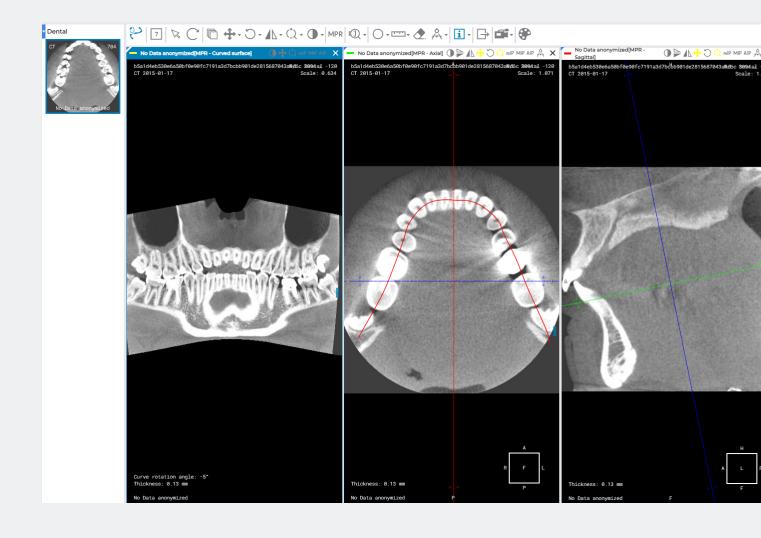
software for healthcare

Inobitec Web DICOM Viewer version 2.8

USER'S MANUAL





The information contained in this User Manual belongs to INOBITEC Software FZ-LLC, Dubai Media City, Building 05, Dubai, UAE, P.O. Box 73030. The Manual is delivered to the users of «Inobitec Web DICOM Viewer» software product exclusively for the purpose of working with this product. No part of the information contained herein can be modified, used for any other purpose or delivered to any third party without the prior written consent of INOBITEC Software FZ-LLC. The INOBITEC Software FZ-LLC reserves the right to alter this Manual without prior notice.

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About this Manual

This User Manual describes the functionality of «Inobitec Web DICOM Viewer» (version 2.8) and provides instructions on how to use this software product.

Accepted Conventions

Names of program interface elements, key names and important notes are printed in **bold**. Image captions are printed in *italics*.

Technical Support

Technical support of «Inobitec Web DICOM Viewer» users is provided by the INOBITEC Software FZ-LLC team. If you apply for technical support, please include the following information in your message:

- the name, version and bitness of the operating system of the computer(s) on which the software server components are installed (you can get this information from your system administrator);
- your operating system version from which a client connects to the server (you can get this information from your system administrator);
- product version. The program version is specified in the **About** section of the user's menu. For details, see Section 1.2.1.

To apply for technical support, or if you have any further questions or comments, please email us at **support@inobitec.com**

About the Product

The «Inobitec Web DICOM Viewer» software product is intended for viewing, analyzing and printing medical data obtained from various DICOM equipment (modality). The product is deployed on diagnostic workstations and integrated with DICOM servers. The Web DICOM Viewer propels the capabilities of diagnostics to a new level and makes it possible to detect pathological conditions timely and efficiently, predict their development and plan their elimination.

For the purpose of establishing a diagnosis, «Inobitec Web DICOM Viewer» software must only be used by properly qualified personnel.

×

Making a diagnosis is not allowed when the Web DICOM Viewer is running on mobile devices.

The «Inobitec Web DICOM Viewer» software product and installer does not:

- collect and transfer confidential user information;
- intercept network traffic;
- show ads;
- send spam;
- show messages not related to work;
- automatically update itself without notifying the user.

After uninstalling you do not need to restore your operating system and browser settings. Uninstalling is free of charge. Uninstalling does not adversely affect the operation of the computer and installed software. Files not related to the Web DICOM Viewer are not removed and changed after uninstalling.

For details on the structure of «Inobitec Web DICOM Viewer» package, its installation, launch, licensing, setting up and integration with DICOM servers, see the **Admin's Manual**.

License Agreement is available from the link inobitec.com/eng/about/webviewerLic/.

Web DICOM Viewer Functionality

INOBITEC Products Comparison Table

Function	Lite	Pro	Web
View flat images with the following options:	\checkmark	\checkmark	\checkmark
rotate, pan, zoom, mirror	\checkmark	\checkmark	\checkmark
change window width and level	\checkmark	\checkmark	\checkmark
change the window width and level based on the image parameters (Full dynamic W/L)	\checkmark	\checkmark	
customization of WL presets	\checkmark	\checkmark	
opening DICOM studies from zip archives	\checkmark	\checkmark	
opening several DICOM studies at the same time	\checkmark	\checkmark	\checkmark
automatic filling tab windows with the series	\checkmark	\checkmark	
synchronization of series windows in case of window level/width alteration, movement, scaling, and scrolling	\checkmark	\checkmark	\checkmark
calibrate image size	\checkmark	\checkmark	\checkmark
show image scout lines on other series images	\checkmark	\checkmark	\checkmark
flat image view with orthogonal projections	\checkmark	\checkmark	
merge series into phases	\checkmark	\checkmark	
measure length, angles	\checkmark	\checkmark	\checkmark
play series images subsequently as a movie	\checkmark	\checkmark	\checkmark
choose image resampling filter in the Flat Images View window	\checkmark	\checkmark	
build freeform contours with the Pencil tool	\checkmark	\checkmark	
add comments, labels and various graphic elements on images	\checkmark	\checkmark	\checkmark
moving comments, measurements, measurement results and various graphic elements on images	\checkmark	\checkmark	\checkmark
measure Cobb angles and intensity at a certain point or in a certain area	\checkmark	\checkmark	\checkmark
export images to a graphic file	\checkmark	\checkmark	
export images to a new series of DICOM images	\checkmark	\checkmark	\checkmark
DSA (Digital Subtraction Angiography)	\checkmark	\checkmark	
applying different CLUTs to images	\checkmark	\checkmark	\checkmark
edit CLUTs	\checkmark	\checkmark	

Function	Lite	Pro	Web
browse DICOM image tags	\checkmark	\checkmark	\checkmark
cerebral Perfusion		\checkmark	
blood flow parameters evaluation on the basis of phase-contrast images		\checkmark	
calcium scoring		\checkmark	
measurement of apparent diffusion coefficient (ADC)		\checkmark	
View three-dimensional tissue reconstruction with the following options:	\checkmark	\checkmark	\checkmark
rotate, pan and zoom the model	\checkmark	\checkmark	\checkmark
view scaled-down multiplanar reconstruction of the series simultaneously (with automatic synchronization)	\checkmark	\checkmark	
cut the outside and inside parts of the model	\checkmark	\checkmark	\checkmark
MIP (Maximum Intensity Projection)	\checkmark	\checkmark	
measure length, angles	\checkmark	\checkmark	
export the model to a graphic file	\checkmark	\checkmark	
export the model to a new series of DICOM images	\checkmark	\checkmark	\checkmark
export of several images obtained by rotating the model to a graphic file	\checkmark	\checkmark	
export of several images obtained by rotating the model to a new series of DICOM images	\checkmark	\checkmark	
switching projection	\checkmark	\checkmark	
add markers and line markers	\checkmark	\checkmark	
delete bone tissue in the manual mode (interactively and automatically)	\checkmark	\checkmark	
delete bone tissue for merged series		\checkmark	
build the surface by the model		\checkmark	
surface construction for the main volume		\checkmark	
surface building for each phase of the main volume in multiphase series		\checkmark	
individual change of parameters for each constructed surface		\checkmark	
measure the shortest path on the surface of a visible volume with the Surface Ruler tool		\checkmark	
import of multiple surfaces		\checkmark	

Function	Lite	Pro	Web
segmentation		\checkmark	
build a curvilinear reconstruction		\checkmark	\checkmark
View multiplanar reconstruction with the following options:	\checkmark	\checkmark	\checkmark
view axial, frontal and sagittal sections of tissue	\checkmark	\checkmark	\checkmark
rotate cutting planes in space	\checkmark	\checkmark	\checkmark
3D view	\checkmark	\checkmark	\checkmark
measure length, angles	\checkmark	\checkmark	\checkmark
add markers and line markers	\checkmark	\checkmark	
build freeform contours with the Pencil tool	\checkmark	\checkmark	
setting modes MIP, MinIP, AIP	\checkmark	\checkmark	\checkmark
set slice thickness	\checkmark	\checkmark	\checkmark
MPR view sync on several monitors	\checkmark	\checkmark	
display of MPR planes on 3D view	\checkmark	\checkmark	
build a section of a spatial model by a random surface		\checkmark	
cross-section view for curvelinear reconstruction		\checkmark	
export sections of any of the planes with a selectable increment to a series		\checkmark	
segmentation		\checkmark	
surface construction for the main volume		\checkmark	
surface building for each phase of the main volume in multiphase series		\checkmark	
individual change of parameters for each constructed surface		\checkmark	
import of multiple surfaces		\checkmark	
contour based segmentation		\checkmark	
displaying the boundaries of the segmented structure mask in MPR		\checkmark	
histogram display for ROI		\checkmark	
mixed RGB colors Fusion		\checkmark	
Diffusion Tensor Imaging (DTI) support		\checkmark	
Virtual endoscopy (viewing the inner surfaces of cavities in tissues) with the following options:		\checkmark	
automatically and manually navigate the inside a cavity		\checkmark	

Function	Lite	Pro	Web
build cavity surfaces		\checkmark	
view scaled-down multiplanar reconstruction of the series simultaneously (with automatic synchronization)		\checkmark	
Merge series with the following options:		\checkmark	
remove bone tissue for DualScan and DualEnergy studies		\checkmark	
build three-dimensional models based on multiple series of images of the same tissue in different modes		\checkmark	
quickly switch to a full-scale view of three-dimensional reconstruction, multiplanar reconstruction and virtual endoscopy of the model built by merging series		\checkmark	
stitch images		\checkmark	
View electrocardiogram with the following options:	\checkmark	\checkmark	\checkmark
measure time intervals and values on graphs	\checkmark	\checkmark	\checkmark
using filters		\checkmark	\checkmark
export images to a new series of DICOM images		\checkmark	\checkmark
Print images on paper or film using a DICOM printer with the following options:	\checkmark	\checkmark	
adding reference images	\checkmark	\checkmark	
exporting pages to the PACS server for printing	\checkmark	\checkmark	
adding logo	\checkmark	\checkmark	
Open studies by dragging and dropping on the program icon and program window	\checkmark	\checkmark	
Open studies in the background	\checkmark	\checkmark	
Transformation indicators in the series window with an opportunity to undo the transformations			~
Open/save file using Native OS dialogs	\checkmark	\checkmark	
Voxel and polygon rendering support on Metal for macOS starting with Version 10.15	~	\checkmark	
Set up the interface and the Web DICOM Viewer functionality with the capability to export and import the settings configuration	\checkmark	\checkmark	
Tool control with the left, the right and the middle mouse buttons	\checkmark	\checkmark	\checkmark
High resolution monitor support	\checkmark	\checkmark	\checkmark
The support of secure connection with TLS	\checkmark	\checkmark	

Function	Lite	Pro	Web
PDF data support and presentation in DICOM files			\checkmark
Structured reports support and presentation			\checkmark
The opportunity to view several types of data in one window simultaneously: flat images, volume reconstructions, MPR, ECG, video, PDF documents and structured reports			\checkmark
Minimize application window to system tray	\checkmark	\checkmark	
Integrated help system	\checkmark	\checkmark	\checkmark
Create series display templates for various modalities	\checkmark	\checkmark	
Write data to CD, DVD and flash cards	\checkmark	\checkmark	
Integration with PACS servers	\checkmark	\checkmark	\checkmark
Store data in a local storage	\checkmark	\checkmark	\checkmark
Work as a PACS server, support C-FIND and C-MOVE	\checkmark	\checkmark	
Save studies to the folder	\checkmark	\checkmark	
Save studies as archives	\checkmark	\checkmark	\checkmark
Integration with third-party systems	\checkmark	\checkmark	\checkmark
Edit patient name and study description on the Local Storage	\checkmark	\checkmark	
Anonymize studies and series with the ability to change DICOM tags	\checkmark	\checkmark	
PLY, OBJ, STL, GLB formats surface export		\checkmark	
PLY, OBJ, STL formats surface import		\checkmark	
Saving the model orientation and projection in spj-projects		\checkmark	
Embedded WebViewer		\checkmark	
Register images		\checkmark	
Record video from viewing windows		\checkmark	
Play video contained in DICOM files			\checkmark

«Pro» and «Lite» are the Inobitec DICOM Viewer editions. This is another product. More information can be found in the Products and Downloads pages.

