not allow the modality to be selected for automatic OSD profile selection
- Text box for selecting a custom modality
- Modality selection for automatic OSD profile selection
- Priority specify the priority for automatic OSD profile selection, the highest priority is number 1

Create a new OSD profile

To create a new profile, choose the "Add" action to create a new empty profile, or choose the "Duplicate" action from the desired profile line, selecting this action will duplicate the assigned OSD profile values. Selecting the "Add" or "Duplicate" action will bring up the "Profile editor" table:

			Profile editor						<u> </u>
rofile name* Default_copy	Patient information	-	Orientation	*	Synchronization indicator	-	Custor	n mod	ality
Patient information	DICOM tag value	_			DICOM tag value				÷
3D orientation visualiser	{InstitutionName}		Resize active viewer		{StudyID} {StudyDescription}				
DICOM tag value	DICOM tag value				DICOM tag value		ECG	CR	CT
Fixed text	Acc no. (AccessionNumber)				(SeriesDescription)		DS	DX	ES
Expression	DICOM tag value						10	ко	MG
Windowing information	{StudyDate}				Series number				
Selected image icon		•				•	MR	NM	OP
Series number	Orientation	^		^	Ruler	^	OT	PR	PT
Image number							PX	RF	SC
Ruler	Value scale				Value scale				
DFOV							ST	SR	US
Value scale	Ruler				Value scale		XA		XC
Multiframe information	Ruler				Orientation				
Waveform info	- Concer				onentation				
Waveform scale		*		Ψ.		*			
Resize active viewer	3D orientation visualiser	•	Ruler	•	DICOM tag value KVP {KVP} kV /	_			
Orientation	SD Orientation Visualiser		Kulei		{XRayTubeCurrent} mA				
Older / newer flag	Waveform info		Orientation		DICOM tag value Gantry tilt:				
Color of localizer lines					{GantryDetectorTilt}°				
Synchronization indicator	Color of localizer lines				DICOM tag value Exposure time:				
Projection and slab thickness editor					{ExposureTime} ms				
Mammo warning		-		-	Echo train length: {EchoTrainLength}	-			

Figure 610: img

In this table, it is possible to edit the profile for displaying selected OSD labels under certain conditions: **Dicompass Gateway**

• Profile name

The "Profile name" text field allows you to name the new created profile. To name it, enter the text in this text field:

Profile name*	
Test profile	

Figure 611: img

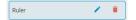
• Layout of OSD labels in the image

This part of the table clearly shows the distribution of the OSD labels in the image into 9 parts:

Patient information PICOM tag value (InstitutionName) PICOM tag value Acc no. (AccessionNumber) PICOM tag value (StudyDate)	▲ ■	Orientation Resize active viewer	•	Synchronization indicator DICOM tag value (StudyID) (StudyDescription) DICOM tag value (SeriesDescription) Series number
Orientation Value scale Ruler Ruler	•		•	Ruler Value scale Value scale Orientation
3D orientation visualiser Waveform info Color of localizer lines	•	Ruler Orientation	•	DicOM tag value KVP (KVP) kV / (KRay Tube/Current) mA DicOM tag value Gantry tilt: (GantryDetectorTilt)* DicOM tag value Exposure time: (Exposure time: (Exposure time) DicOM tag value Exho train length: (Echo Train Length)

Figure 612: img

Individual assigned OSD labels can be "Edited" or "Removed" by hovering the mouse over the selected OSD label and selecting one of the icons:



```
Figure 613: img
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By selecting the "Edit" action, you will display table for editing the OSD label.

9.13.2.4.1 OSD label: DICOM tag value {#viewsettings_DICOM tag value} Text field Value caption specifies the template into which the values of the tags listed in the item Codename values

are added. The position in which the value is replaced is written as empty braces {}. Tag names are entered as an 8-digit hexadecimal number. E.g. we write the Patient's Name tag (0010,0010) as 00100010. The individual values are then separated by a comma. The format sequence_tag[index]/tag is used to enter the sequence. An integer (numbering starts from 0) or an asterisk (*) can be used as an index , which means that we want to use all the descendants of the sequence and connect the values with a comma.

For example, the template can look like this: Name: {}, sex: {} and values 00100010,00100040

If we would like to list all patient identifiers from the sequence, the following settings can be used: Template: Additional patient identifiers: {}, values: 00101002[*]/00100020

9.13.2.4.2 OSD label: Expression {#viewsettings_OSD label Expression} The https://pub.dev/packages/expressio library is used to process expressions. You can also find syntax information on this page.

In addition to the basic syntax, the following features are supported:

- string(value) converts value to a string
- replace(value, pattern, replacement) replaces all matches of the regular expression pattern with replacement in value
- digits(value, digits) converts the value number to a string with a fixed number of decimal places specified in the digits parameter
- nonNull(a,b,c) supports 2-3 parameters and returns the first one that has a non-null value
- time(value) converts the timestamp to a time in the format corresponding to the language setting
- date(value) converts the timestamp to a date in a format corresponding to the language setting
- dateTime(value) converts the timestamp to a datetime in a format that matches the language setting

Example expressions:

time(nonNull(\${(0008,0032)?},\${(0008,0033)?}))
digits(\${(0018,0088)}-\${(0018,0050)},1)

Modalities

This part of the table allows you to select the modalities for which this OSD label profile will be displayed:



Figure 614: img

• OSD label items

This part of the table contains individual OSD labels for assignment to "Layout of OSD labels in the image":

Patient inf	ormation
3D orienta	tion visualiser
DICOM tag	j value
Fixed text	
Expression	ı
Windowing	g information
Selected in	mage icon
Series nun	nber
Image nun	nber
Ruler	
DFOV	
Value scal	e
Multiframe	information
Waveform	info
Waveform	scale
Resize act	ive viewer
Orientation	ı
Older / nev	ver flag
Color of lo	calizer lines
Synchronia	zation indicator
Projection editor	and slab thickness
Mammo w	rarning

Figure 615: img

To add an OSD label, click the left mouse button on the selected OSD label and hold and drag to move it to the selected part "Layout of OSD labels in the image", releasing the mouse button will place the OSD label in the selected part and the editing table will be display (this table may differ according to the selected OSD label):

Aodality						
ECG	CR	CT	DS	DX	ES	
01	КО	MG	MR	NM	OP	
ОТ	PR	PT	PX	RF	_ sc	
ST	SR	US	XA	XC		
) (
Custom r	modality					
Hide	e on small	view				

Figure 616: img

In this table, edit the parameters of the OSD label. After finishing, press the "Save" action to insert the OSD label into the selected section "Layout of OSD labels in the image", press the "Cancel" button to cancel the action.

- Use the "Cancel" button in the "Profile editor" table to cancel the action
- The action "Initial profile layout" will remove the changes made to the OSD labels
- The "Save" action will save the changed "Profile editor" profile

If you have saved a new profile, it will be displayed in the "Label layout" table and can be edited using the "Edit condition", "Edit", "Duplicate", "Set as default", "Remove" tools:

Profile editor					×
+ ADD 🗸 Auto sele	ct OSD profile				
Name	Condition	Priority		Ac	tions
Default			0 /	•	Ð
MG	List of modalities: MG		0 🖊	n 🖻	×
Test profile	List of modalities:		0 /	h	×
ECG	List of modalities: ECG		0 🖊	h	×

Figure 617: img

Ø

Figure 618: img

"Preset windows management" tool is used to define the windowing profiles of selected windows. By

ADD +			
Name •	Windowing information	Hotkey	Actio
Abdomen / Mediastinum	center: 0; width: 350	F2	e ×
Angio	center: 250; width: 800	F7	<i>•</i> >
Bone	center: 480; width: 2,500	F5	<i>•</i> >
Bone 2	center: 400; width: 4,000	F6	e >
Brain	center: 40; width: 80	F4	<i>•</i> >
Lung	center: -500; width: 2,000	F3	/>

selecting this tool, you will display the "Windowing preset" table:

Figure 619: img

This table shows the individual created profiles. The columns contain information about presets: -Name - windowing name - Windowing information - windowing preset center and width value - Hotkey - selected keyboard shortcut for applying the windowing preset - Actions - contains actions for "Edit" and "Remove" the predefined windowing preset

The "Add" action allows you to create a new predefined windowing preset. Select this action to display the "Add" table:

Add		×
Name		
🗆 suv		
Center 512		
Width 1,023		
Hotkey		
	CANCEL	ADD

Figure 620: img

In this table, select the desired parameters for the new profile and then select the "Add" button to add the new profile to the "Window preset" table. The "Cancel" button cancels the action performed. To make the work easier, in this table the values of the center and width of the window are selected according to the currently displayed image, the checkbox "SUV" indicates the use of Standardized Uptake Value.

4

Figure 621: img

9.13.2.6 Tool Selection {#viewsettings_Tool Selection} The "Tool Selection" tool is used to add and edit saved DICOM viewer configurations. Selecting this tool display the "Tool selection" table:

Tools selection							>
ADD +							
Condition	Left mous	Middle mo	Right mou	Left + Righ	Priorit	Default	Actio
List of modalities: ECG	Waveform	Move	Zoom	Magnify gl			1>
List of modalities: CT	Stacking	Move	Windowing	Magnify gl			1>
All modalities	Zoom	Move	Windowing	Magnify gl		~	1
						SA	NE 🗸

Figure 622: img

This table contains:

- **Viewer configuration profiles** individual columns on the profile row contain the selected parameters, the "Actions" column contains "Edit" buttons to call up the profile parameter configuration table and the "Delete" button to remove the selected profile.
