

MASTERCLASS ZA HADRONSKU TERAPIJU

**Planiranje radioterapijskih tretmana u
MatRadu**

Instrukcije

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1. Zadatak

- Korištenje TG119 fantoma
- Planiranje radioterapijskog tretmana pomoću fotona, protona i ugljikovih iona
- Analiza i poređenje dobivenih rezultata

1. Zadatak

- Korištenje TG119 fantoma
- Planiranje radioterapijskog tretmana pomoću fotona, protona i ugljikovih iona
- Analiza i poređenje dobivenih rezultata

1. Učitajte TG119 fantom pomoću opcije Load * .mat (TG119.mat)

The screenshot displays a software interface with a 'Workflow' section on the left. The 'Load *.mat data' button is highlighted with a red arrow. A 'Select File to Open' dialog box is open, showing a file list with 'TG119' selected, also indicated by a red arrow. The dialog box shows the file path 'e0404-matRad-2.1... > e0404-matRad-2957fcc' and the file name 'TG119' in the 'File name' field. The file type is set to 'MAT-files (*.mat)'. The background interface includes various settings for 'Plan', 'Objectives & constraints', and 'Visualization'.

Workflow

Refresh Load *.mat data Load *.COM Import from Bin...

Status: no data loaded

Plan

bixel width in [mm] 5 use M
Gantry Angle in ° 0 3D c
Couch Angle in ° 0 Run
Radiation Mode photons
Machine Generic Stratific
IsoCenter in [mm] 0 0 0 Auto. Run
Fractions 30
Type of optimization none Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.
1				
2				
3				
4				

Visualization

Slice Type of plot inten... GoTo lateral
Beam Plane axial Open 3D-View
Offset Dislav option no option avail...
Show DVH/Q

- plot CT
- plot contour
- plot isolines
- plot dose
- plot isolines labels
- plot iso center
- visualize plan / be...

Select File to Open

Organize New folder

Name	Date modified	Type
standalone	6/19/2019 8:34 AM	File folder
tools	6/19/2019 8:34 AM	File folder
unitTest	6/19/2019 8:34 AM	File folder
vmc++	6/19/2019 8:34 AM	File folder
BOXPHANTOM	6/19/2019 8:33 AM	MAT File
carbon_Generic	6/19/2019 8:34 AM	MAT File
HEAD_AND_NECK	6/19/2019 8:33 AM	MAT File
LIVER	6/19/2019 8:33 AM	MAT File
photons_Generic	6/19/2019 8:34 AM	MAT File
PROSTATE	6/19/2019 8:33 AM	MAT File
protons_Generic	6/19/2019 8:34 AM	MAT File
TG119	6/19/2019 8:34 AM	MAT File

File name: TG119 MAT-files (*.mat)

Open Cancel

min max

Set IsoDose Levels

Viewer Options

None No available

Window Center: 0.5
Window Width: 1.0
Range: 0 1

bone Dose opacity: 0 1

Structure Visibility

no data loaded

Info

v3.0.0
github.com/e0404/mat
About

2. Odaberite fotone kao vrstu zračenja, te odredite jedan ugao snopa (gantry angle)

The screenshot displays the matRad software interface, which is part of the German Cancer Research Center (dkfz) in the Helmholtz Association. The interface is divided into several panels:

- Workflow:** Includes buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, Import from Bin..., and Import Dose. The status is "ready for dose calculation".
- Plan:** Contains input fields for bixel width (10 mm), Gantry Angle (0 degrees), Couch Angle (0 degrees), Radiation Mode (photons), Machine (Generic), IsoCenter (251.3, 236.4, 162.6 mm), # Fractions (30), and Type of optimization (none). It also has radio buttons for "use MC (VMC++) dose calculations", "3D conformal", "Run Sequencing", and "Run Direct Aperture Optimizat...". Stratification Levels are set to 7.
- Objectives & constraints:** A table with columns for VOI name, VOI type, priority, obj. / const., penalty, dose, EUD, volume, and ro. The table contains three rows: Core (OAR, priority 2, square overdosing, penalty 300, dose 25), OuterTarget (TARGET, priority 1, square deviation, penalty 1000, dose 50), and BODY (OAR, priority 3, square overdosing, penalty 100, dose 30). A "save" button is at the bottom right.
- Visualization:** Includes fields for Slice, Beam, and Offset. It has dropdowns for Type of plot (intensity), Plane (axial), and Dislay option (no option avail...). There are buttons for "GoTo lateral" and "Open 3D-View". A list of plot options includes plot CT, plot contour, plot isolines, plot dose, plot isolines labels, plot iso center, and visualize plan / be... The "Show DVH/Q" button is at the bottom.

The central viewing area shows an axial plane at z = 165 mm. The image displays a cross-section of a patient's head and neck with a yellow contour around the target area and a purple contour around the body. A color scale on the right indicates Hounsfield Units from 0 to 60. The x and y axes are labeled in mm, ranging from 50 to 500.

On the right side, there are additional settings for "min max" (1000, 1040), "Set IsoDose Levels", "Viewer Options" (CT (HU), Window: Drapeat, Custom, Window Center: 0.85, Window Width: 1.67, Range: 0.02671, 1.692), "Structure Visibility" (Core, OuterTarget, BODY), and "Info" (v3.0.0, github.com/e0404/mat, About).

3. Pokrenite proračun doze pomoću opcije („Calc. Influence Mx“). Zatim započnite obrnutu optimizaciju klikom na („Optimize“)

The screenshot displays the matRad software interface, which is used for radiation therapy planning. The interface is divided into several panels:

- Workflow Panel:** Contains buttons for 'Refresh', 'Load *.mat data', 'Load DICOM', 'Import from Bin...', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Export', and 'Import Dose'. Red arrows point to the 'Calc. influence Mx' and 'Optimize' buttons. The status below these buttons reads 'Status: ready for optimization'.
- Plan Panel:** Includes input fields for 'bixel width in [mm]' (10), 'Gantry Angle in °' (0), 'Couch Angle in °' (0), 'Radiation Mode' (photons), 'Machine' (Generic), 'IsoCenter in [mm]' (251.3 236.4 162.6), and '# Fractions' (30). It also has radio buttons for 'use MC (VMC++) dose calculations', '3D conformal', 'Run Sequencing', and 'Run Direct Aperture Optimizat...'. A 'Stratification Levels' field is set to 7.
- Objectives & constraints Panel:** Contains a table with the following data:

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no
- Visualization Panel:** Includes 'Slice' and 'Beam' selection, 'Type of plot' (intensity), 'Plane' (axial), and 'Dislay option' (no option avail...). It also has a 'GoTo' field set to 'lateral' and an 'Open 3D-View' button.
- Viewing Panel:** Shows an axial CT scan of a patient's head at z = 165 [mm]. The image displays the brain, eyes, and surrounding structures. A color scale on the right indicates Hounsfield Units from 10 to 60. The 'Viewer Options' panel on the right shows 'CT (HU)' selected, 'Window Center' at 0.85, 'Window Width' at 1.67, and 'Range' from 0.02671 to 1.692. The 'Structure Visibility' panel shows 'Core', 'OuterTarget', and 'BODY' checked.

4. Analizirajte rezultujuću raspodjelu doze




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Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI

Load DICOM Recalc Export

Import from Bin... Import Dose

Status: plan is optimized

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations

Gantry Angle in ° 0 3D conformal

Couch Angle in ° 0 Run Sequencing

Radiation Mode photons Stratification Levels 7

Machine Generic Run Direct Aperture Optimizat...

IsoCenter in [mm] 251.3 236.4 162.6 Auto.

Fractions 30

Type of optimization none

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no

Visualization

Slice Type of plot inten... GoTo lateral plot CT

Beam Plane axial plot contour

Offset Dislay option physicalDose plot isolines

plot dose

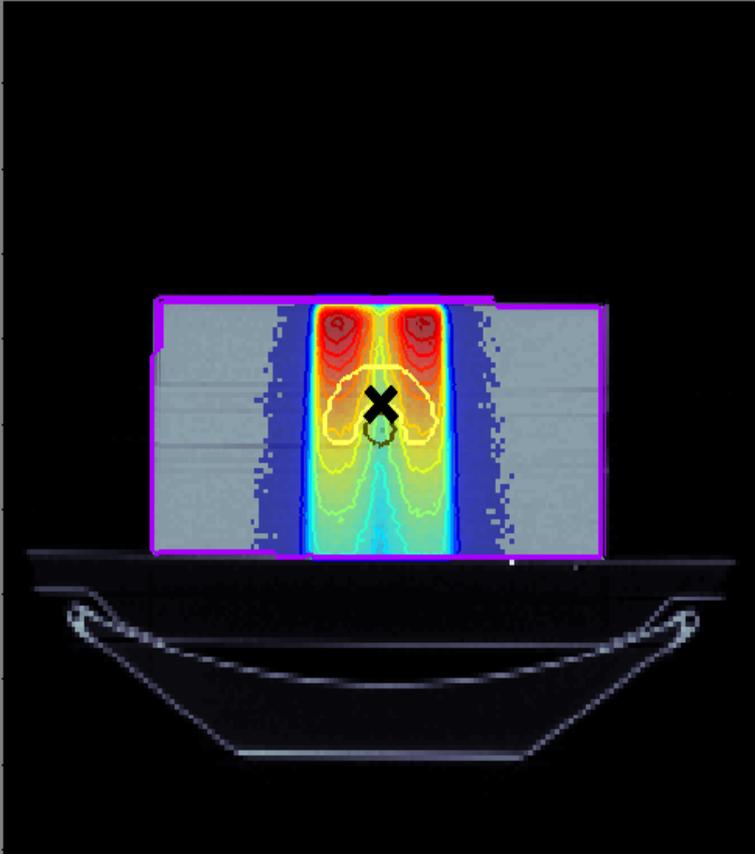
plot isolines labels

plot iso center

visualize plan / be...