Getting Started

To avoid problems with the Web DICOM Viewer, do the following before you start working with the program:

- 1. Examine the browser you are using to work with the Web DICOM Viewer. Do not use keyboard shortcuts that can close the browser (for example Ctrl + W) or interfere with its operation.
- 2. Limit the access to the computers with the program installed in order to protect the patients' personal data and unauthorized deletion of information.
- 3. Do not give your login and password to the system and license keys to unauthorized persons. Store this data in a safe place.
- 4. If you need to download data from other software, check the connection settings by downloading test data.
- 5. Make sure that the data downloaded from the modality can be viewed in the modes chosen.
- 6. To avoid data loss in the event of a failure of the computer on which the storage is located, back up data regularly.

Before starting a procedure that requires the flawless operation of the Web DICOM Viewer, connect the PC to an uninterruptible power supply and check that it is working properly.

Chapter 1

Login to the Web DICOM Viewer

1.1 Login

Open the user console by the link *http://<ip address>:8090*

After opening the page, the login form will be displayed (Fig. 1.1). When a field is empty, its name is displayed in it. To type data in a field, place the cursor in it with a left-click. To move to the next field, press the **Tab** key on the keyboard or use the left mouse button to position the cursor. To go to the previous field, press the keys **Shift + Tab**.

Sign in
User name
User name
Password
Password 🛩
SIGN IN
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Figure 1.1: Login Form

Type the following data:

- type user name to the User name field;
- type password to the **Password** field.

You can get this information from the program administrator. All fields are required. Click the **Sign in** button.

In case of an error, a message appears in the lower right corner.

To escape the program, click the button with the user's initial in the top right-hand corner. E.g., the button for a user named **User** will be displayed as U . Choose **Exit** in the button menu (Fig. 1.2).

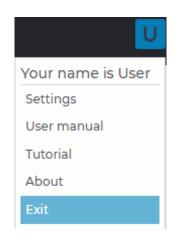


Figure 1.2: Escape the program

1.2 **Tutorial and Usermanual**

When the user first logs in, the window shown in Fig. 1.3 will be displayed.

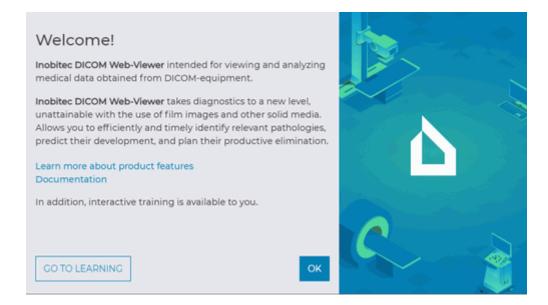


Figure 1.3: Welcome Window

To view the program features, click the **Learn more about product features** link, to open usermanual (pdf file) click the **Documentation** link. To view the tutorial click the **GO TO LEARN-ING** button. To start learning, click the button with the first letter of the user name (e.g. **for User**) and select the **Tutorial** item (Fig. 1.2).

1.2.1 Version of Web DICOM Viewer

To find out the Web DICOM Viewer version, click the button with the first letter of the user name (e.g. **U** for **User**) and select the **About** item (Fig. 1.2).

A window will pop up with the following information about the program (Fig. 1.4):

- Web DICOM Viewer version;
- Link to the website of the software developer;
- Email address of INOBITEC Software FZ-LLC Tectnical Support.

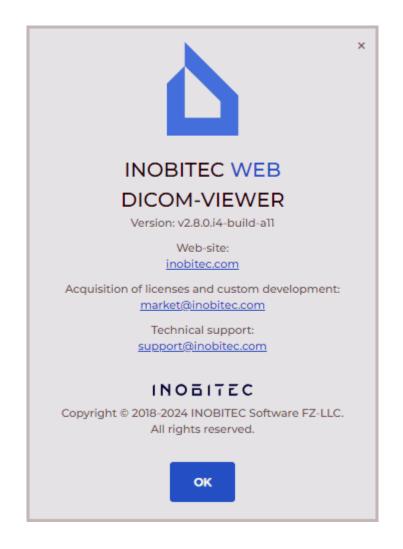


Figure 1.4: Product information

Chapter 2

Web DICOM Viewer Settings

To change settings click the button with the first letter of the user name (e.g.

User) and select the Settings item. The window shown in the Fig. 2.1 will be opened.

for

To apply parameters, click **Apply**, to cancel, click **Cancel**. Always click **Apply** when you change a tab.

2.1 Common settings

Figure 2.1: Common settings

The following parameters are set up in this tab:

- interface theme (light and dark);
- interface language (Russian, English, Portuguese);
- SCU to connect to DICOM Server.

2.2 Hotkeys

The **Hotkeys** tab is used for customizing the tools availability on the toolbar or the context menu, as well as for assigning key combinations to certain tools or actions.

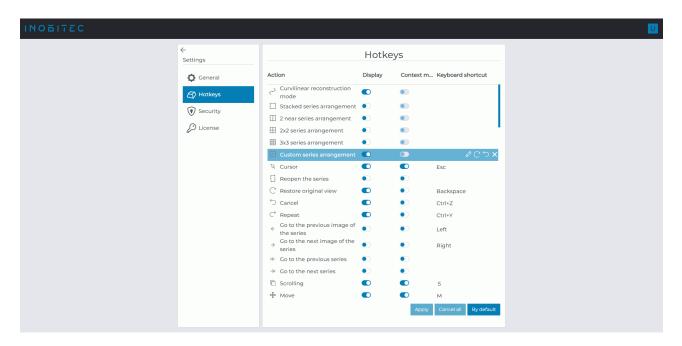


Figure 2.2: Hotkeys

If you want a tool or action to be displayed on the toolbar, put the switch in the **Display** column in the right position. If you want to hide the tool, choose the left position of the switch. If you want a tool to be shown on the context menu, put the switch in the **Context menu** menu column in the right position. If you don't want the tool to be shown, choose the left position of the switch.

In the **Keyboard shortcut** column, you can see the hot keys assigned to the respective tools or actions.

To change or assign a hot key for a selected tool or action, click the **Change hotkey** button and enter the name of the key in the respective field. If the key is already being used for another action, a warning will pop up. To cancel an assigned hot key, click the **Cancel change** button. To restore the default key combination assigned to a tool or action,

click the **Set default settings** button. To delete a hot key assigned to a tool, click the **Remove hotkey** button.

Click **Apply** to apply the settings or **Cancel all** to cancel all the changes. To restore the default tool display settings and hot keys, click the **By default** button.

By default, hot keys are assigned by the administrator for all the users on the **Hotkeys** page of the admin panel.

2.3 Security

The **Security** tab is used to change the user's password.

To change the password, type an old password to the **Old password** field, next type the new password to the **New password** and **Confirm new password** fields. Passwords are hidden during entry.

To save the new user's password, click on the **Update password** button.

2.4 License

If the system is in the personal user licensing mode, the user can:

- See the license key number.
- Delete the license key.
- Activate a new license key.
- See the list of the functions available for the current license key.

Attention! If a license key has been deleted, it cannot be used in future.

If the system is in the competitive licensing mode, in the **License** window, you will only see the list of the functions available for the current license key.

Chapter 3

Program Window Elements

3.1 Toolbar

The Toolbar is shown in Fig. 3.1 (highlighted in red).

IN	0617EC								1
									User storage 🗸
	Name	PID	Sex	DOB	Modality	ACC#	Date	Description	
	Anonymous	Anonymous	0		MR		2019-10-07	L-SPINE	
					Image series				
			15 H	R 15 M	R 15				
Ċ									

Figure 3.1: Toolbar location in the Web DICOM Viewer window

Buttons on the toolbar:



The **Download study** button loads data to the storage.

ò	The Add to archive button downloads studies as archives.
Ū	The Delete study button removes data from the storage.
⊥	The Download from PACS button downloads data to a server.
<u>↑</u>	The Upload to PACS button uploads data to a server.
Q	The Update button searches for studies in the storage or on the server based on the search filter parameters. The list of studies is updated after the search is completed.
∇	The Filter opens the search filter.

Image view buttons on the toolbar:

<u>م</u>	The Image viewer button opens images in the flat image view window.
\$	The Volume reconstruction button opens images in the volume reconstruction window.
₽	The MPR reconstruction button opens images in the multiplanar reconstruction window.
≁∽	The ECG viewer button opens images in the ECG view window.
i.	The Show tags button opens a window for viewing the tags of the series.
Ū	The Remove series button removes the selected series from the storage.

If the action corresponding to a button is not available, the button looks pale and cannot be pressed (inactive). In Fig. 3.2 the only **ECG viewer** button is inactive.



Figure 3.2: The buttons on the toolbar

3.2 Select Data Source

To open studies stored on a DICOM Server or in a local folder, select the source from the drop-down list (Fig. 3.3).

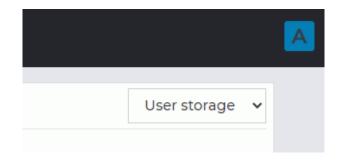


Figure 3.3: Data source panel

3.3 Search filter

To open the panel, click the \checkmark Filter button on the toolbar. The search panel is shown in Fig. 3.4.

Name	Name	nt		PID	
Nume	Name		PID	FID	
Sex	Sex		✓ DOB	YYYY-MM-DD	
Modalit	y	Modality			~
Studi	es da	te			
Exact date			DD		
Interval		YYYY-MM-	DD	YYYY-MM-DD	
O Period		Period			~
 Period 		Period		ply Clear filter C	anc

Figure 3.4: Search panel

The search can be performed by the following parameters:

- patient name;
- patient sex (select from the list);
- patient ID;
- date of birth (is entered manually or select using the calendar, see Section 3.3.1);
- modality;
- study date. You can search by exact date, date interval, or select a period from the list.

