

## Introduction to matRad

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**HIRO**

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supported by:  
German Cancer Research Center (DKFZ)  
Heidelberg University Hospital  
Heidelberg Ion Therapy Center (HIT)  
Medical Faculty Heidelberg

matRad 

**dkfz.**



Research for a Life without Cancer

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## Treatment Planning

- computerized process
- dose is numerically simulated and optimized

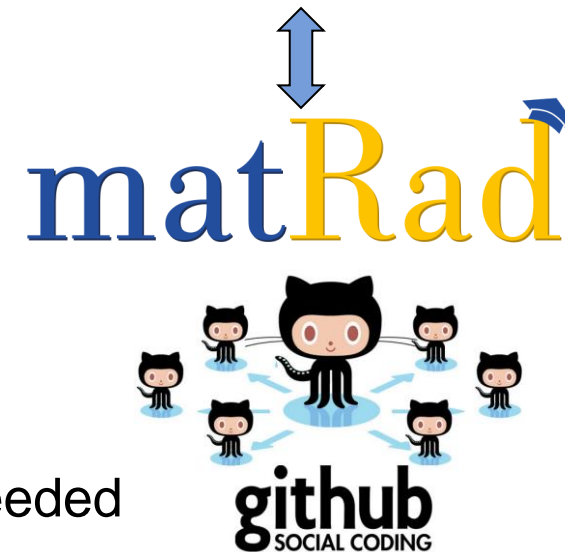
Commercial solutions are closed systems (Black Box)



Research needs flexible, accessible software

Examples for research topics “at the core”:

- Biological Optimization/Planning (RBE, effect, mixed-modality, FLASH)
  - Probabilistic dose calculation & optimization
- low-level access to dose calculation / optimization needed



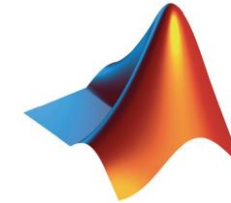
# What is matRad?

- toolkit for three-dimensional intensity-modulated treatment planning for photons, protons and carbon ions
- Entirely written in Matlab & open source
- matRad implements well-established radiotherapy algorithms for **research & education**

## Properties:

- **open-source code, patients and machine files on GitHub**
- **graphical user interface**
- **Non-linear constrained dose optimization (IPOPT)**
- **Import & export functionalities (DICOM, binary formats)**
- **No Matlab? → Octave compatibility & downloadable standalone**

**Why?** Supporting open science, reproducibility and education



[www.matrad.org](http://www.matrad.org)

# Team



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**DFG**

Deutsche  
Forschungsgemeinschaft

BA 2279/3-1  
WA 4707/1-1

## DKFZ Development team

**Niklas Wahl**

Lucas Burigo

Amit Ben Antony Bennan

Jennifer Hardt

Jeremias Kunz

Noa Homolka

**Oliver Jäkel**

## HIT cooperation

**Benjamin Ackermann**

Swantje Ecker

**Malte Ellerbrock**

Andrea Mairani

Thomas Tessonier

**Katia Parodi (LMU)**

## Advisors

Martin Siggel

Peter Ziegenhein

H.-P. Wieser *et al.* (all authors above in **bold**), “Development of the open-source dose calculation and optimization toolkit matRad,” *Med Phys*, vol. 44, no. 6, pp. 2556–2568, 2017, doi: [10.1002/mp.12251](https://doi.org/10.1002/mp.12251).

## Alumni

**Mark Bangert**

**Hans-Peter Wieser**

**Eduardo Cisternas**

Ahmad Neishabouri

Cindy Herman

Thomas Klinge

Verena Böswald

**Henning Mescher**

**Alexander Stadler**

Guisepppe Pezzano

**Lucas-Raphael Müller**

**Hubert Gabrys**

**Silke Ulrich**

Oliver Schrenk

Paul Meder

## Other Contributors

Eric Crhistiansen (Carleton University)

Steven van de Water (PSI)

All Code Contributors: <https://github.com/e0404/matRad/blob/master/AUTHORS.txt>

# More than 25 confirmed institutes somehow working with matRad

Start: 22. Januar 2015

- Github-Fork
- Citations:

