



cvi⁴²

User Manual

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Regulatory Information

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FM 539204

Canada

cmr⁴²: Health Canada device license number: 78347

ct⁴²: Health Canada device license number: 87520

cvi⁴²: Health Canada device license number: 93385

United States of America

cmr⁴²: US FDA 510K number: K082628

ct⁴²: US FDA 510K number: K111373

cvi⁴²: US FDA 510k number: K141480

Australia

ARTG Certificate No.: 177785

Sponsor: Market Access Australia Pty Lt
PO Box 1019,
WAHROONGA, NSW, 2076

European Union



cvi⁴² and cmr⁴² are qualified as a class IIa medical device.
It complies with the requirements of the European Medical
Device Directive 93/42/EEC

EC



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Table of Contents

1	Regulatory	5
2	Measurement Accuracy	5
3	Basic Components and Indication for Use	6
4	Customization	7
5	Viewing.....	8
6	Contour Drawing, Labeling and Measurement Tools	10
7	Workspaces	13
8	Patient Data	14
9	Report Module	15
10	Series Overview	19
11	Study Viewer Module	21
12	Multiplanar Reformatting.....	24
13	4D Viewer	27
14	Flow Module	30
15	LV/RV Function Modules	32
16	Tissue Characterization Module	41
17	T2* Module	44
18	T2 Mapping	45
19	T1 Mapping and ECV Quantification.....	47
20	First Pass Perfusion Module	51
21	Vascular Module	54
22	Tissue Tracking	58
23	4D Flow	64
24	Coronary Module (Auto plaque).....	73
25	Calcium Scoring.....	78
26	Aortic Valve Module	80
27	Femoral Module.....	87
28	Mitral Valve Module	90
29	Trans-Septal Module.....	99
30	Technical Support	101

1 Regulatory

1.1 Regulatory Information

AGENCY	CMR ⁴²	CT	CVI ⁴²
US FDA 510K #	K082628	K111373	
Europe CE Certificate #	CE 539277 (issued by BSI)		
Health Canada Device License #	78347	87520	
Australia TGA ARTG #	177785	pending	
Brazilian ANIVSA			094337/14-1
South Korea KFDA	14-1334	pending	

2 Measurement Accuracy

2.1 Measurement Accuracy

All the measurements in the software application are direct transfer of the original DICOM data acquired as a result of image acquisition process. The expected error for the on-screen measurements will be limited to sub pixel matrix size, which divides up each pixel of the input image into a set of sub pixels, i.e.



Depending on the original pixel dimensions and the number of sub pixels used, the accuracy of our measurements will be limited to each sub pixel dimension.

For example, given an original CT or MR image with an isotropic pixel dimension of 0.4 mm and a sub pixel matrix size of 4 x 4, the length (min/max/average diameter) accuracy will be limited to ± 0.05 mm. The accuracy for area and volume will be dependent on the size and complexity of shape, in addition to the sub pixel matrix size. For example, a 4-pixel diameter circle with isotropic pixel dimension of 0.4 mm and 1 x 1 sub pixel matrix will have an accuracy of ± 0.96 mm².



3 Basic Components and Indication for Use

3.1 Intended Use

cvi⁴² is intended to be used for viewing, post-processing and quantitative evaluation of cardiovascular magnetic resonance (MR) images and cardiovascular computed tomography (CT) images in a Digital Imaging and Communications in Medicine (DICOM) Standard format.

It enables:

- Importing cardiac MR & CT Images in DICOM format.
- Supporting clinical diagnostics by qualitative analysis of cardiac MR & CT images using display functionality such as panning, windowing, zooming, navigation through series/slices and phases, 3D reconstruction of images including multiplanar reconstructions of the images.
- Supporting clinical diagnostics by quantitative measurement of the heart and adjacent vessels in cardiac MR & CT images*, specifically signal intensity, distance, area, volume and mass.
- Supporting clinical diagnostics by using area and volume measurements for measuring LV function and derived parameters cardiac output and cardiac index in long axis and short axis cardiac MR & CT images.
- Flow quantifications based on velocity encodes cardiac MR images including 4D flow analysis
- Tissue characterization of cardiac MR Images**
- Perfusion analysis of cardiac MR Images**
- Supporting clinical diagnostics of cardiac CT images including quantitative measurements of calcified plaques in the coronary arteries (calcium scoring), specifically Agatston and volume and mass calcium scores, evaluation of heart structures including coronaries, femoral, aortic and mitral values.
- Evaluating CT and MR images of blood vessels. Combining digital image processing and visualization tools such as multiplanar reconstruction (MPR), thin/thick maximum intensity projection (MIP), inverted MIP thin/thick, volume rendering technique (VRT), curved planner reformation (CPR), processing tools such as bone removal (based on both single energy and dual energy) table removal and evaluation tools (vessel centerline calculation, lumen calculation, stenosis calculation) and reporting tools (lesion location, lesion characteristics) and key images). The software package is designed to support the physician in conforming the presence or absence of physician identified lesion in blood vessels and evaluation, documentation and follow up of any such lesions.

cvi⁴² shall be used by qualified medical professionals, experienced in examining and evaluating cardiovascular MR or CT images, for the purpose of obtaining diagnostic information as part of a comprehensive diagnostic decision-making process. cvi⁴² is a software application that can be used as a stand-alone product or in a networked environment.

The target population for cvi⁴² is not restricted, however the image acquisition by a cardiac magnetic resonance scanner may limit the use of the software for certain sectors of the general public.

cvi⁴² shall not be used to view or analyze images of any part of the body except the cardiac images acquired from a cardiovascular magnetic resonance scanner.

WARNING:

*Quantitative analysis is dependent on the quality and correctness of the image source data.

WARNING:

**The Tissue Characterization and Perfusion modules are not available in the USA since the FDA does not approve the use of contrast agents for cardiac MR procedures.

4 Customization

- 4.1 How to Define a Protocol..... 7
- 4.2 Customize Tools and Working Modules..... 8

4.1 How to Define A Protocol

The definition of protocols allows for setting up a workflow according to standardized image acquisition protocols.

1. Open the drop-down menu next to Protocol selection 
2. Choose *New* and type in the protocol name. Hitting the *return* key will enter the new name and clear the list
3. Go to *Add* and select your modules
4. Rearrange modules via drag-and-drop
5. Rename a module with a double LMB click on the name (e.g. change "Tissue Characterization" into "Infarct")
6. Delete a module with the *Delete* button (bottom right)
7. Automate protocol selection: If you are using standardized image acquisition protocols, you can link your cvi⁴² protocol to the scanning protocol. Open a standard e.g. AMI sequence, set up a post-processing protocol, open the menu and choose "typical". The next time you open an AMI sequence that exact protocol with all the defined pre-sets will pop up
8. Reset typical studies. This removes all recorded study characteristics for the respective protocol and allows to start selecting typical studies from scratch
9. Save the protocol

4.2 Customize Tools and Working Modules



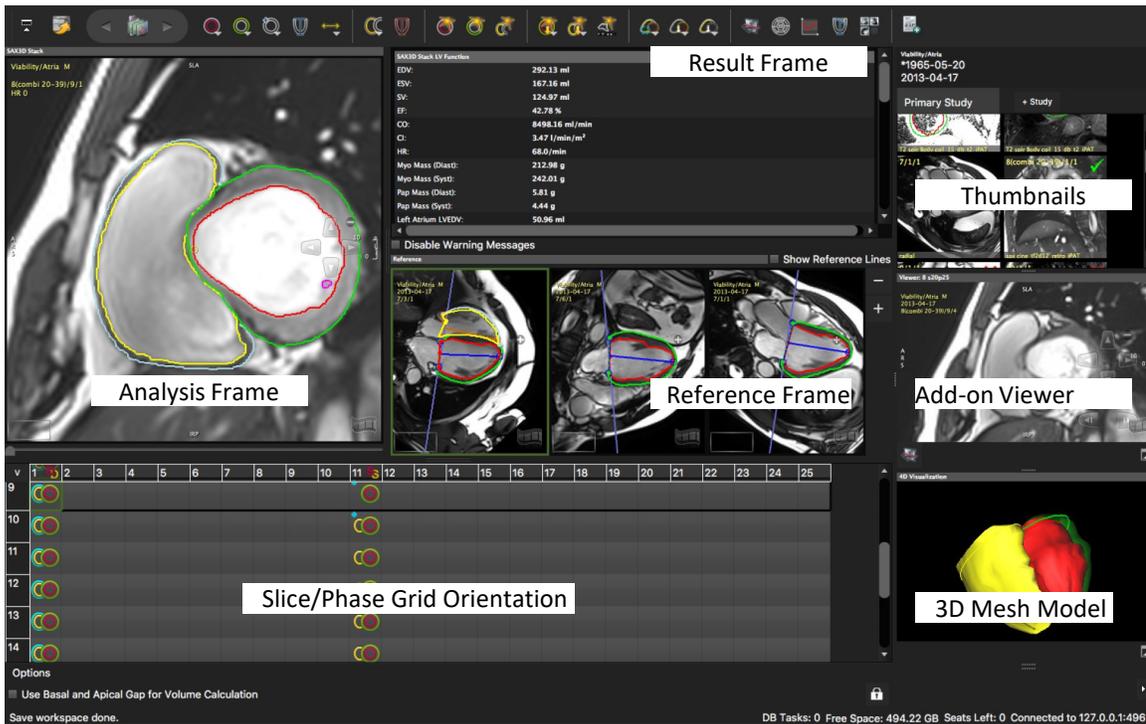
1. In the protocol list, select the module you would like to customize
2. Click on the wrench (bottom left)
3. Everything highlighted in green can be turned on or off. Click the small circular *off* button in the right lower corner of the frame. They will be hidden in all your protocols
4. De-activated functionality is highlighted by a purple frame
5. To reactivate click on the X 
6. To return to default click 
7. Save your selection by clicking the wrench again

5 Viewing

- 5.1 View Ports - Overview 9
- 5.2 Image Editing, Viewer Controls 10
- 5.3 Additional Viewing 10
 - 5.3.1 Floating Viewer 10

5.1 View Ports - Overview

Each module provides viewer frames for image post-processing.
The frames are tagged and prepared for specific image sets (e.g. T2-weighted images)



5.2 Image Editing, Viewer Controls

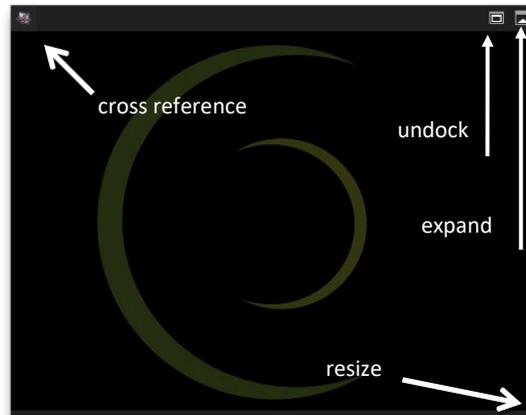
ICON	FUNCTION	MOUSE CLICKS	ALTERNATIVE
	Windowing	Push mouse wheel and drag Auto adjust: double-click MMB Holding shift-key+LMB and drag	Windowing pre-sets: keyboard keys 1-2-3-0 Windowing pre-sets: context menu Define your own pre-set via <i>Edit</i> in the context menu
	Panning	Hold LMB down and pan	
	Zooming	Scroll mouse wheel	Move cursor to the right frame border until symbol appears; LMB drag up/down Context menu: Image/Auto adjust Zoom
	Cine mode		Spacebar
	Navigation	Open on-screen navigation buttons	
	Scroll through phases/slices and series	Hold the shift-key and scroll the mouse wheel	Keyboard arrow keys Customize your mouse wheel: Preferences/Viewer/Mouse Wheel function

5.3 Additional Viewer

5.3.1 Floating Viewer

- Located underneath the thumbnail panel, is an additional detachable viewer that can be undocked and changed in size. It will stay on-screen even when switching to a different module.

-  Expand/Collapse Viewer: Select an image via drag-and-drop from the thumbnail panel
-  Provides a cross reference



6 Contour Drawing, Labeling and Measurement Tools

6.1 Manual Contouring 11

6.1.2 Manual Contour Definition..... 12

6.1.3 Using a Selected Drawing Mode (e.g. by setting points)..... 12

6.1.4 Semiautomatic Contour Detection..... 13

6.1.5 Automatic Contour Detection..... 13

6.1.6 Contour Saving and Deleting 13



- Drawing tools are in the top toolbar or the on-screen pencil box
- Tooltip: All buttons show a brief description on mouse hover

6.1 Manual Contouring

Some tools are hidden by default (as shown below in pink - See Chapter 2)

The toolbar is sorted into tool-sets:



← For the left heart chambers



← For the right heart chambers

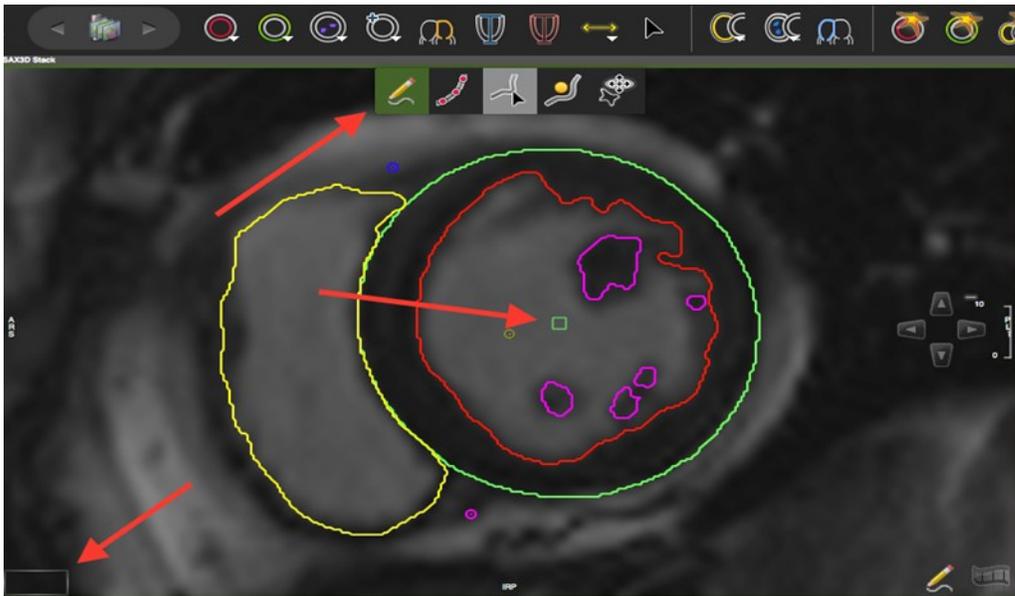


Buttons with a small white triangle indicate that there are more options provided:
 Click and hold the LMB for displaying more options
 To select, drag the cursor to the desired function and release the mouse button

6.1.2 Manual Contour Definition

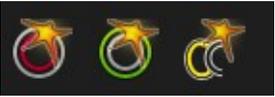
1. Select a tool with a LMB click
2. To draw, click and hold the left mouse button
3. To switch to another tool, simply click on the new tool
4. To drop the tool, double click LMB anywhere within the frame or click ESC

6.1.3 Using a Selected Drawing Mode (e.g. by setting points)



1.  Select a contour (e.g. endocardial)
2. Open the pop-up menu and select the drawing mode:
 -  *Freehand*: default
 -  *Click-Draw-Contour*: sets a series of points for edge definition
 -  *Threshold segmentation*: allows to define borders by dragging with the LMB downwards and sideways until you have found the threshold that best defines your border
 -  *Nudge Contour*: allows to drag part of the contour in a defined direction
 -  *Push Contour*: Shape the contour by pushing the circle
3. Switch back to the freehand mode

6.1.4 Semiautomatic Contour Detection



1. Target your area of analysis (LV or RV) by centering the small yellow circle (target point) in the respective chamber cavity
2. Click the button once to apply a contour detection
3. Click and hold to adjust

6.1.5 Automatic Contour Detection



For automatic contouring based on deep learning please refer to the respective modules (Short 3D, Biplanar, Viewer)

6.1.6 Contour Saving and Deleting

Contour Saving

- Contours are saved automatically

Contour Deletion

- Delete or backspace key
- Context menu
- Shortcut keys (listed in the context menu)

Note: A warning message will be displayed in the results frame (and the exported reports), if:

- Only one phase has been evaluated
- The number of evaluated slices differ between systole and diastole
- Less than 3 slices are used for calculations

At the readers' discretion, warning messages can be turned off. They will re-appear after contour alteration.

WARNING: Automatic Contour Detection

The automatic contour detection provides an initial assumption of contour definitions. It is the responsibility of the user to verify and correct the results.

7 Workspaces

7.1 Workspace Menu

By default, drawn contours are automatically saved and will be reloaded when the patient study is reopened. In addition, the user has the option to:

- *Load Workspace*: Loads a workspace that has been previously stored in a location other than the default location. (“Save Workspace As” option)
- *Save Workspace*: Saves a newly created or updated workspace in the default location
- *Save Workspace As*: Saves the workspace to a different location
- *Load Workspace DICOM*: Loads a workspace that has been previously created as a DICOM Secondary Capture
- *Save Workspace DICOM*: Saves the current workspace as a DICOM Secondary Capture and appends it as an image to the series
- *Import Workspace*: Imports a workspace from a local file system
- *Export Workspace*: Exports a workspace from a local file system
- *Reset Workspace*: Deletes all drawn contours in the current patient study
- *Close Study*

8 Patient Data

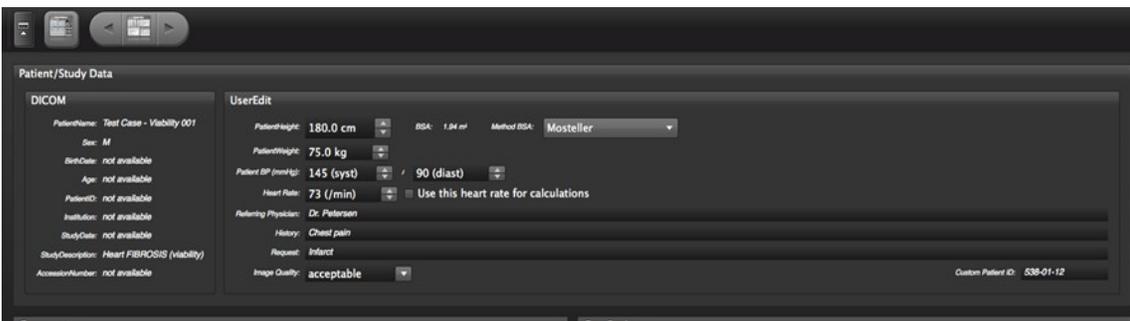
WARNING: Patient Data

The displayed study/patient data is initially derived from the DICOM information if available. Note that editing these values affect the calculations of all modules. It is the responsibility of the user to verify the data before releasing final results.

Note: To change from metric to imperial units go to *Preferences>Appearance>Patient Data Metric*

8.1 Edit Patient Data

1. Edit patient related data necessary for calculation, documentation and reporting
2. Enter patient related comments
3. Create a *Case Review Presentation*: Select your image for review via context menu, Add Current Series to Case Review or ⌘ R.
4. Present Your Case Review: Go to *Overview* and select *Case Review* in the Filter



The screenshot shows a software interface for editing patient data. It is divided into two main sections: 'DICOM' and 'UserEdit'. The 'DICOM' section on the left lists fields like PatientName, Sex, BirthDate, Age, PatientID, Institution, StudyDate, StudyDescription, and AccessionNumber, with many marked as 'not available'. The 'UserEdit' section on the right allows manual input for Patient height (180.0 cm), Patient weight (75.0 kg), Patient BP (145 syst / 90 diast), Heart Rate (73 /min), Referring Physician (Dr. Petersen), History (Chest pain), Request (Infarct), and Image Quality (acceptable). A 'Method BSA' dropdown is set to 'Mosteller'. A 'Custom Patient ID' field at the bottom right contains '538-01-12'.

Note: The software retrieves the heart rate from the first image of a series. A heart rate over 250 bpm will not be displayed. The value remains at 250 bpm.

9 Report Module

9.1	Customize Logo and Header.....	16
9.2	Create a Report.....	16-17
9.3	Update the Report.....	17
9.4	Configure Export Options	18
9.5	Show/Hide Report Fields.....	18

9.1 Customize Logo and Header



Simply type your address and upload a logo

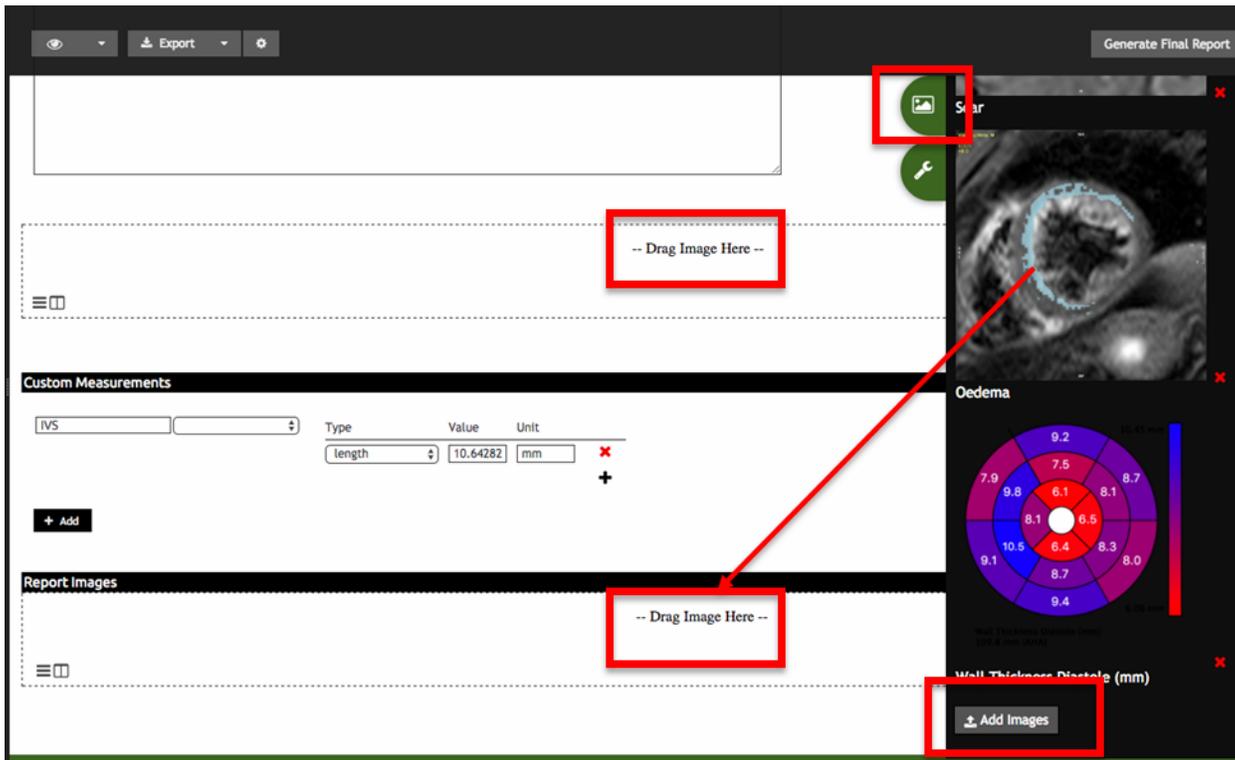
9.2 Create a Report

- *Add to report* will transfer analyses and measurements to the report module
- If gender and birthdate are present, normal values will be classified

	Value	Value / Height	Value / BSA
LVEDV	<input type="text" value="171"/> ml – normal [99 - 199]	96.2 ml/m – normal [60 - 120]	88.34 ml/m ² – normal [53 - 97]
LVESV	<input type="text" value="99"/> ml – increased [17 - 69]	55.57 ml/m – increased [12 - 44]	51.03 ml/m ² – increased [10 - 34]
LVSV	72 ml – normal [68 - 144]	40.63 ml/m –	37.31 ml/m ² – normal
LVEF	42 % – reduced [59 - 83]		
LVCO	5.8 l/min –		3.0 l/min/m ² – Normal [\geq 2.5]
LV Mass	<input type="text" value="178"/> g – increased [74 - 166]	100.19 g/m – increased [47 - 93]	92 g/m ² – increased [42 - 78]
LVESWS	10 ³ N/m ² –		
Global Peak Wall Thickness	<input type="text" value="22.7187"/> mm		

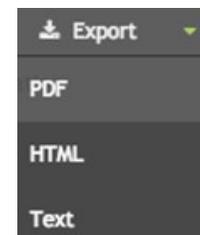
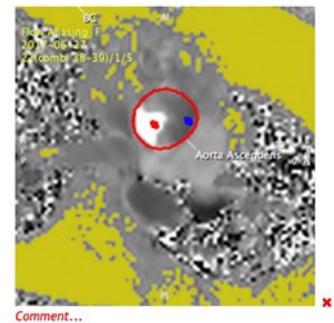


Images collected during the analysis are available in the Image tab and may be added by drag-and-drop into designated “—Drag Image Here—” areas



- *Add Images*: Additional images (e.g. polar maps), can be exported to the desktop and imported into the image pane via the *Add Images* at the bottom of the image pane
- Image captions can be edited from within the image pane, or inline in the report, by clicking on the text directly below the image
- Deleting an image from the report section will return the image to the Image pane

1. Review the report, add findings and write the summary
2. Generate a final report. Once a report has been finalized, a .pdf of this report is stored in settings pane, under History
3. Once the report is done go to *Export* and select the output format. The selected output format will be remembered for future reports



9.3 Update the Report

- Update before finalization, after contour corrections or recalculations: In the module click *Add to Report*. The report should automatically update, it might take a few seconds.
- Update after finalizing the report: Do the same as above and click create addendum.

Note: Any finalized reports or addendums will have a pdf stored under settings pane, *History*.

9.4 Configure Export Options



Click the configuration button (gear icon) to set your Export Preferences:

Export Options ✕

Output Format: **PDF** HTML Text

Indexed Values: Value / Height Value / BSA

PDF, HTML, DICOM

Paper: **US Letter** ↓

Orientation: **Portrait** Landscape

Tissue Diagram: Include If Present
 Polarmaps

Axial Coronary Anatomy: Include If Present

CMR Report Page Theme: **monochrome-light** ↓

Text

Layout: **Tables** Plain

Table Character Width: **70** 80 90 100 110 120

Table Field Padding: **Blank** Dot

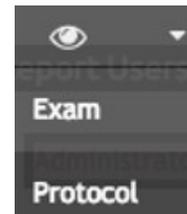
Destination: **Screen** File

Save changes Cancel

9.5 Show/Hide Report Fields

From the drop-down, select what you would like to edit:

- **Exam**: allows to hide sections for the current report (Note that this is the only mode available to non-administrative users and will be used by default)
- **Protocol**: available to administrators, this mode will save the selection to the current protocol



1. Click the eye icon and toggle the field-visibility switch to hide/show report fields. Fields highlighted in red will be hidden from the onscreen and printed/exported reports. Fields highlighted in beige will be visible
2. Click the eye icon again to return to report editing mode

Generate Final Report

Administrator User (admin10001) | Viewer, Reporter, Technician, Analyst | Report Date 27 Feb 2018

Valves and Flow

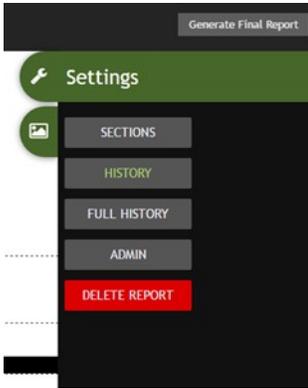
Structure: Pulmonary Artery		Function: [dropdown]		Heart Rate (bpm): 110.7	
Total Forward Flow (ml): 104.8	Forward Flow (ml): 104.8	Backward Flow (ml): 0	Regurgitant Fraction (%): 0	Orifice Area (cm ²): [input]	Gradient: 6.1
Peak Velocity (m/s): 1.2	Regurgitant Fraction (%): 0	Orifice Area (cm ²): [input]	Heart Rate (bpm): 110.7	Gradient: 6.1	

Structure: Aorta Ascendens		Function: [dropdown]		Heart Rate (bpm): 105.6	
Total Forward Flow (ml): 34.9	Forward Flow (ml): 34.9	Backward Flow (ml): 0	Regurgitant Fraction (%): 0	Orifice Area (cm ²): [input]	Gradient: 1.5
Peak Velocity (m/s): 0.6	Regurgitant Fraction (%): 0	Orifice Area (cm ²): [input]	Heart Rate (bpm): 105.6	Gradient: 1.5	

Pulmonary Flow: 11.6 U/min	Pulmonary Artery	Systemic Flow: 3.7 U/min	Aorta Ascendens	Qp/Qs Ratio: 3.1
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9.6 Delete Report

To start with a new report, use the delete report option and the software will create a new report.