The selected search criteria are displayed in the toolbar. To delete a search criterion, click the x symbol next to the criterion or in the corresponding field of the search panel. Click **Apply** to apply parameters, click **Clear filter** to clear the filter or **Cancel** to cancel.

3.3.1 Setting Dates in the Calendar

Dates in the fields **DOB**, **Exact date** and **Interval** can be entered manually (from the keyboard) or by using the calendar in the search panel.

To enter a date using the calendar, left-click in the appropriate field.

Use one of the following methods to specify the date:

1. Use « < » and « > » buttons to choose the month of birth. The months will be spaced at one-month intervals. Use « < < » and « > > » buttons to choose the year of birth. The years will be spaced at one-year intervals. Choose the date (see Figure 3.5).

«	<		lovem er 2021		>	*
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	1	2	3	4	5

Figure 3.5: Calendar drop-down box

Select the area inside the red box (see Figure 3.5). Now, only the current year is displayed (see Figure 3.6). Use « < » and « > » buttons to choose the year with a one-year spacing or « < < » and « > > » buttons to choose the year with a 10-year spacing. Then select the month and the date in the box below.

« ‹	2021	>	»
January	February	Ma	arch
April	May	Ju	ine
July	August	Sept	ember
October	November	Dece	ember

Figure 3.6: Calendar drop-down box. Selecting the month

3. Double-click on the area inside the red box (see Figure 3.5). The current decade will be displayed (Fig. 3.7). Use « < » and « > » buttons to choose the time interval with a 10-year spacing or « < < » and « > » buttons to choose the time interval with a 100-year spacing. Select the year, month and date in the box below.

« <	2021 - 2030	> >>
2021	2022	2023
2024	2025	2026
2027	2028	2029
2030		

Figure 3.7: Calendar drop-down box. Selecting the year

 Click on the area inside the red box once again (see Figure 3.5). The current century will be displayed (see Figure 3.8). Use « < » and « > » buttons to choose the century. Then select the decade, year, month, and date.

¢	2001 – 2100	>
2001 – 2010	2011 – 2020	2021 – 2030
2031 – 2040	2041 – 2050	2051 – 2060
2061 – 2070	2071 – 2080	2081 – 2090
2091 – 2100		

Figure 3.8: Calendar drop-down box. Selecting the decade

To delete the date from the search panel, click the \mathbf{X} symbol in the corresponding field.

3.4 Study Panel

The study panel is shown in Fig. 3.9 (highlighted in red).

D B D ± ± Q ▽ Name	PID	Sex	DOB	Modality	ACC#	Date	Description	User storage
Anonymous	Anonymous	36X	1970-01-01	CT/PT/SR	18	2019-02-07	WB	
				Image series				
et t t et	260 CT 340 PT	263 P	7 263	PT 263	ct 51		A month anonysized	

Figure 3.9: Study panel in the Web DICOM Viewer window

The study panel displays the list of studies stored in the selected location (local storage or PACS server).

3.4.1 Uploading Studies

There are two ways to upload DICOM files to the server:

1. Click the **Download study** button on the toolbar. In the dialog box that pops

up, choose a study or a folder with DICOM studies and click the **Load** button. Click the **Cancel** button to cancel.

 Drag a study or a folder with DICOM studies to the Web DICOM Viewer study panel. When a file or a folder is being dragged, the study panel is replaced by a receive window. Release the mouse button when the cursor is inside the area marked with a dotted line to drop the file or folder.

When uploading studies, you will see a progress bar in the right-hand bottom corner. To interrupt the uploading process, click in the right-hand top corner of the progress bar. If the current DICOM upload process is not completed, no new DICOM files are uploaded by drag-and-drop. In this case, a message is displayed: Wait for download to finish. Try again after the download is complete.

The list of uploaded studies is displayed on the Web DICOM Viewer study panel.

3.4.2 Sort Studies

If two or more studies are displayed, they can be sorted by a certain parameter. To do this, click on the header of the column the parameter is located in. For instance, to sort by date, click on the area highlighted in red in Fig. 3.10.

			User storage 🗸
Modality	ACC#	Date	
MR	0004078851		NEURORADIOLOGIE^Routine
MR/SC	10166710	2020-02-06	Pelvis male
MR/SC	25638412	2019-12-07	Head^Cranial veins

Figure 3.10: Date column header

Next to the column name, there is an arrow showing the sorting order (down arrow — descending , up arrow — ascending). To change the sorting order, click on the column header again. Studies can be sorted only by one parameter at a time.

3.4.3 Downloading Studies

The Web DICOM Viewer allows for downloading studies as archives. The selected study is saved as an archive and may be opened by another viewer program, such as the professional edition of Inobitec DICOM Viewer.

To download a study, proceed as follows:

- 1. On the study panel, select from the list the study you want to download.
- 2. Click the **Add to archive** button on the toolbar. You will see the progress indicator in the right-hand bottom corner while the study is being archived.
- 3. To cancel the archiving process, click the \mathbf{X} button in the right-hand top corner of the progress indicator.

Depending on the browser settings,

- The archive with the study may be saved in the **Downloads** folder, or
- A standard file saving dialog box may pop up.

3.5 Series Panel

The panel is shown in Fig. 3.11 (highlighted in red). Select the study from the study panel to see the list of series for it.

		A
\Box \Box \pm \pm Q ∇	User storage 🕞 👻	
Name PID Sex DOB Modality ACC# Date Descrite Anonymous Anonymous 1970-01-01 C1/P1/SR 18 2019-02-07 WB	ition	
	1 Starting of the second secon	
	iii Mar International Bar John Marin Hannahar Bara 41	
E CT 5001 KM TA WAY. CTAC 3.75 m anonymize B Standard anonymized HB MC anonymized HB MC anonymized TS MC anonymized CT Lungs anonymized Lungs 3 dam anonymize. Dose heard anonym	Dose Report anonymized	

Figure 3.11: Series panel in the Web DICOM Viewer window

3.6 View Images

The Web DICOM Viewer allows you to open a series of one or different studies in all available viewing modes in one tab.

You can open series in the desired mode in 4 ways:



- Double-click on the thumbnails of the series on the series panel. The series will be opened in the **Image viewer** or **ECG viewer** window depending on the modality. If the series cannot be displayed, the DICOM tag viewer window will be opened.
- Right-click on the study and select one of the Open in new tab (the series will be opened in a new tab) or Open in tab -> desired tab. In this case, the first series of the study will be opened in a new window of the tab selected. The series thumbnails will be added to the series panel of the tab.
- Right-click on the series and select one of the Open in 2D, Open in 3D, Open in MPR, Open tags (the series will be opened in a new tab) or Open in tab -> desired tab. In this case, the selected series of the study will be opened in a new window of the tab selected. The series thumbnails will be added to the series panel of the tab.

The series panel will be displayed on the left. It displays studies and thumbnails of the series of the studies opened in this tab. The studies added are highlighted for some time when you go to the tab.

To maximize the study window and display the series thumbnails, left-click on the study on the series panel. You can choose the thumbnails of another study by left-clicking on the name of this study.

To delete a study from the series panel, click the X button in the name of the study. When you delete a study, its series are deleted from the series panel. The windows with the open series from this study are closed. You cannot delete the first study to be opened or its thumbnails. Close the tab with this study to escape.

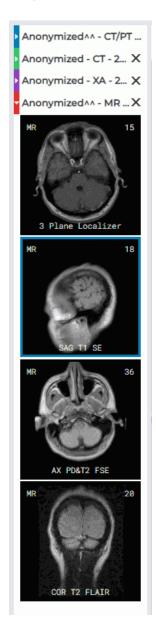


Figure 3.12: Series panel

3.7 View Multiple Series

There are several modes for positioning series windows in the study window. The selected mode is saved when the Web DICOM Viewer is closed.

When you open a new tab, automatic mode is always selected.

If when opening a new series of free cells to accommodate the images is not enough, then the automatic arrangement mode is activated and the required number of cells is added. For example, if only one cell is free, then two more cells are added to open the series in the MPR mode.

3.7.1 Auto Mode

This mode is active by default. If two or more series are open, they will be located in the window and scaled automatically, occupying the entire window. Three series are displayed in Fig. 3.13.

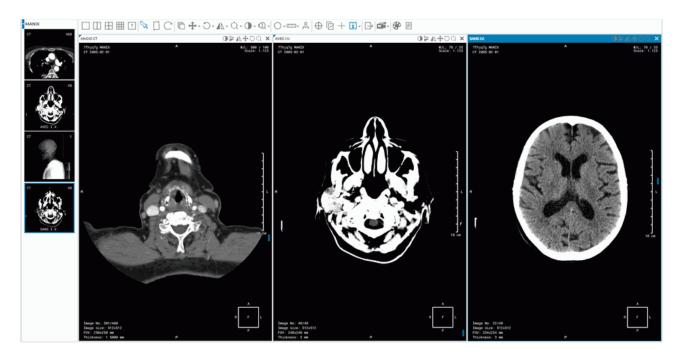


Figure 3.13: Auto Arrangement Mode

3.7.2 Stacked Mode

This tool allows you to receive only left-hand cell in the study window. In this case, the Web DICOM Viewer will request confirmation for closing the current series. If you try to open another series, the Auto Mode will be activated.

To activate this mode, click the

button.

3.7.3 Grid Mode

In this mode the study window is split into parts, and a series window can be opened in each of them.

If all cells in the grid are filled, then auto mode will be activated.

A grid may have from 1 to 25 cells. The maximum number of rows and columns is 5.

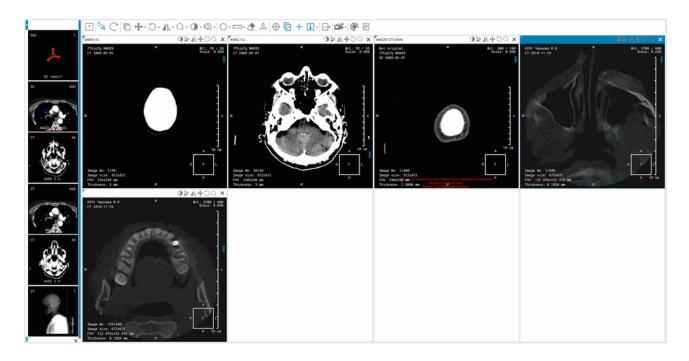


Fig. 3.14 illustrates a $2^{*}4$ grid with three cells filled.

Figure 3.14: Displaying series in a 2*4 grid



To define your own grid configuration, click the

button. drag the cursor over the

panel that opens and select the desired row and column counts (1 to 5). Click the left button in the desired cell to apply the configuration.

?

If the number of already open windows in the tab exceeds the number of cells in the selected configuration, a warning **Are you sure you want to do it? (The number of open series more than the number of cells you want to display)** appears.

To apply a configuration, click **Yes**, to cancel, click **Cancel**. If the number of cells is less than the number of open windows, then windows that do not fit will close. If at least one of the three MPR windows is not enough space, then all three windows for this series will be closed. If the tab has a sufficient number of unoccupied cells, then the newly opened series are reshared in them.

If a configuration from one cell is selected, this is equivalent to activation of the automatic mode.