Filters: is:issue is:open

Issue Title	Issue Number	Opening Date	Author	Labels	Comments
20 Open	186 Closed				
Error with matRad_calcDVH()	#530	opened 8 days ago	fratomtom		4
Error when importing DICOM	#529	opened 12 days ago	fratomtom	bug	15
[Feature Request] Separate dose threshold for matRad_gammaIndex	#515	opened on 14 Jul	wahln	enhancement	
Question about VMAT branches	#499	opened on 31 Mar	chh105	bug, question	55
question about the dij structure	#477	opened on 11 Jan	chh105	enhancement, question	18
[BUG] Color-Scale in GUI wrong after re-optimization with different number of fractions	#472	opened on 28 Dec 2020	wahln	bug	
Sensible structure set resampling for optimization?	#462	opened on 6 Nov 2020	wahln	enhancement, question	
matRad has issues with 2D patients	#324	opened on 23 Jan 2019	wahln	bug, enhancement	



BERRA HOSPITAL HEALTH SERVICES




UNIVERSITÄT HEIDELBERG  
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### Development of the open-source dose calculation and optimization toolkit matRad

Hans-Peter Wieser , Eduardo Cisternas, Niklas Wahl, Silke Ulrich, Alexander Stadler, Henning Mescher, Lucas-Raphael Müller, Thomas Klinge, Hubert Gabrys, Lucas Burigo, Andrea Mairani, Swantje Ecker, Benjamin Ackermann, Malte Ellerbrock, Katia Parodi, Oliver Jäkel, Mark Bangert

*Wieser et al., 2017, Med Phys 44(6)  
- among top 20 downloaded Med Phys  
papers in 2017 -*

- 3D dose calculation (validated)

- Photons: SVD pencil-beam algorithm + sequencing  
MC interface to ompMC (open source)
- Protons: Pencil-Beam algorithm + const. RBE  
MC interface to MCsquare (open source)
- Carbon ions: Pencil-Beam algorithm + biol. effect / RBE

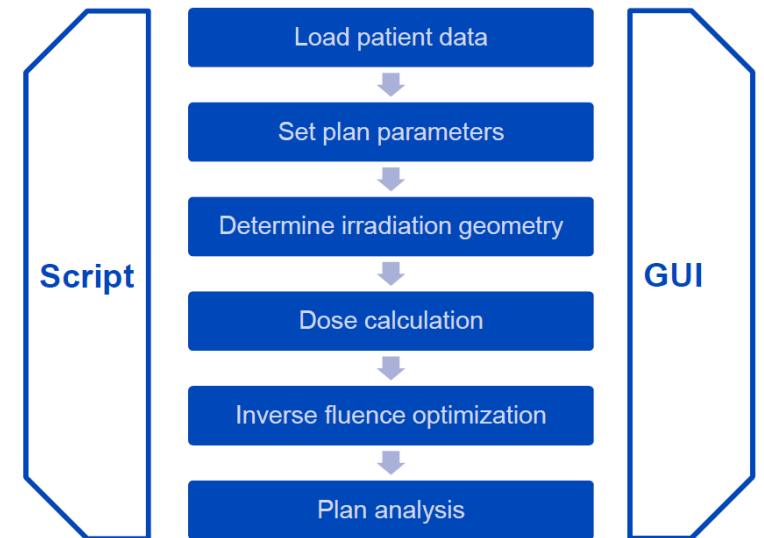
- Base data

- Patient data (CORT data set) & DICOM Import
- Physical (& biological) base data for photon LINAC as well as a proton and a carbon machine