10 Series Overview

10.1	Series Overview	19
	10.1.1 How to Open a Series in One of the Analysis Module	19
10.2	Series Composer	19-20

Intended use:

- Get a quick, in-depth overview of sequences and images
- Choosing image/ series for evaluation
- Series Composer sub-unit allows for recomposing studies

10.1 Series Overview

10.1.1 How to Open a Series in One of the Analysis Mode

1. Drag and drop it into one of the modules in the module pane
- or
2. Open the context menu with a RMB click and select the module from here

10.2 Series Composer

The Series composer allows to generate new manually composed series.

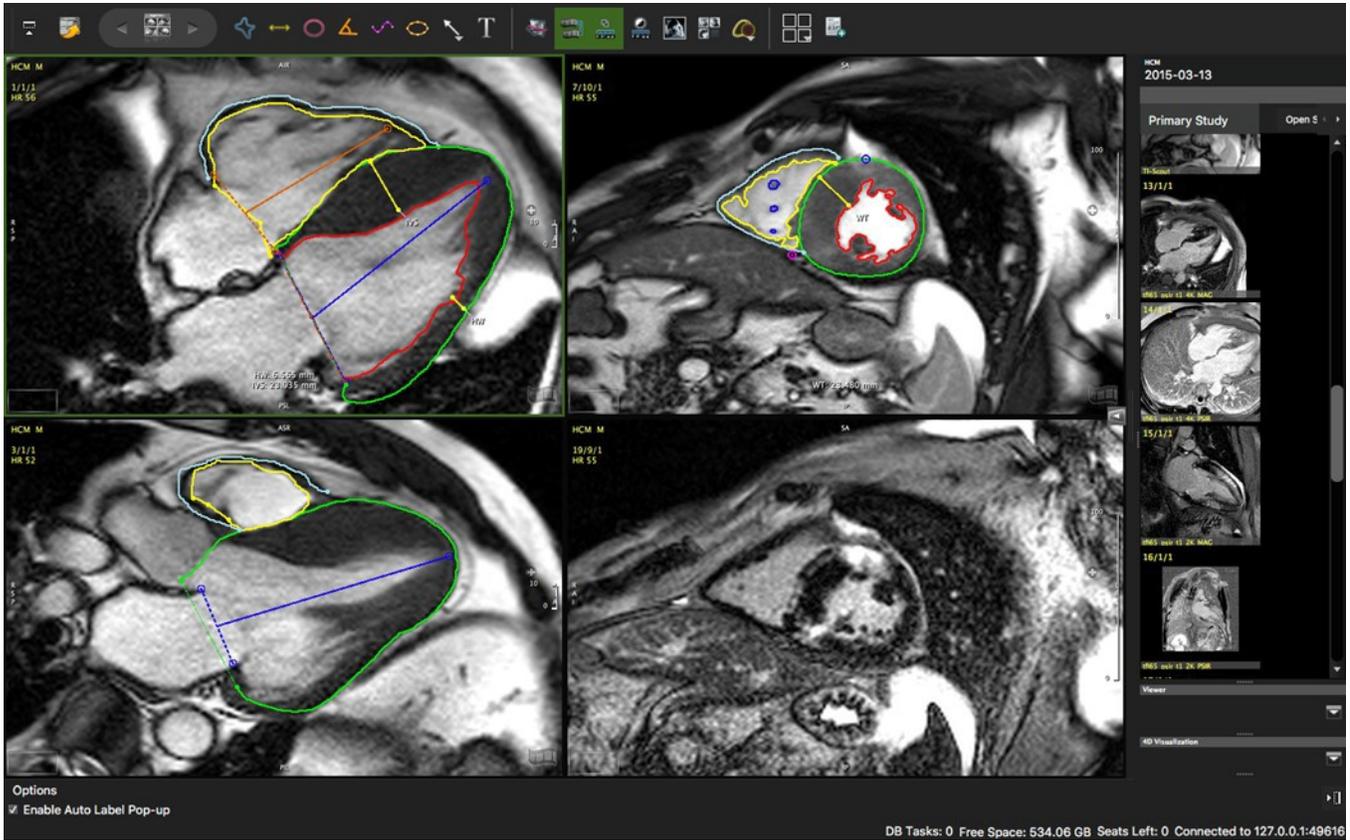


1. Drag a series to the *Browser*, choose slices or phases and drag them into the composer
2. A complete series can be dragged directly to the composer
3. For sequence information (e.g. trigger time or slice location) hover over an image

4. In the *Composer* select the phase/slice or series and use the context menu to remove, insert or sort images
5. Click *Apply* to assemble the newly composed series. The lower window provides a showcase for newly composed series
6. A newly composed series will automatically be appended to the list of series. Manually composed series can be identified by their tag (e.g. 28 (manual 9/1/1))

11 Study Viewer Module

11.1	Custom Measurements.....	21-22
11.2	Compare Follow-up Scans	23

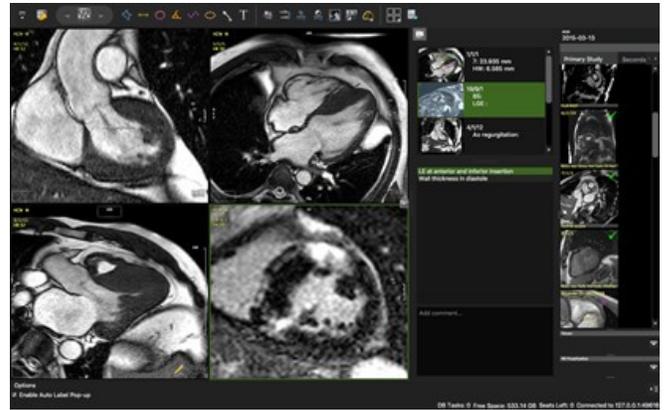


Intended use:

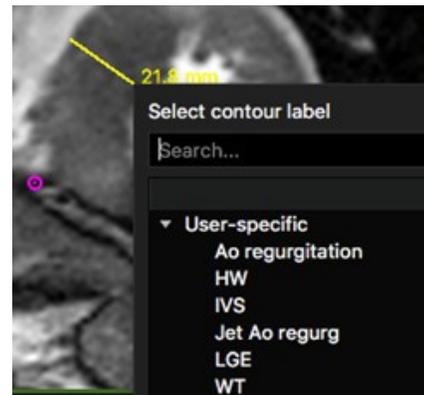
- View and compare multiple sequences next to each other
- View and compare multiple scans of one patient next to each other
- Perform and automatically capture measurements
- Add contour labels and annotations
- Add comments/findings that will be pushed to the report *Findings* section

11.1 Custom Measurements

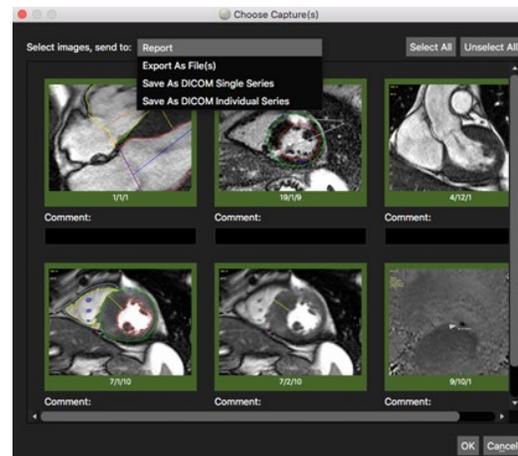
- Open the measurement capture pane



- Perform measurement
- Label the contour
- Image and measurement will be captured automatically in the pane to the right



- Click the small camera to
- Send captures to the report, export or save them
- Optionally add a caption



- *Add Comment* and click enter. Comments will be listed underneath the captures. After adding them to the report, they will show in the findings section



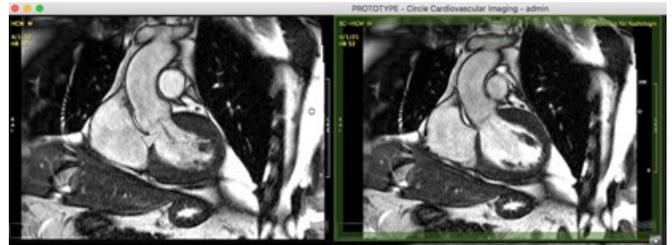
11.2 Compare Follow-up Scans

- Click *Open Secondary Study* in the thumbnail pane
- Select a Filter and type an identifier
- Click on the tabs and drag series in the viewer frame

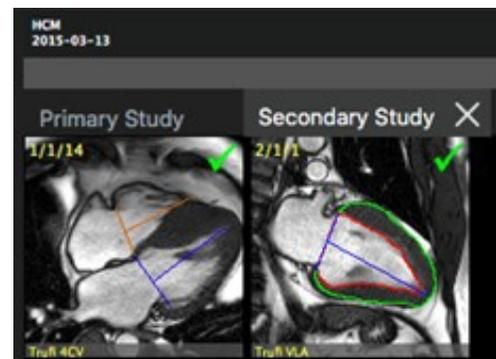


Patient Name
Patient Id
Study Date
Gender
Date of Birth
Modality
Study Description
ALL STUDIES

- The secondary series will be marked by a green frame



- Click the 'X' to unload the series
- It is possible to do a measurement on the secondary series, but they will not be captured



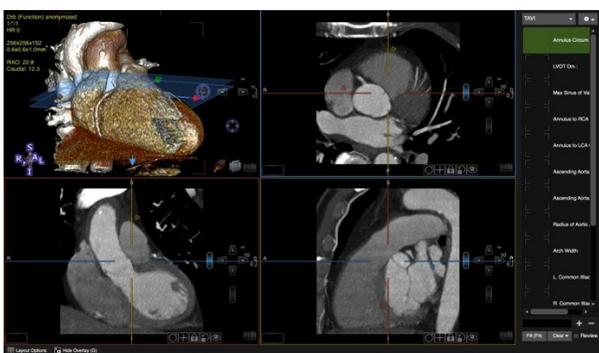
12 Multiplanar Reformatting

12.1	Double Oblique	24
12.1.1	Navigation.....	24
12.1.1.1	Crosshair Navigation.....	24
12.1.1.2	Adjust the Slab-Thickness	25
12.1.2	Reporting and Saving of Image Captures and Measurements.....	26
12.1.2.1	How to Use Multi-Frame Screen Capture (top toolbar)	26
12.1.2.2	How to Save Measurement Captures	26
12.2	Navigation Tab	26

Intended use:

Reconstruct a slice in any position and orientation through the 3D Volume.

The MPR Module provides 2 tabs:



Double Oblique



Free Form Navigation

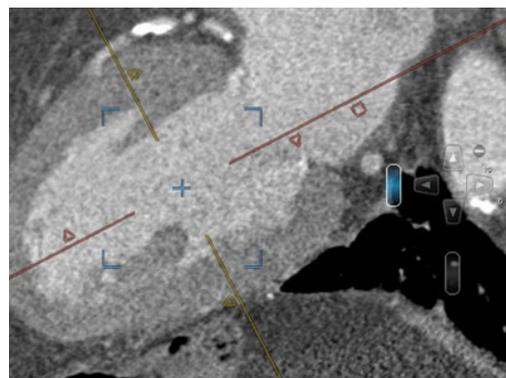
12. 1 Double Oblique

12.1.1 Navigation

By default, a mouse drag in the view port slices through the volume. Alternatively, click the arrow keys.



Use the toolbar button to toggle between panning and slice navigation. Alternatively press the Ctrl (command) key to switch to panning.

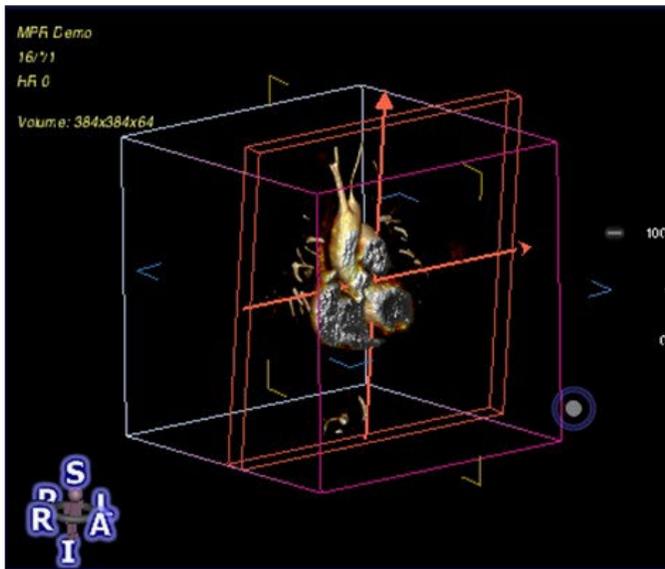
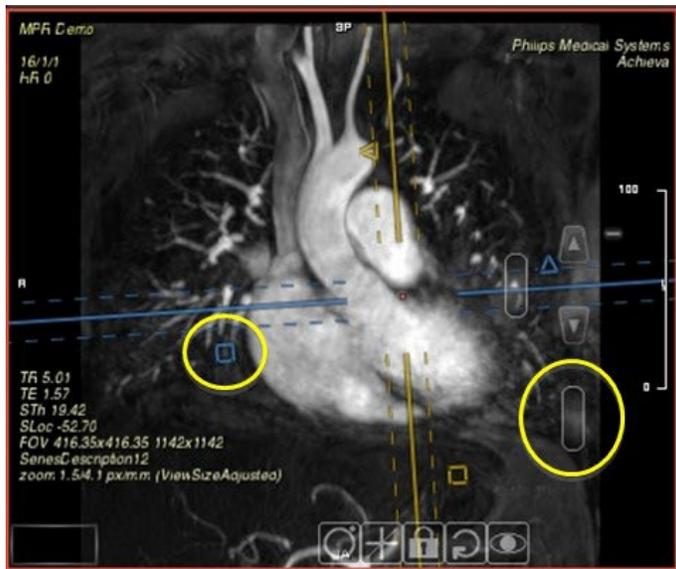


12.1.1.1 Crosshair Navigation

- Two color-coded orthogonal crosshairs within a viewport correspond to the frame with the same color
- Drag the center of the crosshairs to reposition the MPR center. Alternatively, click in a region of interest within the Reference Viewer
- Dragging a crosshair on a triangle will rotate around the MPR center, the lines will always stay orthogonal to each other
- Dragging a crosshair close to the center allows to rotate, and close to the end allows to move it in a vertical/horizontal direction.
- This behaviour will be indicated by icons when hovering over the line
- To move the crosshairs individually, open the context menu and select *Rotate Individual Plane*
- Dragging on a square will adjust the slab thickness

12.1.1.2 Adjust the Slab-Thickness

- A square **icon** next to the crosshairs allows to change the slab thickness by dragging the square
- Another option to change the slab thickness is the **slab-thickness scroll bar** (see screenshot): Dragging the scroll bar with
- LMB increases/decreases the slab thickness using the smallest amount of data increment available for the loaded volume



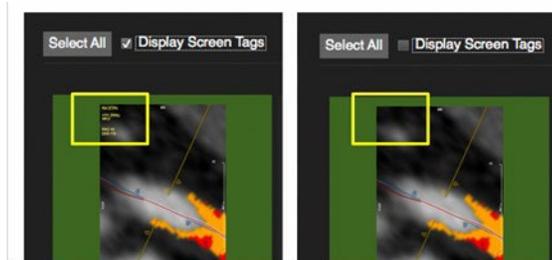
12.1.2 Reporting and Saving of Image Captures and Measurements

12.1.2.1 How to use Multi-Frame Screen Capture (top toolbar)

Add single or multiple images of the current layout to your report.



1. Click the icon in the top toolbar
2. To add images click *Select All* or select single images
3. Choose between displaying or hiding Screen Tags
4. Optionally enter a comment
5. Other options are *Export As Files* and *Save As DICOM*
6. Finally, click the *Add to Report* button
7. Images will show up in the report



12.1.2.2 How to Save Measurement Captures

Add captures to the report.

1. As soon as you have done a measurement the capture will show up in the panel as *Untitled*
2. Drag a label on the capture



3. Click the small camera in the panel to add the capture to the report
4. The captures are part of the workspace and can be reviewed by clicking on the capture
5. Labels will be "auto" saved alongside with the current workflow protocol per user



12.2 Navigation Tab

1. Select the MPR viewer Frame and click the *Navigation* sub-module button
2. Move the MPR center by clicking in your ROI
3. Slice Navigation: left mouse drag or by dragging the slice navigation slider



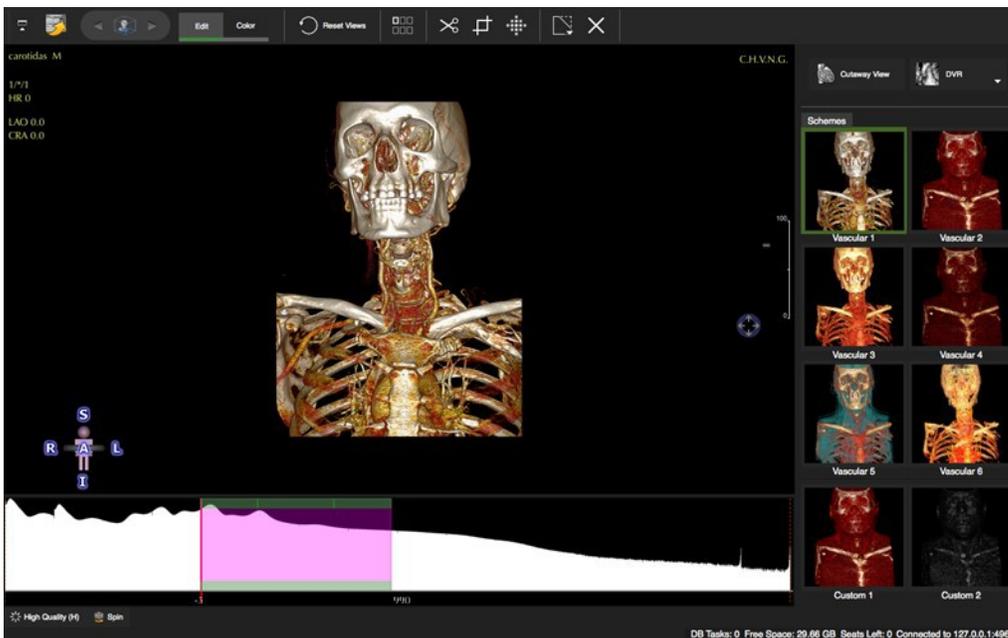
4. Toggle on *Rotate Image* to freely rotate with a mouse drag within the circle
5. When in *Slice Navigation*, you can still freely navigate initiating your mouse drag in the shaded area, of the circle
6. Hold the center of rotation during free-form navigation: Click the "+" at around 11 o'clock or use the modifier key "Shift"
7. When your mouse wheel is set to *Slice Navigation* (Preference->Viewer->Mouse Wheel Function) it is possible to accelerate slice navigation by holding down the Ctrl/CMD key

13 4D Viewer

13.1	Edit.....	27
	13.1.1 Measurement Option	27
	13.1.2 Segmentation	28
	13.1.2.1 How to Do an Automatic Segmentation	28
	13.1.2.2 How to Use Clipping and Cropping	29
13.2	Color	29

This viewer is designed to analyze 3D and 4D data such as MR and CT (cvi⁴² only) angiography.

The *4D Viewer* module provides 2 sub-modules: *Edit* and *Color*



13.1 Edit

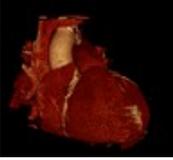
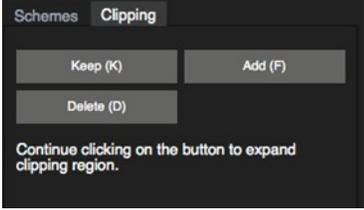
- Drag a 3D, 4D or stacked 2D image data set into the viewer frame.
- The right-side panel offers different rendering options as well as a *Cutaway View*.

13.1.1 Measurement Option

- *Measurement* is an option that can be found in the context menu.
- It allows to set a marker which will indicate the position within the volume that can be viewed across pages and modules. Marker points are automatically saved to a workspace

13.1.2 Segmentation

13.1.2.1 How to Do an Automatic Segmentation

<ul style="list-style-type: none"> • Select the <i>Region of Interest</i> • Select one of four options 	
<p>ROI-Heart:</p> <ul style="list-style-type: none"> • The heart will be segmented automatically 	
<p>ROI - User defined:</p> <ul style="list-style-type: none"> • Activate the Cutaway button (former View Clip) 	
<ul style="list-style-type: none"> • Activate the 3D Region Grow Clip 	
<ul style="list-style-type: none"> • Point the cursor in the middle of your ROI and drag 	
<ul style="list-style-type: none"> • When you are done, go to the clipping pane and click Keep 	
<ul style="list-style-type: none"> • Turn off the grow button 	
<ul style="list-style-type: none"> • Optionally, add a background: Context Menu>Rendering>Background 	
<p>ROI-LV:</p> <ul style="list-style-type: none"> • Repeat steps 1-6 	
<ul style="list-style-type: none"> • No Region of Interest • Will display the original volume 	

13.1.2.2 How to Use Clipping and Cropping

1. Select the tool from the tool bar and select a drawing mode from the tool pop-up menu (default=freehand)



2. *Cut*: Drag the mouse cursor to encircle structures you would like to remove



3. *Crop*: Drag the mouse cursor to encircle things you would like to keep



4. *Grow*: Move the cursor over the region of interest. A black dot will indicate the software has detected a structure that can be segmented. A red square will appear over a structure that has already been segmented. Dragging the cursor downwards will define the region of interest according to the SI threshold. Dragging the cursor sideways will increase the threshold, taking more SI into account.



5. To rotate or pan the volume while in clipping mode, move the mouse over the icons and drag

13.2 Color

- **Ambient** Light: simulates surrounding light
- **Diffuse** Light: light that reflects into all directions
- **Specular** Light: simulates the light reflection on the surface depending on the vantage point
- **Shininess**: increases the highlights

Color Schemes: cvi⁴² provides 6 predefined color schemes (vascular 1-6) and 2 custom schemes

14 Flow Module

14.1	Flow Correction.....	30
14.1.1	How to Do a Flow Correction	30
14.2	Flow Quantification.....	30
14.2.1	How to Do a Flow Analysis.....	30-31
14.2.1.1	How to Display Flow Curves.....	31

Flow analysis (Only through plane velocity encoding)

Intended use:

- Color-coded flow display with semiautomatic contour detection, synchronization and forwarding
- Automatic calculations of flow parameter
- Shunt assessment (Qp/Qs)

14.1 Flow Correction

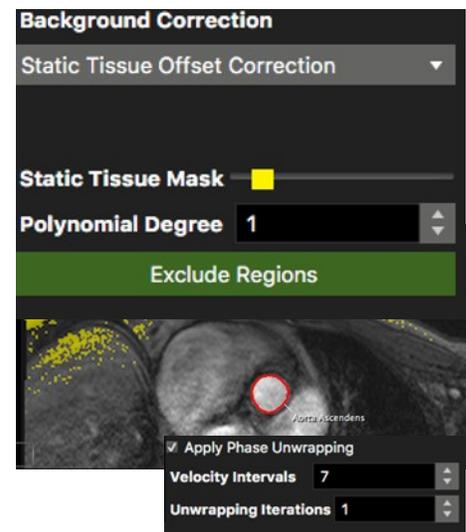
14.1.1 How to Do a Flow Correction

1. Drag the series on a viewer frame. Magnitude and Phase images will be loaded automatically into their respective frames.
2. Draw your ROI contour around the vessel of interest
3. Apply *Corrections* by checking the box.

To change the correction settings, click *Advanced* under *Correction Settings*. This will open the *Background Correction* options:

- **Offset Correction:** incl. *Exclude Region* e.g. spatial aliasing
- **Phantom Correction:** Drag the phantom image in the box
- **Background Correction (BC):** Click on *BC*, draw and forward (use context menu command) a ROI in an area with no flow
- **Apply Phase Unwrapping** (anti-aliasing): Contour the vessel for all phases, then apply phase unwrapping by selecting *Apply Phase Unwrapping*

Any update to the ROIs or background correction settings will reapply both background correction and phase unwrapping.



14.2 Flow Quantification

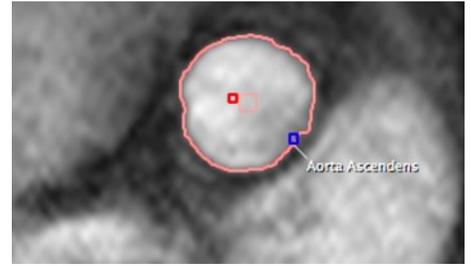
14.2.1 How to Do a Flow Analysis

1.  For color encoding click on the *Toggle Flow Overlay* button
2. To reduce noise, the threshold color overlay can be adjusted;

3.  Define the vessel border
4. Label the contour



5. Forward the contour. The automatic contour detection will grow the
6. contour towards the vessel borders based on the information from all phases. It also forwards the contours to all phases
7. Scroll through phases to ensure the contours are correct
8. *Add to Report*: Flow assessments of different series have to be added individually to the report



14.2.1.1.1 How to Display Flow Curves



1. Use the color-coded checkbox to toggle flow curves on/off
2. Invert Results in case of wrong encoding directions
3. Shift the curve on the time axis for better viewing
4. Adjust the triangles on the bottom of the graph to calculate flow within a set time range (inside ruler)

Measurement	Total	Inside Ruler
Total Forward Volume:	27.52 ml	27.16 ml
Total Backward Volume:	-0.49 ml	-0.06 ml
Total Volume:	27.03 ml	26.31 ml
Regurgitation Fraction:	1.78 %	0.24 %
Vol/min:	2906.43 ml/min	
Vol/min (effective):	2854.72 ml/min	
Heart Rate:	105.60 /min	

5. Hover over a data point to view the numbers
6. A right mouse click in the graph opens a context menu, that offers several display options and an option to export the flow curves



7. Add to report. Flow assessments of different series have to be added individually to the report

15 LV/RV Function Modules

15.1	Bi/Triplanar Module.....	32
	15.1.1 How to Do Bi-triplanar LV Function and Volume Analysis.....	33
	15.1.2 How to Do Automated Atrial Volumetry	33
15.2	Multiple Long LAX Module	33-34
	15.2.1 How to Do LV Function with Multiple Radial Long Axis	34
	15.2.2 How to Create a Ventricular Volume/mass Curve	34
15.3	Short 3D Module.....	35
	15.3.1 Short 3D Interface	35
	15.3.1.1 Thumbnail Grid	36
15.3.2	Global LV function and volume analysis.....	36
	15.3.2.1 How to Do Fully Automatic LV/RV Segmentation	36-37
	15.3.2.2 How to Do Automatic LV Segmentation in a User Selected Phase.....	37
	15.3.2.3 How to Do Auto. Endo. RV Segmentation in a User Selected Phase...37	
	15.3.2.4 How to Do a Mitral/tricuspid Valve Plane Correction.....	38
	15.3.2.5 How to Do Manual Phase Shifting Per Slice.....	38
	15.3.2.6 How to in/exclude Papillary Muscles	39
	15.3.2.7 How to Create and Display a Volume Curve	39
	15.3.2.8 How to Do a Regional Wall Motion Analysis	39-40
	15.3.2.9 How to Do Atrial Volumetry.....	40
	15.3.2.9.1 How to Do Atrial Volumetry in Short/Long Axis Orientation.40	

15.1 Bi/Triplanar Module

Intended use:

- LV function analysis biplanar in 2CV and 4CV or triplanar by adding a short axis view
- Atrial volumetry in one or 2 planes

WARNING: Bi/Triplanar

For the calculations, the diastolic and systolic phase are assumed to be in the same phase of the cardiac cycle in all three views. The volume calculation is done per phase. The smallest volume is assumed to be the systolic volume and the largest volume is assumed to be the diastolic volume. In case a volume can be calculated for one phase only, ESV and EDV will be displayed as equal and no calculations will be performed for derived parameters such as ejection fraction.