- Button "Save" saves the changes made
- Button "Add" selecting "Add" calls up the table for creating a new viewer configuration profile:



Figure 623: img

This table contains:

- Type option to choose for which modality the configuration will be set, you can choose from:
 - All modalities
 - List of modalities
- Modality text field for entering individual modalities, this window is active if you have chosen "List of modalities", enter modalities in this window and separate them with a comma, e.g. CT, MR, PT, etc.
- Priority The priority determines the order in which the tools will be selected. Tools with a priority of 10 will be selected before those with a priority of 1
- Left, Middle, Right mouse buttons drop-down menu for defining the selected tool on the mouse button, in this line, after expanding the menu, click on the selected tool for assignment, move in this menu with the mouse wheel:



Figure 624: img

• Left + right mouse button - drop-down menu for defining the selected tool when the left + right mouse button is pressed at the same time, this menu contains only some tools:



Figure 625: img

• The "Save" button saves the defined viewer configuration, the "Cancel" button cancels the action performed. The changes will be applied at the next login to the system Dicompass Gateway. In case you want to return to the defined viewer configuration while working, you can use the "Select default tools" keyboard shortcut, this keyboard shortcut must be defined first, see paragraph "Shortcuts".

The change to switch to the default tools can also be controlled using the functions located in the "Display toolset" under the "Default tools" tab:

• "Resetting after the active viewer change" - When switching to another panel in the "Image Data Display Window", the currently assigned tools on each mouse button will be replaced by the default tools

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Figure 626: img

9.13.2.7 Position of pinned actions {#viewsettings_Position of pinned actions} The "Position of pinned actions" tools are used to determine the order in which the tools are displayed. Select this tool to call up the "Position of pinned actions" table:

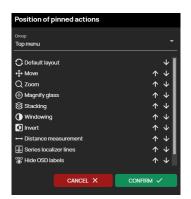


Figure 627: img

In this table, use the "Group" drop-down menu to select the desired area to reposition the tools:

- Top menu tools in "User configurable toolbar"
- Mobile menu tools in the "User configurable toolbar" in mobile mode
- Context menu tools in the "Context menu"

The position of the tool can be changed by clicking on the arrows located to the right of the selected tool.

After making the required changes, choose the "Save" action to save the changed state, the "Close" action cancels the action.

Tools in the "Position of pinned actions" table, you can choose the settings of the display of tools, see the chapter "DICOM viewer tools" paragraph "Display tools".

С

Figure 628: img

9.13.2.8 Restart windows layout {#viewsettings_Restart windows layout} The "Restart windows layout" tool is used to reload the displayed images/series in the "Window for displaying image data". It serves as a "refresh" DICOM viewer.

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Figure 629: img

9.13.2.9 Working set configuration {#viewsettings_Working set configuration} Selecting the "Work Set Configuration" tool brings up the "Work Set Configuration" table, which allows the user to

modify the Viewer work set sidebar, select the viewer work set side position, and switch between the flexible and optimal viewer work set configurations:

Working set config	uration X
Visible working set	Study columns
Position	
Left side 👻	Series columns
	-0 2
Layout type Flexible	Wide study columns
	 5
	Wide series columns
Auto open viewers n	nap on drag rs for auto open viewers map
	2
`	
CAN	CEL SAVE

Figure 630: img

Warning: turning off viewer working set display can be reverted in the viewer configuration tools tab by turning it on in the "Working set configuration" tool.

9.13.3 Browser state {#viewsettings_Viewer state}

The "Browser state" tools group includes the following tools:

lcon	Function	Description
	Load viewer state	View saved DICOM viewer states
	Save viewer state	Saving or sharing edited study, including its measurement, post-processing, localization, etc.
	Save state as hanging protocol	Takes the current view of the DICOM viewer state and inserts these parameters into the configuration of the new hanging protocol

Figure 631: img

9.13.3.1 Load viewer state {#viewsettings_Load viewer state} The "Load viewer state" tool is used to work with and display saved DICOM viewer states. To create a DICOM viewer state, follow the "Save viewer state" paragraph. By selecting this tool, you will display the "Browser states" table:

Browser states				>
Name 🔺	Description	Created	Actions	
test		2022-01-04 15:11	く 🔍 🍵	1
Test1	Test1	2024-04-24 14:42	< 🗖 🔋	

Figure 632: img

In this table, it is possible to filter individual saved DICOM viewer states using the "Enter name" text field.

This table contains saved DICOM viewer states and allows you to perform actions on them:

 Share < - this action allows you to share the saved DICOM viewer state with other users logged into the Dicompass Gateway system. Selecting this action will display the "Manage users" table where you can add users to share the saved state. You can use the check boxes to assign them rights for further sharing "Share" or for possible modification of the status "Edit". The "Save" action activate sharing to users, the "Cancel" action cancels the action:

The second s		Manage	users				
Enter the user login or username test							
vailable users			Selected users				
.ogin name	User name	Add	Login name	User name	Share	Edit	Remove
dw3test1		0 *	test			\checkmark	•
dw3test2		•					
est2		•					
		•					s

Figure 633: img

Comments - the "Comments" action is used for sending messages and writing information about the saved state. By choosing this action, you will display the "Comments" table. Enter text in the "Comment" text field and press the "Add" action - to add a comment. This comment will then be visible to shared users.



Figure 634: img

• Remove - removes the saved DICOM viewer state. By choosing this action, you will be asked to confirm or cancel by the YES/NO action, using the "Question" table. If you confirm the removal of the saved state, you will be informed about the successful removal of the state by information in the footer of the DICOM viewer:

Figure 635: img

Figure 636: img

9.13.3.2 Save viewer state {#viewsettings_Save viewer state} If the work in the browser must be finished and you want to continue it later, it is possible to save all the work for the possibility of reloading. The entire state of the windows, the studies in progress are saved. This status can still be shared with other users, who can add their comments to it. It is a so-called off-line consultation.

By selecting this tool, you will display the "Save browser state" table:

Save browser state		
Name*		
Description		
CANCEL	SAVE	

Figure 637: img

In this table, enter in the "Name" text field the name of the state and "Description" more detailed information about the state. Use the "Save" action to create a new saved DICOM viewer state, use the "Cancel" action to cancel the action. You will be informed about successful saving by information in the DICOM viewer footer:

Figure 638: img

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Figure 639: img

9.13.3.3 Save state as hanging protocol {#viewsettings_Save state as hanging protocol} The "Save state as hanging protocol" tool accepts the current display of the DICOM viewer status and inserts these parameters into the configuration of a new hanging protocol, opening the "Hanging protocol manage" window for possible specification of parameters. More detailed information about the creation of hanging logs can be found in the chapter "My hanging protocols".

9.13.4 Other actions {#viewsettings_Other actions}

lcon	Function	Description
osb ♥	Hide OSD labels	Hiding/displaying OSD labels in individual windows "Image data display window"
(Hide overlay	Hide/show saved measurements in the image
\$	Display shutter	Allows you to hide all pixels that are outside the specified area
ţ	Reverse windows order for HP	Reversing the order of applied hanging protocols on individual monitors
**	Hanging protocols enabled	Disables/enables the option to display hanging protocols
	Image editor	Tool for editing the selected image
8	Help	Product label view and User Guide link
Ð	Logout	Logging out and ending work in the Dicompass Gateway system
Ê	All Studies of Patient	View all studies performed on the selected patient

The "Other Actions" tool group contains the following tools:

OSD

Figure 640: img

9.13.4.1 Hide OSD labels {#viewsettings_Hide OSD labels} The "Hide OSD labels" tool is used to hide/display the OSD labels in individual windows "Image data display window". Select this tool to hide the OSD labels:

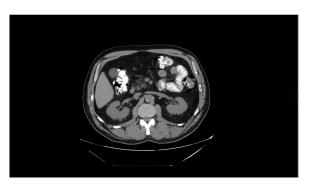


Figure 641: img

To display the OSD labels, select this tool again, the OSD labels will be displayed:

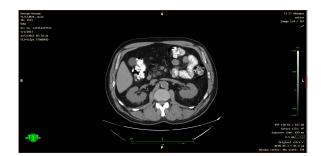


Figure 642: img

Figure 643: img

9.13.4.2 Hide overlay {#viewsettings_Hide overlay} The "Hide overlay" tool allows you to hide/show saved measurements in the image. Selecting this will hide the saved measurements:

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Figure 644: img

To display the overlayed measurements in the image, select this tool again, the saved measurements will be displayed:



Figure 645: img

Warning: The "Hide overlay" tool is used to hide measurements entered directly into the DICOM image information, the so-called overlay. This tool does not hide measurements saved as "PR" - presentation state.