- Inverse planning with new optimization interface

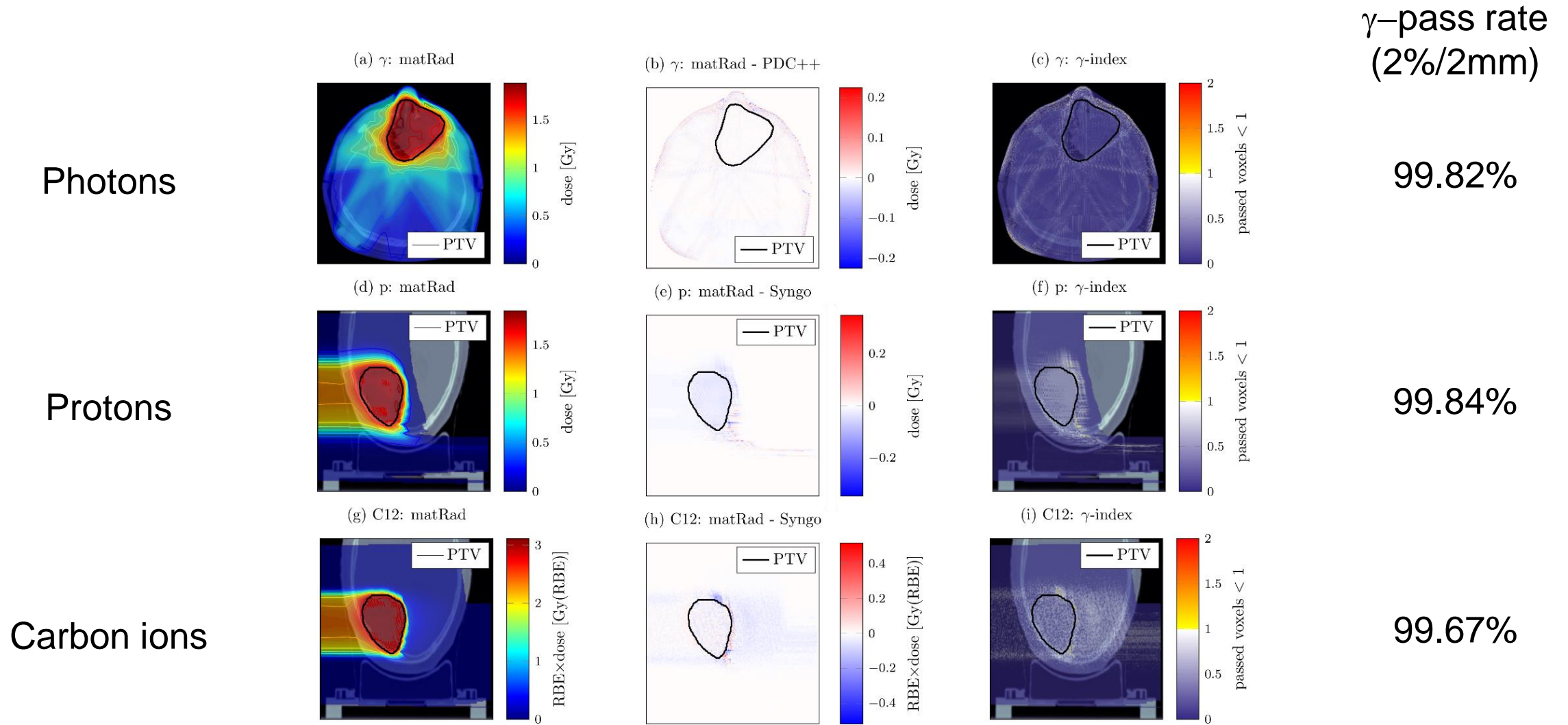
- Photons: Physical dose optimization & DAO
- Protons: + Constant RBE optimization
- Carbon-ions: + RBE (1.1 or variable) or effect optimization

- Scripting & Graphical User Interface





# Validation of dose calculation



Wieser et al. "Development of the open-source dose calculation and optimization toolkit matRad." Medical Physics (2017).

### Workflow

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

Status: plan is optimized

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### Plan

bixel width in [mm]: 5  
 Gantry Angle in °: 90 270  
 Couch Angle in °: 0 0  
 Radiation Mode: protons  
 Machine: Generic  
 IsoCenter in [mm]: 263.3 265.9 124  Auto.  
 # Fractions: 30  
 Type of optimization: const\_RBExD

3D conformal  
 Run Sequencing  
 Stratification Levels: 7  
 Run Direct Aperture Optimization

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### Objectives & constraints

+-	VOI name	VOI type	OP	Function	p	Parameters
-	Rectum	OAR	3	Squared Overdosing	300	$d^{max}$ : 50
-	PTV_68	TARGET	1	Squared Deviation	1000	$d^{ref}$ : 68
-	PTV_56	TARGET	2	Squared Deviation	1000	$d^{ref}$ : 56
-	Bladder	OAR	3	Squared Overdosing	300	$d^{max}$ : 50
-	BODY	OAR	4	Squared Overdosing	100	$d^{max}$ : 30