Endocardial contours must always be on the inside of the epicardial contour, cvi⁴² does not check for that.

Calculations in cvi⁴² are performed based on the current state of contour definition and image selection. For every change the calculations are updated immediately. It is the responsibility of the user to decide whether the stage of contour definitions reflects the intended measurement task.

15.1.1 How to Do Bi-triplanar LV Function and Volume Analysis

1. Drag and drop the series (2CV, 4CV) from the thumbnail panel into their respective frames and adjust viewer properties
2. Select the end-diastolic phase



3. **LAX LV Extent** contour: Set two points in the mitral valve plane, and a third point in the apex to define the length of the ventricle. As soon as the apex point has been placed, the endocardial contour will be detected automatically.

Correct if necessary

4. Draw an epicardial contour
5. Do the same for the end-systolic phase



6. **Segment Myocardial Contours LAX in Slice**: Optionally propagate endo- and epicardial contours for a myocardial segmentation in all phases



7. Add analysis to Report

15.1.2 How to Do Automated Atrial Volumetry

Mono or bi-plane area-length method.

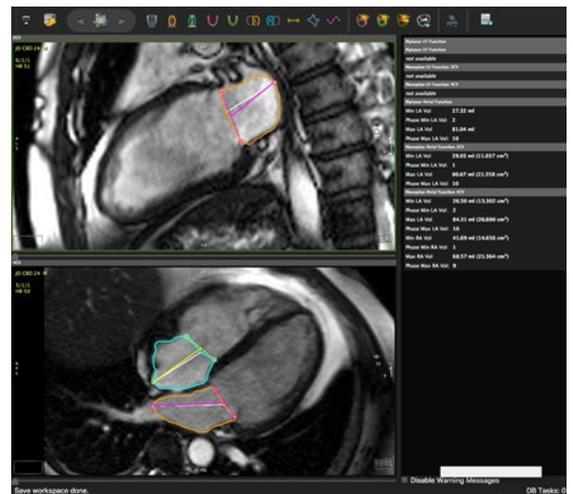
1. Load the 2CV and the 4CV in the respective frames



2. Click Detect LA/RA Contours Current Slice
3. Check contours
4. Minimum and maximum volume will be reported mono- and biplane



5. Add to Report



15.2 Multiple Long LAX Module

Intended use:

LV function volume and mass analysis.

WARNING: Multiple Long

The input for multiple long calculations requires a multi-slice series with long axis slices that are acquired along a central rotation axis. cvi⁴² checks for the presence of these requirements, although it allows some variance to avoid small aberrations being rejected from the analysis. It is the responsibility of the user to verify the cutting line of the slices and decide whether the variance in the rotation axis is acceptable.

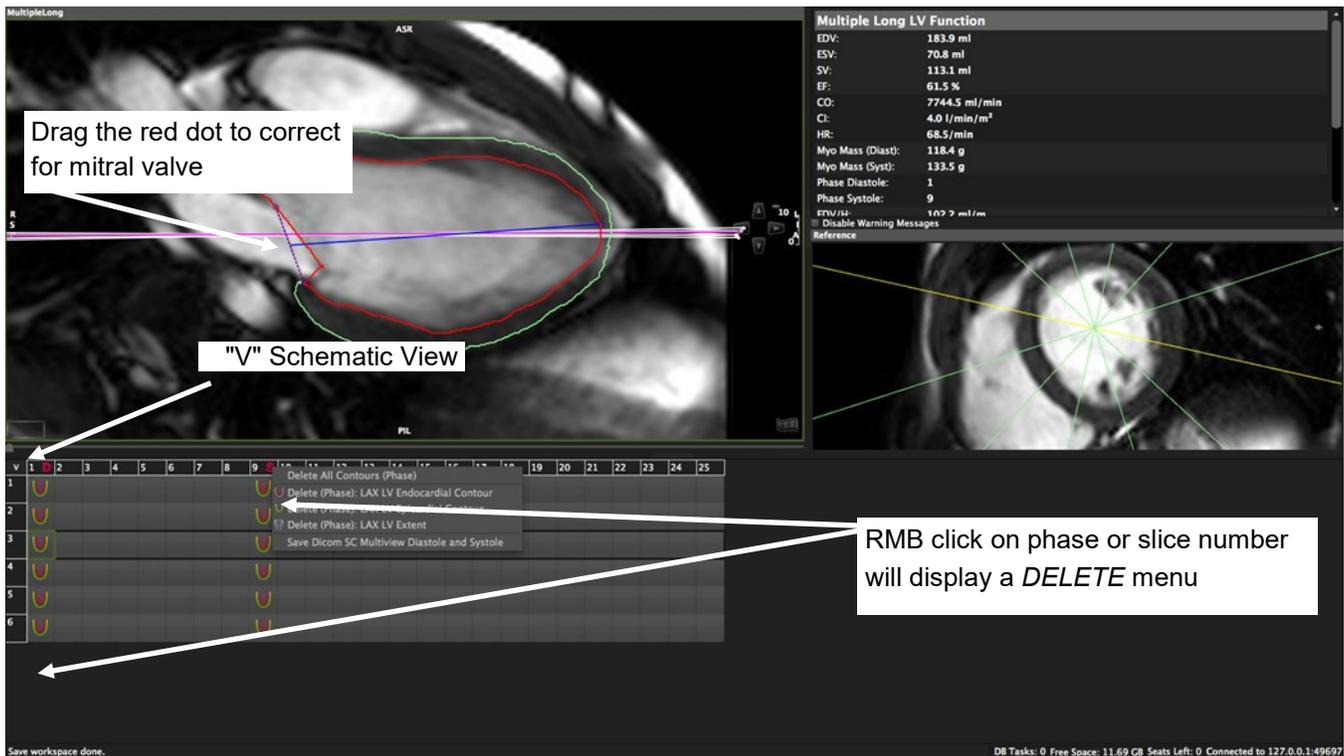
Depending on the shape of the left ventricle the method in use might not be suitable for all evaluations (e.g. severe regional wall motion abnormalities), it is the user's responsibility to verify the suitability of the selected method.

Endocardial contours must always be within (i.e. on the luminal side of) the epicardial contour, cvi⁴² does not check for that.

The volume calculation is performed for every phase. The smallest volume is assumed to be the systolic volume and the largest volume is assumed to be the diastolic volume. If the volume can only be calculated for one phase, ESV and EDV will be displayed as equal and no further calculations are done.

Missing contours are interpolated, if possible.

Calculations are performed based on the current state of contour definition and image selections. For every change in this state the calculations are updated immediately. It is the responsibility of the user to verify that the contours reflect the desired measurements before releasing final results.



15.2.1 How to Do LV Function with Multiple Radial Long Axis

1. Drag the radial long axis series from the thumbnail panel into the respective frame and adjust viewer properties. A perpendicular short axis cine series (if available) will automatically be displayed in the reference window
2. Clicking on the "V" in the left top corner switches to a schematic view to help you find the right slice and phase
3. Visually select diastole and systole



4. Define the extend of the left ventricle: Set two points defining the mitral valve plane and a third point in the apex to define the length of the ventricle, the endocardial contour will be detected automatically. Correct endocardial contour
5. Adjust the valve plane by dragging the red dot



6. Draw an epicardial contour manually or semiautomatic



7. Optionally propagate the contour use the toolbar button *Segment Myocardial Contours LAX in Slice* button
8. Repeat for all slices



9. Add analysis to Report

15.2.2 How to Create a Ventricular Volume/mass Curve

1. Define endocardial and epicardial contours in all phases and slices



2. Click LV volume curve

Peak Ejection and Filling Rate and Peak Wall Thickness can be found in the scientific report (Report Module)

15.3 Short 3D Module

Intended use:

Using the disc-summation technique this module can analyze a stack of parallel images in long, short or axial orientations:

- Global and regional LV function and volume analysis
- Global RV function analysis
- Atrial volumetry

WARNING: SA 3D

The volume calculation is done for phases. The smallest volume is assumed to be in systole and the largest volume is assumed to represent diastole. In case a volume can only be calculated in one phase, ESV and EDV will be displayed as equal and no further calculations are done.

Missing contours are interpolated, if possible.

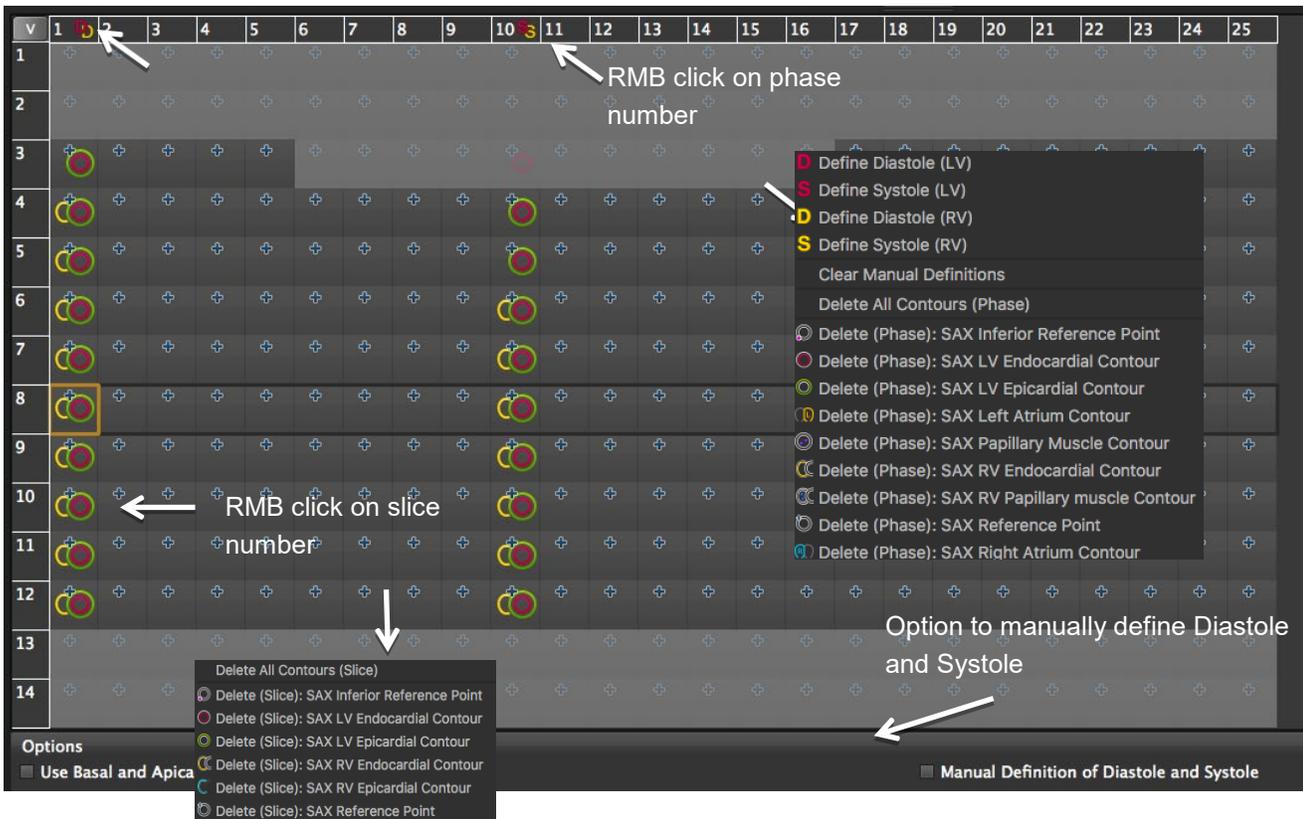
Calculations are performed based on the current state of contour definition and image selections. For every change in this state the calculations are updated immediately. It is the responsibility of the user to verify that the contours reflect the desired measurements before releasing final results.

15.3.1 Short 3D Interface



15.3.1.1 Thumbnail Grid

The thumbnail grid represents images of all slices and phases that are contained in a parallel stack. Optionally, the display can be changed to a schematic view click the **V**.



RMB click on phase number:

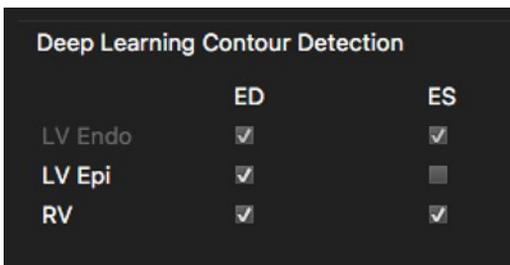
- 'V' toggles between image and contour view
- Right mouse click on a phase or slice number opens a context menu

15.3.2 Global LV Function and Volume Analysis

15.3.2.1 How to Do Fully Automatic LV/RV Segmentation

This option will detect diastole and systole automatically, based on the largest/smallest volume calculated per phase. In addition, it allows the user to set their own default in regard to border delineation per phase:

1. Go to the *Preference > Contours* and set a checkmark



2. Next, program the button: Do a right mouse click and select a function. Step 1 and 2 will be remembered. With the next patient, move right to step 3

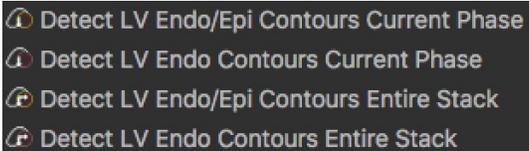




3. Click the button
4. Move the mouse in the analysis frame and click 'm' for a multiview display and review the segmentation in all slices
5. Correct where needed
6. Click Add to Report

15.3.2.2 How to Do Automatic LV Segmentation in a User Selected Phase

1. Select a phase
2. Open the drop-down menu and select the desired function. This step will be remembered for the next study



3. Click the icon

15.3.2.3 How to Do Automatic Endocardial RV Segmentation in a User Selected Phase

1. Select end-diastolic, -systolic phase



- 2.
3. Right-click and select phase or stack. This step will be remembered for the next patient



4. Click the icon

WARNING: Automatic Contour Detection

The expected input for the volume calculation in this module is a series containing parallel slice in the short axis direction.

Endocardial contours must always be inside the epicardial contour. cvi⁴² does not check for that.

In case of missing contours in one slice linear interpolation is done if possible.

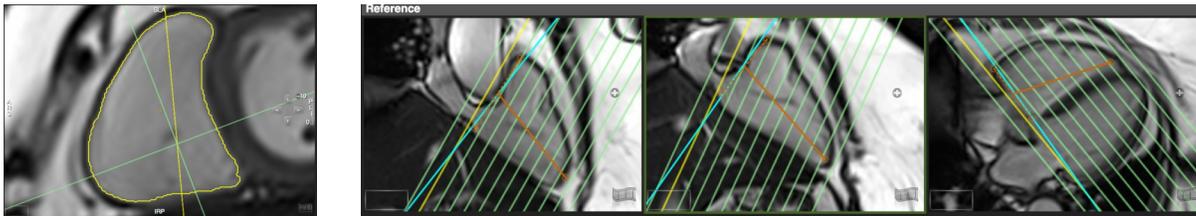
Module calculations in cvi⁴² are in generally done based on the current state of contour definition and image selections. For every change in this state, the calculations are updated immediately. It is the responsibility of the user to decide whether the stage of contour definitions reflects the desired measurement task before releasing final results.

15.3.2.4 How to Do a Mitral/Tricuspid Valve Plane Correction

1. Visually identify the most basal slice of the left ventricle and/or the outflow tract



2. Activate the *Valve plane correction* button to open additional reference windows



3. Using the *Cross-Reference* display, define the plane in two orthogonal orientations

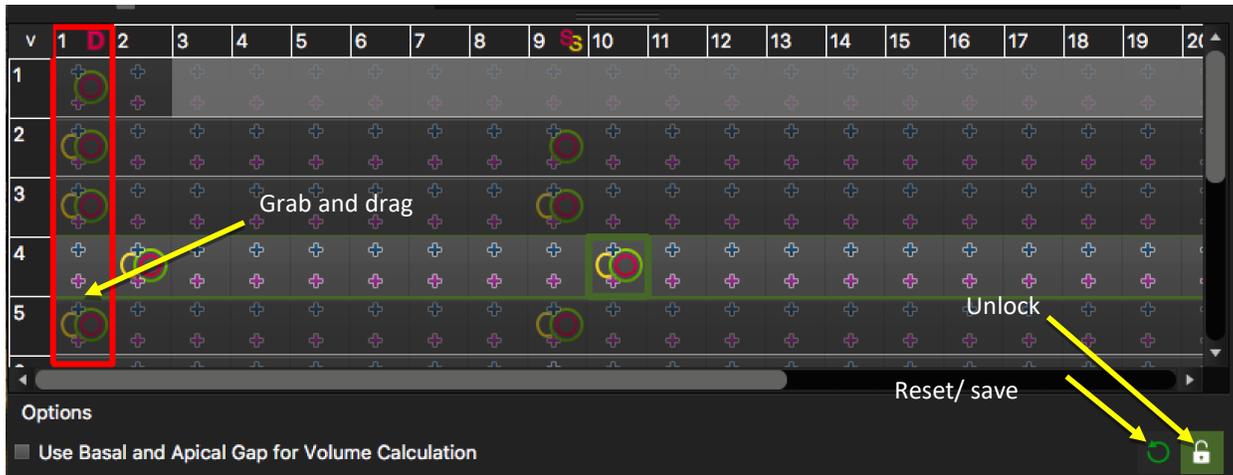
4. The correction will be calculated automatically



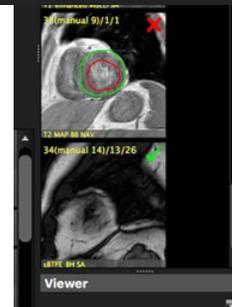
5. Use the *Exclude Area Contour* tool to exclude atrial volume in mitral plane

15.3.2.5 How to Do Manual Phase Shifting Per Slice

In case of mis-triggered slices:



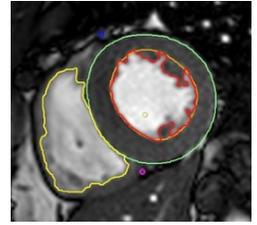
1. Click the lock
2. In the slice, grab the box with correct diastolic phase and shift it into the red box
3. Delete any existing contours that are outside diastolic and/or systolic phase
4. Delineate the myocardium for the newly defined systole and diastole
5. Click the lock again
6. The phase shift will be remembered and is stored in the workspace



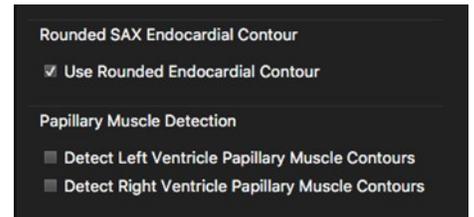
7. Only if you want to run the ML based automatic contour detection after phase shifting, you have to save the shifted series. This will create a manually composed series.
8. Load the new series and run the ML contour detection

15.3.2.6 How to in/exclude Papillary Muscles

By default, the automatic contour detection will include papillary muscles in mass calculations and exclude them from the lumen. This can be changed in the *Preferences*.



-  To exclude papillary muscle in the current image, use the Exclude Papillary tool
- To exclude them when using an automated contour detection method, go to the *Preferences/Contour*:
- Check "*Use Rounded Endocardial Contour*". Using automated edge detection, the software will apply a rounded contour, cutting off trabecular structures
- De-select Detect *papillary muscle contours* to turn off the detection of papillaries within the LV lumen (pink contour)



-  Include papillary muscles in the current image: *SA papillary muscle contour tool*

15.3.2.7 How to Create and Display a Volume Curve

Define endocardial and epicardial contours in all phases.

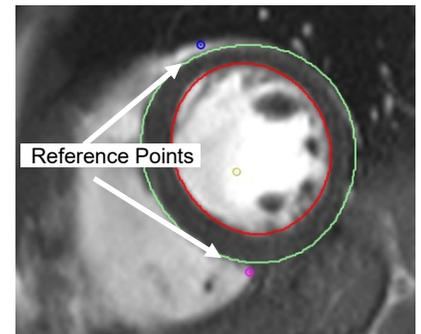


- A click on the volume curve button will display a graph next to the study display frame.
- Peak Ejection Rate and Filling Rate and Peak wall thickness will be reported in the Scientific Report (Report Module)

15.3.2.8 How to Do a Regional Wall Motion Analysis

-  Define the analysis range: Define length of anatomical long axis using the *LA LV Extent Contour* button. (see above)
-  Define reference points for the segmentation: In the analysis window select a mid-ventricular slice and set the points at the anterior and inferior insertion of the RV

Note: The segmentation points don't have to be adjusted in the remaining images.



-  **Display Polar Map:** To view the polar maps click the *View Regional Function* button

- Chords: Range between 2 and 100 chords. (segments will be counted clockwise)
 - AHA segmentation: Volumes per AHA segment will be reported in the *Scientific Report*
- From the menu select wall thickening/motion or wall thickness (Diastole/Systole). The chords in the analysis image reflect the wall thickness measurements (not the segment border)



-  Add analysis to Report. Max. diastolic wall thickness will be reported in the scientific report

5. RMB click allows to export the polar map
6. Repeat for all slices you want to include in the analysis.



7. Add analysis to Report

Note: Make sure to always use the same technique, either in or exclude papillaries and/or delineate trabecle. Left and right SV should match. Follow-up results done with a different contouring technique can mimic deterioration.

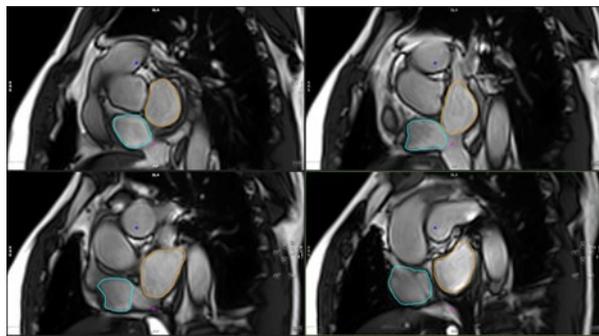
15.3.2.9 How to Do Atrial Volumetry

By default, the atrial contour buttons are hidden. In the protocol pane go to the wrench to edit the interface and activate the buttons.



15.3.2.9.1 How to Do Atrial Volumetry in Short or Long Axis Orientation

1. The atrial volumes will be reported for LV systole and diastole and therefore require the definition for the LV end-systolic and end-diastolic phase.
2. Drag a stack of parallel images, long or short axis, into the analysis frame. If there is no LV analysis applicable, you can manually define the phases. Please refer to the chapter "Manual Definition of Systole and/or Diastole for LV and/or RV"
3. Click on the *SAX Left Atrium Contour* tool to trace the border of the left atrium
4. Use the *SAX right Atrium Contour* tool for the right atrium
5. The software will report ESV, and EDV for the right and left atrium



16 Tissue Characterization Module

16.1	Analysis of LGE and T2 Images	41
	16.1 How to Do Scar and Edema Analysis	41
	16.1.2 How to Do T2 Ratio Analysis	42
	16.1.3 How to Assess Microvascular Obstruction	42
	16.1.4 How to Do Greyzone Analysis	42
	16.1.5 How to Assess Salvaged Area at Risk	42
16.2	Early Enhancement Analysis	42
16.3	Regional LGE and T2 Analysis and Assessment of Transmurality	43
	16.3.1 How to Do Segmental Scar and Edema Analysis	43
	16.3.2 How to Assess Transmurality	43
	16.3.2.1 How to Display a Greyzone Polar Map	43

cvi⁴² provides four adjustable viewer frames for determining various tissue characteristics by CMR methods.

16.1 Analysis of LGE and T2 Images

16.1.1 How to Do Scar and Edema Analysis

1. Drag and drop series into the frames marked “Late Enhancement” and “T2” respectively.
2. Derive existing contours via *Context Menu/Contours/Derive Cardiac Contours*. If no contours are present, draw endocardial and epicardial contours for each slice.



3. *Synchronize Contours*: Provided that T2 and LGE images have the same slice location, contours will automatically be copied to the other frame



4. *Segment Reference ROIs*: Automatically detect a reference ROI. The blue contour will depict the remote myocardium and the pink contour a reference ROI for FWHM algorithm in the enhanced area



5. Toggle the *Overlay Display* to display a color overlay for scar and edema



6. Use the *Exclude Enhancement* contour in case of e.g. artifacts



7. Quantify MVO (see below for more information)



8. Click on *Display Result* to view the analysis



9. Add measurements to report by clicking the *Add to Report* button

16.1.2 How to Do T2 Ratio Analysis

-  Select the *Skeletal Muscle* contour tool and draw a contour within skeletal muscle in every slice. The percentage as well as the SI ratio will be reported on the right bottom corner of the frame

-  Activate the T2 Ratio Overlay
- Repeat for all slices

16.1.3 How to Assess Microvascular Obstruction

-  To quantify a 'No Reflow Area' (depicting MVO - micro vascular obstruction), use the *No Reflow Contour* button
- Roughly draw a contour around the region with low signal

-  Click on *Display Result* to view the analysis

16.1.4 How to Do Greyzone Analysis

- Check the box Greyzone Analysis
- Select a threshold

16.1.5 How to Assess Salvaged Area at Risk

Using the results of LGE and T2 segmentation, calculation will be done automatically by subtracting scar from edema volumes.



You will find the quantified area at risk in the result pane (click on the *Display Result* button).

Note: It will **not** be calculated in case

- slice locations are not matching
- number of slices are not matching
- there is a difference of myocardial volumes > 10%

16.2 Early Enhancement

16.2.1 How to Assess the Early Gd Enhancement Ratio

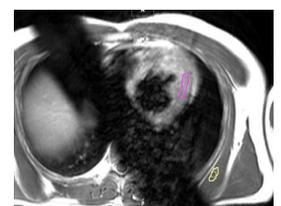
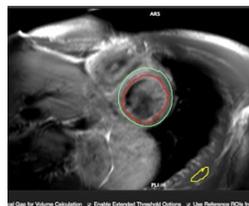
Draw endocardial and epicardial contours in T1 pre and post contrast images.