Viewing

axial plane z = 165 [mm]



min max n 2.342

Viewer Options

Result (i.e. dose)

Window:

Window Center: 1.17

Window Width: 2.34

Range: 0 2.342

jet

Dose opacity: 1

Structure Visibility

- Core
- OuterTarget
- BODY

Info

v3.0.0

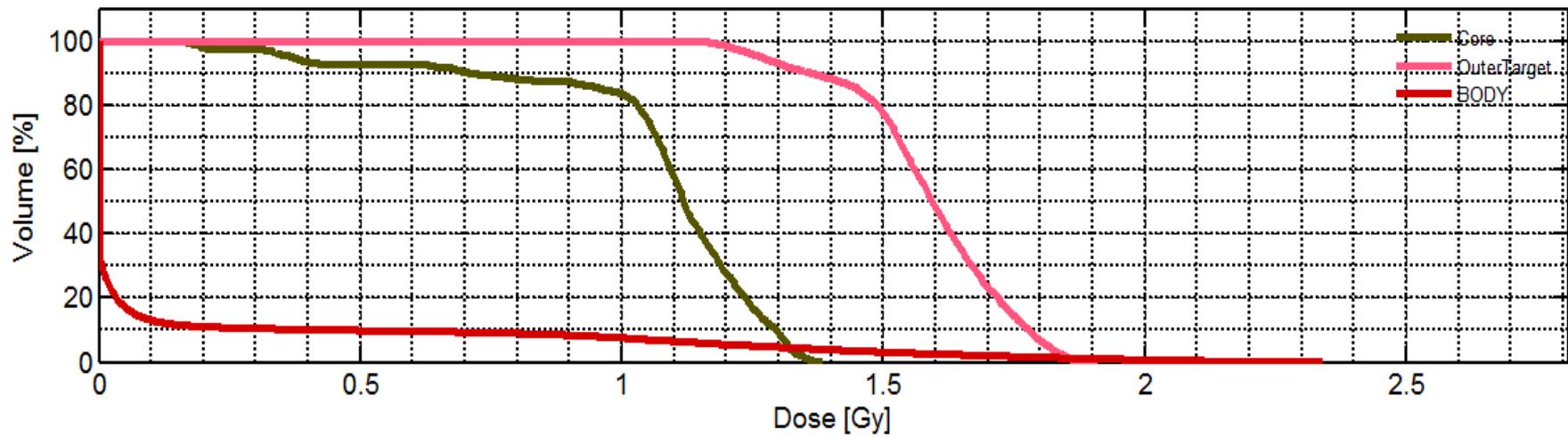
github.com/e0404/mat

5. Spremite rezultat optimizacije putem („Save to GUI“), te prikažite DVH pomoću („Show DVH/QI“).

The screenshot displays the matRad software interface, which is used for radiation therapy optimization. The interface is divided into several panels:

- Workflow:** Contains buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, and Import Dose. A red arrow points to the "Save to GUI" button. The status below indicates "plan is optimized".
- Plan:** Contains various parameters for optimization, including bixel width (10 mm), Gantry Angle (0°), Couch Angle (0°), Radiation Mode (photons), Machine (Generic), IsoCenter (251.3, 236.4, 162.6 mm), # Fractions (30), and Type of optimization (none). There are also radio buttons for "use MC (VMC++) dose calculations", "3D conformal", "Run Sequencing", "Stratification Levels" (7), and "Run Direct Aperture Optimizat...".
- Objectives & constraints:** A table listing VOI names, types, priorities, and constraints.
- Visualization:** Contains options for Slice, Beam, Offset, Type of plot (intensity), Plane (axial), and Display option (physicalDose). A red arrow points to the "Show DVH/QI" button.
- Viewing:** Displays a dose distribution plot for an axial plane at z = 165 mm. The plot shows a color-coded dose distribution with a color scale on the right ranging from 0 to 60 Gy. The plot is overlaid on a CT scan of a patient's head and neck.
- Viewer Options:** Contains options for Result (i.e. dose), Window Preset (Custom), Window Center (1.17), Window Width (2.34), Range (0 to 2.342), and Dose opacity (1).
- Structure Visibility:** A list of structures with checkboxes: Core (checked), OuterTarget (checked), and BODY (checked).
- Info:** Displays the version (v3.0.0) and the GitHub repository (github.com/e0404/mat).

VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1 Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2 OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3 BODY	OAR	3	square overdosing	100	30	NaN	NaN	no



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.4Gy	V_0.9Gy	V_1.4Gy	V_1.9Gy
Core	1.0665	0.2554	1.3860	0.1329	1.3434	1.3187	1.1183	0.3706	0.1988	1	0.9341	0.8727	0	0
OuterTarget	1.5852	0.1536	1.9115	1.0935	1.8453	1.8153	1.5941	1.2663	1.2077	1	1	1	0.8824	0
BODY	0.1443	0.4168	2.3420	0	1.7203	1.2694	0	0	0	1	0.1019	0.0846	0.0393	0

6. Promijenite vrstu zračenja u Proton i ostavite uglove snopa nepromijenjenim

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
 Load DICOM Recalc Export
 Import from Bin... Import Dose

Status: plan is optimized

Plan

bixel width in [mm] 10
 Gantry Angle in ° 0
 Couch Angle in ° 0
 Radiation Mode photons
 Machine photons
 IsoCenter in [mm] protons **←**
 # Fractions carbon
 Type of optimization none Set Tissue

use MC (VMC++) dose calculations
 3D conformal
 Run Sequencing
 Stratification Levels 7
 Run Direct Aperture Optimizat...

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no

save

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Dislay option physicalDose

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

Show DVH/QI

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axial plane z = 165 [mm]

min 0
max 2.342

Set IsoDose Levels

Viewer Options

Result (i.e. dose) Window Doseat Custom
 Window Center: 1.17
 Window Width: 2.34
 Range: 0 2.342
 jet Dose opacity: 1

Structure Visibility

Core
 OuterTarget
 BODY

Info

v3.0.0
 github.com/e0404/mat
 About

7. Pokrenite proračun doze pomoću opcije („Calc. Influence Mx“) i započnite obrnutu optimizaciju klikom na („Optimize“)

Workflow

Refresh Load *.mat data **Calc. influence Mx** Optimize Save to GUI
 Load DICOM Finalc Export
 Import from Bin... Import Dose

Status: 1 ready for optimization 2

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations
 Gantry Angle in ° 0 3D conformal
 Couch Angle in ° 0 Run Sequencing
 Radiation Mode protons Stratification Levels 7
 Machine Generic Run Direct Aperture Optimizat...
 IsoCenter in [mm] 251.3 236.4 162.6 Auto.
 # Fractions 30
 Type of optimization const_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Dislay option physicalDose
 Show DVH/QI

- plot CT
- plot contour
- plot isolines
- plot dose
- plot isolines labels
- plot iso center
- visualize plan / be...

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Viewing axial plane z = 165 [mm]

min max n 2.342

Set IsoDose Levels

Viewer Options

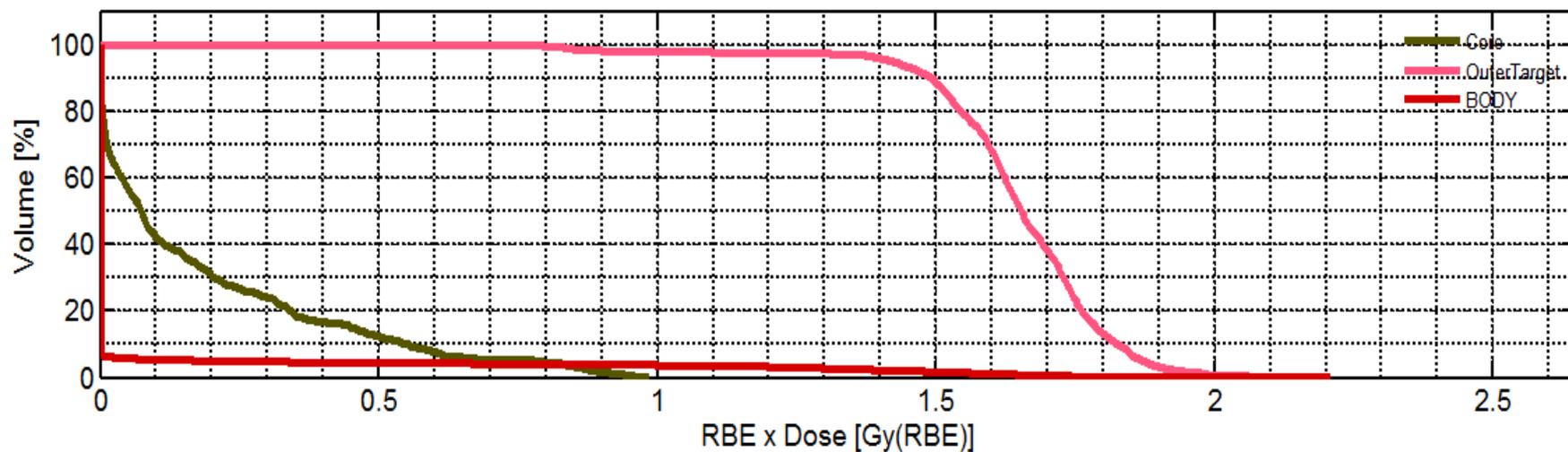
Result (i.e. dose) Window Presest Custom
 Window Center: Window Width: 1.17
 Range: 0 2.34
 jet Dose opacity: 1
 Structure Visibility
 Core
 OuterTarget
 BODY

Info v3.0.0 github.com/e0404/mat About

8. Spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)

The screenshot displays the matRad software interface. The top left shows the workflow area with buttons for 'Refresh', 'Load *.mat data', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Load DICOM', 'Recalc', 'Export', and 'Import dose'. A red arrow labeled '1' points to the 'Save to GUI' button. Below this is the 'Plan' section with various parameters like 'bixel width in [mm]', 'Gantry Angle in °', 'Couch Angle in °', 'Radiation Mode', 'Machine', 'IsoCenter in [mm]', '# Fractions', and 'Type of optimization'. The 'Objectives & constraints' section contains a table with columns for 'VOI name', 'VOI type', 'priority', 'obj. / const.', 'penalty', 'dose', 'EUD', 'volume', and 'ro'. The 'Visualization' section at the bottom left has controls for 'Slice', 'Beam', 'Offset', 'Type of plot', 'Plane', 'Dislay option', and 'Show DVH/QI', with a red arrow labeled '2' pointing to the 'Show DVH/QI' button. The main viewing area shows an axial plane at z = 165 [mm] with a color-coded dose distribution. A color scale on the right indicates 'RBE x Dose [Gy(RBE)]' from 0 to 60. The 'Viewer Options' panel on the right includes 'Set IsoDose Levels', 'Result (i.e. dose)', 'Window: Default', 'Custom', 'Window Center', 'Window Width', 'Range', 'jet', 'Dose opacity', and 'Structure Visibility'.

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro	
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no	+
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no	-
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no	save



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.4Gy	V_0.8Gy	V_1.3Gy	V_1.8Gy
Core	0.1815	0.2396	0.9866	2.0386e-09	0.8909	0.7849	0.0744	2.4933e-05	6.0723e-07	1	0.1682	0.0470	0	0
OuterTarget	1.6449	0.1770	2.1789	0.7475	1.9408	1.8726	1.6533	1.4205	0.9187	1	1	0.9949	0.9722	0.9222
BODY	0.0640	0.2912	2.2101	0	1.4572	0.2364	0	0	0	1	0.0462	0.0405	0.0282	0.0182

Rezultati

- Srednje doze po regionima (Gy):

Region/Zračenje	Fotoni	Protoni
Osjetljivi region	1.0665	0.1815
Vanjska meta	1.5852	1.6449
Tijelo	0.1443	0.0640

- Fotoni daju najveću dozu na površini
- Protoni daju najveću dozu na meti (tumoru) uz maksimalnu zaštitu osjetljivih regiona

9. Pokušajte definisati bolji plan tretmana fotonima pomoću većeg broja uglova snopa (npr. ekvivalentni razmak [0, 72, 144, 216, 288]. Pokrenite proračun doze i započnite obrnutu optimizaciju.