3.8 Cancel Tool Actions

In the title of each window there are pictograms of the tools available for this window (Fig. 3.15). After the tool is applied to this series, its icon is highlighted in yellow (Fig. 3.16). To undo the tool action, left-click on its icon.

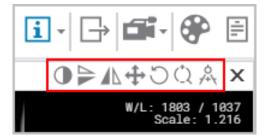


Figure 3.15: Available tools



Figure 3.16: Move and rotate tools are applied

To cancel all the changes made with different tools, click the **C** Restore original view

button on the toolbar.

3.9 Tool Control Buttons

Starting with the version 2.1, you can use the left, the right and the middle mouse button, which allows you to activate up to three tools at the same time. To activate a tool, click on the left-hand side of the button with the mouse button. There are two ways to deactivate a tool:

- 1. Activate another tool with the same button or
- 2. Click on the left-hand side of the tool button with the same mouse button.

On the tool button, you will see a symbol representing the mouse with the respective button selected (see Fig. 3.17).



Figure 3.17: The tools activated with the left, the middle and the right mouse button respectively (from left to right)

If a limited number of clicks is needed to create a graphic object (e.g. three clicks for corner measurements), then the creation process will be completed after these clicks. If you need an unlimited number of points to create a graphic object (e.g. a polygon or an MPR curve), you have to double-click the button with which the tool was activated to complete the creation process.

Whether the tool is active or not, the graphic objects created with this tool can be edited with any mouse button, and the right button click invokes the context menu.

3.10 Context menu

The context menu is available in the DICOM data view window (Fig. 3.18). The menu contains:

- tools available for the window in which it is open;
- series that can be opened in the current window.

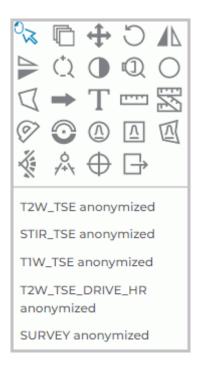


Figure 3.18: Context menu

The tools are activated with the left, right and middle mouse buttons (see Section 3.9).

Chapter 4

View Flat Images



When viewing images, you may see some artifacts that can be attributed to the image compression method. If you see any artifacts, scale the image.



256 shades of gray or color are used to show the tissue density. As there are more than 256 density values, similar densities are represented by the same shade. To differentiate between similar values, decrease the window width and level. Hence, the 256 shades will be used to represent fewer density values.

4.1 View Images in a Series

When a series is opened, its first image is displayed in the series window. There are three ways to switch to other images:

- Roll the mouse wheel up to switch to the previous image or down to go to the next one. One click of the wheel changes the position by one image.
- Use the scroll bar on the right-hand side of the series window. Move the slider along the bar to go to the image required.
- Use the Scrolling tool. Activate the tool by choosing the

button with the left,

right, or middle mouse button. While holding this mouse button, move the cursor down to go to the next image or up — to go to the previous image. You will find the details concerning the use of the tools in Section 3.9.

4.2 Play images

The Web DICOM Viewer allows you to automatically play back images of a series at a certain speed. To set up playback, click on the right part of the **Play** and choose parameters.

- 1. Select the speed (5, 10, 15, 20, or 25 frames per second).
- 2. If you want to play images cyclically, select the **Replay** item. The item will be marked with a flag. To disable the cyclic playback, select this command again. The flag will be removed.

To play the images click on the left side of the **form** button on the toolbar with the left-

/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. To end playback, click on the left side of the button again.

4.3 Zoom. Move. Rotation

4.3.1 Zoom

To zoom an image, you can choose one of the following options:

- 1. Roll the mouse wheel while holding down the Ctrl key. To enlarge the image, roll the wheel forward; to shrink it, roll the wheel backward. The center of the zoomed image will be in the point where the cursor is placed.
- 2. Use the **Zoom** (tool on the toolbar. Click the arrow on the right-hand side of the

Zoom button and choose one of the following options from the drop-down list:

- fixed zoom level: 0.5x, x1, x2, x3, or x4;
- **Custom zoom**. For this, you need to provide the zoom level from 0.001 to 1,000,000,000 in the dialog box. If the value entered is less than 1, the image will be shrunk. If the value is larger than 1, the image will be enlarged;
- **Reset zoom** to fit the image to the screen;
- for simultaneous zooming of all the images in the open windows of the tab, choose Zooming synchronization. The icon representing the active tool will be highlighted in blue. When zooming is completed, click the Zooming synchronization button again to deactivate the tool.
- 3. Activate the **Zoom**, tool by clicking the left-hand side of the (button with the left,

right, or middle mouse button. To work with this tool, use the button with which it was activated. You will find the details concerning the use of the tools in Section 3.9. To zoom the image, move the cursor up or down while holding the mouse button with which the tool was activated. In the window, you will see the point that is the center of the zoomed image. The current zoom level is displayed next to the cursor. When zooming has been completed, click the **Zoom**, button again to deactivate the tool.

4.3.2 Move

To pan an image, activate the **Move** tool by clicking on the \longleftrightarrow button with the left, right,

or middle mouse button. Use the same button to work with the tool. For details on working with tools, see Section 3.9. Drag the image with the mouse, holding the left button.

To restore the initial position of the image, click the arrow on the right-hand side of the **Move** button and select **Position reset**. If you want to change the position of images in all the open windows of the tab, select **Moving synchronization**.

4.3.3 Rotation

To rotate an image, you can choose one of the following options:

1. Use the **Rotation** tool on the toolbar. Click the arrow on the right-hand side of

the Rotation button and choose one of the following options from the drop-down list:

- a fixed rotation angle (a multiple of 90°): 90° clockwise, 180°, or 90° counterclockwise;
- Custom rotation. For this, you need to provide the rotation angle from -1,000 $^{\circ}$ to 1,000 $^{\circ}$ in the dialog box;
- Reset rotation to restore the initial position of the image;
- for simultaneous rotation of all the images in the open windows of the tab, choose Rotation synchronization. The icon representing the active tool will be highlighted in blue. When rotation is completed, click the Rotation synchronization button again to deactivate the tool.
- 2. To rotate an image by an arbitrary angle, click the left-hand side of the Rotation

button with the left, right, or middle mouse button. To work with this tool, use the button with which it was activated. You will find the details concerning the use of the tools in Section 3.9. To rotate the image, move the mouse while holding the button with which the tool was activated. The image is rotated about the center of the window. While the image is being rotated, the current rotation angle is displayed. When rotation is completed, click the **Rotation** button again to deactivate the tool.

4.4 Set Window Level and Width

To change the parameters, activate the Change W/L

tool on the toolbar with the

left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. Move the mouse in the study window, holding the right mouse button

- to increase the window level down;
- to reduce the window level up;

- to reduce the window width left;
- to increase the window width right.

The terms «window width» and «window level» in digital images commonly refer to «contrast» and «brightness» respectively on your computer.

The current window level and width values are displayed in the top right-hand corner of the window.

The contains predefined width and level values for some specific tissues. To select them,

click on the left side of the Change W/L

button. The following modes will be available

in the newly opened window:

- to view abdomen tissues;
- to view bone tissues;
- to view brain tissues;
- to view lung tissues;
- to view chest tissues;
- to view headneck tissues;
- custom mode. The values defined by the user can be edited;
- Reset W/L to set optimal default values;
- W/L synchronization for changing the level and width of all the open windows of the tab simultaneously. The icon representing the active tool will be highlighted in blue. When you have changed the level and width, click the W/L synchronization button again to deactivate the tool.

You cannot edit the values predefined for specific tissues.

4.5 Magnifier

This tool allows you to increase the tissue with a zoom increment of 2 to 100 (Fig. 4.1). To magnify the tissue:

- 1. Activate the **Magnifier** tool by clicking the 1 button.
- 2. Move the cursor to the area you want to make out.
- The radius and the zoom increment of the tool are set in the context menu of the tool.
 To set these parameters, click on the arrow on the right side of the button
 Image: Click on the arrow on the right side of the button Image: Click on the arrow on the right side of

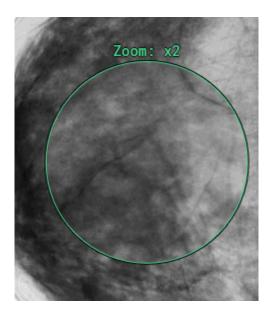


Figure 4.1: Tool Magnifier

4.6 Interpolation Filters

Interpolation may be needed when an image is being transformed in some way. The most common type of transformation is image enlargement.

Depending on the complexity of the image interpolation algorithm, from 1 to 256 adjacent pixels are used for image transformation. The greater is the number of the adjacent pixels included in the process, the more accurate are the enlarged images. This accuracy is achieved at the cost of the time required for image processing.

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When filters are used (except for the Bilinear filter), distortions may occur. If you see any dubious artifacts, select the Bilinear filter.

To change the image interpolation method, you need to refer to the system administrator. Web DICOM Viewer allows for the use of the following filters:

- Bilinear;
- Bicubic;
- Lanczos3;
- Lanczos5.

By default the Lanczos5 filter is active.

4.7 Measurements



For projections (such as x-rays), you have to use the Calibration tool (see Section 4.11) to avoid the mistakes associated with the mismatch between the image size and the real size.



The measuring accuracy is one screen pixel. As screen pixels are smaller than the pixels of the original image, the actual linear measurement error will be equal to the size of the original image pixel. Certain inaccuracy is also associated with density measurements.

The following tools are used to measure various parameters: **Distance measurement ruler**, **Polyline for measuring distances**, **Angle measurement**, **Point value**, **ROI ellipse**, **ROI rectangle**, **ROI polygon**, **Kobb angle**. To select one of these tools, click on the arrow on the right side of the tool selection button. The button will look different depending on the selected tool:

[]	Distance measurement ruler
R	Polyline for measuring distances
Ø	Angle measurement
\odot	Point value
(1)	ROI ellipse
Δ	ROI rectangle
	ROI polygon
×	Cobb Angle

To activate or deactivate the tool that is currently selected, just click on the left side of the tool selection button with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. If some tool is activated, the button is highlighted. The drawn objects will be displayed in the window while it is open, and you can pan, zoom or rotate them together with the image.

4.7.1 Distance measurement ruler

To measure distance in an image:

1. Activate the **Distance measurement ruler I** tool on the toolbar by clicking the left

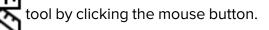
mouse button.

- 2. Click the mouse button on the image to mark the first point and hold down the button.
- 3. Move the cursor over the screen. The distance from the first point to the current point will be displayed next to the line.
- 4. Release the mouse button to fix the current point.
- 5. To cancel an incomplete measurement, press **Esc** on the keyboard.

4.7.2 **Polyline for measuring distances**

To perform polygonal linear measurements:

1. Activate the **Polyline for measuring distances** tool by clicking the mouse button.



tool from the toolbar by clicking the mouse

- 2. Mark the first point on the toolbar by clicking the mouse button.
- 3. Drag the cursor over the screen. The distance from the first point to the current point will be displayed beside the line.
- 4. To fix the current point, click the mouse button.
- 5. Repeat Steps 3 and 4 until the last but one point is fixed.
- 6. Fix the last point by double-clicking the mouse.
- 7. To cancel an incomplete measurement, press **Esc** on the keyboard.
- To move the polygonal measurement point:
- 1. Activate any measurement tool.
- 2. Locate the cursor on the point to be moved.
- 3. Drag the point, holding the mouse button.
- 4. Release the mouse button.