-  If you are not able to define endo and epicardial contours, use the *Myocardial Contour* tool

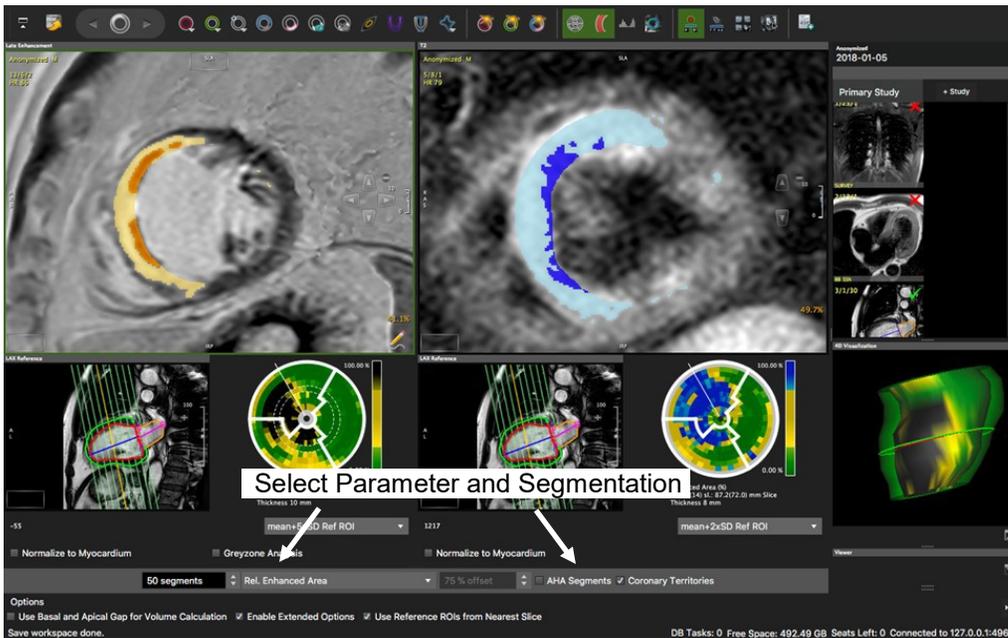
-  Define a skeletal reference
- Repeat for all slices

-  To view analysis results, click the *Display Result* button

-  Add measurements to report using the *Add to Report* button



16.3 Regional LGE and T2 Analysis and Assessment of Transmurality



Toggle Polar Map button

16.3.1 How to Do Segmental Scar and Edema Analysis

1.  Set Segmentation points
2.  Define length of anatomical long axis using the *LAX Extent* contour button: Set 3 points, where the first two define the base, and the third point is set in the apex.
3. Select Parameter from drop-down menu

For the *Enhanced Area* you have the option between custom and AHA segmentation.

16.3.2 How to Assess Transmurality

1. Repeat step 1-2
2. Select *Transmurality* from the drop-down menu at the bottom of the page
3. If you want to display a defined percentage, select *Transmurality Offset* and set the percentage with the arrow keys

16.3.2.1 How to Display a Greyzone Polar Map

1. Repeat steps 1-2
2. Select *Relative Greyzone* from the menu (make sure you have defined a threshold for the greyzone and activated the analysis option)

17 T2* Module

17.1 How to Do T2* Analysis 44

Intended use:

T2* quantification and mapping for myocardial tissue characterization (iron, hemorrhage, thrombus)

17.1 How to Do T2* Analysis

-  For global T2* times, draw an endocardial and an epicardial contour for the assessment of global T2* values

-  Draw a ROI contour in the septal wall for regional T2* values

-  Forward contours to other images within the series or via context menu *Forward Contour*

T2* values will immediately be displayed, as well as the absolute iron content (1.5 Tesla only)

T2* decay curves will immediately be displayed, choose a fitting algorithm (nonlinear=default) and a fitting correction method from the option menu

-  Toggle color map on

-  Create DICOM grayscale map. In addition to the grayscale map a R2 map will be created. You will find both maps combined in one series. They will be appended to your thumbnails

-  Add measurements to your report

18 T2 Mapping

18.1	T2 Measurements.....	45
	18.1.1 How to Do a T2 Analysis	45
	18.1.2 How to Apply a Motion Correction to T2 Series	46
18.2	T2 Map	46
	18.2.1 How to Analyze T2 Maps	46

Intended use:

T2 quantification and mapping for myocardial tissue characterization (edema)

18.1 T2 Measurements

The module consists of two pages:

- T2 Measurements: that allow for the assessment of T2 times
- T2 Map: and the analysis of a T2 map

18.1.1 How to Do a T2 Analysis

-  1. Segment the myocardium for global T2* values. Forward the contours via context menu
-  2. Select one of the ROI contours for the assessment of a specific region of interest. The ROI can be forwarded with the *Forward contour* button 

 - Global and regional T2 times will be calculated immediately
 - The decay curve will be displayed. A right mouse click in the graph allows to change the display or to export the graph. In the option menu choose a fitting algorithm and a fitting correction method
-  3. Add calculations to your report

Note: Automate DICOM map generation by checking the box for *Create map upon loading*. Save this step to your protocol. The next time a T1 sequence will be loaded, the map will be generated automatically

18.1.2 How to Apply a Motion Correction to T2 Series

1. Make sure you have endo- and epicardial contours in all images.



2. Set the Segmentation point

3. Select the image with the best contrast and enter the phase that you would like to use for the registration



4. Click the registration button: The new series will be created and automatically loaded into the respective frames

5. If you decide to use a different phase for registration, reload the original series and repeat steps 1-4

Note: The registration works for points that are on or inside the contours; the image outside of the contours may appear warped or distorted after registration. This is expected, and users should not use any outer regions for analysis

18.2 T2 Map

18.2.1 How to Analyze T2 Maps

1. Drag and drop the map from the thumbnail panel into the respective window



2. Drag a long axis reference into the reference window and define the extent of the left ventricle: Set 3 points, the first two define the base, the third defines the apex and thereby the extent of the LV

3. Derive existing contours from the T2 measurement via context menu > *Derive Contours Phase*. Alternatively, draw new contours



4. Set a segmentation reference point



5. Add analysis to the report. The values can be found in the scientific report

19 T1 Mapping and ECV Quantification

19.1	Module Layout.....	47
19.2	T1 Analysis (native and post contrast).....	47
	19.2.1 How to Load Various Sequences.....	47
	19.2.2 How to Do Global or Regional T1 Analysis	48
19.3	Motion Correction for ECV map generation	48
	19.3.1 How to Do Intensity Based Registration.....	48
	19.3.2 How to Do Feature Based Registration	49
19.4	T1 Map Analysis.....	49
	19.4.1 T1* Maps	49
	19.4.2 How to Analyze the T1 Map	49
19.5	ECV and lambda quantification (λ)	50
	19.5.1 How to Do Segmental ECV and Lambda Quantification	50

Note: Due to a lack of standardization, this module is for research only.

Intended use:

T1 quantification and mapping for myocardial tissue characterization (fibrosis, edema, tissue infiltration).

19.1 Module Layout

Analysis options are organized in pages:

- *T1 Native*: T1 values and recovery curve for global and/or regional myocardium
- *T1 CA*: s.a. post contrast
- *T1 Map Native*: T1 and R² maps and a segmental polar map display (1-100 or AHA)
- *T1 CA*: s.a. post contrast
- ECV/ λ : Polar maps and global/per slice ECV and partition coefficient (λ) quantification

19.2 T1 Analysis (native and post contrast)

19.2.1 How to Load Various Sequences

1. Select the correct sequence in the option menu
2. Load the sequence from the thumbnail panel. If the series is rejected, it could have been due to one of the following reasons:

- The last image of the series is a scanner generated map (Siemens). **Solution:** In the option menu, check the box for *Skip Last Slice Image*
- In the DICOM header the T1 time is encoded in the field for the Trigger Time (Philips). **Solution:** In the option menu, check the box for *Use Trigger Time as Inversion Time*
- During anonymization the information for T1 has been stripped and you get a warning message "T1 times are equal." **Solution:** Go to the patient list module and in the *Extended view* check if the information for the T1 can be found under either *T1* or *Image Comments* (Siemens). If possible, try to re-anonymize the data, preserving the T1 information

Note: Some recent **Siemens** sequences use a correction or scaling factor for the calculation of T1 times. You will find the scaling factor in your Siemens documentation. Enter the factor in the option menu. Alternatively, use Siemens-generated maps.

19.2.2 How to Do Global or Regional T1 Analysis

-  For global T1 values, draw endocardial and an epicardial contours. Forward the contours via context menu *Forward All Contours Slice* (short-cut: ⌘E/Ctrl E)
-  For regional analysis, select one of the ROI contours. This contour can be forwarded via toolbar button 
- Repeat for all slices
- Global, regional T₁ Relaxation times and R²-values will be shown in the result frame
- The T₁ recovery curve will be presented in the frame below. RBM click in the graph opens a *Context Menu* offering display options
-  Create DICOM map
- Automate DICOM map generation: In the option menu, check the box for *Create map upon loading* and save this step to your protocol
- For the post-contrast series, contours can be derived via context menu: *Derive Contours>Derive Contours Cardiac Phase*, will derive previously drawn contours for all slices
-  Add evaluation (images, values and graph) to the report

19.3 Motion Correction for ECV Map Generation

- *Intensity-based* registration is to be used with already motion corrected images and will register between pre- and post-contrast series
- Feature-based registration registers all images (pre- and post-contrast) to a selected image in the native series. Therefore, it is required to have contours in both the pre- and post-contrast series as well as in all slices
- Go to the respective pages and select the motion correction of your choice

19.3.1 How to Do Intensity Based Registration

- Identify motion corrected series in the thumbnail panel
- Drag the series into the respective pages: T1 Native/T1 CA
- Choose a phase with good contrast between endo- and epicardial contour as well as between surrounding tissues in both the native and post contrast series. Avoid images with bright intensities in the outer regions (areas of fat, air regions, etc.)
- In the *Registration Options* enter the phase number:
- Native Registration Phase*: Image in the native series to be the reference image
- CA Registration*: Image in the post contrast series, to be registered with the reference image
-  Click the registration button
- The registration registers the CA Series to the Native Series
- To re-register, reload the initial series and repeat steps 1-6

19.3.2 How to Do Feature Based Registration

This registration can be used on motion and non-motion corrected series. It registers every phase in both the native and post-contrast series to a chosen reference image in the native series.

1. Make sure you have accurately drawn the endo- and epicardial contours, as well as a segmentation point in all images, in pre- and post-contrast series
2. Select a reference image in the native series with the best contrast and enter the phase into *Native Registration Phase*



3. Click the registration button: Two new registered series will be created and automatically loaded into the respective frames. Contours of the reference phase will be forwarded to all registered images
4. If you decide to use a different phase for registration, reload the original series and repeat steps 1-4

Note: The registration works for points that are on or inside the contours; the image outside of the contours may appear warped or distorted after registration. This is expected, and users should not use any outer regions for analysis

19.4 T1 Map Analysis

Either create map in cvi⁴² or load a map that has been created by the scanner software.

19.4.1 T1* Maps

- T1* maps may offer a more accurate estimation of blood pool T1
- T1* maps will be automatically extracted of Molli, shMolli and Molli Oxford sequences

19.4.2 How to Analyze the T1 Map

1. Drag the map from the thumbnail panel to the analysis window
2. Draw or derive contours for a segmental analysis
3. For a color-coded display, select a color map from the drop-down menu or create your own LUT
4. For a segmental polar map display, place an anterior and inferior *SAX Reference* point
5. Define the left ventricular extent in the reference window
6. In the polar map frame, the number of segments per ring can be customized or you can switch to a 16- segment AHA model
7. To exclude zero values (non-signals) you can set an off-set by checking *Exclude Zero Values for Map Measurements*



8. Add evaluations (images, values)

19.5 ECV and Lambda Quantification (λ)

19.5.1 How to Do Segmental ECV and Lambda Quantification

1. Make sure you have contours, segmentation points and a blood-pool contour  in all slices in both *T1 map Native* and *T1Map CA*. The blood pool contour can be drawn either in the *T1* or the *T1** map
2. Enter the hematocrit for the ECV quantification
3.  Add an ECV map
4.  For a polar map display, make sure you have defined the Long Axis Extent in the reference frame of the *T1 Map Native* and *T1 Map CA* page
5. Add evaluation (images, values and graph) to the report 

20 First Pass Perfusion Module

20.1	Multi View Compare Rest and Stress Perfusion	51
20.1.1	How to Do a Visual Perfusion Analysis	51
20.2	Edit Contour, Semi-Quantitative Analysis.....	52
20.2.1	How to Do Semi-Quantitative Perfusion Analysis.....	52
20.2.2	Option Menu	53
20.2.3	How to Analyze Perfusion Curves.....	53
20.3	Analysis, Segmental Analysis.....	53

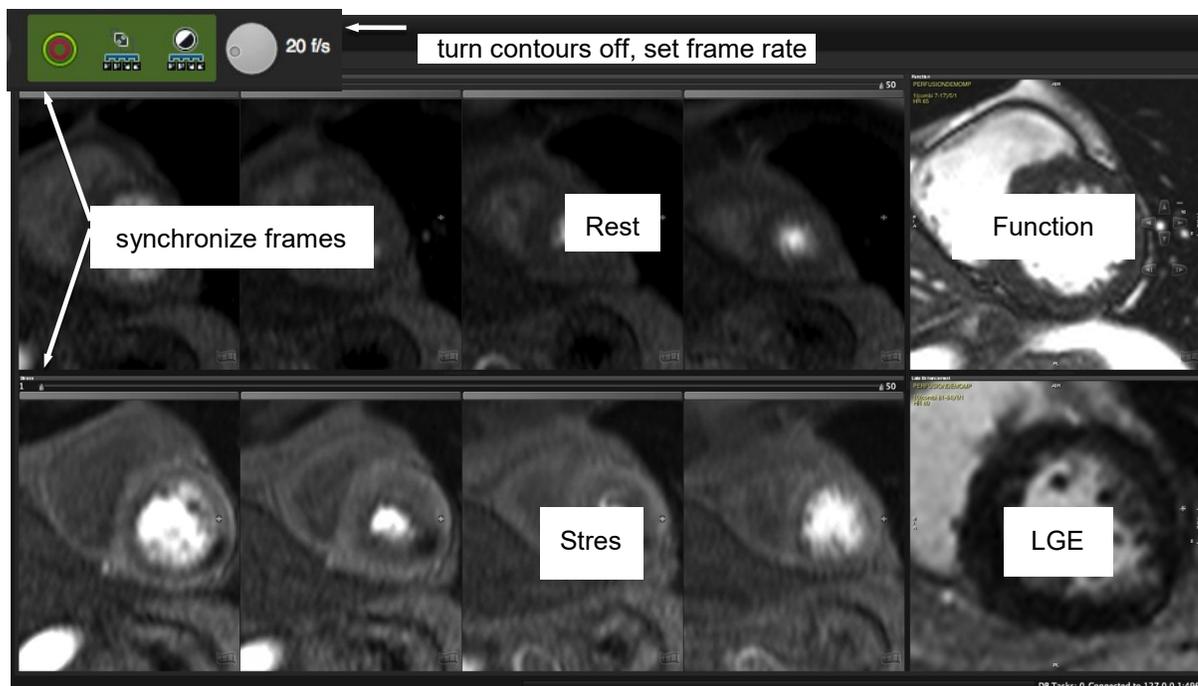
Intended use:

Visual and semi-quantitative assessment of myocardial perfusion

The Perfusion module consists of 3 pages:

- *Multi View*: Compare rest and stress perfusion with scar images and wall motion kinetics
- *Edit Contour*: Semi-quantitative analysis
- *Analysis*: Polar Maps

20.1 Multi View Compare Rest and Stress Perfusion

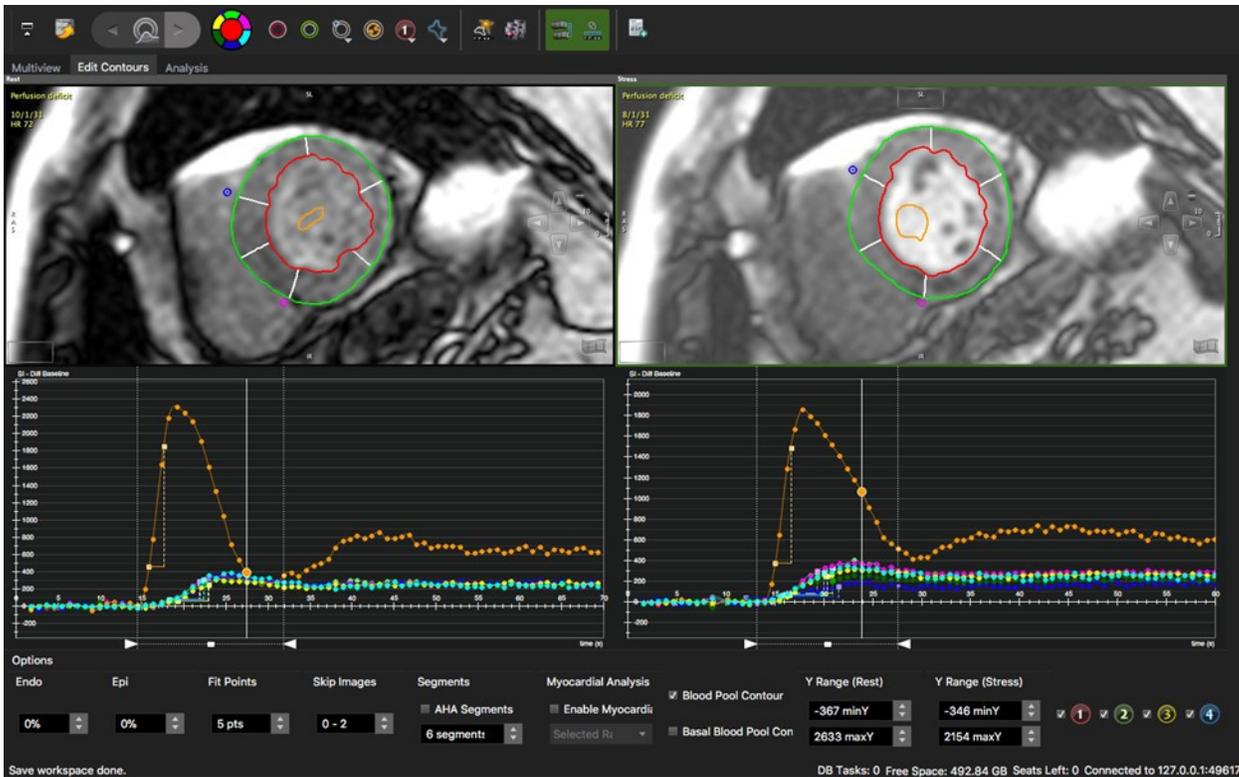


20.1.1 How to Do a Visual Perfusion Analysis

1. Adjust window and zoom for rest and stress series
2. To set the frame rate use the wheel in the toolbar
3. Turn existing contours off (toolbar button)
4. Synchronize the frames for rest and stress by using the slider
5. If applicable, drag Late Enhancement and Function images into their respective frames for comparison
6. Add the *Floating Viewer* for more viewing options

20.2 Edit Contour, Semi-Quantitative Analysis

20.2.1 How to Do Semi-Quantitative Perfusion Analysis



1. Drag the appropriate series in the *Rest* and *Stress* frame and adjust viewer properties
2. Select a phase with good contrast and draw endo- and epicardial contours
3.  Draw a *Blood Pool* contour, (input function) in the brightest, most homogeneous area of the blood pool
4. Check *Use Bloodpool Contour* in the *Options* (underneath the graph)
5.  Set the anterior and inferior segmentation reference points at the insertion of the RV to the epicardial border of the LV
6.  Contour Propagation: Forward all contours, including blood pool contour and segmentation point to the remaining phases
7.  Check and adjust contours for each phase within the analysis range. Use the align contour button to correct for breathing motion
8. Choose between AHA or custom segmentation
9. Define the analysis range (contrast inflow during first pass) by moving the left and right slider at the bottom of the graph. Alternatively, set start and end frame via context menu. This will also limit the frames in the 'Multi View' to the defined range
10. Apply baseline correction via context menu (right-click in the graph)
11. For a special region of interest, there are 4 additional contours in the top tool bar. To view the curve, check the respective ROI in the *Option* menu
12.  Repeat for all slices and add results to the report

20.2.2 Option Menu

- *Endo/Epi*: Exclude sub-endocardial and/or sub-epicardial layers
- *Fit points*: Choose the number of points that will define the slope
- *Skip Images*: you can exclude images from the from baseline
- *Segments*: Choose between custom segments (up to 100) or AHA segmentation
- *Myocardial Analysis*: Analyze different layers
- *Blood Pool contour*: Checked, it requires to define a blood pool contour
- *Basal Blood Pool Contour*: The software only uses the most basal blood pool contour
- Check a box to view the perfusion curve of a drawn ROI

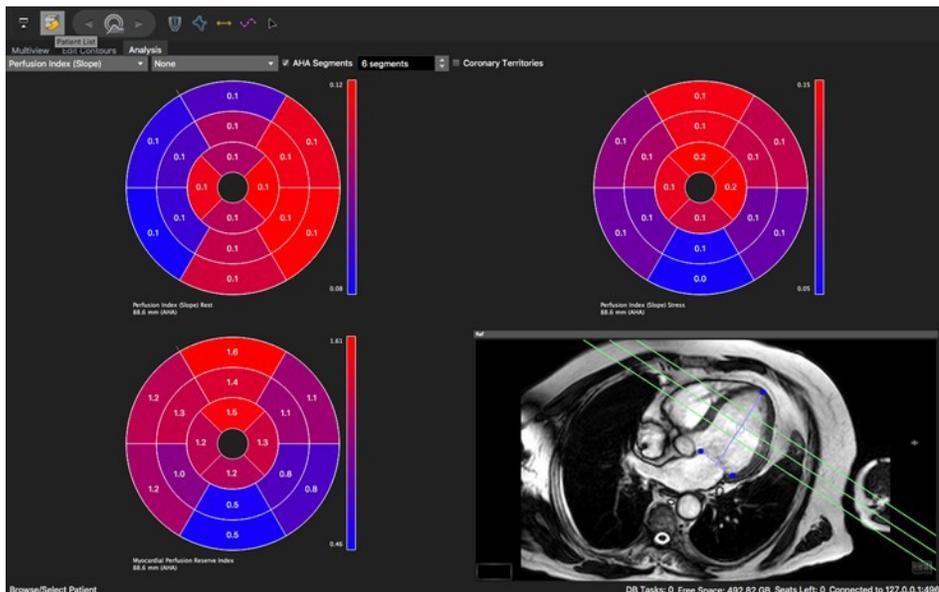
20.2.3 How to Analyze Perfusion Curves

1. To switch to an enlarged view, go to the context menu and check *Enlarged View*



2. Move the mouse cursor over the icon to view the curve of the respective segment
3. Display semi-quantitative parameter such as *upslope* or *time to max.* from the context menu
4. Optionally apply a baseline correction
5. *Exit the Enlarged View* via context menu (uncheck)

20.3 Analysis, Segmental Analysis



How to do a regional perfusion analysis:



1. Define length of anatomical long axis using the *LAX LV Extent Contour* button
2. Select a parameter from the drop-down list
3. Apply a baseline correction

21 Vascular Module

21.1	Centerline page	54
	21.1.1 Main Page	54
	21.1.2 Vessel Lists	55
	21.1.3 Region Growing	55-56
	21.1.4 Edit the Centerline	56
21.2	Cross Section.....	57
	21.2.1 How to Do a Stenosis Assessment.....	57

The module covers two topics:

- Centerline: Centerline extraction of any vessel using orthogonal MPR views
- Cross Sections: Visual and quantitative assessment

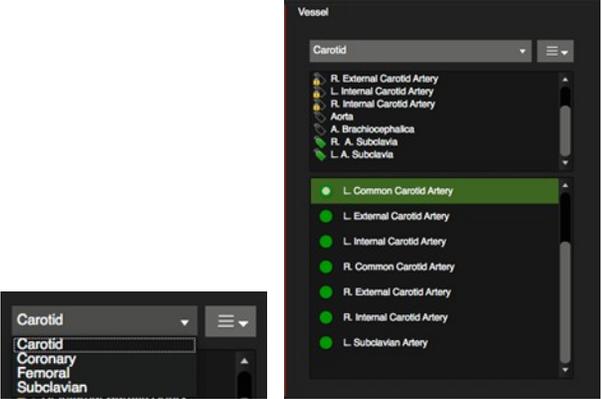
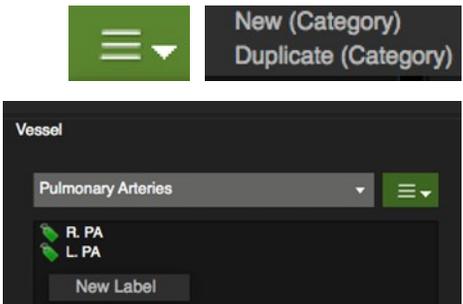
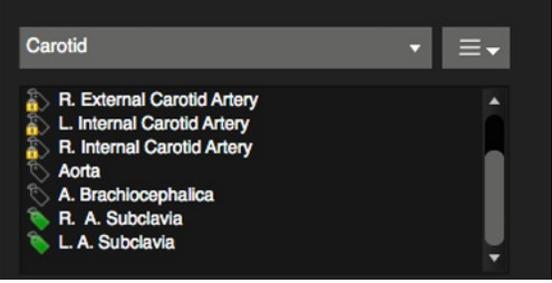


21.1 Centerline Page 21.1.1 Main Page



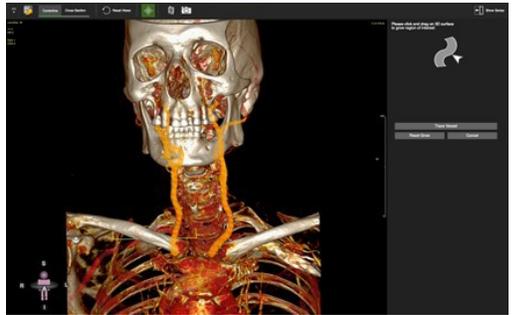
Trace a centerline by setting a start and an end seed point (see Coronary module)

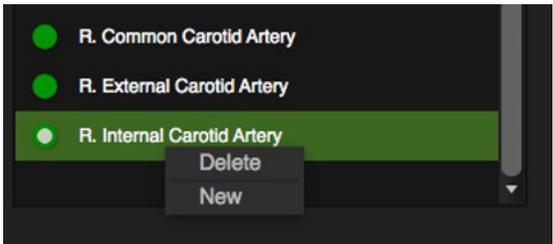
21.1.2 Vessel Lists

<p>Lists of predefined labels for standard measurements. The software comes with lists for:</p> <ul style="list-style-type: none"> • Carotid Arteries • Coronary Arteries • Femoral Arteries • Subclavian Arteries 	
<ul style="list-style-type: none"> • For a new list select <i>New (category)</i>, type a name in the window, click <i>Enter</i> • To add or delete a label, right-click in the background • The selected vessel list will be used sequentially when the user defines centerlines via seed-points or regiongrowing 	
<p>A list can display three label icons:</p> <ul style="list-style-type: none"> • Labels with a lock: predefined • Green: unused • Outlined labels: used 	

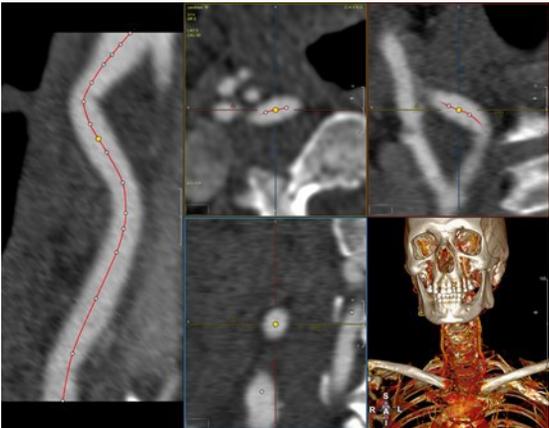
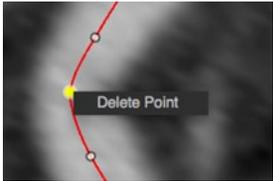
21.1.3 Region Growing

Trace a centerline via region growing.