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
 Load DICOM ReCalc Export
 Import from Bin... Import Dose

Status: ready for dose calculation

Plan

bixel width in [mm] 10
 Gantry Angle in ° 0 72 144 216 288
 Couch Angle in ° 0 0 0 0
 Radiation Mode photons
 Machine Generic
 IsoCenter in [mm] 251.3 236.4 162.6 Auto.
 # Fractions 30
 Type of optimization none Set Tissue

use MC (VMC++) dose calculations
 3D conformal
 Run Sequencing
 Stratification Levels 7
 Run Direct Aperture Optimizat...

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2	OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Disolv option physicalDose

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

Show DVH/QI

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Viewing axial plane z = 165 [mm]

min max 1.902

Set IsoDose Levels

Viewer Options

Result (i.e. dose)
 Window: Breast
 Custom
 Window Center: 0.95
 Window Width: 1.9
 Range: 0 1.903
 jet
 Dose opacity: 1

Structure Visibility

- Core
- OuterTarget
- BODY

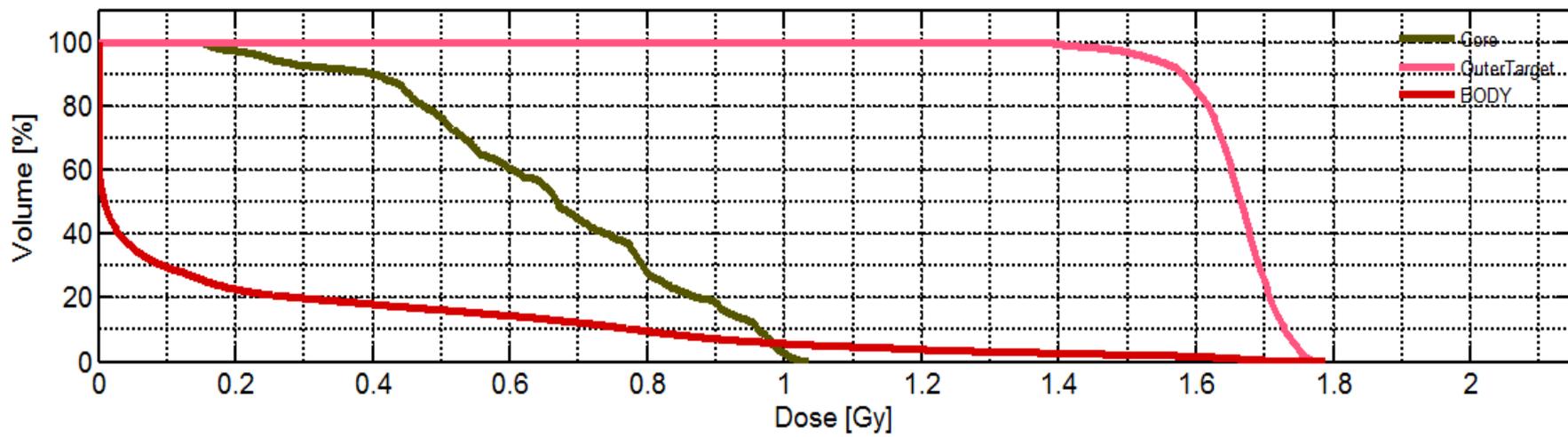
Info v3.0.0 github.com/e0404/mat About

10. Spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)

The screenshot displays the matRad software interface, which is part of the German Cancer Research Center (dkfz) in the Helmholtz Association. The interface is divided into several functional areas:

- Workflow:** Contains buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, and Import Dose. A red arrow points to the "Save to GUI" button.
- Status:** Indicates "plan is optimized".
- Plan:** Lists parameters such as bixel width (10 mm), Gantry Angle (0 72 144 216 288), Couch Angle (0 0 0 0), Radiation Mode (photons), Machine (Generic), IsoCenter (251.3 236.4 162.6), # Fractions (30), and Type of optimization (none). It also includes options for MC dose calculations, 3D conformal, Run Sequencing, Stratification Levels (7), and Run Direct Aperture Optimization.
- Objectives & constraints:** A table defining the optimization goals:

VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1 Core	OAR	2	square overdosing	300	25	NaN	NaN	no
2 OuterTarget	TARGET	1	square deviation	1000	50	NaN	NaN	no
3 BODY	OAR	3	square overdosing	100	30	NaN	NaN	no
- Visualization:** Includes controls for Slice, Beam, and Offset, as well as plot options (intensity, axial, physicalDose) and visualization settings (plot CT, contour, isolines, dose, labels, center, plan). A red arrow points to the "Show DVH/QI" button.
- Viewer:** Shows a dose distribution plot for an axial plane at z = 165 mm. The plot uses a color scale from 0 to 60 Gy. The y-axis ranges from 50 to 500 mm, and the x-axis ranges from 50 to 500 mm. A red arrow points to the "Show DVH/QI" button in the visualization area.
- Viewer Options:** Includes controls for Result (i.e. dose), Window (Custom), Window Center (0.89), Window Width (1.79), Range (0 to 1.79), and Dose opacity (1).
- Structure Visibility:** Lists visible structures: Core, OuterTarget, and BODY.
- Info:** Shows version v3.0.0 and the GitHub repository link qithub.com/e0404/mat.



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.3Gy	V_0.7Gy	V_1Gy	V_1.5Gy
Core	0.6625	0.2176	1.0370	0.1450	1.0030	0.9853	0.6686	0.2460	0.1755	1	0.9265	0.4477	0.0250	0.0000
OuterTarget	1.6563	0.0659	1.7897	1.2866	1.7566	1.7450	1.6652	1.5323	1.4636	1	1	1	1	0
BODY	0.1968	0.3777	1.7897	0	1.5510	1.0629	0.0091	0	0	1	0.1986	0.1230	0.0568	0

Rezultati

- Srednje doze po regionima (Gy):

Region/Zračenje(uglovi)	Fotoni(0)	Protoni(0)	Fotoni (0,72,144,216,288)
Osjetljivi region	1.0665	0.1815	0.6625
Vanjska meta	1.5852	1.6449	1.6563
Tijelo	0.1443	0.0640	0.1968

- Plan tretmana sa više snopova fotona daje bolje rezultate nego sa jednim snopom fotona.
- Najbolji rezultati su dobiveni korištenjem protona.

11. Promijenite cilj optimizacije za poboljšanje plana liječenja protonima. Koristite tabelu („Objectives & constraints“) i dodajte ograničenje (npr. maksimalna doza u meti ili minimalna doza u vanjskoj ciljnoj strukturi).

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
 Load DICOM Recalc Export
 Import from Bin... Import Dose

Status: ready for optimization

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations
 Gantry Angle in ° 0 72 144 216 288 3D conformal
 Couch Angle in ° 0 0 0 0 Run Sequencing
 Radiation Mode photons Stratification Levels 7
 Machine Generic Run Direct Aperture Optimizat...
 IsoCenter in [mm] 251.3 236.4 162.6 Auto.
 # Fractions 30
 Type of optimization none Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Core	OAR	2	max dose constraint	5	NaN	NaN	no	+
2	OuterTarget	TARGET	1	min dose constraint	5	NaN	NaN	no	-
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN	no

save

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Dislay option physicalDose
 Show DVH/QI

- plot CT
- plot contour
- plot isolines
- plot dose
- plot isolines labels
- plot iso center
- visualize plan / be...

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Viewing

axial plane z = 165 [mm]

physicalDose [Gy]

min max n
max 1.789

Set IsoDose Levels

Viewer Options

Result (i.e. dose) Window Preset Custom
 Window Center: 0.89
 Window Width: 1.79
 Range: 0 1.79
 jet Dose opacity: 1

Structure Visibility

- Core
- OuterTarget
- BODY

Info

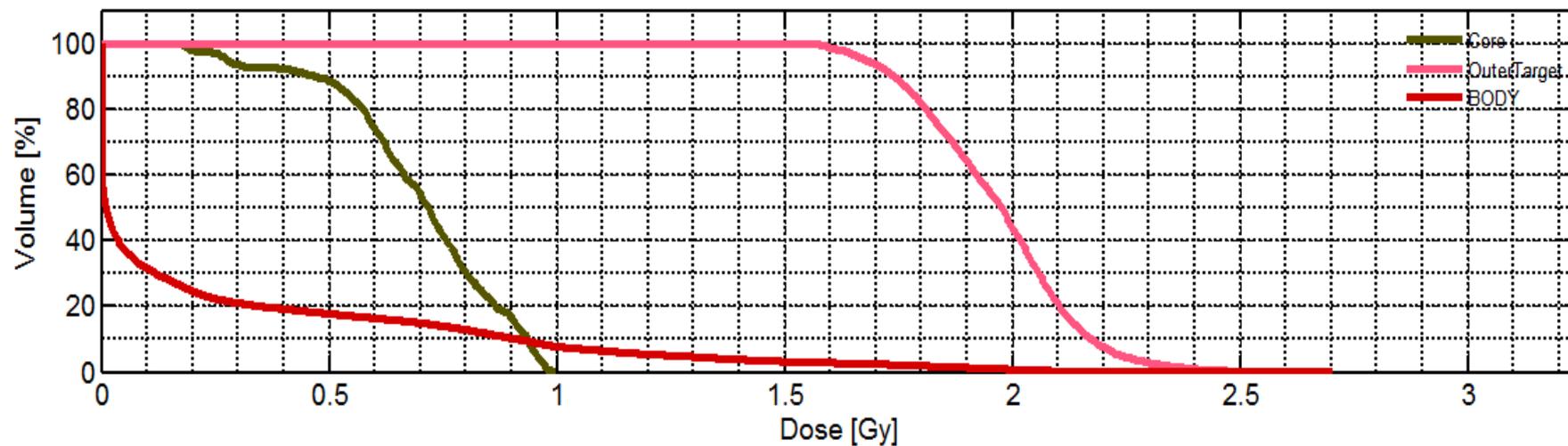
v3.0.0
 github.com/e0404/mat
 About

12. Pokrenite proračun doze pomoću opcije („Calc. Influence Mx“) i započnite obrnutu optimizaciju klikom na („Optimize“). Zatim spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)

The screenshot displays the matRad software interface, which is part of the German Cancer Research Center (dkfz) in the Helmholtz Association. The interface is divided into several sections:

- Workflow:** Contains buttons for 'Refresh', 'Load *.mat data', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Load DICOM', 'Replc', 'Export', and 'Import dose'. Red arrows point to 'Calc. influence Mx', 'Optimize', and 'Save to GUI'. The status below reads 'Status: plan is optimized'.
- Plan:** Includes parameters for 'bixel width in [mm]' (10), 'Gantry Angle in °' (0 72 144 216 288), 'Couch Angle in °' (0 0 0 0), 'Radiation Mode' (photons), 'Machine' (Generic), 'IsoCenter in [mm]' (251.3 236.4 162.6), '# Fractions' (30), and 'Type of optimization' (none). It also has radio buttons for 'use MC (VMC++) dose calculations', '3D conformal', 'Run Sequencing', and 'Run Direct Aperture Optimizat...', and a 'Stratification Levels' dropdown set to '7'.
- Objectives & constraints:** A table with columns for 'VOI name', 'VOI type', 'priority', 'obj. / const.', 'penalty', 'dose', 'EUD', 'volume', and 'ro'.

VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1 Core	OAR	2	max dose constraint	NaN	25	NaN	NaN	no
2 OuterTarget	TARGET	1	min dose constraint	NaN	50	NaN	NaN	no
3 BODY	OAR	3	square overdosing	100	30	NaN	NaN	no
- Visualization:** Includes 'Slice', 'Beam', and 'Offset' controls, 'Type of plot' (intensity), 'Plane' (axial), and 'Dislay option' (physicalDose). A 'Show DVH/QI' button is highlighted with a red arrow.
- Viewing:** Shows an axial plane at z = 165 [mm]. The plot displays a color-coded dose distribution with a central target area (Core) and surrounding organs at risk (OAR). A color scale on the right indicates 'physicalDose [Gy]' from 0 to 60. The axes are labeled 'x [mm]' and 'y [mm]'.
- Viewer Options:** Includes 'Result (i.e. dose)', 'Window: Dose', 'Custom', 'Window Center', 'Window Width', 'Range', 'jet' color map, and 'Dose opacity'.
- Structure Visibility:** Shows 'Core', 'OuterTarget', and 'BODY' as visible structures.
- Info:** Displays 'v3.0.0' and 'qithub.com/e0404/mat'.



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.5Gy	V_1Gy	V_1.6Gy	V_...
Core	0.6974	0.1876	0.9986	0.1704	0.9743	0.9563	0.7189	0.2781	0.1981	1	0.8848	0	0	
OuterTarget	1.9652	0.1732	2.7054	1.5511	2.3409	2.2397	1.9766	1.6761	1.6190	1	1	1	0.9857	
BODY	0.2343	0.4481	2.7054	0	1.7993	1.2658	0.0110	0	0	1	0.1780	0.0784	0.0288	

Rezultati

- Srednje doze po regionima (Gy) dobivene pomoću 5 snopova fotona sa i bez ograničenja:

Region/Zračenje	Bez ograničenja	Sa ograničenjima
Osjetljivi region	0.6625	0.6974
Vanjska meta	1.6563	1.9652
Tijelo	0.1968	0.2343

2. Zadatak

- Plan liječenja bolesti jetre ugljikovim ionima
- Definisanje vlastitog plana liječenja fotonima i protonima
- Analiza i poređenje različitih planova liječenja

1. Učitajte jetru pacijenta pomoću opcije Load *.mat (LIVER.mat)

The screenshot displays the matRad software interface. The top toolbar contains several buttons, with 'Load *.mat data' highlighted by a red arrow. Below the toolbar, the 'Plan' section shows various parameters such as 'bixel width in [mm]' (20), 'Gantry Angle in °' (0 72 144 216 288), and 'Radiation Mode' (protons). The 'Objectives & constraints' section contains a table with the following data:

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	Core	OAR	2	max dose constraint	NaN	25	NaN	NaN
2	OuterTarget	TARGET	1	min dose constraint	NaN	50	NaN	NaN
3	BODY	OAR	3	square overdosing	100	30	NaN	NaN

The 'Select File to Open' dialog box is open, showing a file list with 'LIVER' selected. The file list includes folders like 'standalone', 'tools', 'unitTest', 'vmc++' and files like 'BOXPHANTOM', 'carbon_Generic', 'HEAD_AND_NECK', 'LIVER', 'photons_Generic', 'PROSTATE', 'protons_Generic', and 'TG119'. The 'LIVER' file is highlighted with a red arrow. The dialog box also shows the file name 'LIVER' and the file type 'MAT-files (*.mat)'. The background interface shows the 'Workflow' section with buttons like 'Refresh', 'Load *.mat data', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Load *.COM', 'Recalc', 'Export', 'Import Plan Bin...', and 'Import Dose'. The status bar indicates 'plan is optimized'. The 'Visualization' section at the bottom shows options for 'Slice', 'Beam', and 'Offset', along with 'Type of plot' (intensity), 'Plane' (axial), and 'Display option' (RBEExDose). The 'Structure Visibility' section shows 'Core', 'OuterTarget', and 'BODY' checked. The bottom right corner shows the version 'v3.0.0' and the GitHub repository 'github.com/e0404/mat'.

2. Definišite vlastiti plan liječenja fotonima sa 4-5 uglova snopa

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
 Load DICOM Recalc Export
 Import from Bin... Import Dose

Status: ready for optimization

Plan

bixel width in [mm] 10 1

Gantry Angle in ° 0 180 225 270 315 2

Couch Angle in ° 0 0 0 0 2

Radiation Mode photons

Machine Generic

IsoCenter in [mm] 265.8 296.7 316.4 Auto.

Fractions 30

Type of optimization none Set Tissue

use MC (VMC++) dose calculations
 3D conformal
 Run Sequencing
 Stratification Levels 7
 Run Direct Aperture Optimizat...

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no

save

Visualization

Slice Type of plot inten... GoTo lateral

Beam Plane axial Open 3D-View

Offset Dislay option physicalDose

Show DVH/QI

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

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axial plane z = 317.5 [mm]

min 0
max 3.706

Set IsoDose Levels

Viewer Options

Result (i.e. dose) Window Doseat Custom
 Window Center: 1.85
 Window Width: 3.71
 Range: 0 3.706
 jet Dose opacity: 1

Structure Visibility

- GTV
- Kidney_R
- Kidney_L
- Stomach
- SmallBowel
- LargeBowel
- Cellac
- SMA_SMV
- Liver
- Heart
- SpinalCord
- DoseFalloff

Info v3.0.0
 github.com/e0404/mat
 About

3. Pokrenite proračun doze pomoću opcije („Calc. Influence Mx“) i započnite obrnutu optimizaciju klikom na („Optimize“)

The screenshot displays the matRad software interface, which is used for radiation therapy planning. The interface is divided into several panels:

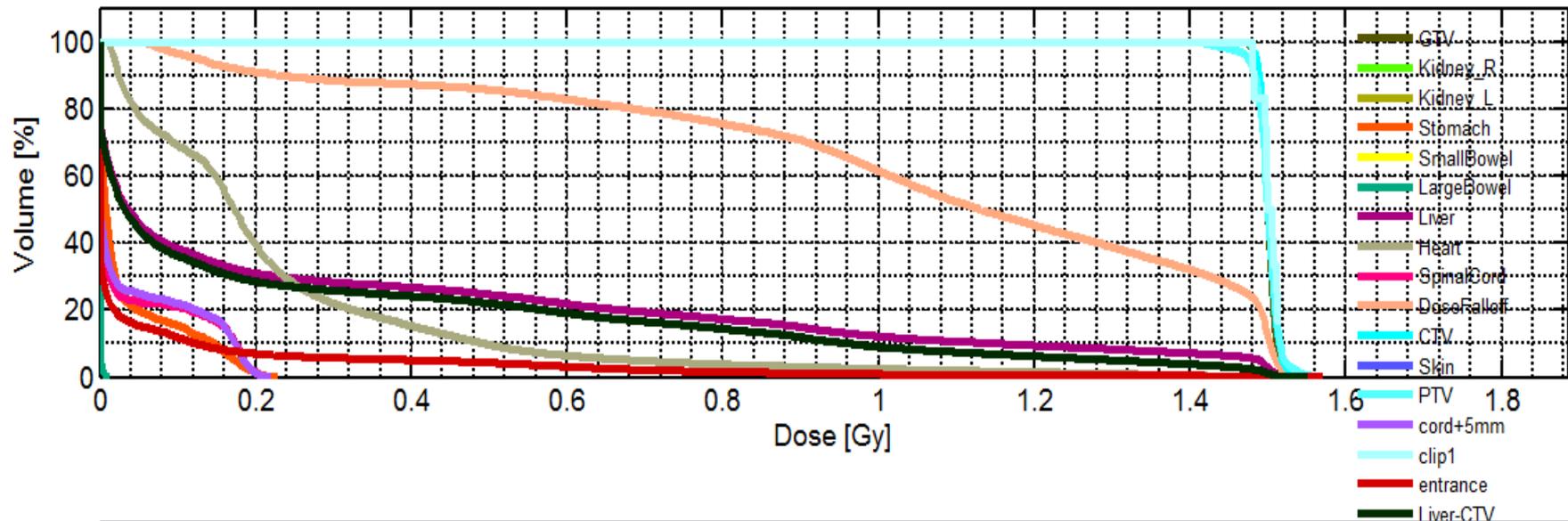
- Workflow:** Contains buttons for 'Refresh', 'Load *.mat data', 'Load DICOM', 'Import from Bin...', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Export', and 'Import Dose'. Red arrows point to 'Calc. influence Mx' (labeled '1') and 'Optimize' (labeled '2'). The status below indicates 'ready for optimization'.
- Plan:** Includes parameters for 'bixel width in [mm]' (10), 'Gantry Angle in °' (0 180 225 270 315), 'Couch Angle in °' (0 0 0 0), 'Radiation Mode' (photons), 'Machine' (Generic), 'IsoCenter in [mm]' (265.8 296.7 316.4), '# Fractions' (30), and 'Type of optimization' (none). It also has options for 'use MC (VMC++) dose calculations', '3D conformal', 'Run Sequencing', 'Stratification Levels' (7), and 'Run Direct Aperture Optimizat...'. A 'Set Tissue' button is also present.
- Objectives & constraints:** A table listing objectives and constraints for different VOI names.
- Visualization:** Shows 'Slice', 'Beam', and 'Offset' settings, 'Type of plot' (intensity), 'Plane' (axial), and 'Dislay option' (physicalDose). It includes a 'GoTo' button set to 'lateral' and an 'Open 3D-View' button. A legend on the right lists various structures to be plotted, such as 'plot CT', 'plot contour', 'plot isolines', 'plot dose', 'plot isolines labels', 'plot iso center', and 'visualize plan / be...'. A 'Show DVH/QI' button is also visible.
- 3D Visualization:** A central window showing an axial CT scan of a patient's head and neck. A color scale on the right indicates 'physicalDose [Gy]' ranging from 0 to 60. The axes are labeled 'x [mm]' and 'y [mm]'. The text 'axial plane z = 317.5 [mm]' is displayed above the image.
- Viewer Options:** A panel on the right side of the 3D view showing 'min' (0) and 'max' (3.706) values, 'Set IsoDose Levels', 'Viewer Options' (Result (i.e. dose), Window, Custom), 'Window Center' (1.85), 'Window Width' (3.71), 'Range' (0 3.706), 'jet' color map, and 'Dose opacity' (1).
- Structure Visibility:** A list of structures to be displayed, including GTV, Kidney_R, Kidney_L, Stomach, SmallBowel, LargeBowel, Cellac, SMA_SMV, Liver, Heart, SpinalCord, and DoseFalloff.
- Info:** A bottom right corner showing the version 'v3.0.0' and the GitHub repository 'github.com/e0404/mat', along with an 'About' button.

4. Spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)

The screenshot displays the matRad software interface, which is used for radiation therapy optimization. The interface is divided into several functional areas:

- Workflow:** Contains buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, and Import dose. A red arrow points to the 'Save to GUI' button, labeled with the number '1'.
- Plan:** Shows various parameters for the optimization, including:
 - bixel width in [mm]: 10
 - Gantry Angle in °: 0 180 225 270 315
 - Couch Angle in °: 0 0 0 0
 - Radiation Mode: photons
 - Machine: Generic
 - IsoCenter in [mm]: 265.8 296.7 316.4 (with 'Auto' checked)
 - # Fractions: 30
 - Type of optimization: none
 - Stratification Levels: 7
 - Options: use MC (VMC++) dose calculations, 3D conformal, Run Sequencing, Run Direct Aperture Optimizat...
- Objectives & constraints:** A table listing constraints for different VOI (Volume of Interest) types:

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no
- Visualization:** Includes controls for Slice, Beam, and Offset. It also has dropdowns for 'Type of plot' (set to 'intentional'), 'Plane' (set to 'axial'), and 'Dislay option' (set to 'physicalDose'). A red arrow points to the 'Show DVH/QI' button, labeled with the number '2'.
- Viewing:** Shows an axial cross-section of a head and neck at z = 317.5 [mm]. A color scale on the right indicates 'physicalDose [Gy]' from 0 to 60. The target area is highlighted in red and yellow, while organs at risk are in blue and green.
- Right Panel:** Contains 'Viewer Options' (Result (i.e. dose), Window: Dose, Custom, Window Center: 0.78, Window Width: 1.57, Range: 0 1.572, jet color map, Dose opacity: 1) and 'Structure Visibility' (listing GTV, Kidney_R, Kidney_L, Stomach, SmallBowel, LargeBowel, Celiac, SMA_SMV, Liver, Heart, SpinalCord, DoseFalloff).



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.3Gy	V_0.6Gy	V_0.9Gy
GTV	1.5000	0.0090	1.5281	1.4727	1.5188	1.5148	1.5002	1.4851	1.4796	1	1	1	
Kidney_R	0	0	0	0	0	0	0	0	0	1	0	0	0
Kidney_L	0	0	0	0	0	0	0	0	0	1	0	0	0
Stomach	0.0342	0.0566	0.2310	0	0.1940	0.1736	0.0082	0	0	1	0	0	0
SmallBowel	0	0	0	0	0	0	0	0	0	1	0	0	0
LargeBowel	2.6018e-04	0.0012	0.0147	0	0.0047	0.0019	0	0	0	1	0	0	0
Celiac	0	0	0	0	0	0	0	0	0	1	0	0	0
SMA_SMV	0	0	0	0	0	0	0	0	0	1	0	0	0
Liver	0.3033	0.4713	1.5526	0	1.5042	1.4889	0.0367	0	0	1	0.2838	0.2190	0
Heart	0.2296	0.2426	1.5232	0.0066	1.1065	0.6913	0.1728	0.0182	0.0141	1	0.2202	0.0650	0
SpinalCord	0.0391	0.0686	0.2167	0	0.1969	0.1856	0	0	0	1	0	0	0

5. Definišite vlastiti plan liječenja protonima s jednim snopom, npr. 315 °, te pokrenite proračun doze pomoću opcije („Calc. Influence Mx“) i obrnutu optimizaciju klikom na („Optimize“)

Workflow

Refresh Load *.mat data **Calc. influence Mx** Optimize Save to GUI
 Load DICOM Export
 Import from Bin... Import Dose

Status: ready for dose calculation

Plan

bixel width in [mm] 10
 Gantry Angle in ° **315**
 Couch Angle in ° 0
 Radiation Mode protons
 Machine Generic
 IsoCenter in [mm] 265.8 296.7 316.4 Auto.
 # Fractions 30
 Type of optimization const_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Displav option physicalDose

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

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Viewing

axial plane z = 317.5 [mm]

physicalDose [Gy]

min max n
 max 1.572

Set IsoDose Levels

Viewer Options

Result (i.e. dose) Window Dose Custom
 Window Center: 0.781
 Window Width: 1.57
 Range: 0 1.572
 jet Dose opacity: 1

Structure Visibility

- GTV
- Kidney_R
- Kidney_L
- Stomach
- SmallBowel
- LargeBowel
- Celiac
- SMA_SMV
- Liver
- Heart
- SpinalCord
- DoseFalloff

Info

v3.0.0
 github.com/e0404/mat
 About

6. Spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)




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min max n
2.501

Set IsoDose Levels

Viewer Options

Result (i.e. dose)
 Window: Prezent
 Custom

Window Center:
 1.25

Window Width:
 2.5

Range: 0 2.501

jet

Dose opacity: 1

Structure Visibility

- GTV
- Kidney_R
- Kidney_L
- Stomach
- SmallBowel
- LargeBowel
- Celiac
- SMA_SMV
- Liver
- Heart
- SpinalCord
- DoseFalloff

Info

v3.0.0
github.com/e0404/mat

About

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize **Save to GUI** Export Import

Load DICOM Recalc

Import from Bin...

Status: plan is optimized 1

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations

Gantry Angle in ° 315 3D conformal

Couch Angle in ° 0 Run Sequencing

Radiation Mode protons Stratification Levels 7

Machine Generic Run Direct Aperture Optimizat...

IsoCenter in [mm] 265.8 296.7 316.4 Auto.

Fractions 30

Type of optimization const_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no

save

Visualization

Slice Type of plot inten... GoTo lateral

Beam Plane axial Open 3D-View

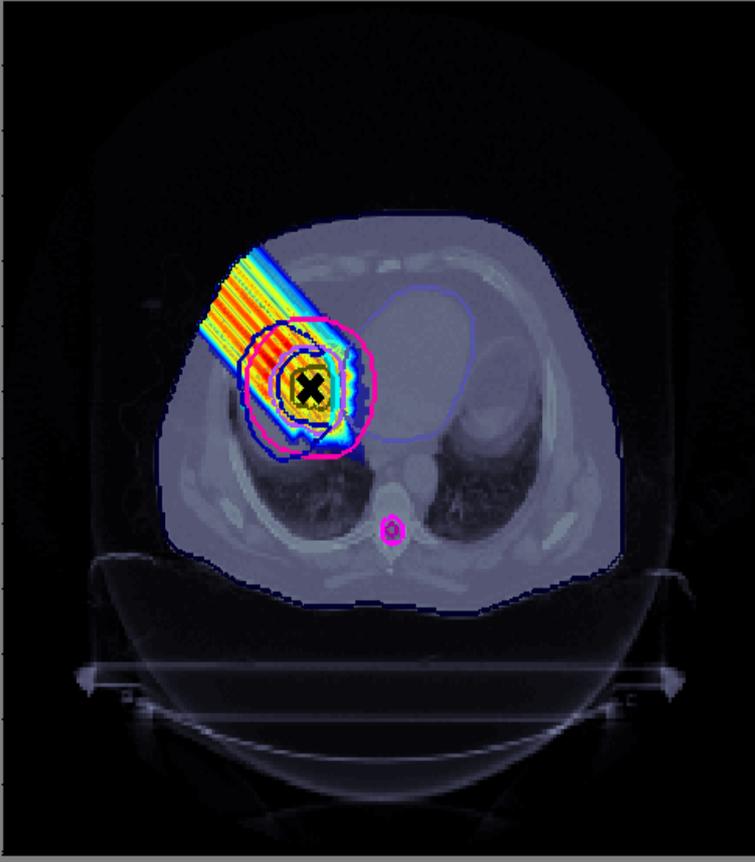
Offset Dislay option RBExDose 2

Show DVH/QI

- plot CT
- plot contour
- plot isolines
- plot dose
- plot isolines labels
- plot iso center
- visualize plan / be...

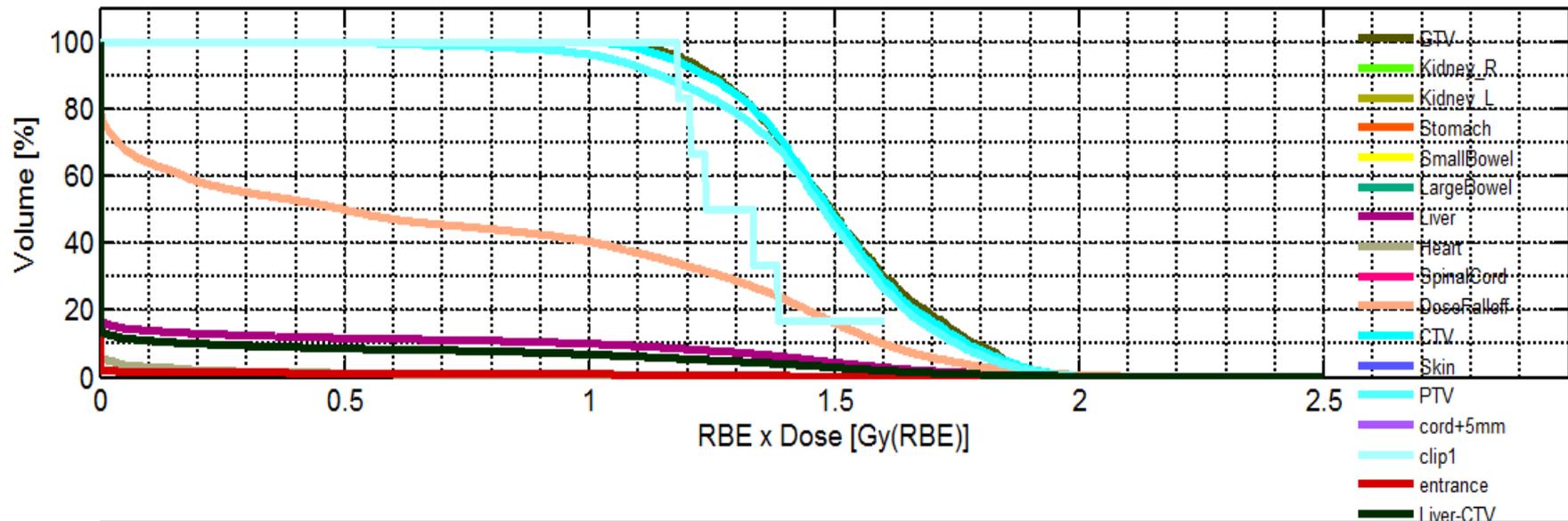
Viewing

axial plane z = 317.5 [mm]



y [mm] 50 100 150 200 250 300 350 400 450 500 550 600 650

x [mm] 50 100 150 200 250 300 350 400 450 500 550 600 650



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.5Gy	V_1Gy	V_1.5Gy
GTV	1.5053	0.1981	2.0110	1.0341	1.8973	1.8506	1.4947	1.1921	1.1231	1	1	1	0
Kidney_R	0	0	0	0	0	0	0	0	0	1	0	0	0
Kidney_L	0	0	0	0	0	0	0	0	0	1	0	0	0
Stomach	0	0	0	0	0	0	0	0	0	1	0	0	0
SmallBowel	0	0	0	0	0	0	0	0	0	1	0	0	0
LargeBowel	0	0	0	0	0	0	0	0	0	1	0	0	0
Celiac	0	0	0	0	0	0	0	0	0	1	0	0	0
SMA_SMV	0	0	0	0	0	0	0	0	0	1	0	0	0
Liver	0.1694	0.4605	2.5011	0	1.6940	1.4688	0	0	0	1	0.1177	0.1008	0
Heart	0.0172	0.1143	1.8597	0	0.2483	0.0195	0	0	0	1	0.0127	0.0050	0
SpinalCord	0	0	0	0	0	0	0	0	0	1	0	0	0