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### Visualization

Slice Selection:  Type of plot: intensity GoTo: lateral  plot CT  
 Beam Selection:  Plane Selection: axial   plot contour  
 Offset:  Display option: RBExDose  plot isolines  
 plot dose  
 plot isolines labels  
 plot iso center  
 visualize plan / beams

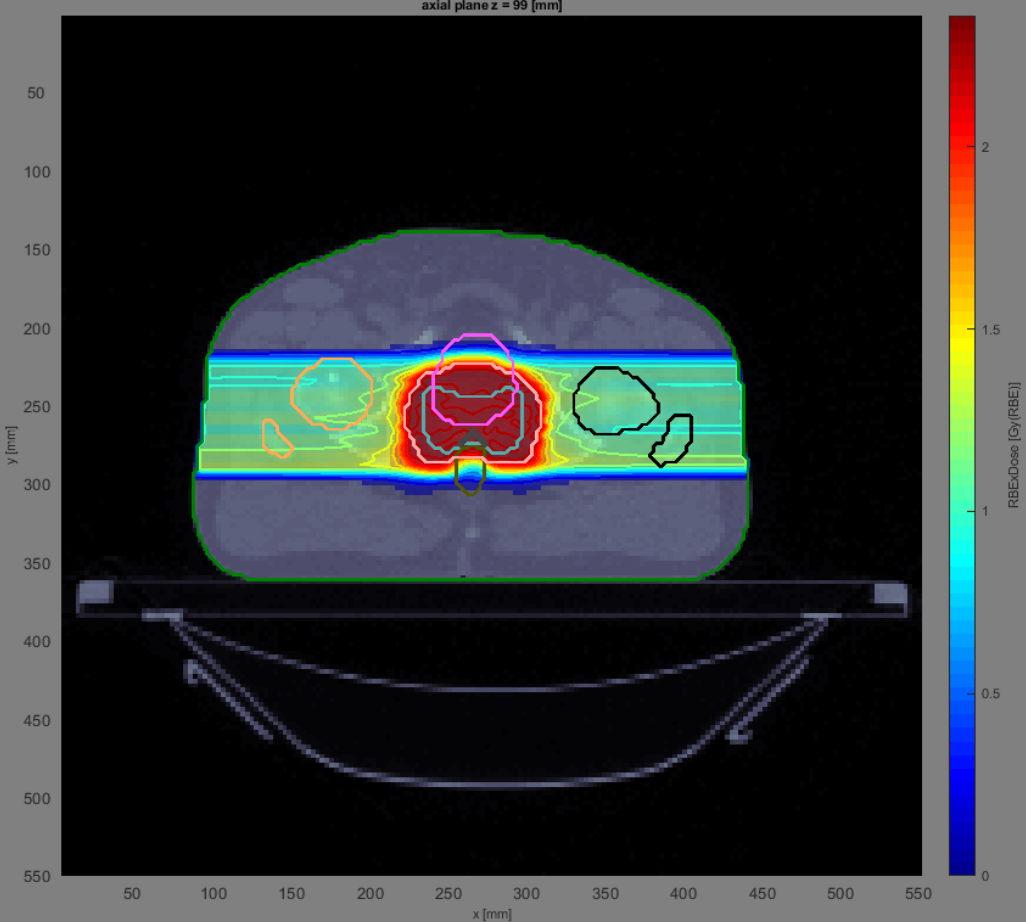



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### Viewing

axial plane z = 99 [mm]



min value: 0  
max value: 2.3596

#### Viewer Options

Result (i.e. dose):   
 Window Preset: Custom  
 Window Center: 1.18  
 Window Width: 2.36  
 Range: 0 2.36  
 jet

Lock Settings  
 Dose opacity: 1

#### Structure Visibility

- Rectum
- Penile\_bulb
- Lymph Nodes
- Rt femoral head
- prostate\_bed
- PTV\_68
- PTV\_56
- Bladder
- BODY
- Lt femoral head