<ul style="list-style-type: none"> • Click the <i>Region Growing</i> button in the top toolbar. The software will open the image in an enlarged view. 	
<ul style="list-style-type: none"> • The top right panel will guide you through the steps 	

<ul style="list-style-type: none"> • Grow the vessel by dragging on the 3D surface • Click <i>Trace vessel</i>, to extract the vessel • 'Alt' key allows to turn, 'Ctrl' (command Mac) allows to pan the image • In the segmented vessel, extract the centerline by placing a start and an endpoint • 'T' toggles the centerline on/off 	
<ul style="list-style-type: none"> • Click Next Vessel to proceed to the next vessel in your list • Right-click to add or delete a centerline. The label will be updated to 'un-used' 	
<ul style="list-style-type: none"> • <i>Done</i> will go back to the Main page 	

21.1.4 Edit the Centerline

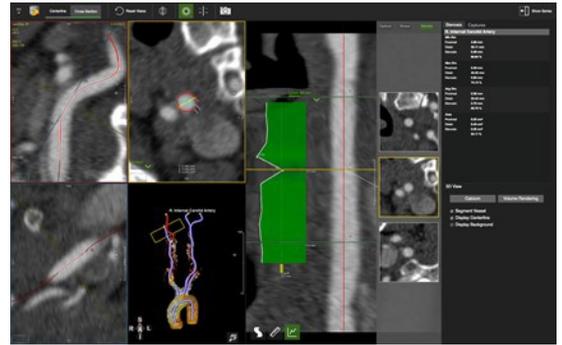
<ul style="list-style-type: none"> • You can move the centerline control points in the CPR or the oblique views • Add control points with a double-click on the centerline 	
<ul style="list-style-type: none"> • Delete a control point with a right mouse click • Rotate around the centerline with a left mouse drag close to the centerline • Pan the image by moving the cursor away from the centerline until the hand symbol appears 	

21.2 Cross Section

Stenosis Assessment

21.2.1 How to Do a Stenosis Assessment

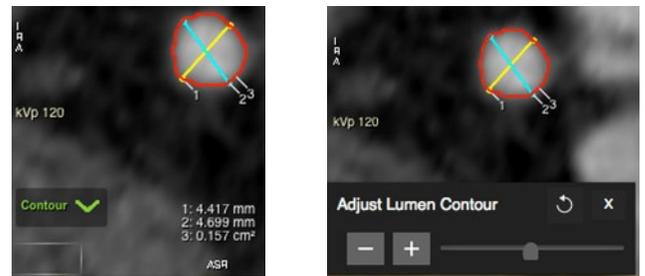
- Select the vessel by clicking on the respective vessel in the reference viewer ('T' – toggles the centerline on/off)
- Start in *Browse* view and drag the yellow line down the vessel
- Switch to *Stenosis* View and place the stenosis marker either by dragging or by double-clicking in the stenosis



- Adjust the plane to accurately display a cross-sectional view



- Move to the MPR view and use the *Auto Vessel Lumen* tool to contour the stenosis and the 2 reference images
- To adjust the threshold, click the green *contour* label. It will open a slider that allows to adjust the contour



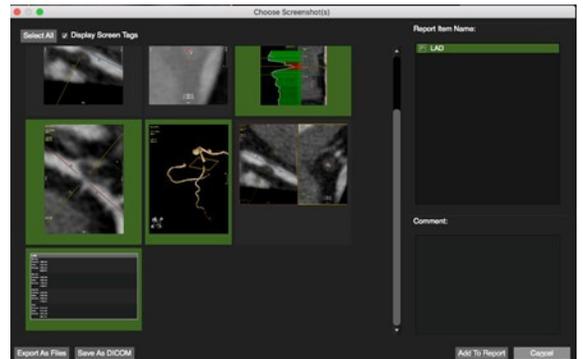
- Stenosis measurements will be filled in automatically

Stenosis	Captures	Avg Dm.
R. Internal Carotid Artery		
Min Dm.		Proximal: 5.28 mm
Proximal:	4.99 mm	Distal: 32.43 mm
Distal:	28.11 mm	Stenosis: 5.70 mm
Stenosis:	5.49 mm	69.76 %
Max Dm.		Area
Proximal:	5.49 mm	Proximal: 0.22 cm ²
Distal:	40.45 mm	Distal: 8.43 cm ²
Stenosis:	5.85 mm	Stenosis: 0.25 cm ²
	74.12 %	94.11 %

- Images can be captured via context menu.



- Use the *Multiframe Capture* to save the stenosis measurements and optionally some images



22 Tissue Tracking

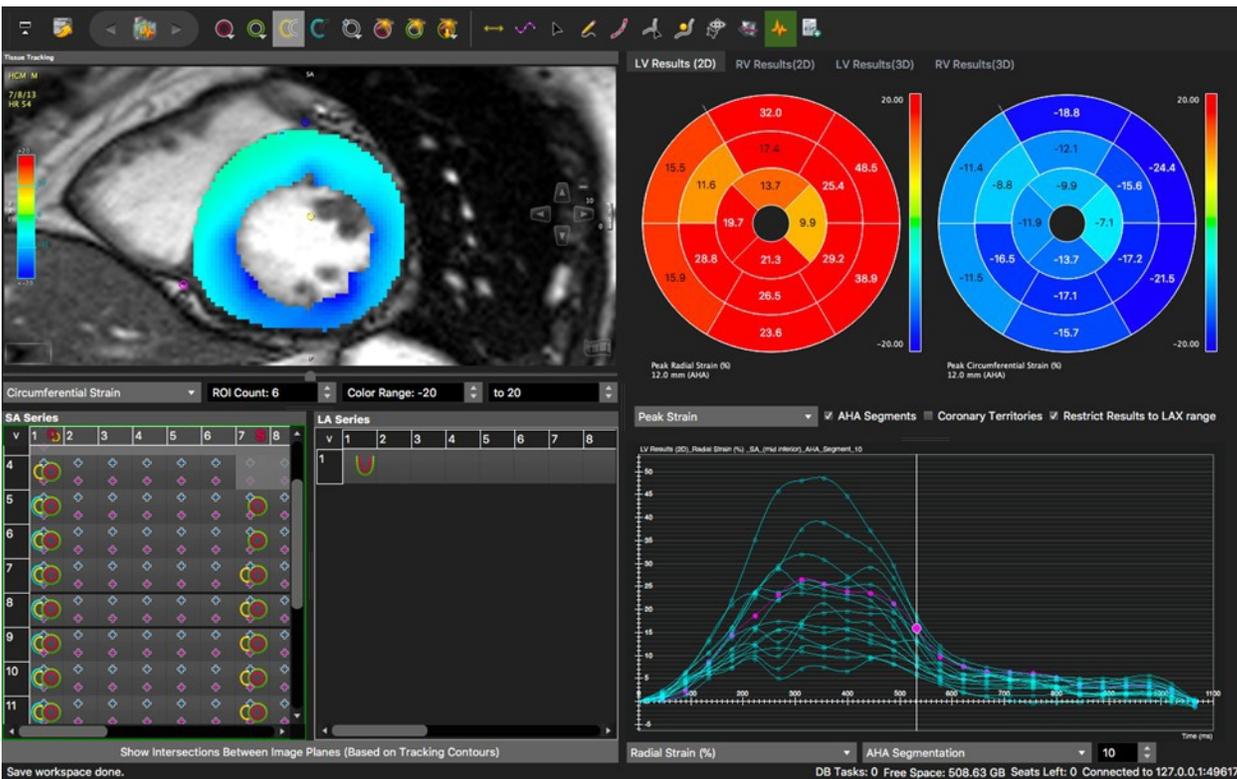
22.1	Strain Analysis	58
22.1.1	How to Do LV 2D Strain Analysis in SAX.....	58-59
22.1.2	How to Do LV 2D Strain Analysis in LAX.....	59-60
22.1.3	How to Do 3D Strain Analysis.....	60-61
22.1.4	How to Do RV Strain Analysis	61
22.2	Performance Check and Adjustments	61
22.3	4D LV Visualization	61
22.3.1	How to View a Strain Cine Loop	61
22.4	Regional LV Analysis	62
22.4.1	How to Display Polar Maps	62
22.4.2	How to Evaluate a Single Segment.....	62
22.5	Diagrams for Strain, Displacement and Torsion	62-63
22.5.1	How to Display Strain and Displacement Diagrams.....	62-63

Intended use:

Assessment of myocardial deformation.

22.1 Strain Analysis

22.1.1 How to do LV 2D Strain Analysis in SAX



1. Load a short axis stack into the analysis frame



2. In a reference phase (most commonly end-diastole) define endo- and epicardial contours for all slices that cover the left ventricle

3. Use open contours for the base



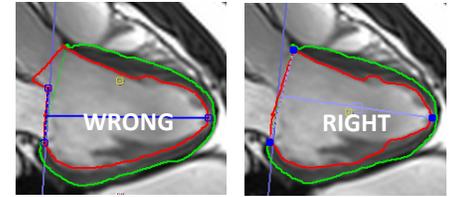
For a regional analysis (polar maps):



4. set the anterior SAX reference points at the insertion of the right ventricle
5. Next, load a long axis series
6. For contour delineation, go to the same phase in which you drew your short axis contours



7. Use the LAX LV extend contour to define the analysis range. The landmark for the basal points is the muscle and not the valve
8. The endocardial contour will be drawn automatically. Correct if necessary. Vice versa, drawing an endocardial contour will automatically place a LAX extend contour.



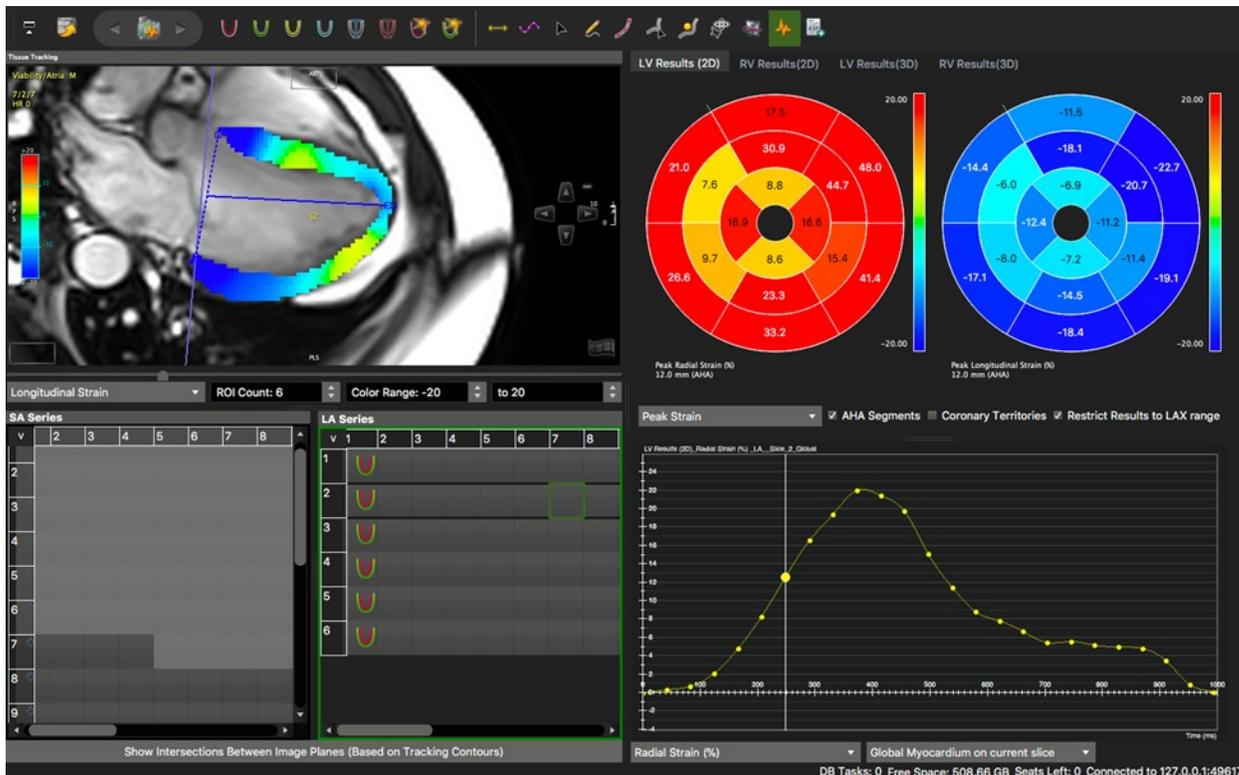
9. Define an epicardial contour

10. Check 'Restrict Results to LAX range'



11. Make sure you have selected the reference phase with the contours and Perform Automatic Strain Analysis.
12. Click the LV Results 2D tab above the polar maps

22.1.2 How to do LV 2D Strain Analysis in LAX



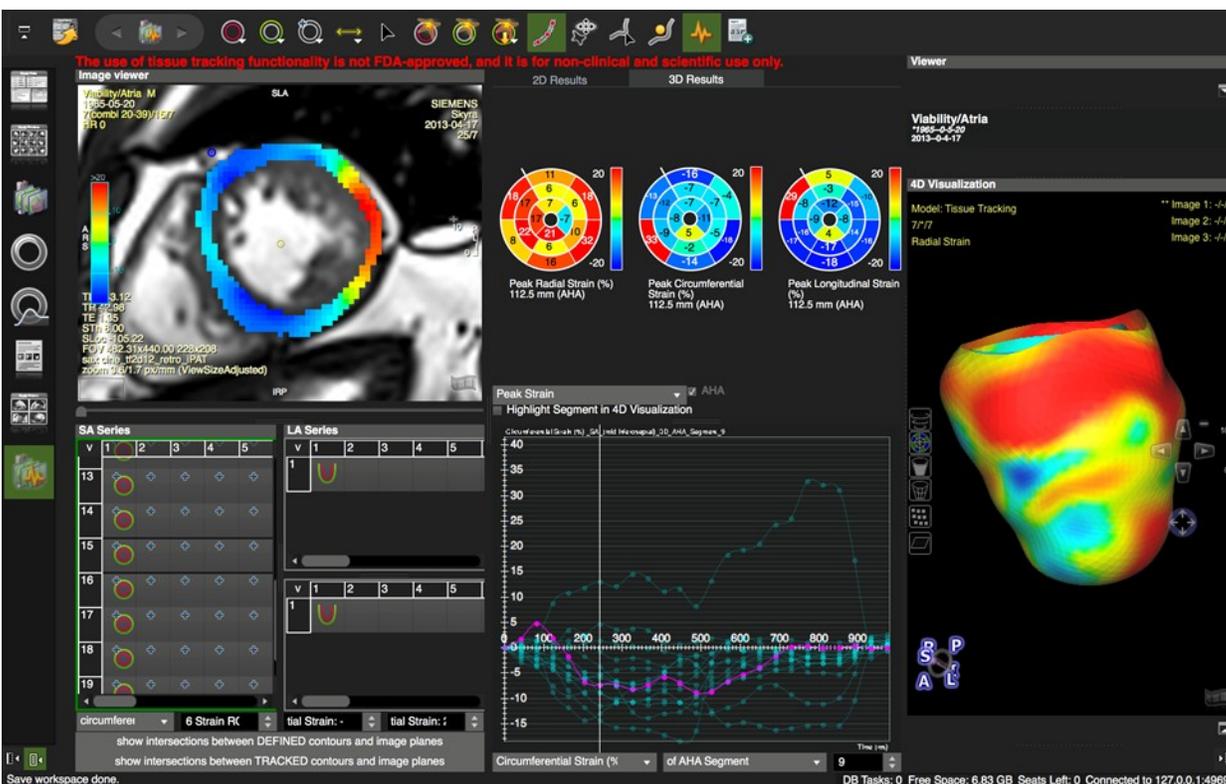
It is recommended to have at least 3 LAX slices in total and one mid-ventricular short axis slice for the generation of a polar map.

1. Load a radial long axis stack or sequentially load all radial long axis series into the viewer

2.  Place a *Long Axis Extend Contour*, which will trigger an endocardial contour in a reference phase or vice versa
3.  Define epicardial LAX contours
4. For the generation of polar maps:
 - Load a short axis slice
 - Draw epicardial and endocardial contours in the SAX image
5.  Place an anterior insertion point for the polar map segmentation
6.  Select the reference phase by clicking on an image within that phase and click the *Perform Automatic Strain Analysis* button. Strain will be automatically computed in all slices, that contain endo- and epicardial contours

22.1.3 How to do 3D Strain Analysis

It is recommended to have at least 3 short and long axis slices and 25 phases.



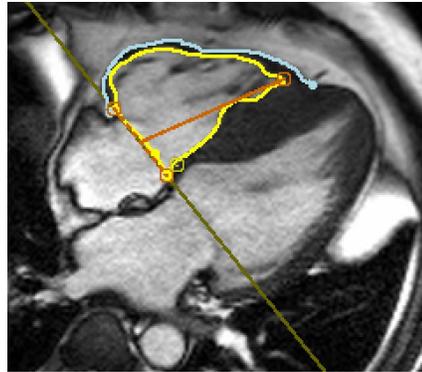
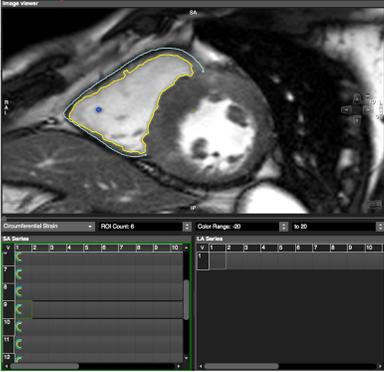
1. Sequentially load short and long axis (either a stack or 3 different radial cines) in the analysis frame
2.  Define endo- and epicardial SAX contours in all slices that that cover the LV. Make sure not to include blood volume
3.  Define the anterior *SAX reference point* at the insertion of the right ventricle in a SAX
4.  In long axis, setting the *LV extend* will trigger an endocardial contour detection and vice versa
5. Correct if needed. Pay attention to exclude papillary muscle



6. Select the reference phase by clicking on an image within that phase and click the *Perform Automatic Strain Analysis* button. Strain will be automatically computed in all slices, that contain endo- and epicardial contours
7. Click the *3D LV Results* tab above the polar maps
8. The software will generate polar maps, strain curves and a 4D contour model based on the tracking results

22.1.4 How to do RV Strain Analysis

1. Load a short and/or long axis stack into the analysis frame
2. In a reference phase, define RV endo- and epicardial contours in all slices that cover the RV



3. For a segmental analysis:
 - Load an orthogonal slice of the RV with the same number of phases
 - Endo- and epicardial contours have to be drawn consistently in the same phase for long and short axis.
 - Add an anterior and inferior insertion point
4. For 3D RV analysis it is recommended to load at least 3 additional orthogonal slices



5. Select the reference phase and click the *Perform Automatic Strain Analysis* button. Strain will be automatically computed in all slices, that contain endo- and epicardial contours.

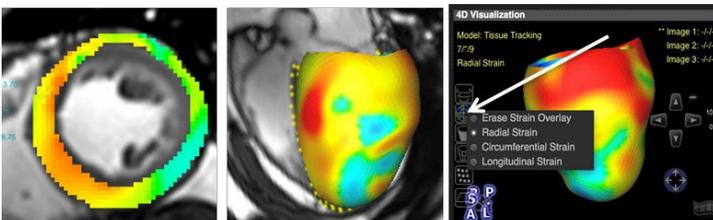
22.2 Performance Check and Adjustments

- Use the overlays to check the performance of myocardial tracking
- To review the parts of the myocardium that have been included into the strain analysis, based on the Restrict to LAX extend, select the Myocardial Points
- If you are not satisfied with the strain borders visible in the overlay, try to re-adjust contours or draw contours in a different phase, then re-run the strain analysis

22.3 4D LV Visualization

22.3.1 How to View a Strain Cine Loop

- After the strain analysis, you can view a cine loop of radial/longitudinal or circumferential strain in the *4D Visualization Viewer*
- Click the film icon or spacebar to run the movie. Cines are synchronized with the image viewer

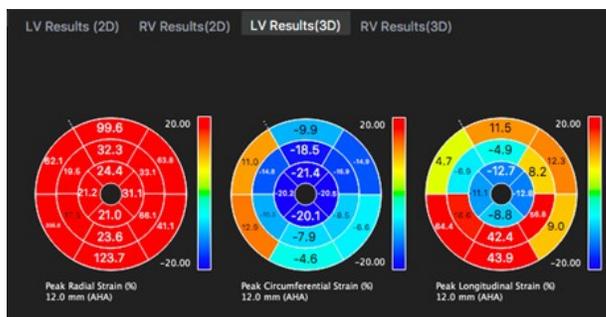


2D Radial Strain

2D/3D Visualization

22.4 Regional LV Analysis

22.4.1 How to Display Polar Maps

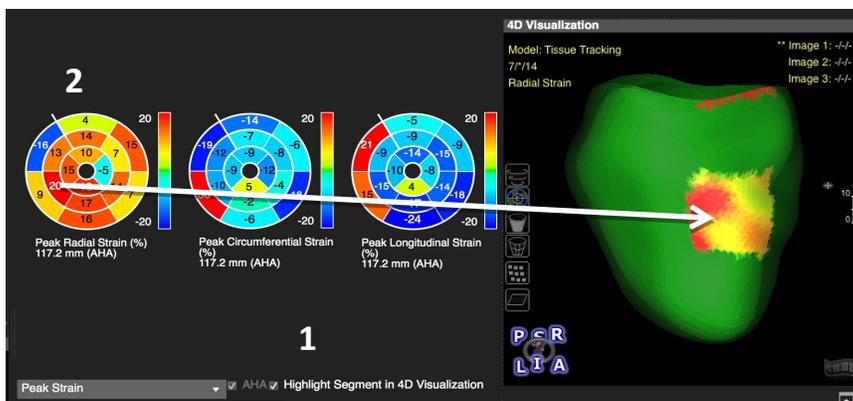


Polar maps display different values, which can be chosen from a drop-down menu.

- To view polar maps for the different strain analyses, click the *2D or 3D Results* tab
- A click on the polar map will automatically synchronize curves, image viewer and 4D viewer to display the selected strain, e.g. circumferential strain
- 2D results can be viewed for a 16-segment model or a defined number (between 6 and 24) of ROIs. To set the number of ROIs per slice, use the drop-down menu located underneath the analysis frame.
- Adjust the color map for minimum and maximum strain using the menu located underneath the LAX thumbnail grid
- The drop-down menu will offer different results

22.4.2 How to Evaluate a Single Segment

1. To view a specific segment in the 4D viewer check the box for *Highlight Segment in 4D Visualization*
2. Double-click a segment in the polar map and view the strain of the selected segment in 4D by clicking the film icon
3. The respective segment in the AHA segment will be highlighted automatically



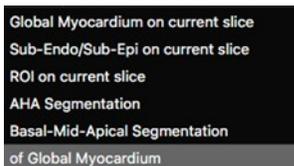
22.5 Diagrams for Strain, Displacement and Torsion

22.5.1 How to Display Strain and Displacement Diagrams

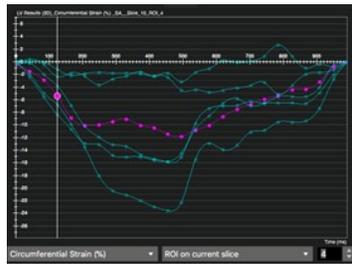
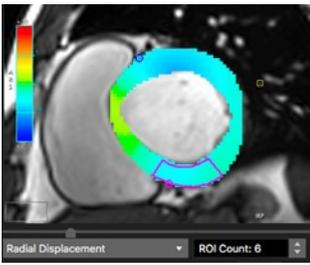
1. Start with selecting the region for which you would like to display the parameter:

- *ROI on current slice*

- The number of ROIs can be in-/decreased (minimum is 6, maximum 24) using the menu located underneath analysis viewer



2. Use the up/down arrows next to *ROI on current slice* to highlight the respective ROI



Torsion:

- If not defined, the software uses the most basal and apical slice to calculate torsion
- To select different slices for the torsion calculation: Right-click on the slice number in the thumbnail grid and select B/A



23 4D Flow

23.1	Data Cropping	64
23.2	Preprocessing	64
	23.2.1 How to Preview 4D Flow	64-65
	23.2.2 How to Do an Offset Correction and Phase Aliasing	65
23.3	Segmentation	66
	23.3.1 How to Segment a Vessel	66
	23.3.2 How to Extract a Centerline	67
23.4	Analysis	67
	23.4.1 How to Visualize Flow in 4D	67-68
	23.4.2 How to Do a Flow Measurements	69
	23.4.3 How to Do a Shunt Calculation Qp/Qs	70
23.5	Advanced 4D Flow Parameter Analysis	70
	23.5.1 How to Assess Wall Shear Stress	70
	23.5.2 How to Do Relative Pressure Mapping	70
	23.5.3 How to Calculate Pulse Wave Velocity	71
	23.5.4 How to Analyze Energy Loss	71
	23.5.5 How to Analyze Ventricular Flow	72

Intended Use:

Preprocessing.

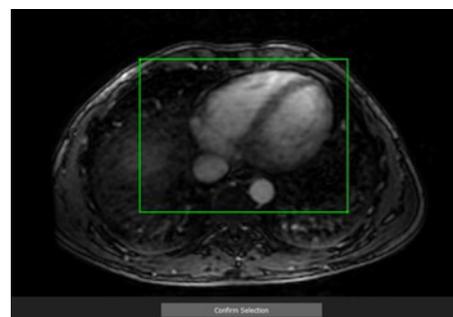
23.1 Data Cropping

1. Load the 4D flow study from the thumbnail pane into the module via double-click or drag-and-drop

- The cropping tool can be used to better define the field of interest and improve performance for big studies (acquisitions with more than 3000 slices)

Note: Once the bounding box is fixed, all the further analysis will be done within the bounding box. If the bounding box is deleted or modified before saving workspace, all the measurements will be changed.

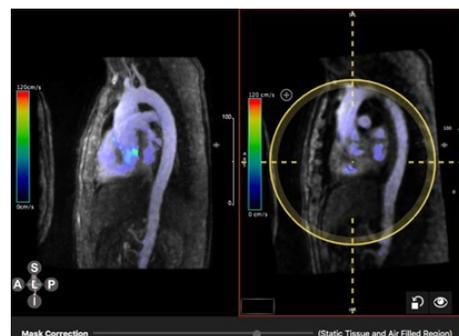
2. Select the box by clicking the green line
3. Grab the green dots at the left top and right bottom corner of the box and drag to confine the region of interest
4. *Confirm Selection*
5. To change the selection, go to *Workspace>Reset Workspace*



23.2 Preprocessing

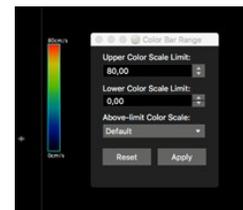
23.2.1 How to Preview 4D Flow

1. Run the cine and scroll through the slices. The left view port displays the time averaged MRA. The right MPR Viewer allows to navigate through the volume (see MPR Viewer)
2. The mask is based on detection of areas with high flow
3. Use the *Mask Correction* slider to adjust the mask



4. Adjust the color bar: LMB click on the bar will open a dialog.

Note: Structures on preview page are independent from the structures on the segmentation page.