7. Napravite tretman ugljikovim jonima sa tačno istim postavkama kao za plan protonskog tretmana

Workflow

Refresh Load *.mat data **Calc. influence Mx** **Optimize** Save to GUI
 Load DICOM **Re** Export
 Import from Bin... Import Dose

Status: ready for dose calculation

Plan

bixel width in [mm]: 10 use MC (VMC++) dose calculations
 Gantry Angle in °: 315 3D conformal
 Couch Angle in °: 0 Run Sequencing
 Radiation Mode: carbon Stratification Levels: 7
 Machine: photons Auto. Run Direct Aperture Optimizat...
 IsoCenter in [mm]: # Fractions: carbon Type of optimization: LEMIV_RBExD

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no

Visualization

Slice: Type of plot: inten... GoTo: lateral
 Beam: Plane: axial Open 3D-View
 Offset: Dislay option: RBExDose

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

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axial plane z = 317.5 [mm]

min 0
max 2.501

Set IsoDose Levels

Viewer Options

Result (i.e. dose)
 Window: Default
 Custom
 Window Center: 1.25
 Window Width: 2.5
 Range: 0 2.501
 jet
 Dose opacity: 1

Structure Visibility

- GTV
- Kidney_R
- Kidney_L
- Stomach
- SmallBowel
- LargeBowel
- Celiac
- SMA_SMV
- Liver
- Heart
- SpinalCord
- DoseFalloff

Info

v3.0.0
github.com/e0404/mat
 About

8. Spremite rezultat optimizacije putem („Save to GUI“) i prikažite DVH pomoću („Show DVH/QI“)

The screenshot displays the matRad GUI interface. The top left contains a 'Workflow' section with buttons for 'Refresh', 'Load *.mat data', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Load DICOM', 'Recalc', 'Export', and 'Import dose'. A red arrow points to the 'Save to GUI' button, labeled with a '1'. Below this is a 'Plan' section with various parameters like 'bixel width in [mm]', 'Gantry Angle in °', 'Couch Angle in °', 'Radiation Mode', 'Machine', 'IsoCenter in [mm]', '# Fractions', and 'Type of optimization'. The 'Objectives & constraints' section contains a table with columns for 'VOI name', 'VOI type', 'priority', 'obj. / const.', 'penalty', 'dose', 'EUD', 'volume', and 'ro'. The 'Visualization' section at the bottom left includes 'Slice', 'Beam', 'Offset', 'Type of plot', 'Plane', 'Dislay option', and 'Show DVH/QI'. A red arrow points to the 'Show DVH/QI' button, labeled with a '2'. The main viewing area shows an axial plane at z = 317.5 [mm] with a color scale for 'RBExDose [Gy(RBE)]' ranging from 0 to 60. The right side features 'Viewer Options' and 'Structure Visibility' panels.

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
Load DICOM Recalc Export
Import dose

Status: plan is optimized

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations
Gantry Angle in ° 315 3D conformal
Couch Angle in ° 0 Run Sequencing
Radiation Mode carbon Stratification Levels 7
Machine Generic Run Direct Aperture Optimizat...
IsoCenter in [mm] 265.8 296.7 316.4 Auto.
Fractions 30
Type of optimization LEMIV_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	ro
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN	no
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN	no

Visualization

Slice Type of plot inten... GoTo lateral plot CT
Beam Plane axial Open 3D-View plot contour
Offset Dislay option RBExDose plot isolines
Show DVH/QI plot dose
plot isolines labels
plot iso center
visualize plan / be...

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axial plane z = 317.5 [mm]

min 0
max 2.123

Set IsoDose Levels

Viewer Options

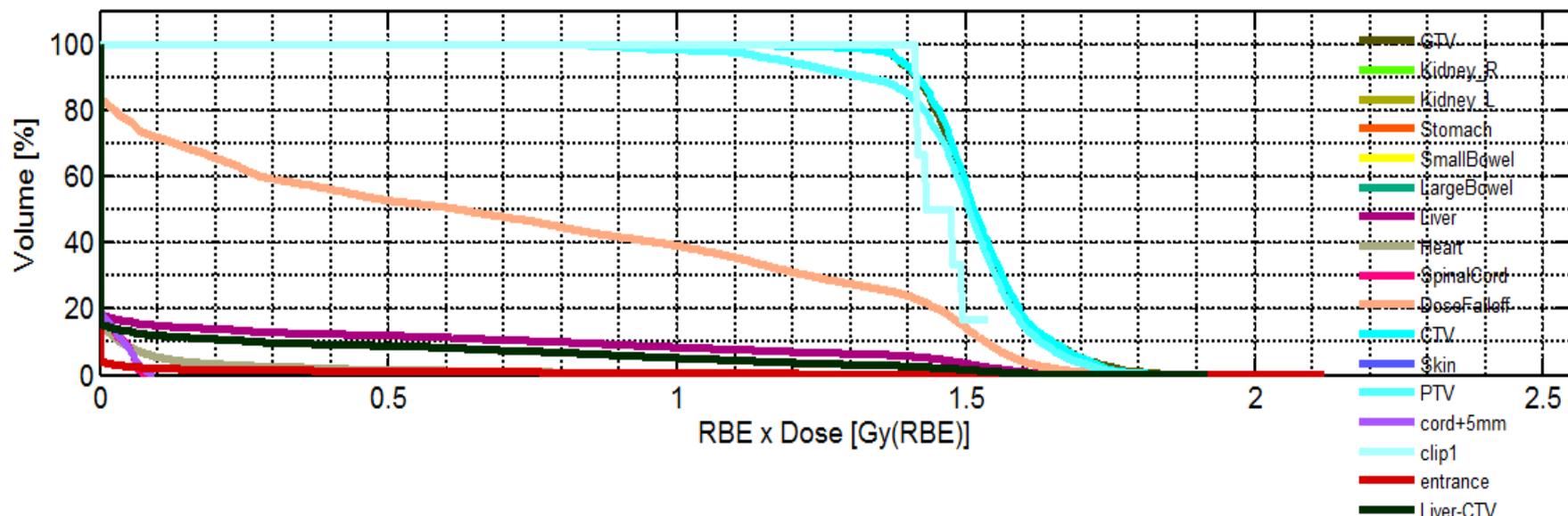
Result (i.e. dose) Window Default Custom
Window Center: 1.06
Window Width: 2.12
Range: 0 2.123
jet Dose opacity: 1

Structure Visibility

- GTV
- Kidney_R
- Kidney_L
- Stomach
- SmallBowel
- LargeBowel
- Celiac
- SMA_SMV
- Liver
- Heart
- SpinalCord
- DoseFalloff

Info

v3.0.0
github.com/e0404/mat
About



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.4Gy	V_0.8Gy	V_1.2Gy
GTV	1.5212	0.0930	1.8920	1.2809	1.7595	1.7032	1.5090	1.3845	1.3641	1	1	1	
Kidney_R	0	0	0	0	0	0	0	0	0	1	0	0	
Kidney_L	0	0	0	0	0	0	0	0	0	1	0	0	
Stomach	0	0	0	0	0	0	0	0	0	1	0	0	
SmallBowel	0	0	0	0	0	0	0	0	0	1	0	0	
LargeBowel	0	0	0	0	0	0	0	0	0	1	0	0	
Celiac	0	0	0	0	0	0	0	0	0	1	0	0	
SMA_SMV	0	0	0	0	0	0	0	0	0	1	0	0	
Liver	0.1570	0.4178	1.9880	0	1.5533	1.4456	0	0	0	1	0.1243	0.1004	0.0000
Heart	0.0277	0.1314	1.8137	0	0.4139	0.1145	0	0	0	1	0.0212	0.0088	0.0000
SpinalCord	0.0077	0.0187	0.0855	0	0.0659	0.0582	0	0	0	1	0	0	

Rezultati

- Srednje doze po regionima (Gy) dobivene pomoću 5 snopova fotona, jednog snopa protona i iona karbona:

Region/Zračenje(uglovi)	Fotoni(0,180,225,270,315)	Protoni(315)	Karbon(315)
GTV	1.5	1.5053	1.5212
Bubrezi	0	0	0
Stomak	0.0342	0	0
Jetra	0.3033	0.1694	0.1570
Srce	0.2296	0.0172	0.0277
Kičmena moždina	0.0391	0	0.0077
CTV	1.5015	1.4981	1.5236
PTV	1.4991	1.4595	1.4868
Koža	0.0568	0.0179	0.0162

3. Zadatak

- Neizvjesnosti u planiranju liječenja
- Planiranje protonske terapije za glavu pacijenta
- Simuliranje greške u postavljanju pacijenta
- Analiza i upoređivanje rezultujućih raspodjela doze

1. Učitajte glavu pacijenta pomoću opcije Load * .mat (HEAD_AND_NECK or ALDERSON.mat)

The screenshot displays the matRad software interface. The main window shows the 'Workflow' section with buttons for 'Refresh', 'Load *.mat data', 'Calc. influence Mx', 'Optimize', 'Save to GUI', 'Load COM', 'Recalc', 'Export', 'Import Dose', and 'Import Bin...'. A red arrow points to the 'Load *.mat data' button. The 'Plan' section shows various parameters like 'bixel width in [mm]', 'Gantry Angle in °', 'Couch Angle in °', 'Radiation Mode', 'Machine', 'IsoCenter in [mm]', '# Fractions', and 'Type of optimization'. The 'Objectives & constraints' section contains a table with columns for 'VOI name', 'VOI type', 'priority', 'obj. / const.', 'penalty', 'dose', 'EUD', and 'volume'. The 'Visualization' section includes 'Slice', 'Beam', 'Offset', 'Type of plot', 'Plane', 'Disolav option', and 'Show DVH/QI'. A 'Select File to Open' dialog box is open, showing a file list with 'HEAD_AND_NECK' selected. The dialog box also shows the file name 'HEAD_AND_NECK' and the file type 'MAT-files (*.mat)'. The background shows the matRad logo and the DKFZ logo (GERMAN CANCER RESEARCH CENTER IN THE HELMHOLTZ ASSOCIATION).

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
Load COM Recalc Export
Import Bin... Import Dose

Status: plan is optimized

Plan

bixel width in [mm] 10
Gantry Angle in ° 315
Couch Angle in ° 0
Radiation Mode carbon
Machine Generic
IsoCenter in [mm] 265.8 296.7 316.4 Auto.
Fractions 30
Type of optimization LEMIV_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	Skin	OAR	2	square overdosing	300	25	NaN	NaN
2	PTV	TARGET	1	square deviation	1000	45	NaN	NaN

Visualization

Slice Type of plot inten... GoTo lateral
Beam Plane axial Open 3D-View
Offset Disolav option RBExDose Show DVH/QI

Select File to Open

File name: HEAD_AND_NECK MAT-files (*.mat)

Open Cancel

2. Dodajte tri ugla za protonske snopove po vašoj želji. Izračunajte i optimizirajte dozu („Calc. Influence Mx“ & „Optimize“).