 \mathbf{i}

Figure 646: img

9.13.4.3 Display shutter {#viewsettings_Displey shutter} The "Display shutter" tool allows you to hide part of the image where appropriate. Turn on this function to hide part of the image:

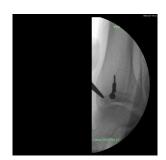


Figure 647: img

Selecting this action again will disable this feature:



Figure 648: img

←→

Figure 649: img

9.13.4.4 Reverse windows order for HP {#viewsettings_Reverse windows order for HP} In case you have set hanging protocols for multiple monitors, the "Reverse windows order for HP" tool reverses the order of applied hanging protocols on the monitors, i.e. that the hanging protocol for the second monitor will be displayed on the first monitor and vice versa.

When you choose this tool, you will be notified by the system of the information: "Reverse windows order for HP: Yes/No".

Figure 650: img

9.13.4.5 Hanging protocols enabled {#viewsettings_Hanging protocols enabled} The "Hanging protocols enabled" tool disables/enables the option to display hanging protocols. In case of deactiva-

tion, the "Hanging protocols" icon will be inactive 📰.

When you choose this tool, the system will alert you with the information: "Hanging protocols enabled: Yes/No".

For more information about displaying hanging protocols, refer to "Viewer Tools" ("Selecting a Hanging Protocol"). The HP settings are described in the section "Hanging Protocols Settings".

Figure 651: img

9.13.4.6 Image editor {#viewsettings_Image editor} The "Image Editor" tool allows the user to create a new edited image, e.g. with a change of side marking, image flipping or rotation.

To edit an image, follow these steps:

- open the selected image for editing in the "Image data display window" and keep the image panel active
- select the "Image Editor" tool to call up the "Image Editor" tab:



Figure 652: img

- the table contains:
 - left column patient's name with ID, study name with date of creation and series name with image value
 - middle column preview of the edited image
 - right column options for image editing and save destination
- to edit the image, use the options in the Right column, which includes:
 - Attributes Change the ability to edit the side labels of the mammography image:



Figure 653: img

- Erase Area - this function adds the option to mark the part of the image to be erased. After selecting this tool, hover over the desired area and click and drag to create a rectangular selection of the area to be deleted:



Figure 654: img

 Add text - if you have selected and marked "Erase area", it is possible to select the function "Add text". Use the "Text" field to enter the desired text, which will then be inserted in the place of the "Delete area" marking:

	Attributes change No change
8	Fraso area Add text Text L
	Flip horizontally Flip vertically Relation
	Target. Archive Archive Archive DPGW@ (CLOUDPACS@127.0.0.1)
	Series description Ankle-joint - edited

Figure 655: img

- Flip horizontally flips the image according to the horizontal axis
- Flip vertically flips the image according to the vertical axis
- Rotation select to rotate the image in degrees "°" using the text field

- select "Target" (archive, station, user), if archive destination is selected, select the chosen "Archive" and name the saved image in "Series description"
- selecting the "CANCEL" action cancels the work performed, selecting the "SAVE" action saves the modified image according to the selected "Target" and adds it to the viewer working set.

For example, the saved image looks like this:

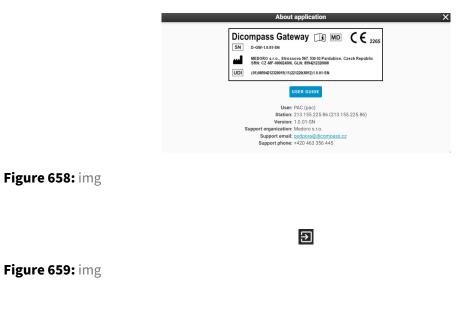


Figure 656: img

?

Figure 657: img

9.13.4.7 Help The "Help" tool is used to display a table containing basic information about the product, with the option of opening this "User guide":



9.13.4.8 Logout The "Logout" tool is used for logging out and ending work in the Dicompass Gateway system.

Selecting this tool will display a page for a possible re-login:



Figure 660: img

By selecting the "Log in again" action, you will be redirected to the Dicompass Gateway system login page.

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Figure 661: img

9.13.4.9 All patient studies {#viewsettings_All patient studies} By selecting the "All patient studies" function, you will call up a table at the bottom of the screen displaying all the performed examinations of the selected patient. For more information, refer to the paragraph "Function All patient studies".

10 Settings {#settings_Global settings}

In this chapter, the third main part will be described, namely the Dicompass Gateway configuration. This dialog can be displayed from the main menu by selecting the category "Settings" located in the upper right corner of the basic registry dialog. Another option for opening the "Settings" dialog is from the DICOM viewer dialog, from the "DICOM viewer settings" tools group and selecting the "Global configuration" tool 📽.

Choose one of these actions to open the "Settings" dialog:

User guide

Dicompass Gateway

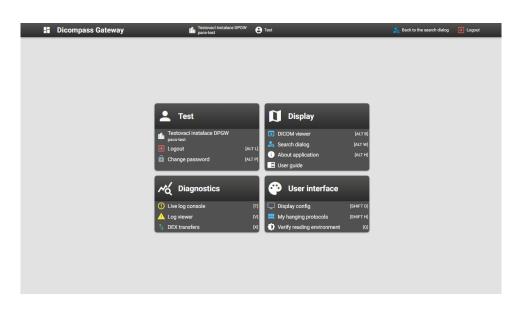


Figure 662: img

In case you opened "Settings" from the basic registry dialog and now want to close it, choose the icon React to the search dataget, which you return to the previous work in the basic registry dialog. In case you opened "Settings" from the DICOM viewer dialog and now want to close it, select the icon by selecting this icon In case you opened "Settings" from the DICOM viewer dialog and now want to close it, select the icon by selecting this icon In case you opened "Settings" from the DICOM viewer dialog and now want to close it, select the icon by selecting this icon In case you opened "Settings" from the DICOM viewer dialog and now want to close it, select the icon by selecting this icon In case you opened "Settings" from the DICOM viewer dialog and now want to close it, select the icon by selecting this icon "In the basic registry dialog."

If you want to return from the configuration tool dialog to the main "Settings" dialog, use the icon

If you want to log out of the DPGW system, select the "Logout" icon Organization.

The main dialog "Settings" contains configuration tools that are located in individual tabs, these are: - User" -
"User" -
"Display" -
"User interface"

Warning: Individual configuration tools are displayed based on the rights assigned to the user. If you need advanced system configuration, contact your administrator. All configuration tools are described in the administrator's manual.

Warning: In the case of a DPGW system located in the cloud, the main dialog "Settings" also contains the tab "Account statistics". This tab contains: - Tariff name - Number of opened studies - Storage usage - Number of users - External share links

10.1 User

-

Figure 663: img

The "User" tool group contains the following tools:

lcon	Function	Description
i∰a	Archive Name	Contains the name of the archive used DPGW
Э	Logout	Logging out of the system DPGW
â	Change password	Changing the password of the currently logged in user

10.1.1 Change password {#settings_Change password}

Figure 664: img

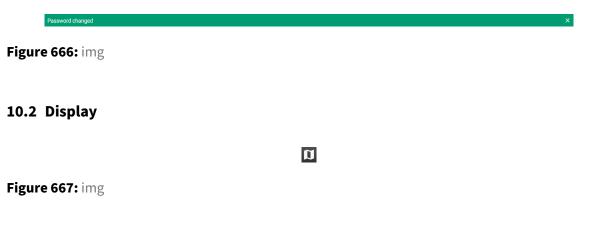
The "Change password" tool allows you to change the password of the currently logged-in user. By selecting this tool, you will display the "Change password" table:

Ô

	Change password	×
Old password*		
New password	*	
Check for a new	v password*	
CANCEL		SAVE

Figure 665: img

By choosing the "CANCEL" action, you cancel the action being performed. By selecting the "SAVE" action, a new password will be applied, according to the specified parameters. You will be informed about a successful password change in the footer of the screen:



The "Display" tool group includes the following tools:

lcon	Function	Description
ត	DICOM viewer	Returning to the DICOM viewer dialog
2	Search dialog	Returning to the basic registry dialog
•	About application	Display information of the DPGW system, including its label
	User guide	Display this user manual

10.2.1 About application {#settings_About application}

Figure 668: img

The "About application" tool is only informative, containing basic information and the system label DPGW. By selecting this tool, you will display "About application" table:

i



Figure 669: img

Close this table by selecting the "X" action.

10.2.2 User guide {#settings_User guide}

Figure 670: img

The "User guide" tool is used to display this user manual. Selecting this tool will display a new web browser window containing this user manual.

10.3 Diagnostics

Figure 671: img

The "Diagnostics" tool group contains the following tools:

lcon	Function	Description
0	Live log console	View the DPGW activity log in real time
	Log viewer	preview and search in already saved system log files DPGW
t _↓	DEX transfers	preview and search received and sent studies using the mDEX inter-hospital exchange network

10.3.1 Live log console {#settings_Live log console}

Figure 672: img

The "Live log console" configuration tool is used to display the DPGW activity log in real time. Using this configuration tool, it is possible to obtain detailed information about the processes performed by the system.

(!)