4.7.3 Angle measurement

To measure an angle:

- 1. Activate the **Angle measurement** button.
- 2. Put two points on the image, clicking the mouse button. The second point is the angle apex.

(0,

- 3. Move the cursor over the screen to set the other side of the angle.
- 4. Click the mouse button to fix the second side of the angle.
- 5. To cancel an incomplete measurement, press **Esc** on the keyboard.

4.7.4 **Point ROI measurement**

To measure intensity at some point:

- 1. Activate the **Point value** (on the toolbar.
- 2. Click on the target point.

The point will be highlighted at the image, and the intensity value will be displayed next to it.

4.7.5 Measure the Intensity Average and Standard Deviation in an Area

To measure the average intensity value and standard deviation in a particular area:

- , ellipse (M), or polygon 1. Activate one of the ROI tools (rectangle Л) from the toolbar by clicking the left/right/middle mouse button.
- 2. To draw a rectangle, press the mouse button in its top left-hand and release on righthand corners.
- 3. To draw an ellipse, just draw the rectangle the ellipse is inscribed in.
- 4. To draw a polygon, click the mouse button to mark each apex and then double-click the mouse to finish.
- 5. To cancel an incomplete measurement, press **Esc** on the keyboard.

The following parameters will be displayed next to the highlighted area:

- minimum intensity value;
- maximum intensity value;
- average intensity value;
- standard deviation;
- area perimeter;
- area space.

4.7.6 Cobb angle meter

To measure a Cobb angle:

1. Activate the **Cobb angle** tool from the toolbar.



- 2. Draw a line that runs along the border of one of the vertebrae. To do this, click the mouse button to set two points through which the line must pass.
- 3. Likewise, construct a line for the second vertebra.
- 4. To cancel an incomplete measurement, press **Esc** on the keyboard.

4.7.7 Drawing Parameters

To set the default drawing parameters for measurement tools:

- 1. Click on the arrow on the right side of the measurement tool selection button.
- 2. Select the item Measurement tool settings.
- 3. In the dialog box that appears, set the color, line thickness and the font size.
- 4. Set length and area units (millimetres or centimetres).
- 5. To apply the settings, click **Yes**. To cancel the settings, click **Cancel**.

To edit the drawing parameters of an existing measurement:

- 1. Locate the cursor on the measurement to highlight a line or a point.
- 2. Right-click the mouse.
- 3. Select the **Set render params...** item from the context menu.
- 4. In the dialog box that appears, set the parameters in the same way as the default drawing parameters.

4.7.8 Move Measurements

To move a measurement:

- 1. Activate any measurement tool.
- 2. Locate the cursor on the measurement to highlight the line or point or to magnify the cross, marking the angular point. Do not locate the cursor on angle points.
- 3. Move the measurement, holding the mouse button.

4.7.9 Moving the measurement data

The measurement data and other text information is provided next to lines, indices and highlighted areas. Sometimes the text is illegible; it may overlap the important parts of the image or other text.

In Web DICOM Viewer, you can move the texts placed next to graphics.

To move the text, activate any measuring or drawing tool. Mouse over the text you need to move. You will see a dotted frame round the text, and the text itself will be in bold. Press the mouse button with which the tool was activated and move it to another spot. For the user's convenience, the text is bound to the graphic element or to the respective image area with a dotted line.

When you move a measurement, the text area is also moved.

4.7.10 Delete Measurements

To delete measurement:

- 1. Locate the cursor on the measurement to highlight a line or a point.
- 2. Right-click the mouse.
- 3. Select the **Delete** item from the context menu.

To delete all annotations and measurements, click the **Delete all user overlay** / button

on the toolbar. In the dialog box that appears, Click **Yes** to delete or **Cancel** to cancel. **Please note that all other graphic objects created manually will be deleted as well.**

4.8 Graphic Label Tools

4.8.1 General

This tool allows you to create graphic labels (as arrows, ellipses, polygons and text fields) on an image. The labels will be applied during the entire session of working with the series. The labels are attached to specific points on the image and panned, scaled or rotated together with the image.

The button design depends on the selected tool.

To activate/deactivate the selected tool, click on the left side of the graphic label tool selection button with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. To select and activate one of the tools, click on the right side of the tool selection button and select the tool.

4.8.2 Graphic Label Tool

To create a label on an image:

1. Activate the Arrow tool. The button will look as follows:



- 2. Click the mouse button at the location the arrow should point to and hold it down.
- 3. Release the mouse where the opposite end of the arrow should be. The pointer will be displayed.
- 4. To cancel an incomplete creation, press **Esc** on the keyboard.

To create a text label on an image:

- 1. Activate the **Text** tool. The button will look as follows:
- 2. Click the mouse button where the text should be located.
- 3. Enter text in the field that appears.

4. Click outside the field.

To create a polygon on an image:

- 1. Activate the **Polygon** tool. The button will look as follows:
- 2. Click the mouse button where the first point of the polygon should be located.
- 3. Move the cursor to the location of the next point. The lines connecting the points of the polygon will be displayed on the screen.
- 4. Click the mouse button to save the point.
- 5. Repeat Steps 3 and 4 until the last but one point is set.
- 6. Double-click the mouse to create the last point. The polygon is complete.
- 7. To cancel an incomplete creation, press **Esc** on the keyboard.

To create an ellipse on an image:

- 1. Activate the **Ellipse** tool by clicking the left mouse button. The button will look as follows:
- 2. To build an ellipse, just build the rectangle the ellipse is inscribed into. Click the mouse button where the first angle of the rectangle should be located and hold it down.
- 3. Move the cursor to the location of the opposite angle. The ellipse will be displayed on the screen.
- 4. Release the mouse button to fix the opposite angle. The ellipse is complete.
- 5. To cancel an incomplete creation, press **Esc** on the keyboard.

4.8.3 Actions with Labels

To edit annotations, activate any annotation tool. The Web DICOM Viewer allows you to perform the following actions with labels:

- Drag. Locate the cursor on the figure line or text and drag the object, holding the left mouse button.
- Drag a point (an apex for an arrow or polygon and a rectangle apex for an ellipse). Locate the cursor on a point and drag the point, holding the left mouse button.
- Delete. Locate the cursor on the label border, right-click the mouse and select the **Delete** item from the context menu.
- Set render params. Locate the cursor on the label border, right-click the mouse and select the **Set render params...** item from the context menu. In the dialog box that appears, set the line color, thickness for a figure and size for text. Click **Yes** to apply the changes or **Cancel** to cancel the actions.

To set the default render parameters, click on the right side of the button and select the **Paint tool settings** command. In the dialog box that appears, set the line color, thickness for a figure and size for text. Click **Yes** to apply the changes or **Cancel** to cancel the actions.

To delete all annotations and measurements, click the **Delete all user overlay** *p* button

on the toolbar. In the dialog box that appears, Click **Yes** to delete or **Cancel** to cancel. **Please note that all other graphic objects created manually will be deleted as well.**

4.9 Synchronize Images

Synchronization is applied if two or more series are opened for the same study. This function allows you to scroll images in several series synchronously. By default the option is disabled.

To enable/disable synchronization, click the **Sync series scroll** button on the toolbar.

4.10 Mirror Image Horizontally/Vertically

To flip image horizontally, click the \square button. To flip image vertically, click on the arrow on the right side of the \square button and select the \Longrightarrow Flip Vertically command. If several

series are opened in the view window, the mirroring is activated for each series independently.

4.11 Calibration sizes

X-ray image sizes are increased as compared with the real size of the tissues (Fig. 4.2).

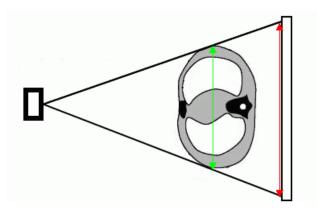


Figure 4.2: The green arrow shows the actual size of the tissues and red arrow shows the size of the image

Hence, the size determined by X-ray modality may be incorrect. If a picture contains an object whose size is known (e.g., a catheter), it is possible to calibrate the image size. To

calibrate the size:

- 1. Click the Reliberation button on the toolbar with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. When the tool is activated, the cursor is displayed as ${}^{igsir}_{igsir}$. To disable the tool, click the $\,\,$ Calibration button again. When the tool is reactivated, the length and the position of the current segment can be edited.
- 2. Mark the first point on the picture by clicking the mouse button and holding it down.
- 3. Drag the cursor over the screen. The calibration interval and its length will be displayed.
- 4. To set the second point, release the mouse button.
- 5. In the dialog box that appears, type the real length of the interval constructed.
 - To calibrate all the images of the series, choose the Series position of the switch. The images from the same study series may differ in the size of pixels. All the images of the series displayed in the current window are calibrated. As a result, the pixel size for all the images of the series is changed to reflect the actual dimensions of the object;
 - To calibrate only the current image, choose the Current image position of the switch.

To calibrate the size, click **Yes**, to cancel, click **Cancel**.

Calibration is factored in for measurements that depend on linear dimensions. After calibration, previously made measurements are automatically adjusted.

Calibration is maintained until the window containing the current series is open. There are two ways to reset calibration:



bration reset:

• left-click on the kicon in the header of the view window (see Section 3.8).

If you reset the calibration, the calibration interval remains in the image.

Visualize Images 4.12

Color look-up tables (CLUTs) are intended for marking pixels characterized by different density with different colors in order to improve perceptibility. The colors you see on the screen do not match the actual colors.

CLUTs button on the toolbar. Select CLUTs in the To change the CLUT click the window that pops up. The current CLUT is marked with a flag. To close the CLUT window click the \mathbf{X} button at the top right of the window.

4.13 Scout Lines

The **Show scout lines** tool is used if two or more series are opened for a particular study. The tool allows you to display the series image scout lines in images from other series if they are synchronized.

The current image scout line is highlighted in green, and the boundary scout lines are highlighted in yellow (Fig. 4.3).

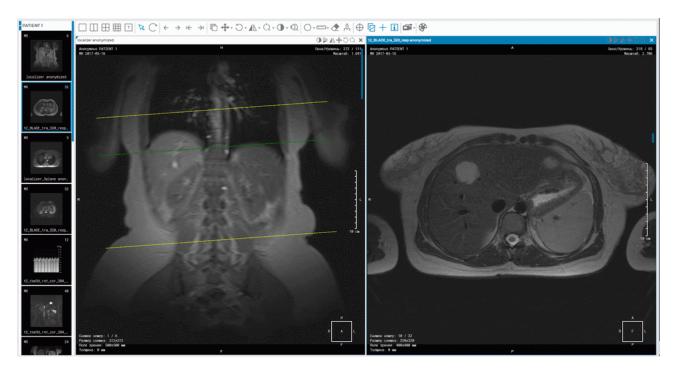


Figure 4.3: The left-hand image scout line is displayed in the right-hand image

4.14 Set Up Viewer Workspace

The **Show/hide annotations** tool allows you to display or hide the information about an image in the series windows. The settings are applied to all the series opened in the current tab. By default, the tool is active. To activate/deactivate the tool, click the **Show/hide annotations**



button on the toolbar.