Workflow

Refresh Load *.mat data **Calc. influence Mx** **Optimize** Save to GUI
 Load DICOM Export
 Import from Bin... Import Dose

Status: ready for dose calculation

Plan

bixel width in [mm] 10
 Gantry Angle in ° 90 180 270
 Couch Angle in ° 0 0 0
 Radiation Mode protons
 Machine Generic
 IsoCenter in [mm] 250.4 205.3 138.5 Auto.
 # Fractions 30
 Type of optimization const_RBExD Set Tissue

use MC (VMC++) dose calculations
 3D conformal
 Run Sequencing
 Stratification Levels 7
 Run Direct Aperture Optimizat...

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	PAROTID_LT	OAR	1	square overdosing	100	25	NaN	NaN
2	PAROTID_RT	OAR	1	square overdosing	100	25	NaN	NaN
3	PTV63	TARGET	2	square deviation	1000	63	NaN	NaN
4	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN

Visualization

Slice Type of plot inten... GoTo lateral
 Beam Plane axial Open 3D-View
 Offset Dislay option

plot CT
 plot contour
 plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...

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Viewing axial plane z = 140 [mm]

min 1024
max 3071

Set IsoDose Levels

Viewer Options

CT (HU)
 Window: Breast
 Custom
 Window Center: 1.27
 Window Width: 2.53
 Range: 0.00324 2.531
 bone
 Dose opacity: 1

Structure Visibility

- BRAIN_STEM
- BRAIN_STEM_PR
- CEREBELLUM
- CHIASMA
- CTV63
- GTV
- LARYNX
- LENS_LT
- LENS_RT
- LIPS
- OPTIC_NRV_LT
- OPTIC_NRV_RT

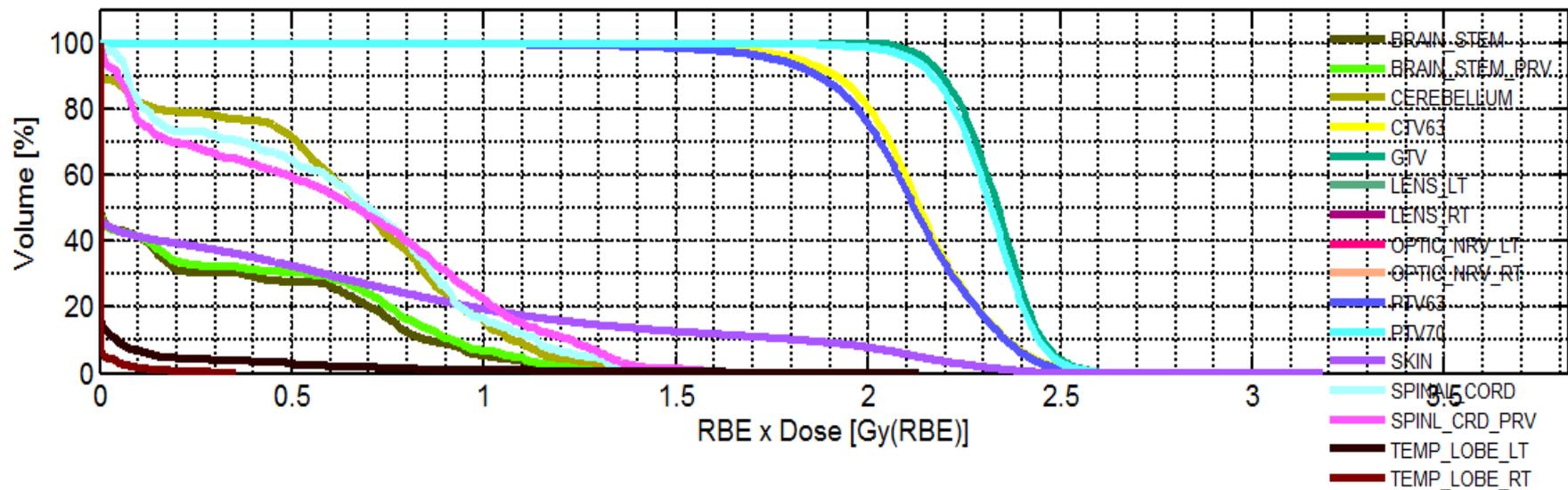
Info
 v3.0.0
 github.com/e0404/mat
 About

3. Analizirajte rezultat (doza i DVH) i sačuvajte („Save to GUI“)

The screenshot displays the matRad software interface, which is used for radiation therapy planning. The interface is divided into several functional areas:

- Workflow:** Contains buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, and Import dose. A red arrow points to the "Save to GUI" button.
- Status:** Indicates "plan is optimized".
- Plan:** Includes input fields for biixel width (10 mm), Gantry Angle (90, 180, 270 degrees), Couch Angle (0, 0, 0 degrees), Radiation Mode (protons), Machine (Generic), IsoCenter (250.4, 205.3, 138.5 mm), # Fractions (30), and Type of optimization (const_RBExD). It also has checkboxes for "use MC (VMC++) dose calculations", "3D conformal", "Run Sequencing", "Stratification Levels" (7), and "Run Direct Aperture Optimizat...".
- Objectives & constraints:** A table listing various VOI (Volume of Interest) names, types, priorities, and associated constraints.
- Visualization:** Includes controls for Slice, Beam, Offset, Type of plot (intensity), Plane (axial), and Dislay option (RBExDose). A red arrow points to the "Show DVH/QI" button.
- Viewing:** Shows an axial plane at z = 140 mm. The visualization includes a color scale for RBExDose [Gy(RBE)] ranging from 0 to 60. The x and y axes are labeled in mm.
- Viewer Options:** Includes "Set IsoDose Levels", "Result (i.e. dose)", "Window Center", "Window Width", "Range", "jet" color map, and "Dose opacity".
- Structure Visibility:** A list of anatomical structures with checkboxes for visibility, including BRAIN_STEM, CEREBELLUM, CHIASMA, CTV63, GTV, LARYNX, LENS, LIPS, and OPTIC NRV.
- Info:** Displays version v3.0.0 and the GitHub repository link github.com/e0404/mat.

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	PAROTID_LT	OAR	1	square overdosing	100	25	NaN	NaN
2	PAROTID_RT	OAR	1	square overdosing	100	25	NaN	NaN
3	PTV63	TARGET	2	square deviation	1000	63	NaN	NaN
4	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.6Gy	V_1.2Gy	V_1.9Gy
BRAIN_STEM	0.2645	0.3831	1.5408	0	1.1597	1.0153	0.0030	0	0	1	0.2649	0.0167	
BRAIN_STEM_PRV	0.2906	0.4099	1.5754	0	1.2980	1.0952	0.0016	0	0	1	0.2896	0.0251	
CEREBELLUM	0.6355	0.3774	2.0785	0	1.3512	1.1661	0.6933	0	0	1	0.5998	0.0469	7.3233
CHIASMA	0	0	0	0	0	0	0	0	0	1	0	0	
CTV63	2.1304	0.1945	3.1861	0.9407	2.4868	2.4230	2.1346	1.8175	1.6587	1	1	0.9973	0.9973
GTV	2.3305	0.1036	2.7047	1.9940	2.5353	2.4898	2.3381	2.1496	2.0935	1	1	1	1
LARYNX	0.9230	0.4283	1.9861	0.2391	1.8607	1.7473	0.8058	0.3375	0.2819	1	0.7891	0.2585	0.2585
LENS_LT	0	0	0	0	0	0	0	0	0	1	0	0	
LENS_RT	0	0	0	0	0	0	0	0	0	1	0	0	
LIPS	0.0157	0.0412	0.2352	1.1603e-35	0.1705	0.1231	5.8836e-06	4.7064e-25	6.6316e-30	1	0	0	
OPTIC_NRV_LT	0	0	0	0	0	0	0	0	0	1	0	0	

4. Simulirajte grešku u postavljanju pacijenta: Uklonite kvakicu u checkboxa za auto-izocentar i odredite novi izocentar. Preračunajte dozu klikom na dugme („Recalc“)

The screenshot displays the matRad software interface, which is part of the German Cancer Research Center (dkfz) in the Helmholtz Association. The interface is divided into several functional areas:

- Workflow:** Contains buttons for Refresh, Load *.mat data, Calc. influence Mx, Optimize, Save to GUI, Load DICOM, Recalc, Export, Import from Bin..., and Import Dose. A red arrow labeled '2' points to the 'Recalc' button.
- Plan:** Includes input fields for 'bixel width in [mm]' (10), 'Gantry Angle in °' (90 180 270), 'Couch Angle in °' (0 0 0), 'Radiation Mode' (protons), 'Machine' (Generic), 'IsoCenter in [mm]' (260 220 150), and '# Fractions' (30). A checkbox for 'Auto' is checked, with a red arrow labeled '1' pointing to it. Other options include 'use MC (VMC++) dose calculations', '3D conformal', 'Run Sequencing', 'Stratification Levels' (7), and 'Type of optimization' (const_RBExD).
- Objectives & constraints:** A table listing various objectives and constraints for the treatment plan.
- Visualization:** Controls for 'Slice', 'Beam', and 'Offset', along with 'Type of plot' (intentional), 'Plane' (axial), and 'Dislay option' (RBExDose). A 'Show DVH/QI' button is also present.
- Viewing:** Shows an axial plane at z = 140 [mm]. The visualization includes a color scale for 'RBExDose [Gy(RBE)]' ranging from 0 to 60. A red 'X' marks the isocenter, and a red circle highlights a specific region.
- Viewer Options:** Includes 'Result (i.e. dose)', 'Window Preset' (Custom), 'Window Center' (1.59), 'Window Width' (3.19), 'Range' (0 to 3.186), 'jet' color map, and 'Dose opacity' (1).
- Structure Visibility:** A list of anatomical structures with checkboxes, including BRAIN_STEM, CEREBELLUM, CHIASSMA, GTV, LARYNX, LENS_LT, LENS_RT, LIPS, OPTIC_NRV_LT, and OPTIC_NRV_RT.
- Info:** Displays the version 'v3.0.0' and the GitHub repository 'github.com/e0404/mat'.

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume	
1	PAROTID_LT	OAR	1	square over dosing	100	25	NaN	NaN	+
2	PAROTID_RT	OAR	1	square over dosing	100	25	NaN	NaN	
3	PTV63	TARGET	2	square deviation	1000	63	NaN	NaN	-
4	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN	
5	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN	



Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI

Load DICOM Recalc Export

Import from Bin... Import Dose

Status: plan is optimized

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations

Gantry Angle in ° 90 180 270 3D conformal

Couch Angle in ° 0 0 0 Run Sequencing

Radiation Mode protons Stratification Levels 7

Machine Generic Run Direct Aperture Optimizat...

IsoCenter in [mm] 260 220 150 Auto.