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| 5                                  | Live log console                                                                                           | Testovací instalace DPGW (CentOS8)<br>pace-test8 | <b>O</b> TEST                                                                                                  | 👔 Back to the viewer 🔁 Logou       | ıt                           |
|------------------------------------|------------------------------------------------------------------------------------------------------------|--------------------------------------------------|----------------------------------------------------------------------------------------------------------------|------------------------------------|------------------------------|
|                                    | er.handler.ContextHandler.doHandle(ContextHan                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | er.handler.ScopedHandler.nextScope(ScopedHand<br>let.ServletHandler.doScope(ServletHandler.jav             |                                                  |                                                                                                                |                                    | x ╤ q ↓ II                   |
|                                    | er.session.SessionHandler.doScope(SessionHand                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | er.handler.ScopedHandler.nextScope(ScopedHand                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | r.handler.ContextHandler.doScope(ContextHand                                                               |                                                  |                                                                                                                |                                    |                              |
|                                    | er.handler.ScopedHandler.handle(ScopedHandler                                                              |                                                  |                                                                                                                |                                    |                              |
| at org.eclipse.jetty.serve         | er.handler.HandlerWrapper.handle(HandlerWrapp                                                              | er.java:122)                                     |                                                                                                                |                                    |                              |
| at org.eclipse.jetty.rewri         | ite.handler.RewriteHandler.handle(RewriteHand                                                              | ller.java:301)                                   |                                                                                                                |                                    |                              |
| at org.eclipse.jetty.serve         | er.handler.HandlerWrapper.handle(HandlerWrapp                                                              | er.java:122)                                     |                                                                                                                |                                    |                              |
|                                    | er.handler.gzip.GzipHandler.handle(GzipHandle                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | er.handler.HandlerCollection.handle(HandlerCo                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | er.handler.HandlerWrapper.handle(HandlerWrapp                                                              | er.java:122)                                     |                                                                                                                |                                    |                              |
|                                    | er.Server.handle(Server.java:563)                                                                          |                                                  |                                                                                                                |                                    |                              |
|                                    | er.HttpChannel.lambda\$handle\$0(HttpChannel.ja                                                            |                                                  |                                                                                                                |                                    |                              |
|                                    | er.HttpChannel.dispatch(HttpChannel.java:762)<br>er.HttpChannel.handle(HttpChannel.java:497)               |                                                  |                                                                                                                |                                    |                              |
|                                    | er.HttpConnection.onFillable(HttpConnection.i                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | ostractConnection\$ReadCallback.succeeded(Abst                                                             |                                                  |                                                                                                                |                                    |                              |
|                                    | illInterest.fillable(FillInterest.java:100)                                                                |                                                  |                                                                                                                |                                    |                              |
|                                    | electableChannelEndPoint\$1.run(SelectableChan                                                             | nelEndPoint.iava:53)                             |                                                                                                                |                                    |                              |
|                                    | .thread.strategy.AdaptiveExecutionStrategy.ru                                                              |                                                  | (, java:421)                                                                                                   |                                    |                              |
| at org.eclipse.jetty.util.         | thread.strategy.AdaptiveExecutionStrategy.co                                                               | nsumeTask(AdaptiveExecutionStra                  | ategy.java:390)                                                                                                |                                    |                              |
|                                    | .thread.strategy.AdaptiveExecutionStrategy.tr                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | .thread.strategy.AdaptiveExecutionStrategy.la                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    | .thread.ReservedThreadExecutor\$ReservedThread                                                             |                                                  | /a:411)                                                                                                        |                                    |                              |
|                                    | thread.QueuedThreadPool.runJob(QueuedThreadP                                                               |                                                  |                                                                                                                |                                    |                              |
|                                    | thread.QueuedThreadPool\$Runner.run(QueuedThr                                                              | eadPool.java:1077)                               |                                                                                                                |                                    |                              |
| at java.base/java.lang.Thr         |                                                                                                            |                                                  |                                                                                                                |                                    |                              |
|                                    | <pre>Lib.QRException: A-ASSOCIATE-RJ[result: 1 - r<br/>lib.RetrieveTools.openAssociation(Unknown Sou</pre> |                                                  | service-user, reason: 1 - no-reason-give                                                                       | n J                                |                              |
|                                    | lib.QueryTools.startQuery(Unknown Source)                                                                  | ince)                                            |                                                                                                                |                                    |                              |
|                                    | igin.dw.archivesearch.SearchCallable.process0                                                              | wony(Unknown Sounce)                             |                                                                                                                |                                    |                              |
|                                    | .gin.dw.archivesearch.SearchCallable.processo                                                              |                                                  |                                                                                                                |                                    |                              |
|                                    |                                                                                                            |                                                  |                                                                                                                |                                    |                              |
|                                    | ugin.dw.archivesearch.SearchCallable.call(Unk                                                              |                                                  |                                                                                                                |                                    |                              |
| at java.base/java.util.con         | current.FutureTask.run(FutureTask.java:264)                                                                |                                                  |                                                                                                                |                                    |                              |
| at java.base/java.util.con         | ncurrent.ThreadPoolExecutor.runWorker(ThreadP                                                              | oolExecutor.java:1136)                           |                                                                                                                |                                    |                              |
|                                    | ncurrent.ThreadPoolExecutor\$Worker.run(Thread                                                             | PoolExecutor.java:635)                           |                                                                                                                |                                    |                              |
| 1 common frames omitte             |                                                                                                            |                                                  |                                                                                                                |                                    |                              |
|                                    | AssociateRJ: A-ASSOCIATE-RJ[result: 1 - rejec                                                              | ted-permanent, source: 1 - serv                  | /ice-user, reason: 1 - no-reason-given]                                                                        |                                    |                              |
|                                    | coder.nextPDU(PDUDecoder.java:182)                                                                         |                                                  |                                                                                                                |                                    |                              |
|                                    | iation\$2.run(Association.java:566)                                                                        |                                                  |                                                                                                                |                                    |                              |
| 3 common frames omitte             | ed<br>⊃-2360443] admin org.medoro.dpgw.web.pLugin.d                                                        | k. ReleventStudiesSeculate [30]                  |                                                                                                                |                                    |                              |
|                                    | 5-2360443] admin org.medoro.dpgw.web.plugin.d<br>5-2360291] admin org.medoro.dpgw.core.blapi.i             |                                                  |                                                                                                                | untion couring is dischlod by polo |                              |
|                                    | adErrorCache] org.medoro.dpgw.core.slapi.i                                                                 |                                                  |                                                                                                                | action saving its disabled by Fole | [DISADL_OSCK_CONFIG_SAVE]. 1 |
|                                    | adErrorCache] org.medoro.dpgw.core.service.s                                                               |                                                  |                                                                                                                |                                    |                              |
| Toto of to donselou, dos info [rea | activities of graceoros apgas coresservices                                                                | Service halfer in                                | and a second |                                    |                              |

#### Figure 673: img

The configuration tool "Live log console" contains several tools to facilitate the work. These tools are located in the upper right corner of the dialog "Live log console": - "Clear log console" - with this tool it is possible to clear the console - "Set log filter" - selecting this tool will display a line for possible console log filtering:

| TRACI - Thread | Logger | Filter | RESET FILTER SET LOG FILTER | × <del>=</del> ۹ ↓ Ⅱ |
|----------------|--------|--------|-----------------------------|----------------------|
|----------------|--------|--------|-----------------------------|----------------------|

#### Figure 674: img

• "Log search" • selecting this tool will display a line for a possible search in the console log:

| Level 👻 Filter | RESET FILTER SEARCH |  | × = ¢, v |
|----------------|---------------------|--|----------|
|----------------|---------------------|--|----------|

#### Figure 675: img

- "Autoscroll" selecting this tool will always display the last line of information in the log console with autoscrolling
- "Pause log events" 🛄 choose this tool to pause the information listing in the log console lines

# **10.4** Log viewer {#settings\_Log viewer}

#### Figure 676: img

The "Log viewer" configuration tool is used for previewing and searching in already saved log files and thus allows you to obtain more detailed information about the operation of the DPGW system. Selecting this tool display the "Log viewer" dialog:

| Log viewer                                                                                                                          | Testovaci instalace DPGW (CentOS8)<br>pacs-test8 | E TEST                                              | 🖪 Back to the viewer 🗧 Logout |                                |
|-------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------|-----------------------------------------------------|-------------------------------|--------------------------------|
| at org.eclipse.jetty.server.handler.HandlerWrapper.handle(H                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.server.handler.gzip.GzipHandler.handle                                                                         |                                                  |                                                     |                               | dpgw.log 🔍 C' 💻                |
| at org.eclipse.jetty.server.handler.HandlerCollection.handl                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.server.handler.HandlerWrapper.handle(H                                                                         | andlerWrapper.java:122)                          |                                                     |                               |                                |
| at org.eclipse.jetty.server.Server.handle(Server.java:563)                                                                          |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.server.HttpChannel.lambda\$handle\$0(Htt<br>at org.eclipse.jetty.server.HttpChannel.dispatch(HttpChanne        |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.server.HttpChannel.handle(HttpChannel.                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.server.HttpConnection.onFillable(HttpC                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.io.AbstractConnection\$ReadCallback.suc                                                                        |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.io.FillInterest.fillable(FillInterest.                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.jo.SelectableChannelEndPoint\$1.run(Sel                                                                        |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.util.thread.strategy.AdaptiveExecution                                                                         |                                                  | ev.java:421)                                        |                               |                                |