23.2.2 How to do an Offset Correction and Phase Aliasing

<ul style="list-style-type: none"> The <i>Preview</i> button toggles between magnitude image visualization and three-dimensional velocity magnitude visualization 	
<ul style="list-style-type: none"> <i>Mask Overlay</i> displays the static- tissue (yellow) and air-filled (blue) mask overlays. 	
<ul style="list-style-type: none"> Use the <i>Mask Correction</i> slider to adjust the regions. <p>Note: Static tissue mask can go outside of the bounding box, air filled region only shows sup inside of the box and is limited by the bounding box.</p>	
<ul style="list-style-type: none"> Run the cine and scroll through the slices to detect artifacts, such as aliasing. 	
<ul style="list-style-type: none"> <i>Check Offset Correction:</i> The static tissue is used to detect an offset. 	<p>This approach works in two steps. The initial step detects the static tissue by comparing the normalized temporal variation of a voxel to a certain threshold. The next step applies fitting procedure to the static tissue to derive a correction function for the velocity offset.</p>
<ul style="list-style-type: none"> Once the offset correction has been performed, the estimated offset field can be visualized for all 3 dimensions, by selecting <i>Show Correction Field</i> in the top toolbar. 	
<ul style="list-style-type: none"> In case of aliasing, check 'Phase Unwrapping' 	
<ul style="list-style-type: none"> Click <i>Apply Processing</i> Green font indicates preprocessing has been applied 	

23.3 Segmentation



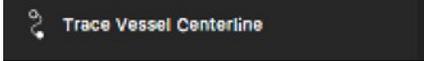
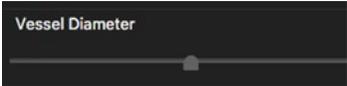
Intended use:

- Segmentation
- Centerline extraction

23.3.1 How to Segment a Vessel

<p>1. <i>Show Magnitude Image</i> toggles between magnitude and phase-contrast display. The magnitude allows you to run a cine in the background.</p>	
<p>2. Activate the <i>Display Mask Overlay</i> button.</p>	
<p>3. Use the <i>Mask Threshold</i> slider to adjust the segmentation.</p>	
<p>4. Use cut and crop to refine the MRA segmentation in the PC MRA</p> <ul style="list-style-type: none"> • Define one or several masks with, or without creating a centerline • Right-click to add a new or delete structure • Double-click on a title to rename • Right-click on the existing vessels to delete 	

23.3.2 How to Extract a Centerline

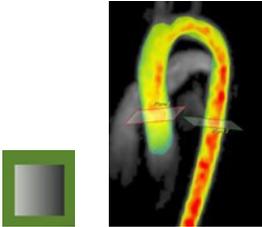
<p>1. Click <i>Trace Centerline</i></p>	
<p>2. Set a start and an endpoint in the PC image</p> <p>3. Select if you are about to trace a small or large vessel (coronaries are small, everything larger is classified as large).</p> <p>4. Adjust the control points in any view port</p> <p>5. Rotate around the centerline with a left mouse drag next to the line in the CPR view.</p> <p>Note: Only the spline and control points within the slab range is visible.</p>	
<p>6. Once you have created several masks and/or centerlines you can easily switch back and forth between the vessels by simply clicking on the vessel title</p>	
<p>7. Use the <i>Vessel Diameter</i> slider to grow the mask from the centerline</p>	

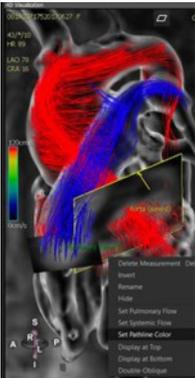
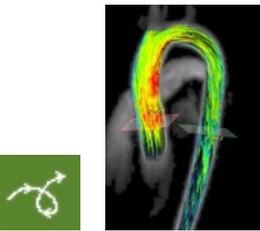
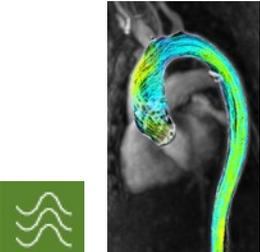
23.4 Analysis

Intended Use

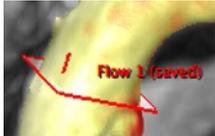
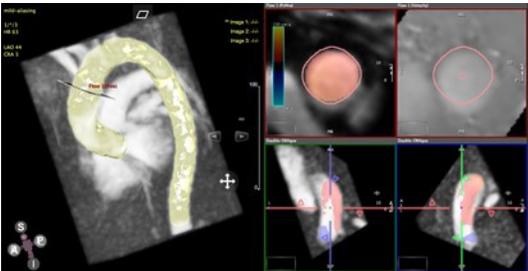
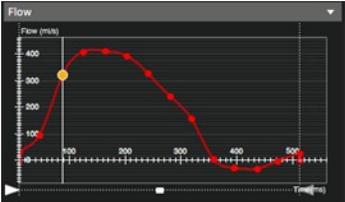
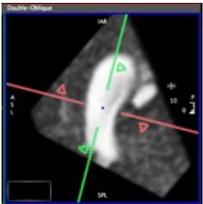
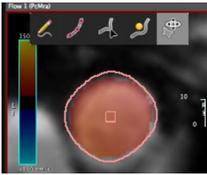
- Flow visualizations
- Quantitative Flow analysis

23.4.1 How to Visualize Flow in 4D

<p>1. Select a flow visualization from the top tool bar</p>	
<p>2. Start the cine and adjust the frame rate</p>	
<ul style="list-style-type: none"> • Velocity Visualization 	

<ul style="list-style-type: none"> • Vector Visualization: Displays speed and direction. Change the density of the vectors between 1-20 lines/ml 	
<ul style="list-style-type: none"> • Pathline (Planes) Visualization: Evolution of 3D blood flow over one or more heartbeats. Note: User needs to check accuracy of contours before observing the particle tracing. • Pathline density can be adjusted via slider heartbeats. Requires a centerline. • Pathline of a specific phase: By default, pathlines are emitted continuously. Use the slider on the bottom of the page to scroll through the phases to identify the onset of e.g. a jet. 	
<ul style="list-style-type: none"> • Pathline color: Illustrate complex flow dynamics in congenital heart disease. <p>Note: Single color map mode does not indicate velocities.</p> <ol style="list-style-type: none"> 1. Click on the flow plane and select <i>Set Pathline Color</i> from the context menu 2. To save and add a second plane, click <i>Add Measurement</i> 3. Position the free plane and set a color 4. Again, save the plane by clicking <i>Add Measurement</i> 5. Run cine to visualize the custom color-coded flow 	
<ul style="list-style-type: none"> • Pathline (Mask) Visualization: Display pathlines of particles emitted from all areas within the segmented mask. Select phase and density of the pathlines 	
<ul style="list-style-type: none"> • Streamline Visualization: 3D velocity fields at a specific temporal phase. Change the density of the streamlines. 	

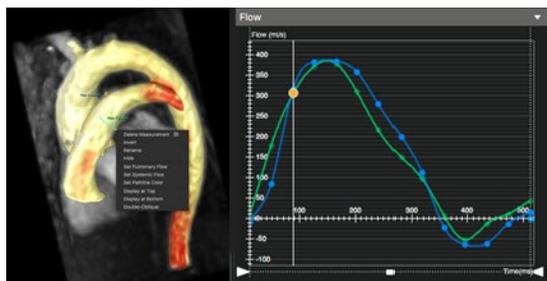
23.4.2 How to do a Flow Measurement

<p>1. Select a vessel from the list</p>																									
<p>2. Place a plane in your ROI. Planes will follow a center-line, if present. (toggle on/off the centerline display with short-cut key 'T')</p> <p>Note: The flow direction of the plane is indicated by an arrow. The direction can be changed via context menu > <i>Invert</i>.</p>																									
<ul style="list-style-type: none"> • Magnitude and Velocity images will be displayed next to the MRA. • Maximize the view: Space-bar-click on a view-port •  Optionally, activate a color-coded flow overlay to the cross-sectional images • <i>Velocity</i> and <i>Vectors</i> can be displayed color-coded in cross-sectional views 																									
<ul style="list-style-type: none"> • A contour will be drawn and propagated automatically, and a flow curve will be displayed instantaneously 																									
<p>3. Adjust the double oblique views to accurately display cross-sectional images</p>																									
<p>4. Run a cine to check for correct edge detection</p> <p>5. Single-click the contour to adjust it. Note: The modified contour is only forwarded to the end of the cycle.</p>																									
<ul style="list-style-type: none"> • Different contouring modes, such as <i>Point-Click</i> or <i>Thresholding</i> will be available by dragging the mouse on the top position tag 																									
<ul style="list-style-type: none"> •  Add a measurement triggers the flow calculation: Net, peak and regurgitant flow will be calculated automatically. • As soon as there are more than 2 planes, pulse wave velocity will be calculated. 	<table border="1" data-bbox="1127 1640 1515 1829"> <thead> <tr> <th>Measurement</th> <th>Net Flow (ml/cycle)</th> <th>Peak Vel. (cm/s)</th> <th>Regur. (%)</th> </tr> </thead> <tbody> <tr> <td>Flow 1</td> <td>98.98</td> <td>120.00</td> <td>2.50</td> </tr> <tr> <td>Aorta Ascendens(Qs)</td> <td>116.12</td> <td>131.00</td> <td>5.59</td> </tr> <tr> <td>Flow 3</td> <td>62.03</td> <td>116.00</td> <td>14.77</td> </tr> <tr> <td>Aort Decedens</td> <td>59.24</td> <td>103.00</td> <td>15.57</td> </tr> <tr> <td>Flow 6</td> <td>49.04</td> <td>109.00</td> <td>8.66</td> </tr> </tbody> </table> <p>Pulse Wave Velocity (m/s): 3.93312</p>	Measurement	Net Flow (ml/cycle)	Peak Vel. (cm/s)	Regur. (%)	Flow 1	98.98	120.00	2.50	Aorta Ascendens(Qs)	116.12	131.00	5.59	Flow 3	62.03	116.00	14.77	Aort Decedens	59.24	103.00	15.57	Flow 6	49.04	109.00	8.66
Measurement	Net Flow (ml/cycle)	Peak Vel. (cm/s)	Regur. (%)																						
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Aort Decedens	59.24	103.00	15.57																						
Flow 6	49.04	109.00	8.66																						
<ul style="list-style-type: none"> • To label a contour right-click on the plane in the PC MRA, or right-click on the flow measurement in the report, select <i>Rename</i> from the menu 	<ul style="list-style-type: none"> Rename Hide Delete Measurement Set Pulmonary Flow Set System Flow 																								

23.4.3 How to do a Shunt Calculation Qp/Qs

Other than in 2D Flow, planes must be set in the same structure.

1. Segment the vessels (no centerline) and move to the analysis page
2. Place a plane in the aorta and in the pulmonary artery
3. Scroll through the phases and correct contours if necessary



4. Open the context menu by clicking on the respective plane, e.g. in the ascending Aorta and select *Set Systemic Flow*



5. Add measurement

	Net Flow	Peak Vel.	Regur.
	(ml/cycle)	(cm/s)	(%)
Flow 7[Qp]	78.94	75.00	2.88
Flow 9[Qs]	81.40	132.00	7.04
Qp/Qs	0.969762		

6. Place a second plane in the Pulmonary artery and select *Set Pulmonary Flow*



7. Add measurement

23.5 Advanced 4D Flow Parameter Analysis

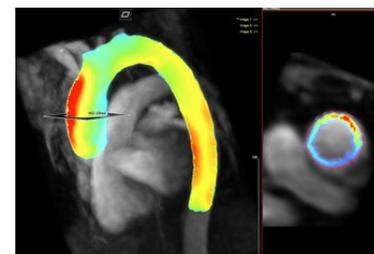
Note: A separate advanced license needs to be installed in order to use the ventricular flow analysis.

23.5.1 How to Assess Wall Shear Stress

cvi⁴² automatically calculates axial and circumferential wall shear stress as soon as a plane is set, and a contour is defined (*Flow* menu).



1. To visualize 3D wall shear stress, activate the button in the top tool bar and run the cine
2. The stress curve will be updated as you drag the plane along the centerline



3. Add a measurement to the report

4. WSS components will be calculated automatically and the curves and their respective measurements can be selected from the *Flow* Menu

	Avg Axial WSS	Avg Cir WSS
	(Pa)	(Pa)
Flow 3	0.22	0.13
Flow 4	0.08	0.18

23.5.2 How to do Relative Pressure Mapping

The software will compute a relative pressure field, giving the pressure difference between any two points. This will be translated into a colored pressure map.

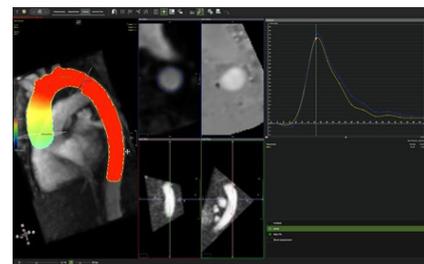


1. Activate the Pressure Visualization button
2. You will be given a reference and a free plane. Position them in your ROI.



3. Add a measurement

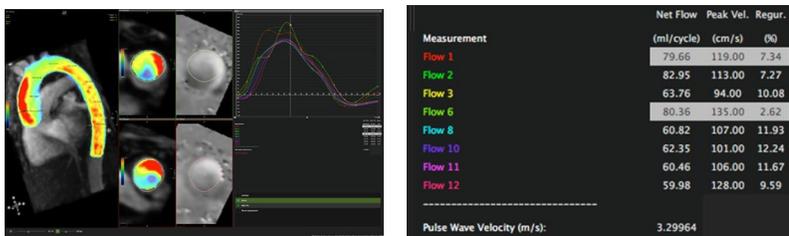
4. Pressure curves from the measurement and the free plane will be displayed in a graph. The relative pressure difference to the reference plane will be reported



	Max Pressure	Avg Pressure
	(mmHg)	(mmHg)
RPM 2	26.47	8.95

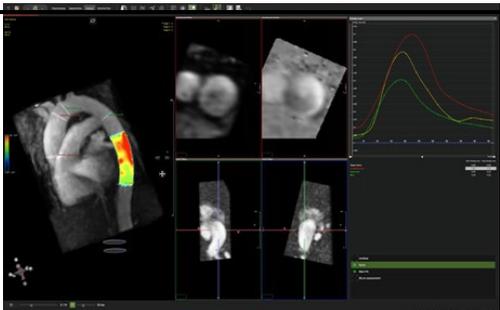
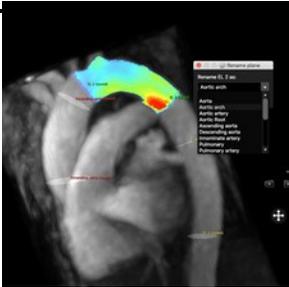
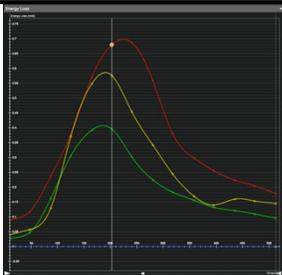
23.5.3 How to Calculate Pulse Wave Velocity

1. Pulse wave velocity (PWV) is a biomarker directly related to vessel stiffness that has the potential to provide information on early atherosclerotic disease burden
2. Pulse wave velocity calculation requires a centerline for the vessel structure with at least two flow measurement planes
3. Once two measurements have been provided (The limit is 12 planes) the system automatically calculates and reports PWV



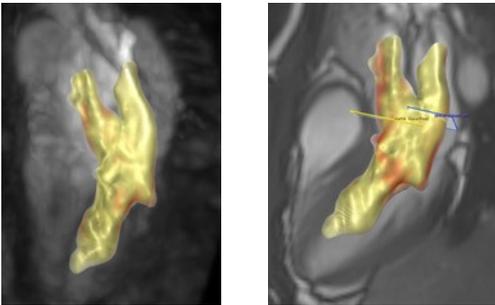
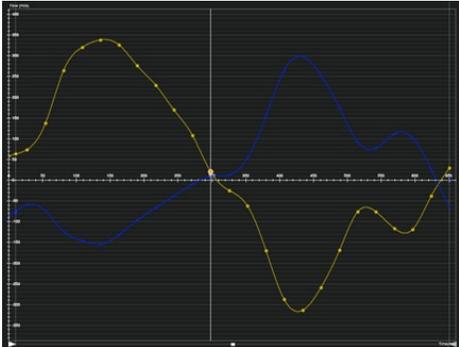
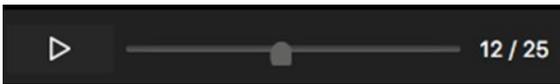
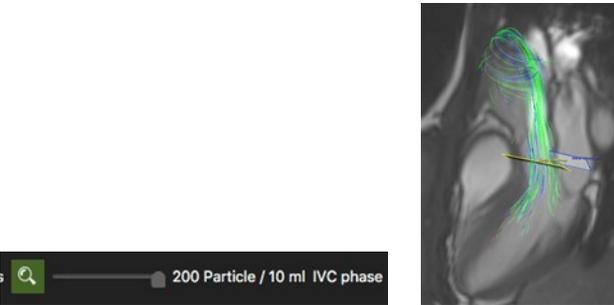
23.5.4 How to Analyze Energy Loss

Note: Not for clinical use.

<ol style="list-style-type: none"> 1.  Activate the button 2. Define a region by placing 2 planes 3. Add a measurement 													
<ol style="list-style-type: none"> 4. Hovering over the plane will display the color overlay 5. Label the region by either right-clicking on the variable in the report or the plane. Choose a description from the menu if present 													
<ol style="list-style-type: none"> 6. Maximum and average energy loss over time will be reported and the curves per region (limited to 6) are displayed 	<table border="1"> <thead> <tr> <th>Region Name</th> <th>Max Energy Loss (mW)</th> <th>Avg Energy Loss (mW)</th> </tr> </thead> <tbody> <tr> <td>EL 1</td> <td>0.69</td> <td>0.36</td> </tr> <tr> <td>EL 2</td> <td>0.40</td> <td>0.20</td> </tr> <tr> <td>EL 3</td> <td>0.58</td> <td>0.26</td> </tr> </tbody> </table> 	Region Name	Max Energy Loss (mW)	Avg Energy Loss (mW)	EL 1	0.69	0.36	EL 2	0.40	0.20	EL 3	0.58	0.26
Region Name	Max Energy Loss (mW)	Avg Energy Loss (mW)											
EL 1	0.69	0.36											
EL 2	0.40	0.20											
EL 3	0.58	0.26											

23.5.5 How to Analyze Ventricular Flow

Note: A separate advanced license needs to be installed in order to use the ventricular flow analysis.

<ol style="list-style-type: none"> 1. <i>Segmentation Tab</i> <ul style="list-style-type: none"> • Clip the ventricle, e.g. via <i>3D Region Growing</i> (context menu or modifier key Ctrl/command + G) 2. <i>Ventricular Flow Tab</i> <ul style="list-style-type: none"> • Drag in a 3-chamber-view, run the cine and locate the valve • Place aortic and mitral valve plane 																												
<ol style="list-style-type: none"> 3. Move through the phases and based on flow curves, determine IVR (isovolumetric relaxation) and IVC (isovolumetric contraction). <p>Note: User needs to check the accuracy of the contours before exporting the results. The IVC phase is the time point that ventricle is full of blood and all the valves are closed and start to contract with constant volume. The IVR phase is the time point that the aortic and mitral valves are closed after end systole and ventricle starts to relax with the constant volume.</p> 																												
<ol style="list-style-type: none"> 4. Read the phase from the slider (underneath the 4D Viewer: 12) 																												
<ol style="list-style-type: none"> 4. Enter the phases in the respective fields. <p>Starting from these phases, the software will track forward and backward pathlines and will separate them into four different flow components: direct flow, retained inflow, delayed ejection flow, and residual volume. Note: The accuracy of the path line for each component depends on user experience to segment the ventricle, place the position of the mitral and aortic valves and set up appropriate IVC and IVR phase number.</p> 																												
<ol style="list-style-type: none"> 6. Click the <i>Flow Component Analysis</i> button in the toolbar 7. Start the cine 8. Adjust the number of particles 																												
<ol style="list-style-type: none"> 9. Read the report 	 <table border="1"> <tbody> <tr> <td>Direct Flow:</td> <td>63.86</td> <td>%</td> </tr> <tr> <td>Delayed Ejection:</td> <td>22.59</td> <td>%</td> </tr> <tr> <td>Retained Inflow:</td> <td>11.14</td> <td>%</td> </tr> <tr> <td>Residual Volume:</td> <td>2.41</td> <td>%</td> </tr> <tr> <td colspan="3">-----</td> </tr> <tr> <td></td> <td>Net Flow</td> <td>Peak Vel. Regur.</td> </tr> <tr> <td>Measurement</td> <td>(ml/cycle)</td> <td>(cm/s) (°)</td> </tr> <tr> <td>Aortic Valve</td> <td>60.47</td> <td>160.00 2.28</td> </tr> <tr> <td>Mitral Valve</td> <td>43.94</td> <td>73.00 59.50</td> </tr> </tbody> </table>	Direct Flow:	63.86	%	Delayed Ejection:	22.59	%	Retained Inflow:	11.14	%	Residual Volume:	2.41	%	-----				Net Flow	Peak Vel. Regur.	Measurement	(ml/cycle)	(cm/s) (°)	Aortic Valve	60.47	160.00 2.28	Mitral Valve	43.94	73.00 59.50
Direct Flow:	63.86	%																										
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Measurement	(ml/cycle)	(cm/s) (°)																										
Aortic Valve	60.47	160.00 2.28																										
Mitral Valve	43.94	73.00 59.50																										

24 Coronary Module (Auto plaque)

- 24.1 Landmark Page 73
- 24.2 Centerline Page 73
 - 24.2.1 How to Define a Centerline via Region Growing 74
 - 24.2.2 Define a Centerline via Seed Points 74
 - 24.2.3 Edit the Centerline 74
 - 24.2.4 Save Options 76
 - 24.2.5 3D View 76
- 24.3 Cross Section 75
 - 24.3.1 How to Do a Stenosis Assessment 75-76
 - 24.3.2 How to Use the Centerline Graph 76
 - 24.3.3 Endoluminal View and a Fly Through 76
- 24.4 Plaque Analysis 76
 - 24.4.1 How to Perform a Plaque Analysis 77

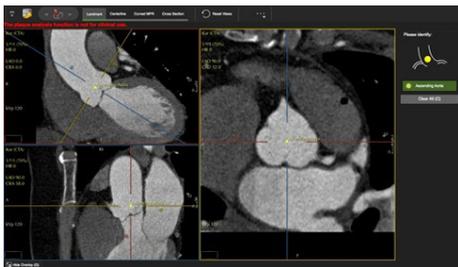
The Coronary Module includes the following pages:

- Landmark: aids the coronary segmentation
- Centerline: Create a centerline for CPR
- Cross Section: Evaluation

Auto-plaque assessment requires a separate license.

24.1 Landmark Page

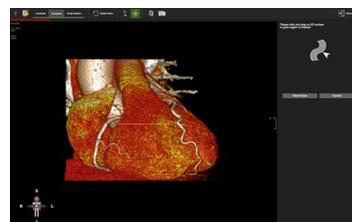
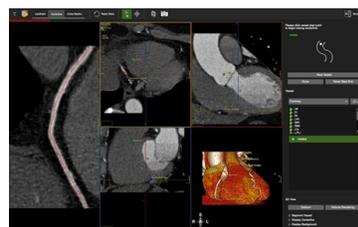
Placing a Landmark in the ascending aorta will support the centerline generation.



24.2 Centerline Page

- Create a centerline via 3D-Region-Growing or setting seed points (start and end points)
- Using the landmark point, the system will automatically segment the heart
- The thresholding page will be the default, if you prefer to create a centerline by setting seed points, exit the page

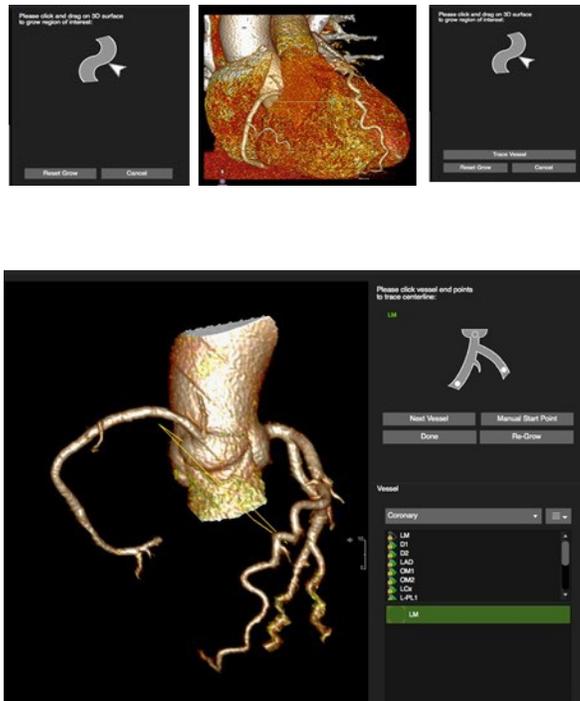
by clicking 3D-Region-Growing in the tool bar



24.2.1 How to Define a Centerline via Region Growing

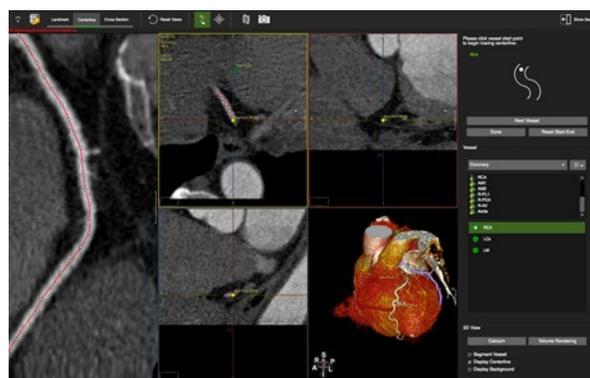
The Grow Region Interface provides a guide diagram to indicate the next user action.

1. The *Region Growing* tool is activated by default. Define a single vessel or the entire coronary tree.
2. Drag on the 3D surface to define a vessel or a vessel tree
3. The modifier key 'alt' will rotate, Ctrl/ (command for Mac) will pan the volume
4. Click Trace Vessel when you are done
5. *Reset Grow* if you are not satisfied
6. Using the landmark, the system will automatically place a start point
7. To relocate the start point click *Manual Start Point*
8. The software has a predefined list of coronaries and will automatically move to the next vessel in the list after setting an end-point
9. Move to the next list item by clicking *Next Vessel*.
10. To relabel a centerline capture, drag the appropriate label on the capture with the green dot
11. Add/Delete a new label with a right mouse click in the background



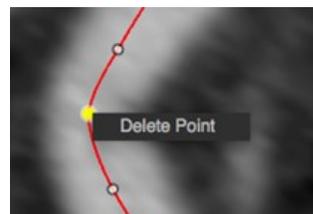
24.2.2 Define a Centerline via Seed Points

1.  Move to the centerline page
2.  Activate the seed point
3. In the axial view locate the origin of the vessel
4. Set a start, optionally intermediate and an end point



24.2.3 Edit the Centerline

1. Shift, add (single left-click), delete (right-click) control points in the CPR or the oblique views
2. Rotate around the centerline with a left mouse drag close to the centerline
3. Pan the image by moving the cursor away from the centerline until the hand symbol appears



24.2.4 Save Options

- Save rotational images



- Multi Frame Captures (See MPR Viewer)



24.2.5 3D View

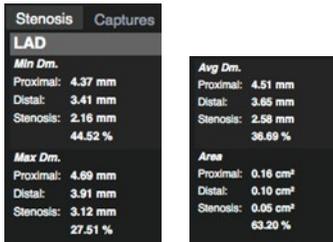
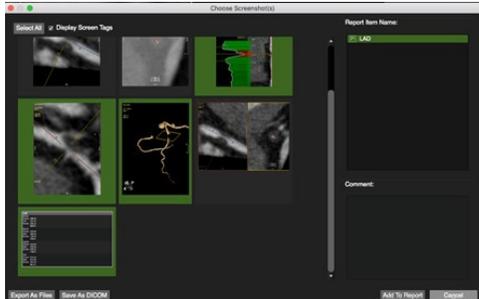
<p>1. Select Calcium or Volume Rendering</p>	
<p>2. Segment Vessel: check it will segment vessels with all centerlines</p> <ul style="list-style-type: none"> • Display/hide centerline • Display/hide background 	

24.3 Cross Section

The page is designed to visually and quantitatively assess the vessels with existing centerlines.