You can adjust the visibility of:

- 1. Orientation cube;
- 2. Orientation letters;
- 3. Render annotations;
- 4. Tag values.
- 5. Rulers.

To customize the Annotations text size, click the arrow on the right side of the **Show/hide** annotations **b**utton, select the **Label settings** command and in the dialog that opens,

enter a value. Click **Yes** to apply the changes or **Cancel** to cancel.

4.14.1 Orientation Cube

The **Orientation Cube** is located in the bottom right-hand corner (Fig. 4.4). The markers on the cube show the side the image is viewed from:

- A Anterior
- R Right
- L Left
- P Posterior
- F Foot
- H Head.

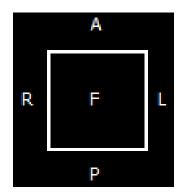


Figure 4.4: Orientation cube

If «F» is written on the cube face, it means that the image is viewed from below; if «R» is written beside a rib, it means that the patient's right side is on this side of the image.

4.14.2 Scale

A graduated scale is displayed on the right side of the window.

4.14.3 Image Information

The image information is displayed in the bottom left-hand corner. It may include the following parameters, depending on the study type:

• Image No — current image number in the series/total number of images in the series;

Show tags

- **Image size** image size (pixels);
- FOV (field of vision) image size (millimeters);
- Thickness interval between slices.

4.14.4 Patient Information

The information about the patient is displayed in the top left-hand corner.

4.15 Viewing PDF documents

Web DICOM Viewer has several modes for viewing PDF documents.

To print a document or copy the text to a text editor, open a PDF document as a text. There are three methods to do that:

- 1. Double-click with the left mouse button on the document thumbnail on the series panel. By default, the document will be opened in a new window as a text.
- 2. Select the PDF document thumbnail on the series panel and click the **viewer** button.
- 3. Open the context menu by right-clicking on the PDF document thumbnail, and choose **Open PDF as text**.

It is more convenient to view PDF documents containing graphs, charts, and pictures as images. If you open a PDF document as an image, you can use **Zoom**, **Move** and **Rotation** tool (see Section 4.3).

To open a PDF document as an image, open the context menu by right-clicking on the PDF document thumbnail, and choose **Open PDF as image**.

There are two ways to view the tags of the series containing a PDF document:

- 1. Select the PDF document thumbnail on the series panel and click the button.
- 2. Open the context menu by right-clicking on the PDF document thumbnail, and choose **Open tags**.

If you choose **Open in tab** -> **desired tab** in the context menu, the study will be added to the series panel in the tab you have selected.

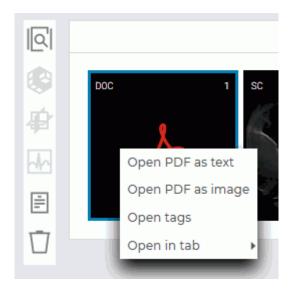


Figure 4.5: Context menu

Figure 4.6 shows a PDF document opened as an image/as a text and the DICOM tags of the series in the Web DICOM Viewer tab.

000 1	× ΩC+∆4 tragente	A DC X	A report.			x
لم			Name Tag	VR	VM	Lengt Text h
Al report	<section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><text><text><section-header><section-header><list-item><list-item><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></list-item></list-item></section-header></section-header></text></text></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header>	<section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header><section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header></section-header>	File-format meta-header (rfer: Li the Endine Explicit) The Endine Explicit) File-MetaInformatic 0002,000 nGroupLength File-MetaInformatic 0002,000 mGroupLength MediaStorageSOPCI0002,000 asSUID TransferSyntaxUID 0002,001 sanceUID SopClessUID 0008,001 SOPClessUID 0008,001 SOPClessUID 0008,001 SOPClessUID 0008,003 SeriesDate 0008,002 StudyDate 0008,003 SeriesDate 0008,003 SeriesDateScription 0008,003 SopStudyDescription 0008,003 SopStudyDescription 0008,003 SopStudyDescription 0008,003 SeriesDescription 0008,003 S	0 UL 1 OB 2 UI 3 UI 2 UI 3 UI 3 SH 5 CS 5 UI 5 CS 6 UI 1 DA 0 DA 0 DA 1 DA 0 TM 1 TM 1 DA 0 DA	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
Exported Series OAT		1 1	PatientID 0010,002 PatientBirthDate 0010,003 PatientSex 0010,004 StudyInstanceUID 0020,000	D LO D DA D CS d UI	1 0 0 1	4 0 26 1.2.410.200051.1547288 816

Figure 4.6: Tab Web DICOM Viewer with an open PDF document

4.16 View DICOM Tags

4.16.1 DICOM Tags. General Information

A DICOM file presents data as a sequence of elements containing information about the image (graph etc.), where the last element has the image (graph etc.) itself. Each element comprises several parts, including:

- Name;
- **Tag** a unique element identifier comprised of a pair of values represented in the hexadecimal number system and determining the group number and the element number;
- Data type (VR) a line showing the data type abbreviation (two symbols);
- The Value Multiplicity of a data element (VM) specifies the number of values that may be encoded in the value field of the data element;
- Data field length;
- **Data text** the information comprised in the element, including the image (graph etc.) itself.

If the **Text** field contains information that cannot be represented in the form of a text (e.g. an image or graph), then the respective column will read: **Binary content. Hidden.**

If the tag is standard, its name will be shown. If the tag is reserved by a certain modality manufacturer or there is an error in its name, then the field will read: **Unknown Tag & Data.** To learn about the data contained in this tag, you will need to study the documents for the modality used in the study.

4.16.2 DICOM Tag Viewer Window

There are four ways to view DICOM tags:

- 1. Select the series thumbnail on the series panel and click the **Show tags** button on the viewer mode panel. The DICOM tag viewer window will be opened in a new tab.
- 2. Right-click on the series thumbnail and select the **Open tags** option in the context menu. The DICOM tag viewer window will be opened in the new tab.
- 3. Right-click on the series thumbnail on the series panel and select the **Open tags** option in the context menu. The DICOM tag viewer window will be opened in the current tab.
- 4. To view the DICOM tags of an open series, click the **Show tags**

button on the

toolbar. The DICOM tag viewer window for the current series will be opened in a pop-up box.

The DICOM tag viewer window is shown in Fig. 4.7.

🚈 😑 Charset: cp-1251 (windows-1251) 🔻 Full						Q. 0028 1/14 🔻 🔺
ame	Tag	VR	VM	Length	Text	
▼ meta-header (xfer: Little Endian Explicit)						
FileMetaInformationGroupLength	0002,0000	UL	1	4	170	
FileMetaInformationVersion	0002,0001	ов	1	2	Bynary content. Hidden.	
MediaStorageSOPClassUID	0002,0002	UI	1	26	1.2.840.10008.5.1.4.1.1.2	
MediaStorageSOPInstanceUID	0002,0003	UI	1	28	1.2.410.200051.1543504284.0	
TransferSyntaxUID	0002,0010	UI	1	18	1.2.840.10008.1.2	
ImplementationClassUID	0002,0012	UI	1	28	1.2.276.0.7230010.3.0.3.5.4	
ImplementationVersionName	0002,0013	SH	1	16	OFFIS_DCMTK_354	
▼ data-set (xfer: Little Endian Implicit)						
ImageType	0008,0008	CS	3	22	ORIGINAL\PRIMARY\AXIAL	
SOPClassUID	0008,0016	UI	1	26	1.2.840.10008.5.1.4.1.1.2	
SOPInstanceUID	0008,0018	UI	1	28	1.2.410.200051.1543504284.0	
StudyDate	0008,0020	DA	1	8	20181129	
SeriesDate	0008,0021	DA	1	8	20181129	
AcquisitionDate	0008,0022	DA	1	8	20181129	
ContentDate	0008,0023	DA	1	8	20181129	
StudyTime	0008,0030	тм	1	6	181108	
SeriesTime	0008,0031	ТМ	1	6	181108	
AcquisitionTime	0008,0032	тм	1	6	181108	
ContentTime	0008,0033	ТМ	1	6	181108	
AccessionNumber	0008,0050	SH	0	0		
• Modality	0008,0060	CS	1	2	ст	
Manufacturer	0008,0070	LO	1	8	PointNix	
InstitutionName	0009 0090	10	1	24	Пиатиостика на Понинском	

Figure 4.7: DICOM tag viewer window

To go to the tags of another image, move the slider along the horizontal scroll bar at the top of the window. Alternatively, you can scroll the mouse wheel when the cursor is on the slider line.

The tag display options may be changed from the drop-down list on the toolbar:

- Full. Set by default. All the columns with tag parameters are displayed in the window;
- **Compact**. All the columns except the column with the tag name are displayed as one (Fig. 4.8). Next to the name of the column, you will see the navigation buttons you can

use to go to the previous	←	and the next	\rightarrow	column.
---------------------------	---	--------------	---------------	---------

When the window is reduced (\leq 768 pixels) e.g. when series are arranged in a grid, or when a mobile device is being used, the tag view mode is automatically switched to **Compact** type. If the window size is >768 pixels, the tag view mode is automatically switched to **Full** type.

		V4
🔚 😑 📔 cp-1251 (windows-125 💙 Compact	✔ Q 0028	1/14 🔻 🔺
lame	\leftarrow	Tag -
▼ meta-header (xfer: Little Endian Explicit)		
FileMetaInformationGroupLength	0002,0000	
FileMetaInformationVersion	0002,0001	
MediaStorageSOPClassUID	0002,0002	
MediaStorageSOPInstanceUID	0002,0003	
TransferSyntaxUID	0002,0010	
ImplementationClassUID	0002,0012	
ImplementationVersionName	0002,0013	
▼ data-set (xfer: Little Endian Implicit)		
ImageType	0008,0008	
SOPClassUID	0008,0016	
SOPInstanceUID	0008,0018	
StudyDate	0008,0020	
SeriesDate	0008,0021	
AcquisitionDate	0008,0022	
ContentDate	0008,0023	
StudyTime	0008,0030	
SeriesTime	0008,0031	
AcquisitionTime	0008,0032	
ContentTime	0008,0033	
AccessionNumber	0008,0050	
Modality	0008,0060	
Manufacturer	0008,0070	

Figure 4.8: Reduced DICOM tag viewer window

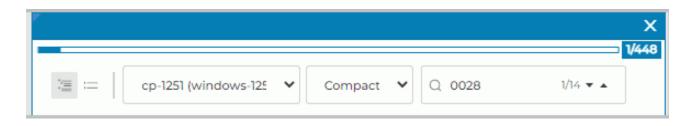
To copy text from the DICOM Tag Viewer Window into a text editor, highlight the selected text open the context menu of the browser by right-clicking on the highlighted text and choose **Copy**.

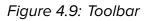
To close the DICOM tag viewer window click the \mathbf{X} button at window.

button at the top right of the

4.16.3 Toolbar

The toolbar is placed in the upper part of the window (Fig. 4.9).