| at org.eclipse.jetty.util.thread.strategy.AdaptiveExecution                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.util.thread.strategy.AdaptiveExecution                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.util.thread.strategy.AdaptiveExecution                                                                         | Strategy.lambda\$new\$0(AdaptiveExecutionSt      | trategy.java:139)                                   |                               |                                |
| at org.eclipse.jetty.util.thread.ReservedThreadExecutor\$Res                                                                        |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.util.thread.QueuedThreadPool.runJob(Qu                                                                         |                                                  |                                                     |                               |                                |
| at org.eclipse.jetty.util.thread.QueuedThreadPool\$Runner.ru                                                                        | n(QueuedThreadPool.java:1077)                    |                                                     |                               |                                |
| at java.base/java.lang.Thread.run(Thread.java:833)                                                                                  |                                                  |                                                     |                               |                                |
| Caused by: org.medoro.dpgw.dicom.lib.QRException: A-ASSOCIATE-RJ[re                                                                 |                                                  | <ul> <li>service-user, reason: 1 - no-re</li> </ul> | ason-given]                   |                                |
| at org.medoro.dpgw.dicom.lib.RetrieveTools.openAssociation(                                                                         |                                                  |                                                     |                               |                                |
| at org.medoro.dpgw.dicom.lib.QueryTools.startQuery(Unknown                                                                          |                                                  |                                                     |                               |                                |
| at org.medoro.dpgw.web.plugin.dw.archivesearch.SearchCallab                                                                         |                                                  |                                                     |                               |                                |
| at java.base/java.util.concurrent.FutureTask.run(FutureTask                                                                         |                                                  |                                                     |                               |                                |
| <pre>at java.base/java.util.concurrent.ThreadPoolExecutor.runWor at java.base/java.util.concurrent.ThreadPoolExecutor\$Worker</pre> |                                                  |                                                     |                               |                                |
| 1 common frames omitted                                                                                                             | .run(Inreadrooitexecutor.java:655)               |                                                     |                               |                                |
| Caused by: org.dcm4che3.net.pdu.AAssociateRJ: A-ASSOCIATE-RJ[result                                                                 |                                                  |                                                     |                               |                                |
| at org.dcm4che3.net.PDUDecoder.nextPDU(PDUDecoder.java:182)                                                                         |                                                  | rvice-user, reason, i - no-reason                   | r-grvenj                      |                                |
| at org.dcm4che3.net.Association\$2.run(Association, java:566)                                                                       |                                                  |                                                     |                               |                                |
| 3 common frames omitted                                                                                                             |                                                  |                                                     |                               |                                |
| 2023-01-10 08:51:57,483 INFO [web-2360443] admin org.medoro.dpgw.we                                                                 | b.plugin.dw.RelevantStudiesServlet: [10]         | relevant studies found, [1] erro                    | ons                           |                                |
| 2023-01-10 08:51:57,837 INFO [web-2360291] admin org.medoro.dpgw.co                                                                 |                                                  |                                                     |                               | DISABLE USER CONFIG SAVE]. Ign |
| 2023-01-10 08:52:00,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 | service.ServiceExecutor: Service name=[re        | eadErrorCache] is starting                          |                               |                                |
| 2023-01-10 08:52:00,003 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 | service.ServiceExecutor: Service_name=[re        | eadErrorCache] has finished                         |                               |                                |
| 2023-01-10 08:52:30,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:52:30,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:53:00,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:53:00,004 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:53:30,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:53:30,003 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:53:46,031 INFO [digiQueueClean-1] org.medoro.dpgw.dig                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:54:00,002 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 |                                                  |                                                     |                               |                                |
| 2023-01-10 08:54:00,003 INFO [readErrorCache] org.medoro.dpgw.core.                                                                 | service.ServiceExecutor: <u>Service_name=[re</u> | eaderrorcache hàs finished                          |                               |                                |

# Figure 677: img

The configuration tool "Log viewer" contains several tools to facilitate the work. These tools are located in the upper right corner of the "Log viewer" dialog: - "Log file name" deputed - by choosing this tool it is possible to open selected log file, e.g.:

| dpg  | w.log               |
|------|---------------------|
|      | prouting.log        |
| brov | wser.log            |
| hl7- | in.log              |
|      | out.log             |
| pref | etch.log            |
| stdo | out.log             |
| dpg  | w.20220801_0.log.gz |
|      | w.20220731_0.log.gz |
|      | w.20220730_0.log.gz |
|      | w.20220729_0.log.gz |
|      | w.20220728_0.log.gz |
|      | w.20220727_0.log.gz |
|      | w.20220726_0.log.gz |
|      | w.20220725_0.log.gz |
|      | w.20220724_0.log.gz |
|      | w.20220723_0.log.gz |
|      | w.20220722_0.log.gz |
|      | w.20220721_0.log.gz |
| dpg  | w.20220720_0.log.gz |
| brow | wser.202207.log.gz  |

# Figure 678: img

The text in the tool icon is displayed according to the name of the currently opened log file.

• "Log search" - selecting this tool will display a line for a possible search in the "Log viewer":

| Dicompass Gateway                                                                                                                                                                                                                                                   | User guide     |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------|
|                                                                                                                                                                                                                                                                     |                |
| Level 👻 Filter RESET FILTER SEARCH                                                                                                                                                                                                                                  |                |
| at org.eclipse.jstty.server.handler.Handler#rapper.handle(Handler#rapper.java:122)<br>at org.eclipse.jstty.server.handle.eng.zip.GripHundler.handle(GripHundler.java:822)<br>at org.eclipse.jsty.server.handle.HandlerCollection.handle(HandlerCollection.java:131) | dpgw.log Q C 👱 |

Figure 679: img

- "Refresh log file" 🖸 "Log viewer" does not display information in real time, it is only used for reading saved information, so use "Refresh log file" tool to obtain latest information
- "Download log" by selecting this tool, you will start downloading the open log in the "Log viewer" dialog to the station's local storage in .log format, it is o text file with selected log

# **10.5 DEX transfers {#settings\_DEX transfers}**

# Figure 680: img

The "DEX transfers" tool is used to preview and search for data received and sent via the mDEX interhospital network. Select this tool to display the search dialog for:

 $|\uparrow_{\downarrow}|$ 

- Received data
- Sent data

These dialogs contain a history of sent/received data and filtering using text fields: - Sender/receiver - sending/receiving healthcare facility - Patient name - Patient identification - ID - Object identification - UID study

|                    |                     | DEX trar                                         | nsfers  | ŵ                | CloudPACS @ PAC<br>cloudpacs1 @ ADMIN | ISTRATOR & Back to the search diak                   | ; 🗃 Logout            |              |           |         |
|--------------------|---------------------|--------------------------------------------------|---------|------------------|---------------------------------------|------------------------------------------------------|-----------------------|--------------|-----------|---------|
|                    |                     |                                                  |         |                  | RECEIVED DATA                         | FORWARDED DATA                                       |                       |              |           |         |
| ender              |                     | Patient<br>test                                  | name    |                  | F                                     | Patient identification                               | Object identification |              |           |         |
| ransfer start 🔻 👘  | Transfer end        | Sender                                           | City    | Patient name     | Patient identification                | Object Identification                                |                       | Object count | Size      | Actions |
| 023-10-20 09:13:40 | 2023-10-20 09:13:40 | I - An nemocnice v 3m + p.o.                     | Second  | NOHA testovaci   | HE SHOELD AS                          | 1.3.6.1.4.1.20744.3.1.2.2.12.1328101243577.516569133 |                       | 5            | 5.5 MB    | •       |
| 023-07-24 12:49:10 | 2023-07-24 12:49:13 | Plicní ambulance Plicini Plicní Plicné ambulance | rines.  | TEST             | menter in                             | 1.2.840.113564.1921681090.202306121327183281         |                       | 7            | 101.23 MB | 0       |
| 023-07-24 12:41:40 | 2023-07-24 12:41:40 | Plicní ambulance Philippe - Internetion          | Piece . | TEST TEST        | DOM: N HOM                            | 1.2.840.113564.1921681209.202307211148299211         |                       | 1            | 15.58 MB  | 0       |
| 023-07-24 12:40:54 | 2023-07-24 12:40:55 | Plicni ambulance Phillippi - IDC/ s              | Berg    | TEST TEST        | AT DECISION                           | 1.2.840.113564.1921681209.202307211148299211         |                       | 1            | 15.58 MB  | •       |
| 023-02-20 17:50:10 | 2023-02-20 17:50:10 | remocnice i stateme                              | and the | TESTOVACI POKUS2 | TOTAL NO.                             | 1.2.826.0.1.3680043.8.1053.6.1637051840682.25.282572 |                       | 1            | 407.56 kB | Ο       |

# Figure 681: img

For more information about the sent/received study, use the "eye" action on the right side to call up the study detail dialog:

# User guide

## Dicompass Gateway

|       | <b>B</b> DEX transfers                                            | the CloudPACS 😕 | PAC ADMINISTRATOR                                                             | ack to the search dialog | E Logout               |
|-------|-------------------------------------------------------------------|-----------------|-------------------------------------------------------------------------------|--------------------------|------------------------|
| 1 - 6 |                                                                   |                 | Transfer start                                                                |                          | Transfer end           |
| 2 *   | "patient": {                                                      |                 | 2023-10-20 09:13:40                                                           |                          | 2023-10-20 09:13:40    |
| 3     | "name": "noha^testovaci",                                         |                 |                                                                               |                          |                        |
| 4     | "ident": "141514151415",                                          |                 | Sender                                                                        |                          | Receiver               |
| 5     | "sex": "F",                                                       |                 | Boullá nemocnice - (p.o.                                                      |                          | Medoro s.r.o.          |
| 6     | "birthDate": "2000-00-00 00:00",                                  |                 |                                                                               |                          |                        |
| 7 ~   | "study": {                                                        |                 | Oty                                                                           |                          | City                   |
| 8     | "uid": "1.3.6.1.4.1.20744.3.1.2.2.12.1328101010577.516569133",    |                 | Sec.                                                                          |                          | Pardubice              |
| 9     | "description": "Noha P",                                          |                 |                                                                               |                          |                        |
| 10    | "accessionNumber": "12184646",                                    |                 | Patient name                                                                  |                          | Patient identification |
| 11    | "studyDate": "2023-10-10 00:00",                                  |                 | NOHA testovaci                                                                |                          | 141514151415           |
| 12 *  | "series": {                                                       |                 |                                                                               |                          |                        |
| 13 ~  | "1.3.6.1.4.1.20744.3.1.2.2.13.1328101243577.516637837": {         |                 | Direct count                                                                  |                          | Stre                   |
| 14    | "modality": "SR",                                                 |                 | S                                                                             |                          | 5.5 MB                 |
| 15    | "imageCount": 1,                                                  |                 | •                                                                             |                          |                        |
| 16    | "size": 937                                                       |                 |                                                                               |                          |                        |
| 17    | },<br>"1.3.46.670589.30.39.0.1.966169727528.1697719504144.1": {   |                 | Object identification<br>1.3.6.1.4.1.20744.3.1.2.2.12.1328101243577.516569133 |                          | Object type<br>DICOM   |
| 18 *  |                                                                   |                 | 1.3.8.1.4.1.20744.3.1.2.2.12.1328101243577.51656913                           |                          |                        |
| 20    | "modality": "SR",<br>"description": "Radiation Dose Information", |                 |                                                                               |                          |                        |
| 20    | "imageCount": 1,                                                  |                 | Domain                                                                        |                          |                        |
| 22    | "size": 5684                                                      |                 |                                                                               |                          |                        |
| 22    | 5126 . 3064                                                       |                 |                                                                               |                          |                        |

# Figure 682: img

Use the "Back" action to return to the "DEX Transfers" dialog.