Fractions 30

Type of optimization const_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	PAROTID_LT	OAR	1	square overdosing	100	25	NaN	NaN
2	PAROTID_RT	OAR	1	square overdosing	100	25	NaN	NaN
3	PTV63	TARGET	2	square deviation	1000	63	NaN	NaN
4	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN

save

Visualization

Slice Type of plot inten... GoTo lateral

Beam Plane axial Open 3D-View

Offset Disolv option physicalDose

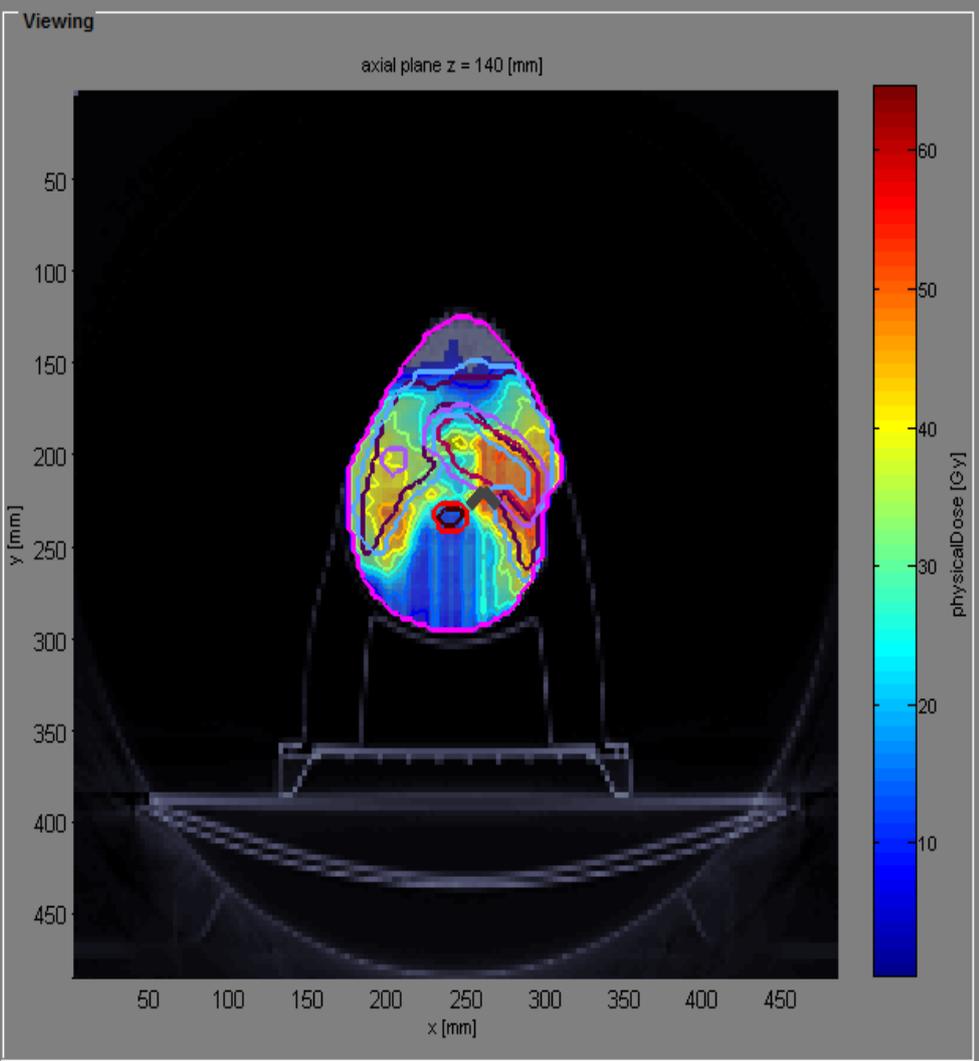
Show DVH/Q!

- plot CT
- plot contour
- plot isolines
- plot dose
- plot isolines labels
- plot iso center
- visualize plan / be...

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min 0
max 3.186

Set IsoDose Levels

Viewer Options

Result (i.e. dose)

Window: Doseat

Custom

Window Center: 1.59

Window Width: 3.19

Range: 0 3.186

jet

Dose opacity: 1

Structure Visibility

- BRAIN_STEM
- BRAIN_STEM_PR
- CEREBELLUM
- CHIASMA
- CTV63
- GTV
- LARYNX
- LENS_LT
- LENS_RT
- LIPS
- OPTIC_NRV_LT
- OPTIC_NRV_RT

Info

v3.0.0

github.com/e0404/mat

About

5. Podešavanjem opcije “Slice” pronađite izocentar te analizirajte i uporedite rezultirajuću raspodjelu doze (doza i DVH)

Workflow

Refresh Load *.mat data Calc. influence Mx Optimize Save to GUI
 Load DICOM Recalc Export
 Import from Bin... Import Dose

Status: plan is optimized

Plan

bixel width in [mm] 10 use MC (VMC++) dose calculations
 Gantry Angle in ° 90 180 270 3D conformal
 Couch Angle in ° 0 0 0 Run Sequencing
 Radiation Mode protons Stratification Levels 7
 Machine Generic Run Direct Aperture Optimizat...
 IsoCenter in [mm] 260 220 150 Auto.
 # Fractions 30
 Type of optimization const_RBExD Set Tissue

Objectives & constraints

	VOI name	VOI type	priority	obj. / const.	penalty	dose	EUD	volume
1	PAROTID_LT	OAR	1	square overdosing	100	25	NaN	NaN
2	PAROTID_RT	OAR	1	square overdosing	100	25	NaN	NaN
3	PTV63	TARGET	2	square deviation	1000	63	NaN	NaN
4	PTV70	TARGET	1	square deviation	1000	70	NaN	NaN

Visualization

Slice inten... GoTo lateral plot CT
 Beam Plane 1 axial Open 3D-View plot contour
 Offset Dislay option physicalDose plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / be...
 Show DVH/QI

matRad dkfz. GERMAN CANCER RESEARCH CENTER IN THE HELMHOLTZ ASSOCIATION

axial plane z = 150 [mm]

physicalDose [Gy]

min 0 max 3.186

Set IsoDose Levels

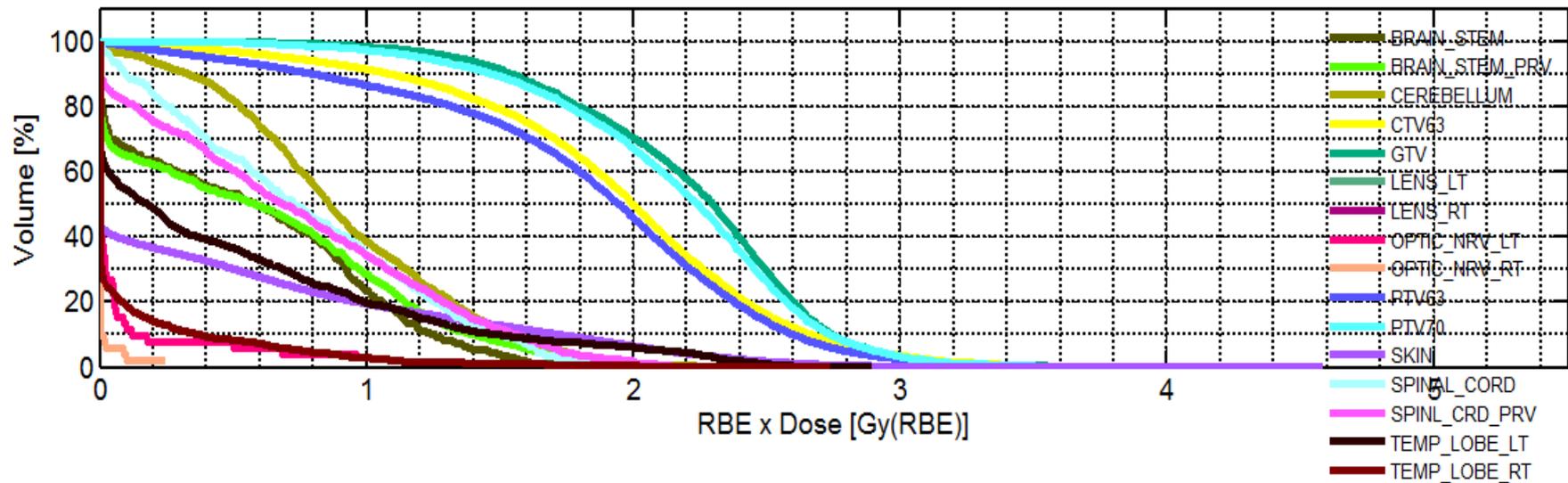
Viewer Options

Result (i.e. dose) Window Doseat Custom
 Window Center: 1.59
 Window Width: 3.19
 Range: 0 3.186
 jet Dose opacity: 1

Structure Visibility

- BRAIN_STEM
- BRAIN_STEM_PR
- CEREBELLUM
- CHIASMA
- CTV63
- GTV
- LARYNX
- LENS_LT
- LENS_RT
- LIPS
- OPTIC_NRV_LT
- OPTIC_NRV_RT

Info v3.0.0 github.com/e0404/mat About



	mean	std	max	min	D_2	D_5	D_50	D_95	D_98	V_0Gy	V_0.9Gy	V_1.8Gy	V_2.7Gy
BRAIN_STEM	0.5784	0.5092	1.8823	0	1.5814	1.4499	0.5847	0	0	1	0.3294	0.0048	
BRAIN_STEM_PRV	0.6153	0.5759	2.3528	0	1.8157	1.6326	0.5786	0	0	1	0.3519	0.0240	
CEREBELLUM	0.9112	0.4681	2.5823	0	1.9049	1.7408	0.8620	0.1636	0.0032	1	0.4588	0.0373	
CHIASMA	0.2487	0.2353	0.8091	0.0071	0.7505	0.6536	0.2672	0.0169	0.0118	1	0	0	
CTV63	1.9376	0.6348	4.0525	0.0093	3.1482	2.8966	1.9997	0.7051	0.3469	1	0.9282	0.6441	0.0001
GTV	2.2150	0.4918	3.9825	0.4100	3.1008	2.8992	2.2980	1.3330	1.0648	1	0.9886	0.7991	0.0001
LARYNX	0.5702	0.3493	1.7209	0.0422	1.5158	1.2552	0.4717	0.1262	0.1006	1	0.1769	0	
LENS_LT	0	0	0	0	0	0	0	0	0	1	0	0	
LENS_RT	0	0	0	0	0	0	0	0	0	1	0	0	
LIPS	0.0064	0.0261	0.2268	0	0.0963	0.0371	8.7893e-18	0	0	1	0	0	
OPTIC_NRV_LT	0.0775	0.2143	0.9674	0	0.9571	0.5805	7.5343e-04	0	0	1	0.0385	0	

Rezultati

- Srednje doze po regionima (Gy) dobivene pomoću tri snopa protona, sa i bez pomijeranja:

Region/Izocentar	Bez pomijeranja	Sa pomijeranjem
Moždano stablo	0.2645	0.5784
Mali mozak	0.6355	0.9112
CTV63	2.1304	1.9376
GTV	2.3305	2.2150
Leće (L,D)	0,0	0,0
Koža	0.4682	0.4555
Optički nerv (L,D)	0,0	0.0775, 0.0092
Kičmena moždina	0.6268	0.7466
PTV63	2.1092	1.8369
PTV70	2.3102	2.1671

Hvala na pažnji :)