24.3.1 How to do a Stenosis Assessment

<ol style="list-style-type: none"> 1. Start in <i>Browse</i> view and drag the yellow line down the vessel 2. Switch to <i>Stenosis</i> View and place the stenosis marker either by dragging or by double-clicking in the stenosis 	
<ol style="list-style-type: none"> 3.  Move to the MPR view and use the <i>Auto Vessel Lumen Contour</i> to contour the stenosis and the 2 references images 4. To adjust the threshold, click the green contour label. It will open a slider that allows to adjust the contour 	

<p>5. Stenosis measurements will be filled in automatically</p>	
<p>6. Use the <i>Multiframe Capture</i> to save the measurements and images</p>	

24.3.2 How to Use the Centerline Graph



Centerline Graph displays min./max./average lumen diameter, lumen area and tortuosity. It will automatically detect the minimum and maximum point along the centerline graphs.

Options:

- *Lumen* menu, offering display options
- Cut off: Drag the yellow handle to set a cut off



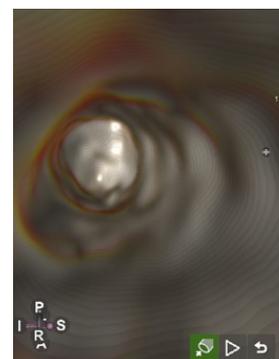
- The ruler measures a custom defined distance



24.3.3 Endoluminal View and Fly Through



1. In the reference viewer click the *Endoluminal view*
 2. Drag the stenosis marker in the CPR viewer
 3. Click the *Play* button, for a fly through animation
 4. Export via *Context Menu>Export>Export Video Immersion*
- The context menu offers viewing, rendering, saving, export and more options



24.4 Plaque Analysis

WARNING: Auto-Plaque

The use of the *Auto-Plaque* is not FDA approved and is for non-clinical, scientific use only.

25 Calcium Scoring

25.1	Toolbar.....	78
	25.1.1 How to Do Calcium Scoring.....	78-79

Calcium Scoring is a method to quantify the plaque load of the coronary arteries.

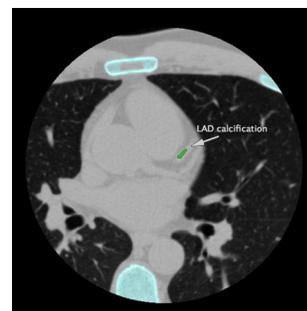
Post processing for Calcium Scoring requires the detection of calcium and assignment to a coronary.

25.1 Toolbar

- The toolbar provides 4 buttons to identify Calcium in the different coronary arteries
- Ca Mass calibration factor: Using a phantom with calcium inserts of known mass allows for determination of a calibration factor for different MDCT scan protocols

25.1.1 How to do Calcium Scoring

1. Open the on-screen navigation buttons or use your keyboard arrow keys to scroll through the slices

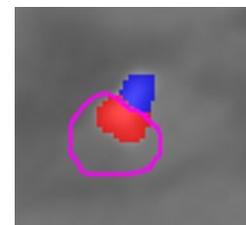


2.  Toggle the Overlay Display and scroll through the volume

3. When you have detected Ca, select the respective coronary ROI-button and simply click in the calcified area

4. To re-do the region selection, click again

5.  To reset all region selections, use the *Reset Selection* button in the toolbar



6.  To assign calcification to two different coronaries:
7. Split the lesion with the Ca Scoring Selection Contour
8. Select the first matching ROI (e.g. LM) and click into the respective section
9. Then, select the second ROI (e.g. LAD) and click into the other part of the lesion

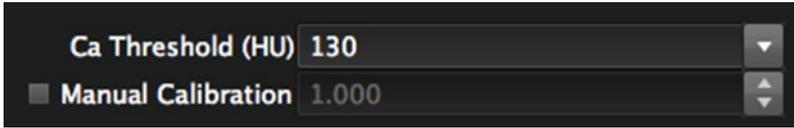
10.  Add the analysis to your report

Threshold

- Set the Threshold (default 130) (on the bottom of the reporting panel, underneath the Percentiles)

Calibration Factor

- The Calibration Factor will be retrieved from the DICOM header or can be entered manually (Calibration Factors are specific to scanner, protocol and patient weight)



Agatston Classification

- Volume, Mass and Agatston Score and Agatston Classification determined automatically

Percentiles

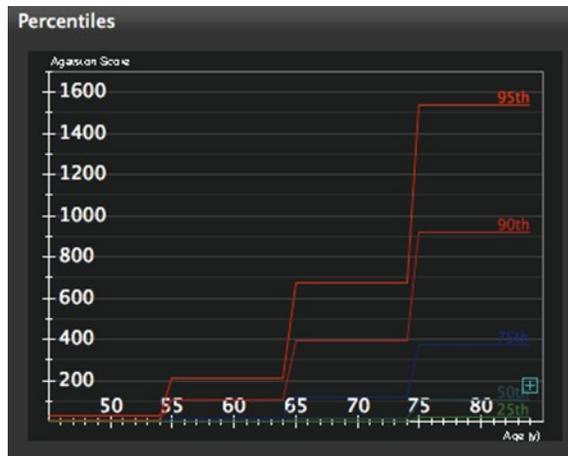
- To determine the Percentiles of the patients Agatston Score per coronary, select the ethnic group

Ca Scoring Report

Artery	Volume mm ³	Mass mg	Score Agatston
LM	0.00	0.00	0.00
LAD	75.44	27.55	100.59
CX	0.00	0.00	0.00
RCA	54.44	19.51	72.59
Total	129.88	47.06	173.18

Ca Threshold: 130.00
Ca Mass Calibration Factor: 1.00

Agatston Score Classification:
Moderate Coronary Artery Calcification



26 Aortic Valve Module

26.1	Landmark page	80-81
26.2	Annulus Page	81
	26.2.1 Measurements	81-82
	26.2.2 Reporting options	82
	26.2.2.1 Capture Measurements and Images	83
	26.2.2.2 Standard Report.....	83
26.3	Aorta Fluoro Page	83
26.4	Calcium Page	84
	26.4.1 Control Feature	84
	26.4.2 How to Detect Calcium.....	84-85
26.5	Valve in Valve	85
26.6	Apical Page	86

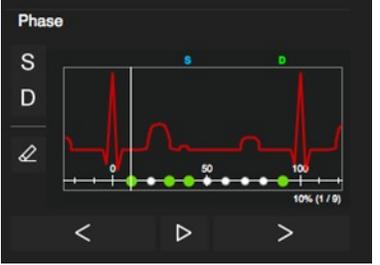
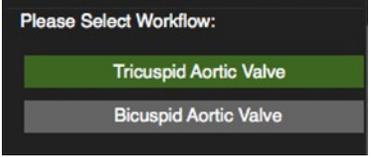
Intended Use

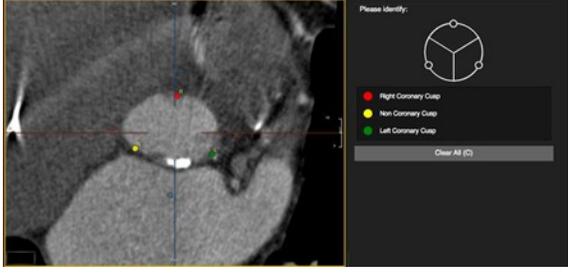
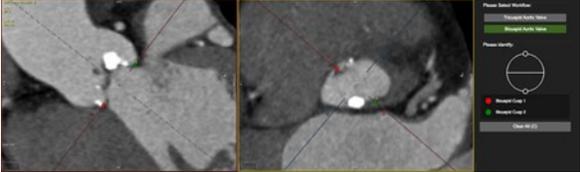
cvi⁴² – The Aortic Valve module offers a structured workflow for pre-procedural planning of trans catheter aortic valve replacement.

26.1 Landmark page

Landmarks will define the annulus plane.

If measurements already exist, the module will skip the landmark page and will open the Annulus page.

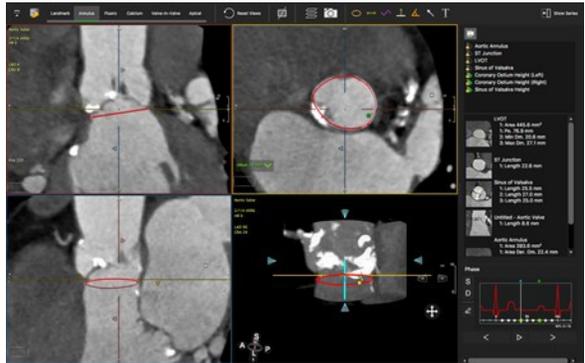
<p>1. Open the side panel and load the data via drag-and-drop or double-click</p>	
<p>2. Start with selecting a phase for assessment</p> <ul style="list-style-type: none"> Optionally set a bookmark for <i>Systole</i> and <i>Diastole</i> 	
<p>3. Select Workflow:</p> <ul style="list-style-type: none"> Choose between a bi- or tricuspid valve 	

<p>4. Identify the cusps (tricuspid)</p> <ul style="list-style-type: none"> In random order, place three points at the hinge of the right, left and non-coronary cusp. The software will identify and label the cusp points 	
<p>5. Identify the cusps (bicuspid)</p> <ul style="list-style-type: none"> Clicking on one of the hinge point markers will automatically rotate the red crosshair to display the point in the respectively color-coded orthogonal viewer frame Modifier key 'M' hides the cross-reference lines If necessary, the marker can be adjusted with a mouse drag in either frame 	

26.2 Annulus Page

26.2.1 Measurements

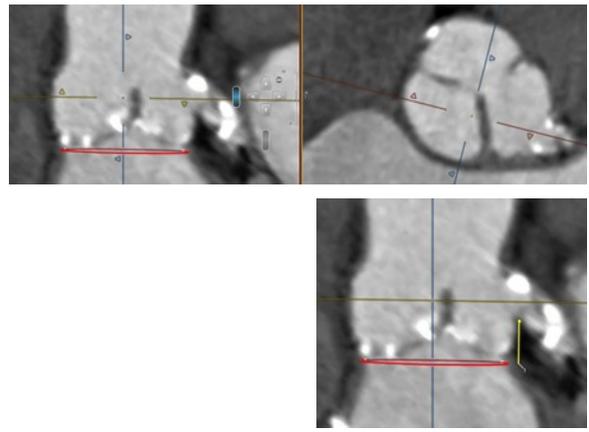
Aortic Annulus

<ol style="list-style-type: none">  Use the <i>Perimeter</i> tool and set control points around the annulus. You can maximize the view with a spacebar-click, or by clicking in the top right corner of the view port To edit the perimeter, single-click on the control points. Single-click in the background to disengage The annulus will be the only measurement that is labelled automatically 	
<p>Ancillary measurements:</p> <ul style="list-style-type: none"> 'Reset View' button in the toolbar will take you back to the annulus plane 	
<ul style="list-style-type: none"> Tools are in the top tool bar and the pencil box in the left bottom corner of the viewer frame. Measurements can be done in any viewer frame 	
<ul style="list-style-type: none"> Measurements and images are captured automatically Drag the appropriate label from the predefined list on the capture To review a measurement, simply click on the capture Right-click to delete the capture. Alternatively, use the delete key 	

4. Ostium Height measurement

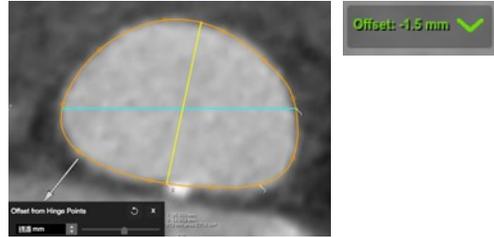


- Using the *Ostium Height* measurement tool, drop a point at the height of the ostium with a left mouse click
- Drag the label on the measurement capture
- Repeat for the other coronary artery



5. LVOT measurement

- Open the small *Offset* box in the bottom left corner (see white arrow), and use the slider to scroll down into the LVOT
- Using the perimeter tool, measure at narrowest portion of the LV outflow tract



26.2.2 Reporting Options

26.2.2.1 Capture Measurements and Images

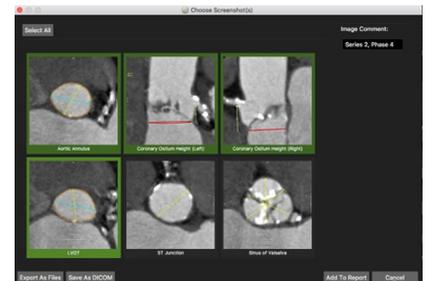
- Worklist measurements, labelled with the tags provided, will automatically be entered into an infographic (see below) in the report module.
- Measurements changes will be updated.



Measurement Captures (Report Panel):



- Activating the measurement capture camera will allow to add respective pictures to the report.

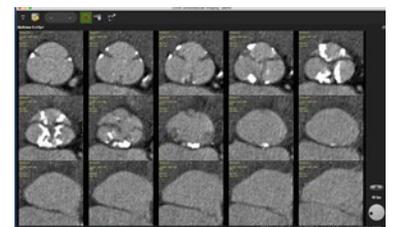


Multi Frame Screen Captures (Tool Bar):

- Captures images of the current layout e.g. corresponding cross-sectional images of a particular plane or measurement.



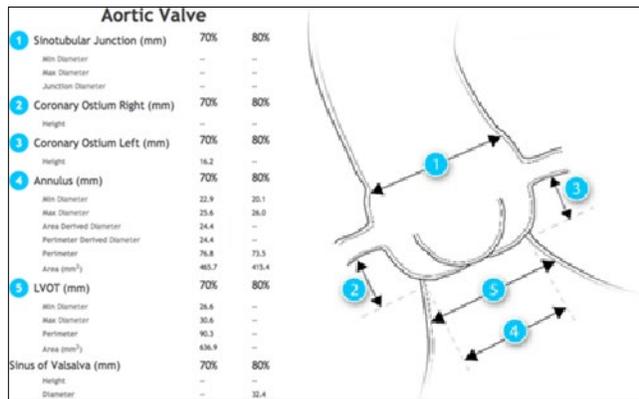
- Save *Multi stack Images* (toolbar) as a secondary DICOM capture.
- Individual frames can be added to the report by selecting the *Add to Report* option in the context menu.



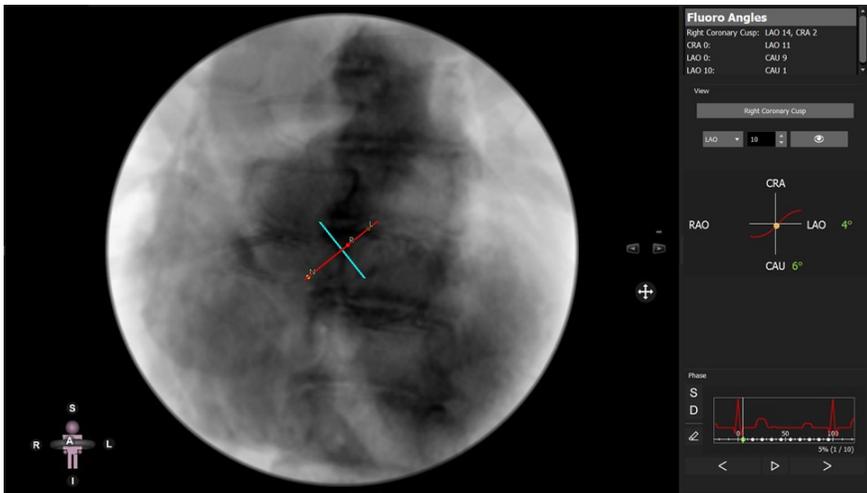
26.2.2.2 Standard Report

Upon opening the default report module:

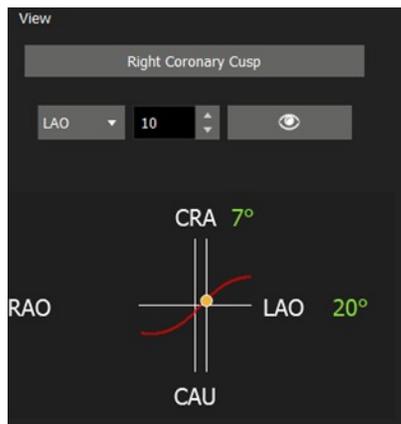
- Review the infographic and related measurements



26.3 Aorta Fluoro Page



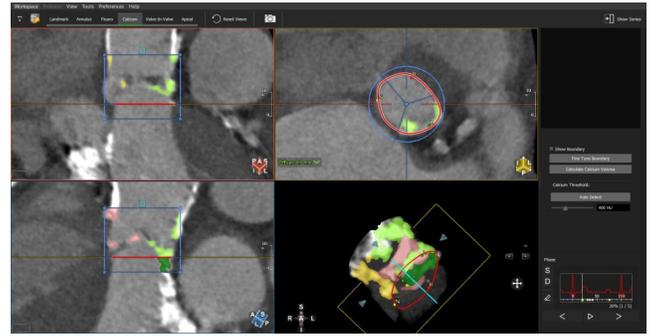
Each point on the “s-curve” will return an angio angle perpendicular to the aortic valve plane.



These angles can be used as suggested orientation for C-arm during the intervention, due to the fact that the conditions during the CT scan are not identical to those during the procedure, angles may vary.

26.4 Calcium Page

- 3 MPR and a 3D Viewer
- Auto detect or manually adjust the calcium threshold
- Cusp specific color-coded calcium overlay
- Calcium volume report per cusp, in aorta and LVOT



26.4.1 Control Feature

1. Mouse drag in the long axis will rotate around the valve center
2. Mouse drag in the short axis will scroll up/down the Aorta/LVOT
3. Press the Ctrl (command in Mac) key to pan the image in the view port

26.4.2 How to Detect Calcium

<ol style="list-style-type: none"> 1. In the valve view, scroll up to the sinus to display the commissures 2. The calcium color corresponds to the nadir point colors. Adjust the center and move the arms of the cusp divider, so that the colors are properly assigned to a cusp 	
<ol style="list-style-type: none"> 3. Drag horizontal lines to increase or decrease height 4. Drag control points to increase or decrease width 	
<ol style="list-style-type: none"> 5. Use the fine tune option to include or exclude certain structures, e.g. to exclude the coronaries. Once the fine tune option is enabled drag the control points in the short and long axis views 	<p>Fine Tune Boundary</p>
<ol style="list-style-type: none"> 6. Using the <i>Auto Detect</i> button, the software will automatically place a sample area (blue square) above the annulus in the log axis view. From that sample, a threshold will be determined to auto detect calcium. (According to the formula: average HU + 4.0 * standard deviation) 7. Alternatively, drag the slider to manually change the threshold for HU 	<p>Calcium Threshold: Auto Detect 485 HU</p>

8. In the report pane click *Calculate Calcium Volume* for a calcium report



9. Click on the camera to add the Calcium Volume and images to your report

Calcium Volume	
Aortic Valve Region Calcium:	1213.5 mm ³
- Left Coronary Cusp:	305.8 mm ³
- Right Coronary Cusp:	387.0 mm ³
- Non-Coronary Cusp:	520.7 mm ³
Upper LVOT Calcium:	44.0 mm ³
- Below Left Coronary Cusp:	28.8 mm ³
- Below Right Coronary Cusp:	0.0 mm ³
- Below Non-Coronary Cusp:	15.2 mm ³

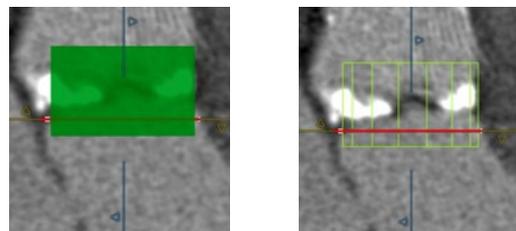
26.5 Valve in Valve

- Device simulation
- Additional measurements



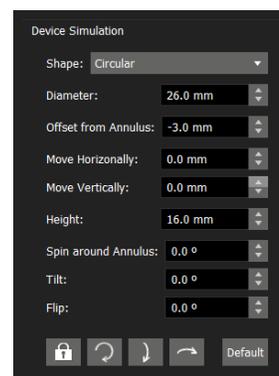
For device simulation choose between:

-  Display Device intersection
-  Display Device Mesh



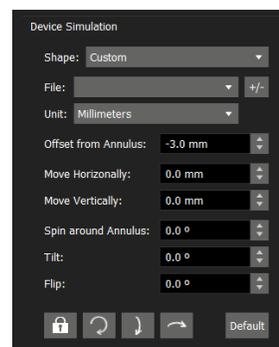
• In the report pane choose Circular shape and use the options provided to position the device.

-  Lock device position across different phases.



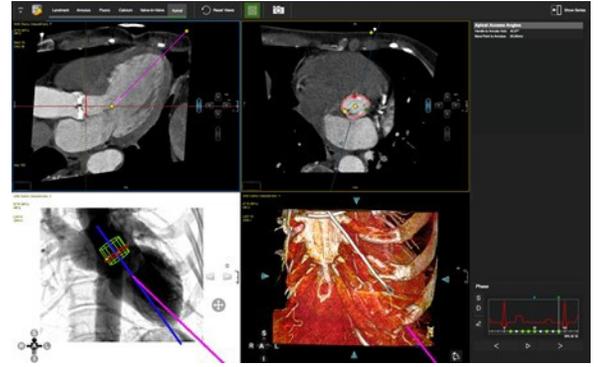
Custom Shape:

- Allows to import vendor specific image files
- Change units from metric to imperial

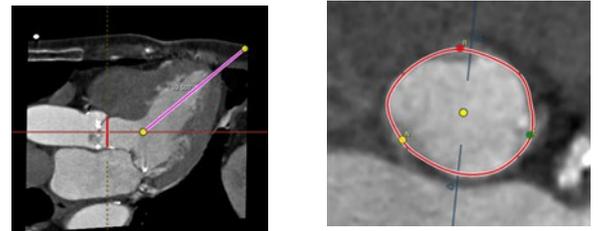


26.6 Apical Page

- Apical access planning



- A red line in the top left viewer indicates the central axis perpendicular to the annulus (trajectory), the ideal access path.
- Rotate around the trajectory line to display the apex



- The purple line can be adjusted to the manual access route
- Adjust the yellow control point that slides along the red trajectory line, to accommodate for bending of the manual access route

Apical Access Angles

Handle to Annular Axis: 45.26°
Bend Point to Annulus: 25.00mm

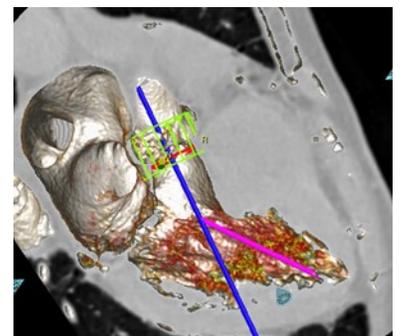
- Record the intercostal space

-  Assess the course of the coronary arteries



Display Device Mesh:

-  Toggles to an endoluminal view to display the device
- The context menu allows to change viewing and/or clip-pings

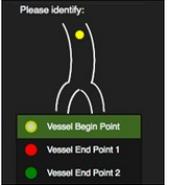


27 Femoral Module

27.1	Landmark Page	87
27.2	Centerline Page	87
	27.2.1 Edit the Centerline.....	88
	27.2.2 Assessment of the Puncture Site.....	88
27.3	Cross Section	88
	27.3.1 Reporting	89

The Femoral Module allows to assess the access for cardiac interventions.

27.1 Landmark Page

<ul style="list-style-type: none"> Segment the aorta by clicking on the abdominal aorta. Optionally, skip the segmentation 	
<ul style="list-style-type: none"> Setting landmarks will assist the software to create centerlines 	
<p>Using the MPR and/or volume view set:</p> <ul style="list-style-type: none"> Point 1 in the aorta abdominalis Point 2 in the left femoral artery Point 3 in the right femoral artery To Delete a point right click on them and select "Delete Object" Move to the Centerline Page 	

27.2 Centerline Page

- Edit centerline control points

- Assess the calcium load

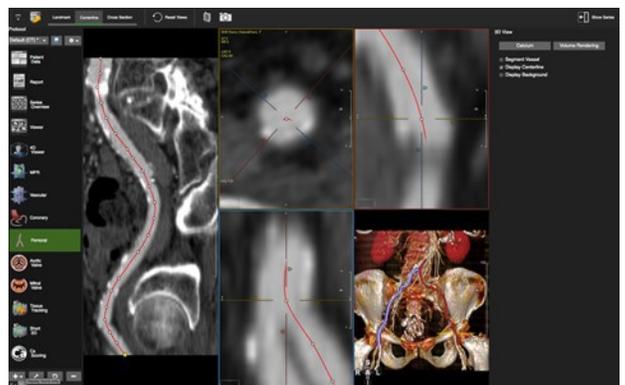
Based on the landmarks a centerline is generated automatically.

- 'T' toggles the centerline on/off

- Select the vessel simply by clicking on the vessel of interest in the reference viewer

- Make sure the centerline is not crossing calcium.

The software will dynamically calculate a calcium threshold for automatic vessel lumen assessment based on the centerline points



27.2.1 Edit the Centerline

- Use the up/down arrow keys on the keyboard to navigate along the control points
- Add control points with a double-click on the centerline
- Delete a control point with a right mouse click on the point
- Rotate around the centerline with a left mouse drag close to the centerline
- Pan the image by moving the cursor away from the centerline until the hand symbol appears

27.2.2 Assessment of the Puncture Site

Use the *3D View* menu in the right-hand panel to:

- Hide the centerline
- Segment the vessels
- Add a background to visualize the femur heads
- Assess Calcium by switching to a *Calcium View*
- Enlarge the view by clicking the space bar key



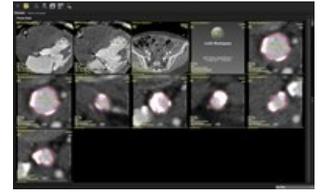
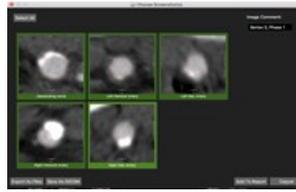
27.3 Cross Section

<ol style="list-style-type: none"> 1. Display the centerlines in the reference viewer 2. Select a vessel by clicking on the centerline in the reference image 	
<ol style="list-style-type: none"> 3. Open the centerline graph. The graph is based on the diameter of the best fitting circle inside the lumen, excluding calcium. The calcium threshold is calculated dynamically based on centerline points. Therefore, during centerline generation, attention should be paid that the centerline is not crossing any calcium 4. Drag the yellow cursor line along the vessel. The diameter is displayed continuously on the right side of the cursor line 	
<ol style="list-style-type: none"> 5. Measurements will be done automatically as soon as the centerline graph is activated, and you halt the cursor line 6. A contour can be drawn manually. In that case, the assumption is made that the contour is drawn excluding calcium and no circle will be fitted 7. Take captures via context menu 8. Use the predefined tags to label them 9. Click on the captures to review axial and orthogonal views 10. Captures will be displayed as cross-sectional images right of the CPR 	

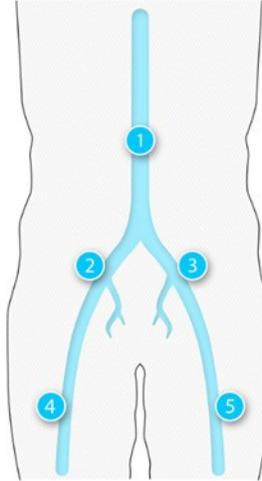
27.3.1 Reporting

Click the camera button in the reporting pane to:

- Add images to the report
- Export them in different formats
- Save the in DICOM format



Measurements labeled as described in the previous step will end up automatically in the infographic of the report.