# **10.6** User interface {#settings\_User interface}

# P

## Figure 683: img

The "User interface" tools group contains the following tools:

| lcon | Function                      | Description                                                                                |
|------|-------------------------------|--------------------------------------------------------------------------------------------|
| Ţ    | Display config                | Configuring display of DPGW on the monitors connected to the workstation                   |
| ***  | My hanging<br>protocols       | Editing and management of hanging protocols stored on the logged in user                   |
| Ø    | Verify reading<br>environment | Indicative verification of the suitability of the environment for the purpose of diagnosis |

# 10.6.1 Display config {#settings\_Display config}

#### Figure 684: img

The configuration tool "Display config" is used to configure the display of DPGW monitors connected to the workstation. Select this tool to open the "Display config" dialog:

Ţ

| 🚦 Display config                                    | Testovací instalace DPGW pacs-test | e Test                                                   | ack to the search dialog | 🔁 Logout   |
|-----------------------------------------------------|------------------------------------|----------------------------------------------------------|--------------------------|------------|
|                                                     | Search dialog                      | Main viewer window                                       |                          |            |
| Search dialog<br>Short example of text. Font size 1 |                                    | Main viewer window<br>Short example of text. Font size 2 |                          |            |
| DEFAULT LAYOUT                                      |                                    |                                                          | s                        | AVE LAYOUT |

# Figure 685: img

The "Display config" configuration tool contains the following areas:

# 1. Search dialog

The area of the search dialog contains tools to configure the display of the basic registry dialog:

- "Fullscreen" option to display in fullscreen mode, this tool can be used if the "Set from system default" tool is disabled
- "Display" 📑 option to choose Light, Dark, Grayscale display mode:



#### Figure 686: img

• "Font size" 🖆 - option to choose the displayed font size:

|                                                  | Main viewer                        |  |
|--------------------------------------------------|------------------------------------|--|
| Search dialog Short example of text. Font size 1 | window                             |  |
|                                                  | Short example of text. Font size 4 |  |
|                                                  |                                    |  |

# Figure 687: img

• "Set from system default" - if this tool is active, it takes over the monitor screen resolution settings from the workstation operating system settings. By default, this tool is active, if you want to change the resolution and monitor position of the search dialog display, deactivate this tool. Deactivating the tool will display the "Edit" action:



## Figure 688: img

Selecting the "Edit" action 2 will open the monitor selection window:

| Si Display config | A laterated | ×                 | 🕹 har i ta anti-bég 🛛 📓 lapet |  |
|-------------------|-------------|-------------------|-------------------------------|--|
|                   |             |                   | Ð                             |  |
| Search dialog     |             | Main viewer windo | m -                           |  |
| ***               |             | ***               |                               |  |
|                   |             |                   |                               |  |
|                   |             |                   |                               |  |
|                   |             |                   |                               |  |
|                   |             |                   |                               |  |

Figure 689: img

Select the selected monitor.

- "Show search dialog on same / another display" / Solution
   by deactivating this tool, the "Main viewer window" will be opened in a separate Internet browser window, disable this tool if the workstation has at least two monitors connected. Deactivate this tool to display the "Edit" action in the "Main viewer window".
- 2. Digitalization

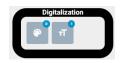


Figure 690: img

The "Digitization" contains tools to configure the display of the "Digitization" dialog: - "Display" 🖻 - "Font size" 💼

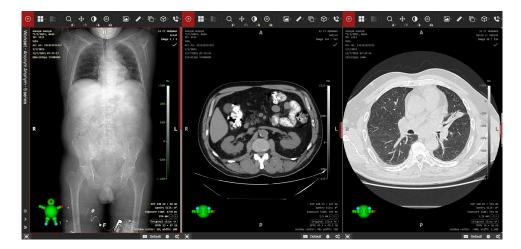
3. Main viewer window



# Figure 691: img

The image of the main browser window contains tools to configure the display of the browser dialog:

- "Fullscreen"
- "Display"
- "Font size" 📑
- "Display multiplicity" by selecting this tool it is possible to divide the area of the "Window for displaying image data" into multiple windows, up to three :



# Figure 692: img

#### 4. Add display

In the case of a multi-monitor workstation, it is possible to add separately opening DPGW system to individual monitors. To add another display, click on the "Add display" action:



#### Figure 693: img

According to the individual monitor resolution or system settings, there are 2 types of assignment of browser windows to the desired monitor:

1. Selecting the "Add monitor" icon will open a window for selecting a monitor:



#### Figure 694: img

Then select the desired monitor by clicking in the appropriate field to assign the viewer window.

2. Selecting the "Add monitor" icon will open a web browser pop-up window, which you drag to the upper left corner of the desired monitor, on which the added viewer window will be opened when the DICOM viewer is opened:



#### Figure 695: img

After selecting one of the options, the "Display 1" monitor is added to the "Display Settings" dialog:

#### **Dicompass Gateway**

| Display config                                   | Testovací instalace DPGW pacs-test | e Test                                                  | ack to the search dialog | E Logout  |
|--------------------------------------------------|------------------------------------|---------------------------------------------------------|--------------------------|-----------|
| Search dialog                                    | Main viewer wi                     | ndow<br>17<br>1<br>1<br>1<br>1<br>1<br>1<br>1<br>1<br>1 | <u>+</u>                 |           |
| Search dialog Short example of text. Font size 1 |                                    | Main viewer windo                                       | w                        |           |
| ★ ★ ★<br>48 34 24                                |                                    |                                                         |                          |           |
| DEFAULT LAYOUT                                   |                                    |                                                         | SA                       | VE LAYOUT |

# Figure 696: img

The "Display 1" display contains the following tools: - "Edit" display editor for configuring the display layout - "Preview font size" - shows a preview of the text size - "Remove" - select this tool to remove the set display

#### 5. Save layout

With the action "Save layout" you save and apply the configured display settings of the displays. Select this action to display the "Question" table:



# Figure 697: img

The action "Default layout" will return to the default configuration of monitor display settings. By selecting this action, you will be notified of the information in the footer of the screen:

Figure 698: img

To make it easier and more comfortable to work with the DPGW system, use the "DICOM viewer extension" add-on. This add-on ensures automatic placement of dialogs on individual set monitor display screens. If this add-on is not installed and you have multiple monitors set up, individual dialogs will only open on one monitor and the user will then manually drag them to the selected monitors.