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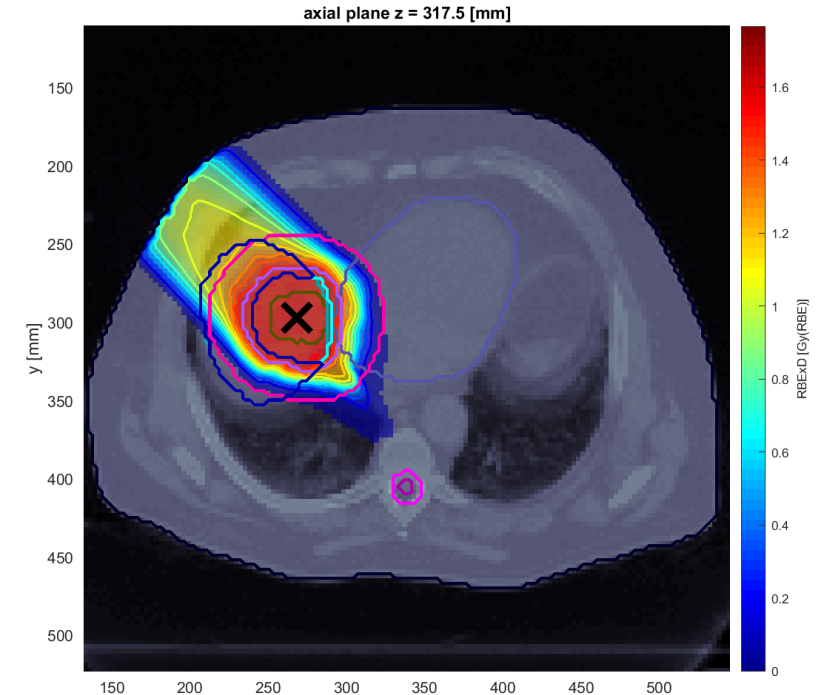
#### Info

v2.10.0 "Blaise"  
(master-c22da7d2)  
www.matRad.org



## Development & Research Branches

- Helium base data (physical & biological)  
`-dev_varRBErobOpt`
- Robust / probabilistic optimization & uncertainty quantification  
`-dev_varRBErobOpt`
- Variable RBE & effect for protons  
`-dev_varRBErobOpt`
- New GUI (Object-oriented & modular, Octave compatible)  
`-dev_classGUI`
- Extended MC interfaces  
`-dev_MonteCarlo`
- Superiorization  
`-research/superiorization`
- External contributions:
  - VMAT `-dev_VMAT(Eric Christiansen)`
  - optimization `-dev_exactOpt (Steven van de Water)`





[www.matrad.org](http://www.matrad.org)



# matRad

## Data IO

DICOM  
\*.nrrd, \*.mha, \*.vtk  
CERR  
VOXELPLAN

## Dose calculation

Photons  
SVD pencil beam  
ompMC interface  
Particles  
IMPT pencil beam  
MCSquare interface  
[TOPAS interface](#)  
Analytical probabilistic modeling

## Analysis & visualization

GUI CT & dose distribution browser  
Dose statistics  
DVHs

## Dose optimization

Fluence and experimental direct aperture optimization  
IPOPT <https://projects.coin-or.org/lpopt>  
Matlab's proprietary fmincon  
Superiorization  
Objectives: Quad. dose deviation, mean dose, EUD, DVH  
Constraints: Min, max, mean dose, EUD, DVH  
Xia, Engel, Siochi MLC sequencer  
[Robust and stochastic optimization](#)  
[Variable RBE optimization for protons](#)  
Coverage based optimization  
Analytical probabilistic modeling  
VMAT

## Base data

Patient data (CT & RTSS)  
Photon pencil beam base data  
→ <https://github.com/e0404/photonPencilBeamKernelCalc>  
Generic proton and carbon ion pencil beam base data  
Carbon ion biological base data (LEM IV)  
[Helium pencil beam base data](#)  
[Helium biological model](#)

**Thank you for your attention!**

Features in [blue](#) are available on development branches

## Exercise to hand in:

- Choose one of the datasets:  
BOXPHANTOM, TG119, LIVER, HEAD\_AND\_NECK, PROSTATE
- Compare the three modalities in matRad
  - Export dose slices and dose-volume histograms (DVH)
    - Create a multi-beam IMRT photon plan (5-9 beams)
    - Create a proton IMPT plan (1-3 beams)
    - Create a carbon ion plan (1-2 beams)