Tools:

1	The Expand all button shows the tag tree.
:=	The Collapse all button is used to collapse the tag tree.
	The Charset drop-down menu allows for the selection of the character set used for displaying data. By default, the character set provided for in the DICOM file is used. If the character set is not mentioned in the DICOM file, windows-1251 character set is used by default. For details see Section 4.16.6.
	The View mode drop-down list allows for the selection the Full or Compact view mode of the tag window. By default the Full mode is activated. When the window is resized, the view mode changes automatically. For details see Section 4.16.2.
Q	Search panel. To perform a search, print some text in the Search field. For details see Section 4.16.5.
\sim	The Next element button. The button is used to go to the next search result.
^	The Previous element button. The button is used to go to the previous search result.
√343	The image indicator shows the current image number and the total number of images in the series. The number is assigned to the image by the modality, and it may be different from the sequence number of the image shown in the Image viewer window.
←	The Switch to previous column button shows the DICOM tags for the previous column in the series.
\rightarrow	The Switch to next column button shows the DICOM tags for the next column in the series.

4.16.4 Tag Tree

The tag tree is placed under the toolbar. If several tags belong to the same group, the group is unfolded and the tags are shown. Groups can be nested, one inside another. To unfold a group, click on the button to the left of its name, to collapse a group, click on the button. To expand all the groups, click on the Expand all button on toolbar; to collapse all the groups, click on the := Collapse all button.

4.16.5 Search

You can perform search by any data contained in the element. To perform search, enter the text you need to find in the **Search...** field. The elements with fully or partially matching values will be highlighted in DICOM tag viewer window. In the search field, you can see the number of the current match and the total number of matches found. To go to the next element found, click the **V Next element** button, to go to the previous element, click the **A Previous element** button.

4.16.6 Changing the Character Set

In the viewer window, the character set provided for in the DICOM file is used for the tags. When working with corrupted DICOM files with invalid character sets, the tags cannot be read. If the character set is not provided in the DICOM file, **cp-1251 (windows-1251)** character set is used by default. Web DICOM Viewer interface provides an opportunity for forced change of the character set.

The **Charset** dropdown list provides for the following character sets:

- cp-1251 (windows-1251);
- latin-1 (ISO-8859-1);
- latin-2 (ISO-8859-2);
- latin-3 (ISO-8859-3);
- latin-4 (ISO-8859-4);
- latin-5 (ISO-8859-9);
- cyrillic (ISO-8859-5);
- arabic (ISO-8859-6);
- greek (ISO-8859-7);
- hebrew (ISO-8859-8);
- utf-8 (utf8).

Choose the character set you need by left-clicking on the respective option.

4.17 Viewing Structured Reports

4.17.1 General Information about Structured Reports

Structured reports (SR) are used for transfer and storage of medical documents. Such documents include both, standard reports in free format and structured data. In the latter case, the accuracy, clarity and usefulness of the documents are improved.

A structured report contains information on the patient and the study performed. Structured reports series have SR modality.

4.17.2 Structured Reports Viewer Window

There are two ways to view a structured report:

- 1. Double-click on the series thumbnail placed on the series panel.
- 2. Right-click on the series thumbnail and choose the **Open in SR** option on the context menu.

The structured reports viewer window is shown in Fig. 4.10.

Dose Record anonymized X								
Charset: windows-1251 V								
X-Ray Radiation Dose Report								
Patient	Study	Report Status						
Patient's name: Anonymous	Study Date: 2019-02-07	Completition Status: COMPLETE						
Patient ID: Anonymous	Study ID: 38	Verification Flag: UNVERIFIED						
Patient's Birth Date: 1970-01-01	Accession Number: 18							
Patient's Sex: -	Referring Physician's Name: Anonymous							
1. Procedure reported: Computed Tomography X	-Ray							
2. Observer Type: Device								
3. Device Observer UID: 1.2.840.113619.6.290								
4. Device Observer Name: pt01	4. Device Observer Name: pt01							
5. Device Observer Manufacturer: GE Medical Systems								
6. Device Observer Model Name: Discovery 710								
7. Start of X-ray Irradiation: 2019-02-07 09:42:12								
8. End of X-ray Irradiation: 2019-02-07 09:56:05								
9. Scope of Accumulation: Study								
1. Study Instance UID: 1.2.840.113619.2.290.3.2831198077.686.1549511545.44								

Figure 4.10: Structured Reports Viewer Window

General information on the patient is provided at the top of the structured reports viewer window. It is arranged in three columns: **Patient**, **Study** and **Report Status**. When the window is reduced (≤768 pixels) e.g. when series are arranged in a grid, or when a mobile device is being used, all the columns with general information are displayed as a vertical option box

(see Fig. 4.11). If the window size is >768 pixels, the structured reports viewer window is automatically switched to full screen mode. To get the details, left-click on the **Patient**, **Study** or **Report Status** option.

Dose Record anonymized				×					
				N					
windows-1251	~	Q No	1/5 🔻 🔺						
X-Ray Radiation Dose Report									
Patient	Pa	atient's name: An	ionymous						
- actoric	Pa	Patient ID: Anonymous							
Study	Pa	Patient's Birth Date: 1970-01-01							
Report Status	Pa	atient's Sex: -							
1. Procedure	repor	ted: Computed T	omography X-Ray						
2. Observer T	Observer Type: Device								
3. Device Ob:	Device Observer UID: 1.2.840.113619.6.290								
4. Device Ob	Device Observer Name: pt01								
5. Device Ob	Device Observer Manufacturer: GE Medical Systems								
6. Device Ob	Device Observer Model Name: Discovery 710								
	2	diation: 2010 02							

Figure 4.11: Reduced structured reports viewer window

In the lower part of the window, structured information is displayed. To copy text from the structured report viewer window into a text editor, highlight the selected text open the context menu of the browser by right-clicking on the highlighted text and choose **Copy**.

To close the structured reports viewer window, click the \mathbf{X} button in the right-hand top corner.

4.17.3 «Search» Field

You can perform search for any data contained in the element. To perform search, enter the text you need to find in the **Search** field. The elements that match the character combination at least partially will be highlighted. In the search field, you can see the number of the current match and the total number of matches found. To go to the next element, click the **V Next result** button. To go to the previous element, click the **Previous result** button.

If the window is reduced, search can only be performed for the visible part of the information — the column displayed in the upper part of the window and the structured information in the lower part.

4.17.4 Changing the Character Set

The character set provided for in the DICOM file is used for structured reports. When the user is working with corrupted DICOM files with invalid character sets, the information from the report becomes inaccessible. If the character set is not provided in the DICOM file **cp-1251** (windows-1251) character set is used by default.

Web DICOM Viewer interface provides an opportunity for forced change of the character set.

The **Charset** dropdown list provides for the following character sets:

- cp-1251 (windows-1251);
- latin-1 (ISO-8859-1);
- latin-2 (ISO-8859-2);
- latin-3 (ISO-8859-3);
- latin-4 (ISO-8859-4);
- latin-5 (ISO-8859-9);
- cyrillic (ISO-8859-5);
- arabic (ISO-8859-6);
- greek (ISO-8859-7);
- hebrew (ISO-8859-8);
- utf-8 (utf8).

Choose the character set you need by left-clicking on the respective option.

4.18 Export series

When exporting, the image is stored in the local storage in a new series of the same study.

To export, click the **Export** button.

Chapter 5

Volume Reconstruction



In the volume reconstruction window, a diagnosis may only be established on the basis of analysis of the series with CT, MR, MG, or XA modality.

The Web DICOM Viewer supports 16-bit image format. If the quality of the original image is higher, some of the information will be lost.

5.1 Model Positioning Tools

To activate/deactivate the tool click the tool button with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9.

The Move tool allows you to model holding down the mouse button. To activate/deactivate

tool click the \longleftrightarrow button on the toolbar.

The Rotation tool allows you to model holding down the mouse button. To activate/deacti-

vate tool click the 💭 button on the toolbar. Note that the initial point of the cursor does not

affect the rotation. If the cursor is moved vertically, the image is rotated round the horizontal axis. If the cursor is moved horizontally, the image is rotated round the vertical axis. That is, the image follows the cursor. If the cursor moves along a slanting line, the image slants the same way.

The **Zoom** tool allows you to model holding down the mouse button. To activate/deactivate tool click the **()** button on the toolbar.

5.2 Model Playing

The Web DICOM Viewer allows you to automatically rotate a volume model around one of the vertical axes relative to the center. The tool is described in Section 4.2.

5.3 **Cutting Tools**

There are cutting tools on the side panel. To activate/deactivate the tool click the tool button with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9.

5.3.1 **Polygonal Cut**

The tool allows you to build a polygon or an area with smooth borders that the cut area is projected on.

To build a polygon:

- 1. Activate the **Polygonal Cut (v**) tool.
- 2. Click the mouse button to set the apices of the polygon marking the figure to be cut, except the last point.
- 3. Fix the last point by double-clicking the mouse.
- 4. To cancel an incomplete building, press **Esc** on the keyboard.

The part of the model to be projected on the area will be cut.

To undo this action, click the Cancel 5 button; to redo the action, click the Repeat



Inverse Polygonal Cut 5.3.2

The tool is similar to the previous one, except that the part of the model not to be projected on the area will be cut. To activate/deactivate the tool, click on the arrow on the right side of



Chapter 6

Multiplanar Reconstruction (MPR)

6.1 View Images

There are two ways to open a series in the MPR reconstruction mode:

- select a study series on the series panel and click the MPR reconstruction button;
- open the context menu for the series thumbnail by right-clicking and select the Open in MPR option.

Fig. 6.1 illustrates the multiplanar reconstruction window.

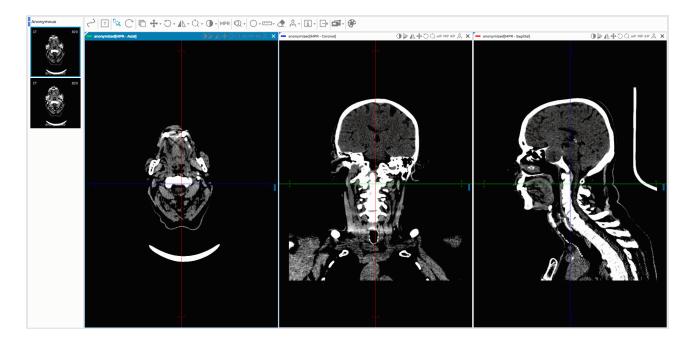


Figure 6.1: Multiplanar reconstruction tab

The series will be opened in a new tab. There will be three image viewing windows: one with the axial plane, one with the frontal plane, and one with the sagittal plane.

View images in the MPR windows is similar to view flat images. To move to the another image, rotate the mouse wheel or use the scroll bar on the right side of the window.

To expand any section plane window of the **MPR reconstruction** tab and restore its size, double-click with the left mouse button on the window header.

6.2 **Reconstruction Modes. Slice Thickness**

6.2.1 Render Modes

To switch between modes, the MPR modes MPR button on the toolbar is used. On the

button, you can see the name of the mode. To change the mode, click the **MPR modes** button and select the mode in the window that pops up (see Fig. 6.2).