You can find the add-on in the "Extension" of the Chrome internet browser, by opening the "Chrome Web Store" and typing "DICOM viewer extension" into the search field:



#### Figure 699: img

#### 10.6.2 My hanging protocols {#settings\_My hanging protocols}

#### Figure 700: img

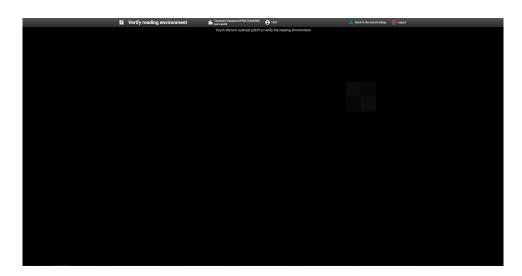
The "My hanging protocols" configuration tool is used for editing and managing hanging protocols stored on the logged in user. For more information about configuration hanging protocols, go to the chapter "Setting Hanging Protocols".

#### **10.6.3 Verify reading environment {#settings\_Verify reading environment}**

0

#### Figure 701: img

The configuration tool "Verify reading environment" is used for indicative verification of the suitability of the environment for the purpose of diagnostics. Select this tool to open the "Verify reading environment" dialog:



# Figure 702: img

In this dialog, click the low-contrast check box to verify the suitability of the environment for diagnostic purposes. If you click on the low-contrast box, you will be notified of successful verification:

|                                    | × |
|------------------------------------|---|
| Verification was successfully done |   |
| ок                                 |   |
|                                    |   |

Figure 703: img

By selecting the "OK" action, you will be returned to the main "Settings" dialog.

If you click outside the low-contrast box, you will be notified of a failed verification:



# Figure 704: img

After selecting the "OK" action, the configuration tool "Verify reading environment" will still be opened with the option of clicking on the low-contrast box.

Warning: Please note that this quality check is only indicative. To verify the quality and suitability of the environment for diagnosis, use safety technical checks and calibration of diagnostic monitors as medical devices.

If set by the administrator, periodically validation of diagnostic monitors can be used. You will be prompted for this validation by a pop-up table with the option to perform the verification, or the option

to postpone this verification with a certain number of postponements, after the completion of the postponement deduction, it will be necessary to perform the verification. Verification will be required every time Dicompass Gateway is run for the first time on a given station. The performed validations is recorded in the "Display device validation audit" tool and for individual stations in the administrator section "Settings".

# **11** Setting Hanging Protocols {#hp\_Setting Hanging Protocols}

Hanging protocol (HP) is used to display images with parameters provided that the specified conditions are met.

This user guide contains basic HP settings, created and saved per logged in user. If advanced HP configuration is required, contact your DPGW system administrator. Advanced HP configuration is described in the Administrator's Guide.

To edit and manage HP, display "Hanging protocols manage" dialog using the configuration tool "My hanging protocols" located in the main "Settings" dialog:

Figure 705: img

Select this tool to open the "Hanging protocols manage" dialog:

| Hanging protocols manage | test.dicompass.cloud<br>Test Dicompass | 📻 Back t | o the viewer 🛛 🛃 Logo |
|--------------------------|----------------------------------------|----------|-----------------------|
| + C                      |                                        |          |                       |
| Name or description      |                                        |          |                       |
| Name •                   | Description                            | Active   | Actions               |
| 77                       |                                        |          | ∕≡                    |
| CT 2                     |                                        |          | ∕∎                    |
| Test                     | Test                                   |          | /=                    |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
|                          |                                        |          |                       |
| BACK                     |                                        |          |                       |

# Figure 706: img

The dialog "Hanging protocols manage" contains:

- "Add HP" action 🛨 by choosing this tool it is possible to create a new HP
- "Reload" action selecting this tool will refresh opened dialog "Hanging protocols manage", , which serves similarly to the "refresh" of the Internet browser
- Text field "Name or description" This text field is used to search for selected created HP
- Field with created HPs here you will find HPs created by the logged in user. This field contains:
  - HP name and description
  - Active option to choose whether the given HP should be displayed/hidden in the DICOM viewer DPGW
  - Actions the action column contains icons for editing HP "Edit" and icon for removing HP "Remove"

| Name + | Description | Active       | Actions |
|--------|-------------|--------------|---------|
| ст     |             | $\checkmark$ | ∕≡      |
| CT 2   |             |              | /=      |

#### Figure 707: img

• "Back" action Action action this action will return you to the main "Settings" dialog

HP can be created in two ways, by clicking the "Add HP" action <sup>+</sup> from the "Hanging protocols manage" dialog, or by selecting the "Save state as hanging log" I from the "DICOM viewer settings" tools group tab in the DPGW DICOM viewer.

# 11.0.1 Creating a new HP "Add HP" {#hp\_Creating a new HP by Add HP}

#### Figure 708: img

To create a new HP, select the "Add HP" action + from the "Hanging protocols manage" dialog. This will redirect you to the new HP dialog box:

+

|      | Hanging protocols manage         | test.dicompass.cloud<br>Test Dicompass | B PAC | Back to the viewer | 🔁 Logout |
|------|----------------------------------|----------------------------------------|-------|--------------------|----------|
| Name | •                                | Description                            |       |                    |          |
| ST   | UDY WINDOW JSON                  |                                        |       |                    |          |
| Stu  | dy query                         |                                        |       |                    |          |
| Grou | p meets all 💌 of conditions.     |                                        |       |                    |          |
|      | Tag 👻 00000000 is not            | empty 👻 .                              |       |                    |          |
| Act  | ions                             |                                        |       |                    |          |
| No   | selected actions (i)             |                                        |       |                    |          |
| Œ    | ) Lock view                      |                                        |       |                    |          |
| Œ    | Lock view and windowing          |                                        |       |                    |          |
| Œ    | Completely lock view             |                                        |       |                    |          |
| Œ    | Automatic lock                   |                                        |       |                    |          |
| Œ    | Automatic lock without windowing |                                        |       |                    |          |
| Œ    | Localizer lines                  |                                        |       |                    |          |
| Œ    | Mammography image alignment      |                                        |       |                    |          |
| BAC  | κ                                |                                        |       |                    | SAVE     |

#### Figure 709: img

In the header of the dialog, it is possible to name the new HP using the "Name" text field and describe more detailed information about the HP in the "Description" text field.

The dialog contains three tabs for entering parameters: - "Study" - contains parameters for the HP application - "Window" - contains HP display parameters - "JSON" - configuration file in JSON format "Back" action action will redirect you to the "Hanging protocols manage" dialog

The "Save" action save is used to save new HP to the logged-in user.

**11.0.1.1 Study tab {#hp\_Study tab}** The "Study" tab contains parameters for the HP application according to the specified conditions:

|      |                  | f conditions.     |         |         |
|------|------------------|-------------------|---------|---------|
|      | Та               | g 👻 0000000       | 0 is no | t empty |
|      | + 🗉              |                   |         |         |
| \cti |                  |                   |         |         |
| ACU  | ons              |                   |         |         |
| No : | selected actions |                   | (i)     |         |
| Ð    | Lock view        |                   |         |         |
| €    | Lock view and w  | indowing          |         |         |
| Ð    | Completely lock  | view              |         |         |
| €    | Automatic lock   |                   |         |         |
| -    | Automatic lock v | without windowing |         |         |
| Ð    |                  |                   |         |         |
| Ŭ    | Localizer lines  |                   |         |         |

# Figure 710: img

The study tab is divided into two areas:

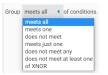
#### 1. Study query

This area contains conditions for initiation HP and is organized into a tree of condition groups, think of it as a file system where each file is located in a specific folder. The tree of groups and conditions can look like this, for example:

| Group |             | ▼ of conditions. |                |
|-------|-------------|------------------|----------------|
|       |             | Tag 👻 [tag]      | is not empty 💌 |
|       |             | Tag 🔻 [tag]      | is not empty 🔻 |
|       |             | Tag 🔻 [tag]      | is not empty 💌 |
|       | +           | •                |                |
| Tag   | ▼ [tag]     | is not empty 🔻 . |                |
| Tag   | ▼ [tag]     | is not empty 🔻 . |                |
| Group | ▼ meets all | ▼ of conditions. |                |
|       |             | Tag 🔻 [tag]      | is not empty 💌 |
|       | +           | <b>E</b>         |                |
| Tag   | ▼ [tag]     | is not empty 💌 . |                |
| Tag   | ▼ [tag]     | is not empty 💌 . |                |

#### Figure 711: img

The main group, or any subgroup, may have certain conditions for initiation HP. These conditions can be displayed by clicking on the text "meets all" in the text "Group meets all of conditions.":



#### Figure 712: img

To add a new condition, select the "Add condition" action 🛨.

To add a new subgroup, select the "Add group" action 🧧.