Femoral Access	
Phase	0
1 Descending Aorta	
Diameter (mm)	16.3
2 Right Iliac Artery	
Diameter (mm)	9.0
3 Left Iliac Artery	
Diameter (mm)	9.5
4 Right Femoral Artery	
Diameter (mm)	6.4
5 Left Femoral Artery	

28 Mitral Valve Module

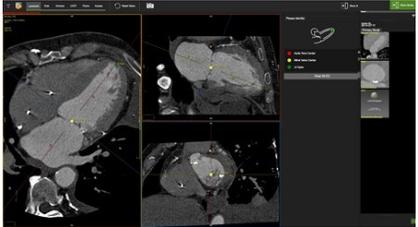
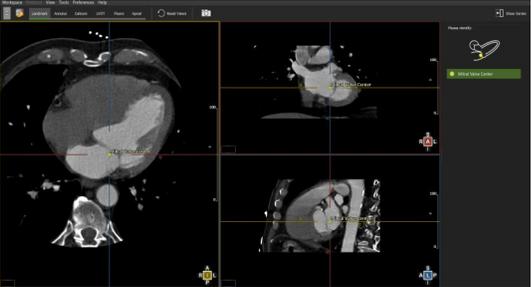
28.1	Landmark page.....	90
28.2	Annulus Page	91
	28.2.1 How to Define the Mitral Annulus (MA) Plane	91
	28.2.2 How to Define the Mitral Annulus (MA) Plane - Advanced	91-92
	28.2.3 Guide Wire Simulation.....	93
28.3	Calcium Page.....	93
	28.3.1 Control Feature.....	93
	28.3.2 Three Chamber View	94
	28.3.3 Quantify the Calcium Load	94
28.4	LVOT Page	94
	28.4.1 LVOT Assessment.....	95
	28.4.2 Device Simulation	96
28.5	Fluoroscopy Page	96
	28.5.1 Optimal Viewing Curve	96
28.6	Apical Page.....	97-99

Intended Use

The *Mitral Valve* module offers a structured workflow for pre-procedural planning of transcatheter mitral valve replacement (TMVR).

28.1 Landmark Page

Set landmark for the assisted generation mitral valve plane

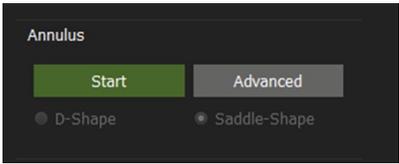
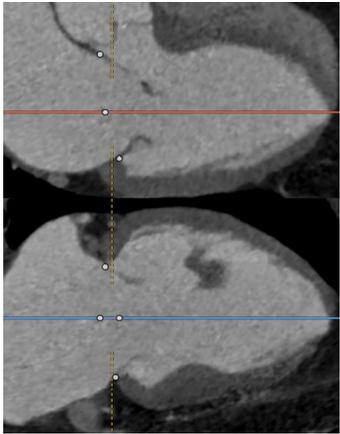
<ul style="list-style-type: none"> Click <i>Show Series</i> and load the series from the thumbnail pane. 	
<ul style="list-style-type: none"> If the software detects contours or measurements, the module will open in the <i>Annulus</i> page. Phases with measurements can be identified by the green dots. 	
<ul style="list-style-type: none"> The Aortic Root center will be detected automatically. Slice through the images by dragging the mouse in the axial image, to display the mitral valve center. Click to set the second landmark. The LV Apex will be placed automatically. Use the crosshairs and swing around the apex to check for correct positioning. Simply drag the points to reposition. 	

28.2 Annulus Page

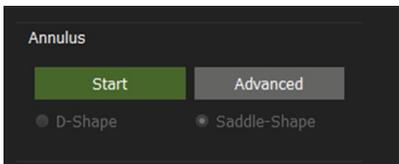
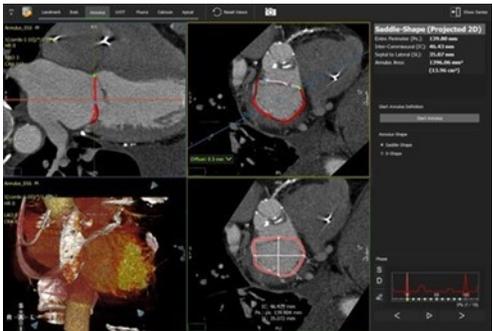
Definition and automated measurement of the valve plane for saddle and D-shaped annulus.

Note: When needed adjust the mitral valve center point and apex point.

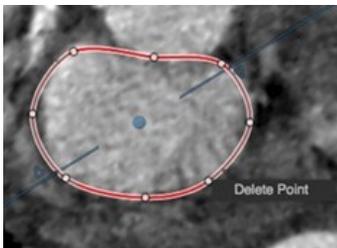
28.2.1 How to Define the Mitral Annulus (MA) Plane

<ol style="list-style-type: none"> 1. Select a phase 2. Press start 	
<ol style="list-style-type: none"> 3. Define 4 control points on the longitudinal views 	

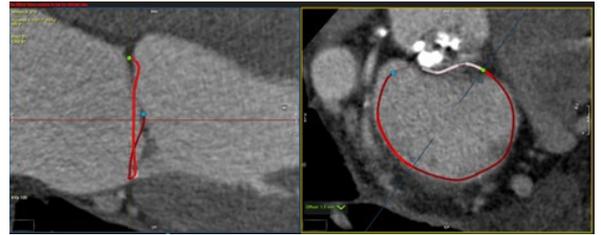
28.2.2 How to Define the Mitral Annulus (MA) Plane - Advanced

<ol style="list-style-type: none"> 1. Select a phase 2. Start the workflow 3. Press "advanced" 	
<ol style="list-style-type: none"> 4. Define control points: <ul style="list-style-type: none"> • The annular contour will be segmented within the 3D space by placing 16 seeding points at the insertion of the valve leaflet • After each click, the software automatically rotates around the red trajectory line by 22.5° • The short axis view is synchronized, displaying the plane that intersects trajectory line and last defined control point • After 16 clicks, the entire circumference of the annulus will be identified and is shown in all 4 viewports 	

Note: Both the regular and the advanced Annulus can be edited

<ol style="list-style-type: none"> 5. Edit the Annulus <ul style="list-style-type: none"> • Double-click to toggle control points on/off • Drag to shift points • Left-click to add points • Right-click to delete points • To redo the annulus, restart the annulus definition by clicking the bar again 	
--	---

- Exiting the editing mode, allows to adjust the trigones to the LVOT



- Measurements are done automatically

Saddle-Shape (Projected 2D)

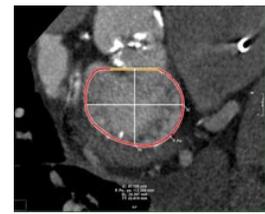
Entire Perimeter (Pe.):	140.74 mm
Inter-Commissural (IC):	46.82 mm
Septal to Lateral (SL):	37.49 mm
Annulus Area:	1487.69 mm ² (14.88 cm ²)



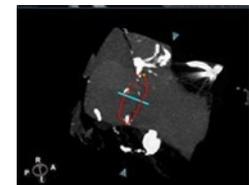
- **Annulus Shape**
 - Saddle-Shape
 - D-Shape
- Switch between saddle and D-shape

D-Shape (Projected 2D)

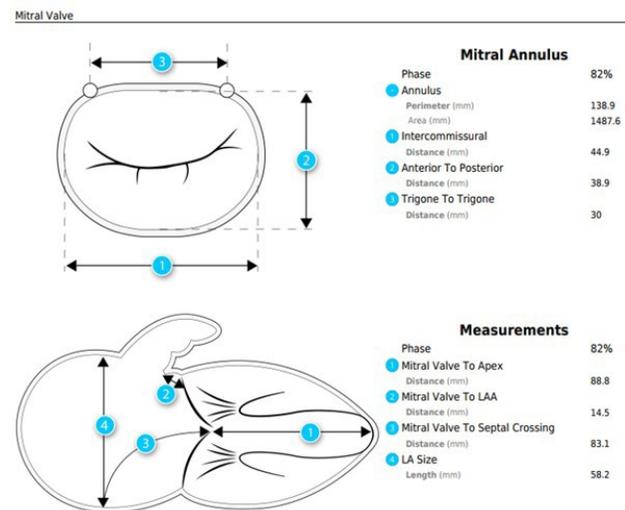
Posterior Perimeter (P. Pe.):	107.69 mm
TT Distance (TT):	31.31 mm
Posterior Per. + TT:	138.99 mm
Inter-Commissural (IC):	46.76 mm
Septal to Lateral (SL):	37.21 mm
Annulus Area:	1452.09 mm ² (14.52 cm ²)



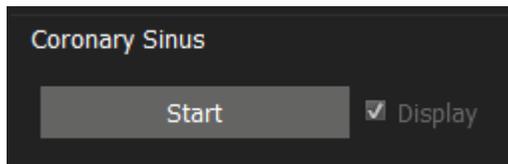
- In the *4D Visualization* viewer adjust the cylindrical clipping cone to define the annular region. Visually inspect the calcification around the valve



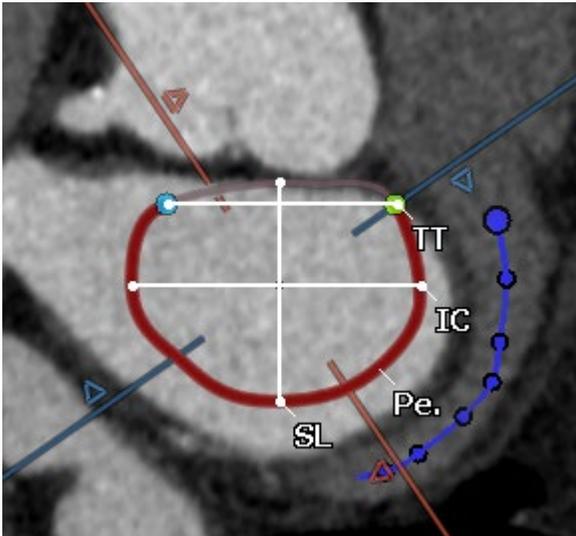
- Labeled measurements will be shown in the infographic in the report



28.2.3 Guide Wire Simulation

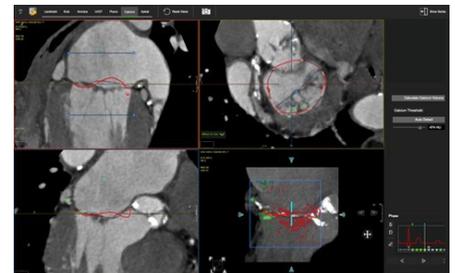


1. Press **Start** to begin defining the Coronary Sinus.
2. Scroll through the data set in the short-axis view or rotate around the horizontal axis in the long-axis view and click to add key points along the sinus.
3. Double-click on the last point to finish.



28.3 Calcium Page

- Auto detect or manually adjust the calcium threshold
- Calcium will be displayed in a color-coded calcium overlay
- Calcium volume will be reported per leaflet

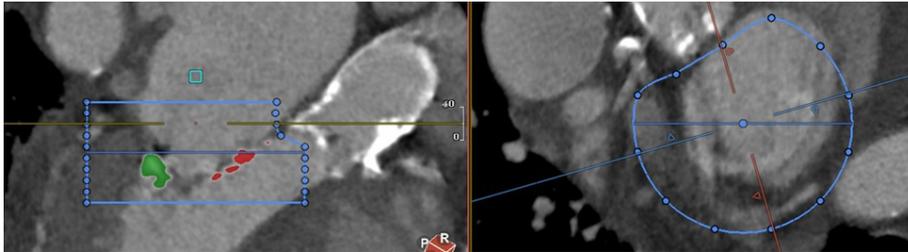
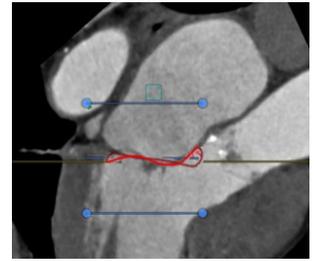


28.3.1 Control Feature

- Mouse drag in the long axis will rotate around the valve center
- Mouse drag in the short axis will scroll up/down along the trajectory line
- Press the Ctrl (command in Mac) key to pan the image in the view port

28.3.2 Three Chamber View

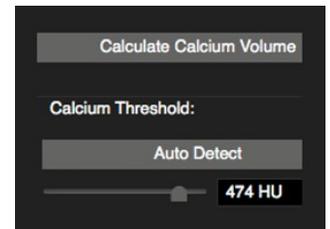
- The area of interest (cylinder) can be in- or decreased by dragging the blue lines up/down or left/right in the long axis viewer (top left).
- The small square in the atrium on top of the line indicates the sample used to calculate the calcium threshold
- Use the fine tune button to include or exclude certain areas



28.3.3 Quantify the Calcium Load

- In the report pane click *Calculate Calcium Volume* for a calcium report
- You can customize the calcium threshold. Click calculate Calcium Volume again

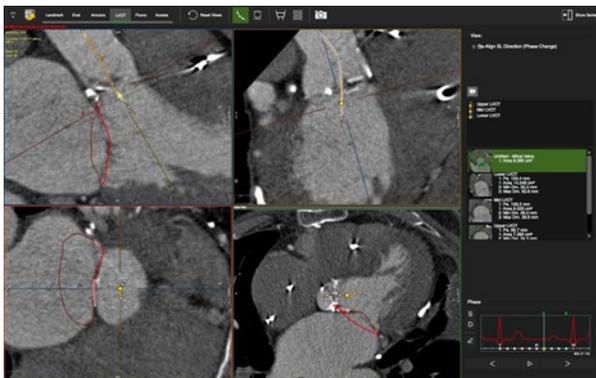
Calcium Volume	
Mitral Valve Region Calcium:	423.7 mm ³
- Anterior Region:	218.0 mm ³
- Posterior Region:	205.7 mm ³



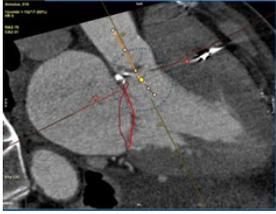
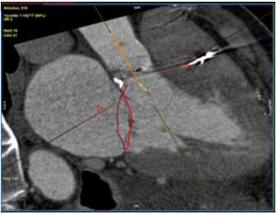
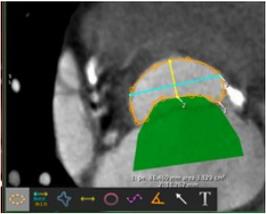
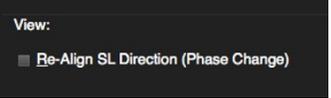
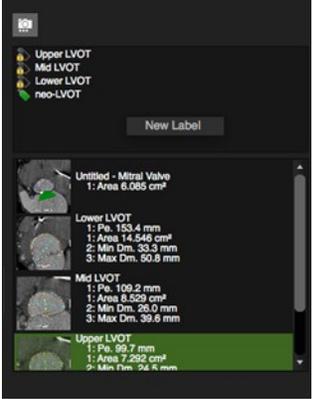
-  Click on the camera to add the calcium Volume and images to your report.

28.4 LVOT Page

- Assessment of the LVOT
- Device simulation
- Planimetric measurement of LVOT and neo-LVOT

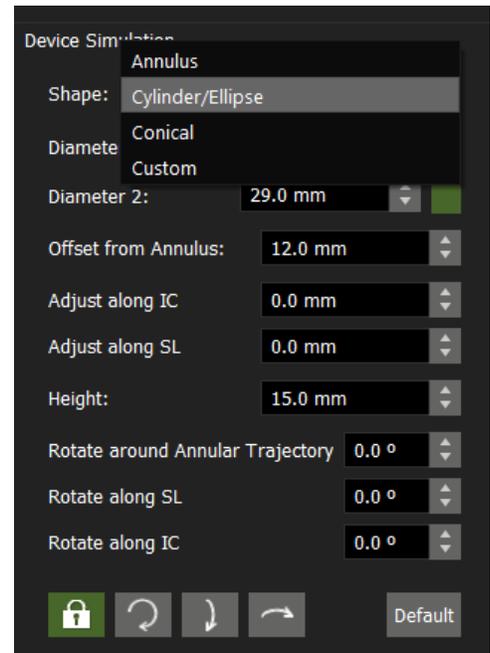


28.4.1 LVOT Assessment

<ol style="list-style-type: none"> 1.  Activate the Centerline Control Points 2. Place points in the Aorta and the LVOT 																										
<ol style="list-style-type: none"> 3. De-activate the centerline points and use the green bar to scroll along the centerline 																										
<ol style="list-style-type: none"> 4. Using the measurement tools provided in the tool box (white box in the lower left corner), measure the upper-, mid-, and lower LVOT 																										
<ol style="list-style-type: none"> 5. Use the phase diagram to inspect the LVOT in different phases. Optionally, realign the view to SL direction with every phase change 																										
<ol style="list-style-type: none"> 6. Label the measurements by dragging the predefined tags on the capture 7. Right-click next to the labels, to add a new label 8. For review, simply click on a capture 	 <table border="1" data-bbox="1032 1213 1218 1444"> <thead> <tr> <th>Segment</th> <th>1: Pe</th> <th>1: Area</th> <th>2: Min Dm</th> <th>3: Max Dm</th> </tr> </thead> <tbody> <tr> <td>Untitled - Mitral Valve</td> <td></td> <td>6.085 cm²</td> <td></td> <td></td> </tr> <tr> <td>Lower LVOT</td> <td>153.4 mm</td> <td>14.549 cm²</td> <td>33.3 mm</td> <td>60.9 mm</td> </tr> <tr> <td>Mid LVOT</td> <td>109.2 mm</td> <td>8.029 cm²</td> <td>26.0 mm</td> <td>39.6 mm</td> </tr> <tr> <td>Upper LVOT</td> <td>99.7 mm</td> <td>7.292 cm²</td> <td></td> <td></td> </tr> </tbody> </table>	Segment	1: Pe	1: Area	2: Min Dm	3: Max Dm	Untitled - Mitral Valve		6.085 cm ²			Lower LVOT	153.4 mm	14.549 cm ²	33.3 mm	60.9 mm	Mid LVOT	109.2 mm	8.029 cm ²	26.0 mm	39.6 mm	Upper LVOT	99.7 mm	7.292 cm ²		
Segment	1: Pe	1: Area	2: Min Dm	3: Max Dm																						
Untitled - Mitral Valve		6.085 cm ²																								
Lower LVOT	153.4 mm	14.549 cm ²	33.3 mm	60.9 mm																						
Mid LVOT	109.2 mm	8.029 cm ²	26.0 mm	39.6 mm																						
Upper LVOT	99.7 mm	7.292 cm ²																								

28.4.2 Device Simulation

1.   Turn on the Device Mesh or Device Intersection buttons
2. Choose a shape from the drop-down list
3. Optionally import vendor specific device image files
4. Adjust the device position
5. Make additional measurements, e.g. neo-LVOT
6.  Lock device position across different phases:



28.5 Fluoroscopy Page

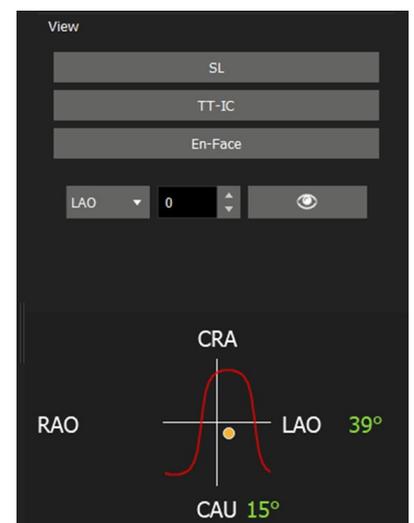
- Display images along the optimal viewing curve
- Capture C-arm angles

28.5.1 Optimal Viewing Curve

1. Display views perpendicular to the annulus

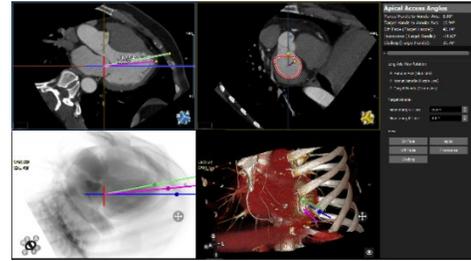


2. Use the pre-set buttons or drag the cursor line over the curve t



28.6 Apical Page

Assessment of trans apical approach to the mitral structure. Visualization of angles, catheter path, relationship to ribcage, pericardial structures, papillary muscles, and mitral apparatus.



1. Displays intraprocedural catheter angles.

2. Select the rotation angles:
 - Perpendicular axis to mitral plane (blue line)
 - Left ventricle axis (purple line)
 - Delivery sheath axis (green line)

3. Positions delivery sheath.

4. Buttons for default fluoroscopic and volume views.

Apical Access Angles

Manual Handle to Annular Axis: 23.56°

Target Handle to Annular Axis: 8.61°

Off Table (Target Handle): 34.54°

Transverse (Target Handle): -19.31°

Clocking (Target Handle): 19.29°

Long Axis View Rotation:

- Annulus Axis (Blue Line)
- Manual Handle (Purple Line)
- Target Handle (Green Line)

Target Handle:

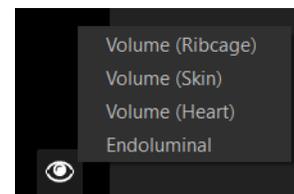
Move along SL Line:

Move along IC Line:

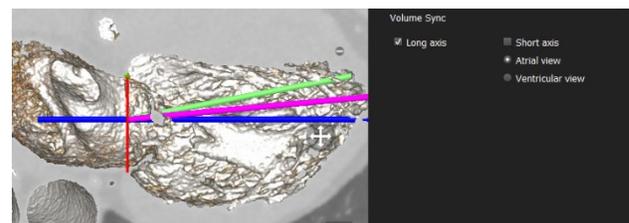
View

En-Face	Apical
Off Table	Transverse
Clocking	

5. Switch between different visualization techniques.



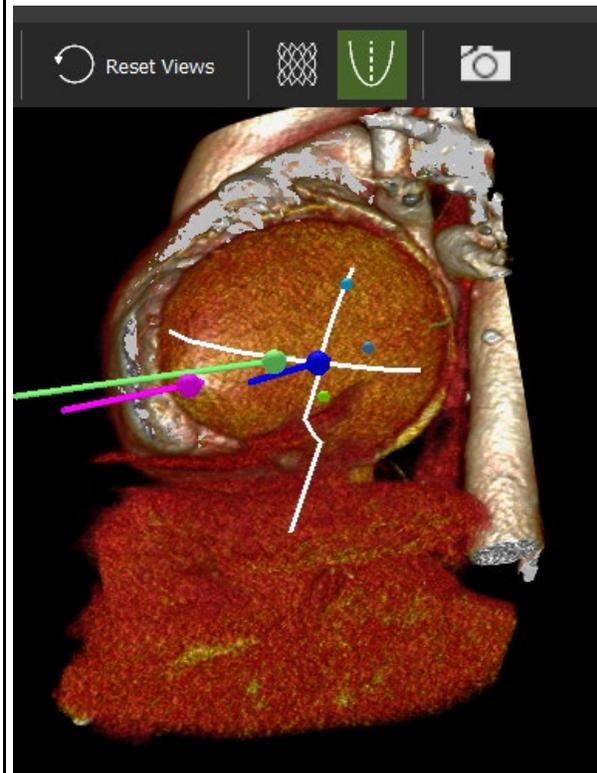
6. When the endoluminal view is active, select the "Volume Sync" options to synchronize the endoluminal view with the long or the short axis MPR.



7. Toggle virtual valve on or off



8. When "Volume(heart)" is selected, use the IC/SL projection button to project the IC and SL lines on the surface of the heart.



29 Trans-Septal Module

- 29.1 Contours page 99
 - 29.1.1 Finding the Fossa Ovalis 99
 - 29.1.2 Marking the Vena Cava Inferior and Superior and Other Structures 100

Intended Use

This module allows the user to gain a better understanding of the relationships of different parts of the anatomy and how they can appear on a fluoroscopic view during the actual trans septal interventions.

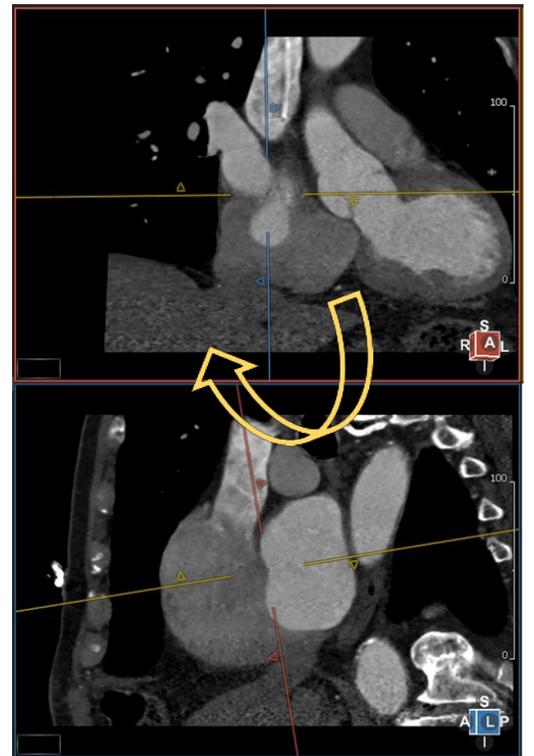
29.1 Contours Page

29.1.1 Finding the Fossa Ovalis

1. On the axial image locate the interatrial septum
2. Translate up and down until the thinnest part of the atrial septum is shown
3. Align the red intersection line with the septum
4. In the lateral view (lower left viewport) align the red intersection line with the septum
5. The resulting view (left upper view will be oriented en-face the Fossa Ovalis)



6. In the en-face view rotate the intersection lines to verify in the orthogonal views that the center of the crosshair is in the middle of the Fossa
7. Click the Fossa Ovalis button and then click on the center of the crosshair in the en-face view to mark the Fossa
8. When needed, resize the annotation that indicates the Fossa

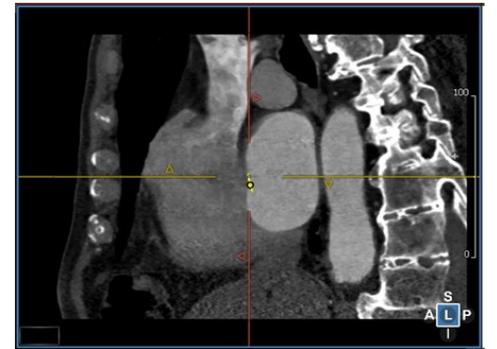


29.1.2 Marking the Vena Cava Inferior and Superior and Other Structures

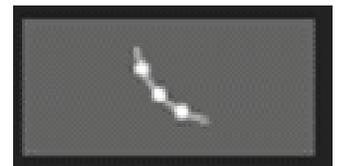
1. Reset views
2. In the lateral viewport translate until the IVC is visible



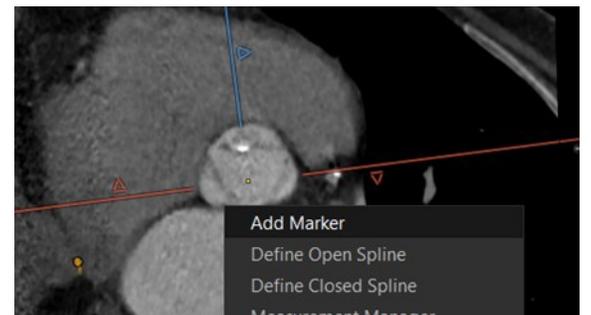
3. Place the crosshair on the IVC
4. On the axial image, the IVC contour can be traced
5. Do the same for the SVC



6. Use the catheter feature to draw a spline to indicate the catheter path or use the automatic option for an automated suggestion.



7. Use markers to identify other anatomical markers, like the aorta
- In case the Mitral annulus has been detected in the Mitral workflow, the annotation will be transferred to the septal crossing workflow



30 Technical Support

30.1 Technical Support

For technical questions please contact our team by phone or e-mail:

North America

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F: +1 403 338 1895

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Circle Cardiovascular Imaging B.V.

Europe Support Phone: + 31 (800)265 8982

Report a problem: support@circlecvi.com

Website: www.circlecvi.com