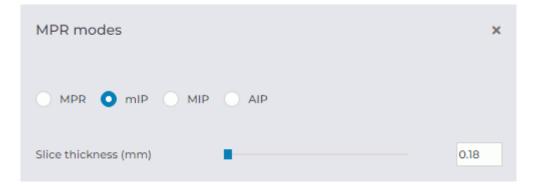


Figure 6.2: The window for reconstruction mode selection

Four render modes are available:

- 1. The MPR mode. Sections are available for viewing. Set by default.
- 2. The **MIP** mode . A slice of a particular thickness is viewed instead of a section. A point with maximum intensity in the slice is projected on each point in the image. For details on how to set the thickness, see the next section.
- 3. The **mIP** mode . Similar to the previous mode, but points with minimum intensity are projected on the image.
- 4. The **AIP** mode. Similar to the previous mode, but the intensity of each point equals the average intensity of the points projected on this point in the image.

6.2.2 Slice Thickness

In the **MIP**, **mIP** and **AIP** render modes, the slice thickness is set using the **MPR modes** window (Fig. 6.2). Click the **MPR modes** button and type the value or move the slider to change the value. In the projection window, the slice borders are marked by dotted lines.

6.3 Working with Orthogonal Planes

Activate the **Cursor** tool with the left/right/middle mouse button. To continue work

with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9. By default the tool is activated by the left button. To move the cutting planes to the target place, click on this place or move the cursor holding down the left button.

To rotate a plane while maintaining orthogonality:

- 1. Locate the cursor on the radial bidirectional arrow on the plane line (see red arrows in Fig. 6.3). The bidirectional arrow will be highlighted.
- 2. Move the cursor, holding the mouse button to rotate the plane by the required angle. The images on the other planes will change.
- 3. Release the mouse button to fix the current position of the plane.

To undo the rotation of all planes, click the **Restore original view** button on the toolbar

. Note that all the other changes made with the tools used for editing images will also be canceled.

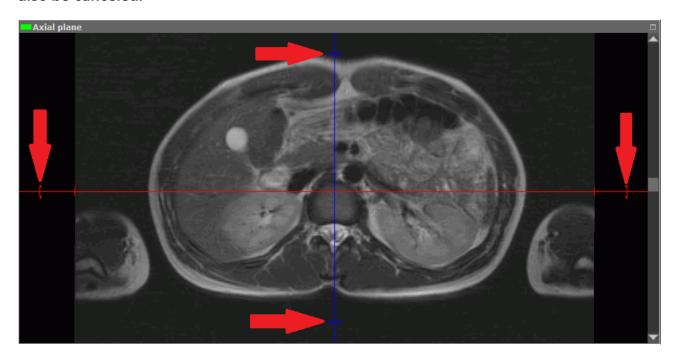


Figure 6.3: Rotate an orthogonal plane

6.4 Curvilinear Reconstruction. Build Surface

Curvilinear reconstruction is a section of tissues by a curvilinear surface whose configuration is set by the trajectory passing through the middle of this surface. This trajectory is called a **curve**. To activate the Curvilinear reconstruction mode, click the **Curvilinear reconstruction mode** button with the left, right, or middle mouse button. To work with this tool, use the button with which it was activated. You will find the details concerning the use of the tools in Section 3.9. When the tool is activated, the cursor looks like

click the **Curvilinear reconstruction mode** button again. When the tool is reactivated, the current curve may be edited.

To build a surface:

- 1. Select the plane in which it would be convenient to start building a curve.
- 2. Roll the mouse wheel to select the slice where the curve will be located.
- 3. Fix the first point on the image by clicking the mouse button with which the tool is activated.
- 4. If necessary, move to another slice by rolling the mouse wheel.
- 5. Move the mouse to select the location of the next point. Fix the point by clicking the mouse button with which the tool is activated. A red curve with a thickness of 1 pixel is drawn between the points.
- 6. Repeat Step 4 and 5 until the last but one point is fixed. While building the curve, the current section will be displayed in the **MPR Curved surface** window (Fig. 6.4).
- 7. Fix the last point by clicking the mouse twice.
- 8. If necessary, correct curve. For details on how to correct, see the section 6.4.1.

To cancel the curve that has not been completed, press **Esc** on the keyboard.

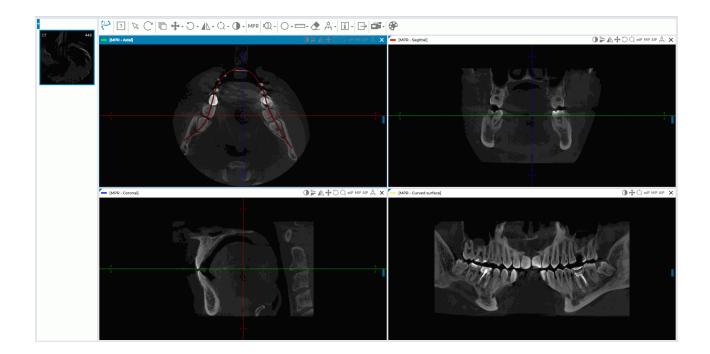


Figure 6.4: Reconstruction by a curve

In the left-hand bottom corner of the window, in the **Curve rotation angle** field, you see the current value of the section rotation angle (in degrees).

There are two ways to rotate a section with a surface in the **MPR - Curved surface** window:

- by rotating the mouse wheel. When you scroll up, the section is rotated by a positive angle. When you scroll down, the section is rotated by a negative angle. One click of the mouse wheel changes the rotation angle by one degree;
- with the help of the scroll bar on the right-hand side of the window. Move the slider up to rotate the section by a positive angle or down to rotate it by a negative angle.

A section with a surface built around the curve may be rotated within the range of -180° to $+180^{\circ}$. Cyclic rotation is available when the **Play**tool is activated with the active **Replay** command. For details see Section 4.2.

To escape the curvilinear reconstruction mode, close the **MPR - Curved surface** window or one of the section plane windows.

6.4.1 Actions with a Curve

Actions with a curve can be performed if the **Curvilinear reconstruction mode** tool, paint tools, or measuring tools are active.

The following actions are available:

- moving a point. Mouse over a point. The point will be highlighted. Move the point while holding the mouse button with which the Curvilinear reconstruction mode tool, paint tool, or measuring tool was activated;
- **deleting a curve**. Mouse over the curve or a point on the curve. Right-click and select **Delete**.

6.5 Measurements

The measurements in the windows of the projections of cutting planes are conducted in the same way as for flat images (see Section 4.7).

The following tools cannot be used in the **MPR - Curved surface** window:

- Curvilinear reconstruction mode;
- Reopen the series;
- Rotation;
- Flip Horizontally/Vertically;
- Magnifier;
- measurement tools (Distance measurement ruler, Polyline for measuring distances, Angle measurement, Kobb angle);
- Synk by point;
- Export;
- Show tags;
- Calibration.

For **ROI ellipse**, **ROI rectangle** and **ROI polygon** tools, the area and perimeter values are not displayed in the **MPR - Curved surface** window.

6.6 Annotations

Annotations are described in Section 4.8.

6.7 Calibration

The procedure of image size calibration in the MPR reconstruction mode is similar to the procedure of image size calibration in the flat image viewer mode, except that you can only calibrate all the images of the series. The procedure of image size calibration is described in Section 4.11.

Chapter 7

ECG Viewer

7.1 View Graphs

If the graphs do not fit on the screen, then:

- pan them horizontally with the mouse, holding the left button;
- pan them vertically by rolling the mouse wheel.

7.2 Toolbar

The toolbar is at the top of the tab (Fig. 7.1).



Figure 7.1: Toolbar

7.2.1 ECG Speed

Two speed values are available: 25 and 50 millimeters per second. The default speed is 50 millimeters per second. The required value can be set using the **«Horizontal Scale ECG»**

7.2.2 Scale

Two scale values are available: 10 and 20 millimeters per millivolt. The default scale is 10 millimeters per millivolt. The required value can be set using the **«Vertical Scale ECG»** -1-1

button.

7.2.3 Length Interval

To select a length interval:

1. Activate the Horizontal Measurement + tool on the toolbar with the left/right/mid-

dle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9.

- 2. Move the mouse along the graph to find the point to start with (it does not matter if the point is located on the left or on the right).
- 3. Click the mouse button to fix the point.
- 4. Move the mouse along the graph to locate the second point. The interval length value will be displayed against the interval background.
- 5. Click the mouse button to fix the point.

To move the interval, locate the cursor on it and drag it holding the left mouse button. Only one interval can be built at a time.

To hide the interval, deactivate the Horizontal Measurement tool.

7.2.4 Value Interval

To select a value interval:

- 1. Activate the **Vertical Measurement** tool on the toolbar with the left/right/middle mouse button. To continue work with this tool, use the button with which the tool was activated. To learn more about tool control, see Section 3.9.
- 2. Move the mouse over the graph to locate the point to start with (it does not matter if the point is located at the top or at the bottom).
- 3. Click the mouse button to fix the point.
- 4. Move the mouse over the graph to locate the second point. The value matching the interval will be displayed against the interval background.
- 5. Click the mouse button to fix the point.

To move the interval, locate the cursor on it and drag it, holding the mouse button. Only one interval can be built at a time.

To hide the interval, deactivate the **Vertical Measurement** tool.

7.2.5 Set Up Coordinate Plane

To change the color scheme, click the **CLUTs** button on the toolbar and in the

CLUTs dialog box select a color scheme. Two color scheme options are available: **classic** (set by default) and **green**.

7.3 Frequency Filters

If an ECG is recorded with distortions, use filters to correct the graphs.



Graphs are changed when filters are applied. To establish a diagnosis, you need to use the original data.

These filters remove the components those frequency is higher than 35 Hz (the 35Hz

Low pass filter 35 Hz button) or 75 Hz (the **75Hz** Low pass filter 35 Hz button). To apply the

filter, click the corresponding button. To cancel filtering, click the corresponding button again. Only one filter can be applied at a time.

To remove power line interference from the ECG signal, use the Notch filter. To apply the filter, click the **Notch filter 50 Hz** button. The tool has current value 50 Hz. If you need

to set a different value, click on the arrow on the right side of the **Notch filter 50 Hz** button and select the **Custom notch filter** item. In the **Render params** dialog box set the value of the frequency in your power supply. To apply the settings, click **Yes**. To cancel the settings, click **Cancel**.

To cancel filtering, click the button again.

To remove the walks of the baseline of the graph, use a High pass filter. To apply the filter,

click the **-Apr High pass filter** button. To cancel filtering, click the button again.

7.4 Export series

When exporting, a series is saved in the filtered form. The series is stored in the local storage in the same study. To export, click the **Export** button.

Chapter 8

Licensing

8.1 Personal User Licenses

A personal user license is registered in the user's account on the **User settings** (see Section 2.4).

The license key activated for a certain user creates one personalized connection providing for continuous access to the system. The user with a personal license has constant access to the system.

8.2 Concurrent licenses

A concurrent license is a license limiting the number of users who can work in the system simultaneously.

The Web DICOM Viewer server automatically provides an unoccupied connection to the user when he/she logs in. The connection provided belongs to the shared unoccupied connection pool. The shared connection pool is a list of unoccupied concurrent connections. The connection stays active during the work session, which ends after the user logs out.

Attention! If there are not enough concurrent connections, the user will be able to work in the system only after another user log out and an unoccupied connection appears. Thanks for selecting our product! The INOBITEC Software FZ-LLC team is constantly working to improve it. We will be grateful for any feedback, comments and suggestions how to enhance the product functionality, user-friendliness and visualization quality.

We wish you success in your work!