By hovering the mouse over the added condition, it is possible to include it in a new group with the "Wrap with group" action, or remove it with the "Remove" action:



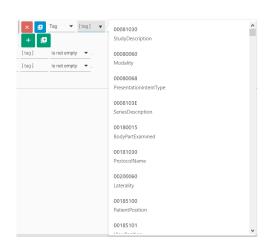
# Figure 713: img

The value of the condition can be changed by clicking on the "Tag" field and choosing the selected range of values:



# Figure 714: img

It contains the following areas: - Tag - the option to choose a value according to the selected DICOM tag of the study, by clicking in the second field [ tag ] you will display a scrolling menu for choosing the selected DICOM tag:



#### Figure 715: img

Click in the third field, in this case "is not empty" you can choose a value condition:

| 🗙 📴 Tag 🔻 [tag] | is not empty 🔹 .                                                                                  |
|-----------------|---------------------------------------------------------------------------------------------------|
| + 🛛             | contains<br>contains (case sensitive)<br>equals<br>equals (case sensitive)<br>is empty<br>is null |
|                 | is not empty                                                                                      |
|                 | meets a regular expression                                                                        |

Figure 716: img

By choosing a value condition other than "is empty", "is null" and "is not empty", a fourth text field [ value ] will be displayed, in which you can enter the required parameter:



# Figure 717: img

Here is an example of the selected HP display condition for the CT modality, according to the CT Modality DICOM tag:



#### Figure 718: img

• Groovy - option to choose a value according to the Groovy script, enter the script value in the text field:



#### Figure 719: img

- Group by selecting this area, you change the condition to a group
- Exists by selecting this area, you will place the condition in the condition initiation state if it exists:



Figure 720: img

#### 2. Actions

The second area in the create new HP dialog is the "Actions" area. This area contains the possible application of tools to the created HP. You can find a description of the individual tools in the "DICOM viewer tools" chapter.

The applicable tools are contained in this area and can be added by clicking on the row of the selected tool:

| Actio | ons                            |     |   |
|-------|--------------------------------|-----|---|
| ⊗     | Lock view                      | =   | ٥ |
| ۲     | Localizer lines                | =   | ٥ |
| Ð     | Lock view and windowing        |     |   |
| Ð     | Completely lock view           |     |   |
| Ð     | Automatic lock                 |     |   |
| Ð     | Automatic lock without windowi | ing |   |
| Ð     | Mammography image alignmen     | t   |   |

# Figure 721: img

Click on the row of the added tool to remove this tool from the added list.

Click and drag the icon to change the order of the tool in the added list.

Click the cicon to go to the "JSON" tab and view the selected tool.

**11.0.1.2 Window tab {#hp\_Window tab}** The "Window" tab contains tools for configuring the display of DICOM viewer layouts and assigning HP information. Go to this tab after completing the settings of the "Study" tab.

| Hanging protocols manage | test.dicompass.cloud PAC | 🖪 Back to the viewer 🔁 Logout |
|--------------------------|--------------------------|-------------------------------|
| Name*<br>Test            | Description<br>test      |                               |
| STUDY WINDOW JSON        |                          |                               |
| Windows count            |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
|                          |                          |                               |
| BACK                     |                          | SAVE                          |

# Figure 722: img

To set the HP display in this tab, proceed as follows:

• choose the number of monitor screens on which the DPGW DICOM viewer dialogs will be opened and press the "Add" icon:

| Windows count<br>1 | • | + |
|--------------------|---|---|
| 1                  |   |   |
| 2                  |   |   |
| 3                  |   |   |
| 4                  |   |   |

#### Figure 723: img

• this has created the HP display configuration and it is possible to continue working with it.

Now the "Window" tab now contains three subfolders "Basic data", "Series", "Layout":

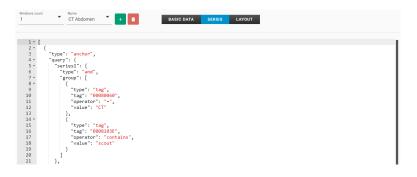
#### 1. Basic data



#### Figure 724: img

In this subfolder, type in the text fields - Name - naming HP, the name will be displayed in the DICOM viewer DPGW - Icon - assignment of an icon to HP (icons correspond to Google fonts and can be found here: https://fonts.google.com/icons, after entering their name, the selected icon will appear) - Group - naming the HP group and then adding it to the HP groups displayed in the DICOM viewer DPGW

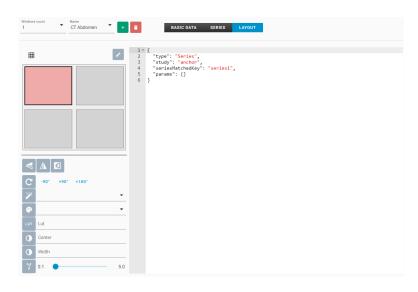
# 2. Series



# Figure 725: img

The "Series" part of the JSON configuration file is displayed in this subfolder. This is an advanced settings management, please contact your administrator to configure it.

#### 3. Layout

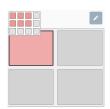


#### Figure 726: img

This subfolder contains tools for displaying series in a DPGW DICOM viewer dialog. This subfolder contains:

• Panels layout:

To layout panels, click the 🔳 icon and select the layout of the panels:



# Figure 727: img

This layout can be entered manually by clicking the  $\boxed{}$  icon and changing the number of "rows" and the number of "cols":



#### Figure 728: img

• Tool assignment:

By selecting a specific panel in "Panels layout", it is possible to assign tools to it that will be applied to selected panel:

|     | Δ               |     |
|-----|-----------------|-----|
| C   | -90° +90° +180° |     |
| ¥.  |                 | •   |
| ۲   |                 | •   |
| LUT | Lut             |     |
| 0   | Střed           |     |
| •   | Šířka           |     |
| Ŷ   | 0.1             | 5.0 |

Figure 729: img

You assign a tool by clicking on the tool icon and assigning its value, e.g.:



#### Figure 730: img

Information on individual tools can be found in the "DICOM viewer tools" chapter.

• JSON config

In this area of the "Layout" subfolder, a section of the JSON configuration file containing the selected tools of the selected HP panel is displayed. This is an advanced settings management, please contact your administrator to configure it. Example:

| 1 -                               | {                              |
|-----------------------------------|--------------------------------|
| 2                                 | "type": "Series",              |
| 3                                 | "study": "anchor",             |
| 4                                 | "seriesMatchedKey": "series1", |
| 5 -                               | "params": {                    |
| 2<br>3<br>5 *<br>6<br>7<br>8<br>9 | "rotate": 90,                  |
| 7                                 | "flipVert": true,              |
| 8                                 | "filter": "SHARPEN",           |
| 9                                 | "filterRate": 1,               |
| 10                                | "windowCenter": 150,           |
| 11                                | "windowWidth": 100,            |
| 12                                | "gamma": 1.6,                  |
| 13                                | "lut": 10                      |
| 14                                | }                              |
| 15                                | }                              |
|                                   |                                |

Figure 731: img

**11.0.1.3 JSON tab {#hp\_JSON tab}** The "JSON" tab contains the HP settings configuration file and allows its editing. This is an advanced settings management, please contact your administrator to configure it. Example:

| 1 - {                                    |                         |  |  |
|------------------------------------------|-------------------------|--|--|
| 2                                        | "active": true,         |  |  |
| 3                                        | name": "Bricho",        |  |  |
| 4                                        | 4 "id": "00011",        |  |  |
| 4<br>5 *<br>6 *<br>7<br>8 *<br>9 *<br>10 | s "matcher": {          |  |  |
|                                          | 6 "anchorstudyQuery": { |  |  |
| 0.1                                      | anchorscubyquery : {    |  |  |
| /                                        | "type": "and",          |  |  |
| 8 -                                      | "group": [              |  |  |
| 9 -                                      | {                       |  |  |
| 10                                       | "type": "tag",          |  |  |
| 11                                       | "tag": "00081030",      |  |  |
| 12                                       | "operator": "equals",   |  |  |
| 13                                       | "value": "Bricho"       |  |  |
| 14                                       |                         |  |  |
| 15 -                                     | p.                      |  |  |
|                                          |                         |  |  |
| 16                                       | "type": "tag",          |  |  |
| 17                                       | "tag": "00080060",      |  |  |
| 18                                       | "operator": "equals",   |  |  |
| 19                                       | "value": "CT"           |  |  |
| 20                                       | 3                       |  |  |
| 21                                       |                         |  |  |
| 22                                       |                         |  |  |
| 23                                       |                         |  |  |
| 23                                       |                         |  |  |
|                                          |                         |  |  |

Figure 732: img

# 11.0.2 Creating a new HP "Save state as hanging log" {#hp\_Creating a new HP by Save state as hanging log}

The second option to create a new HP is to select the "Save state as hanging protocol" tool I from the "DICOM viewer Settings" tools group tab in the DICOM viewer – PRODUCT-SHORT-NAME–.

To create a new HP using this tool, first edit the panel layout in the DICOM viewer DPGW and assign the selected series. Then select the "Save state as hanging protocol" tool. This will take you to the new HP configuration dialog. If you want to edit a new HP, proceed according to the paragraph "Creating a new HP"Add HP"", with the only difference being that the values are pre-filled according to the selected state of the DICOM viewer DPGW.

# **11.0.3 Error conditions {#hp\_Error conditions}**

If the configuration was not set correctly, you will be warned about this by the information in the footer of the new HP configuration dialog, after pressing the "Save" action:

| Invalid query |  |
|---------------|--|
|               |  |

Figure 733: img

In this case, the invalid configuration area will also be highlighted, e.g.:



# Figure 734: img

If all required fields are not filled in, these fields will be highlighted and new HP cannot be saved.

In the case that the HP modification using JSON is not valid, you will be warned about this by the information in the footer of the new HP configuration dialog, after pressing the "Save" action:

The configuration file is not a JSON!

# Figure 735: img

For an invalid line of JSON code, it will be marked as:



Figure 736: img