

Version 3 of the open-source radiotherapy dose calculation and treatment

matRad

Beyond nominal dose calculation and optimization

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Group Radiotherapy Optimization @ DKFZ

Embedded in the Department of Medical Physics in
Radiation Oncology (Prof. Jäkel)

- 1 PostDoc
- 5(+2) PhD students
- 1 M. Sc. Student
- 1 B. Sc. Student
- 1 HiWi

Website: www.dkfz.de/radopt

Research Focus:

- **(Beyond) dose calculation**

Biological dose calculation for particle therapy, nanodosimetry, uncertainty quantification, etc.

- **Treatment planning / optimization techniques**

*Robust & probabilistic optimization, mixed-modality optimization, (N)TCP-based planning,
FLASH/dose-rate optimization*

- **Data Science & AI/ML**

Treatment outcome prediction, AI-driven dose prediction

- Developing/Maintaining the open-source planning toolkit matRad

www.matRad.org



Group Picture: G. Stanic, C. Sepulveda, R. Cristoforetti, T. Becher, S. Facchiano, J. Hardt, A.B.A. Bennan, N. Wahl
Not in picture: P. Stammer, T. Ortkamp, L. Bucher



www.dkfz.de/radopt

dkfz.

Outline

- What & Who is matRad?
- Previous releases & development
- The Coming Release (v3)
- Going further beyond dose & future developments
- How can you get started with matRad?

Treatment Planning

- is a computerized process in which the dose is numerically simulated and optimized

Commercial solutions are closed systems (Black Box)

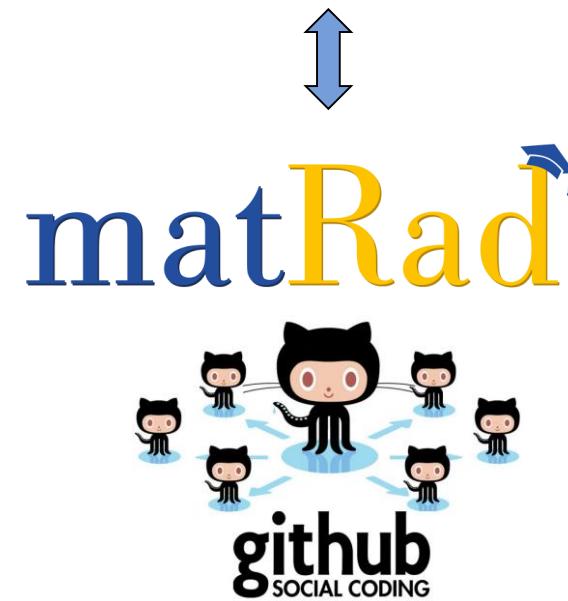


Research needs flexible, accessible software

“Basic” research, e.g.,

- Biological Optimization/Planning
(RBE, effect, mixed-modality, FLASH)
- “Beyond” dose calculation & optimization (probabilistic, nanodosimetric, etc.)

needs low-level access to dose calculation / optimization!

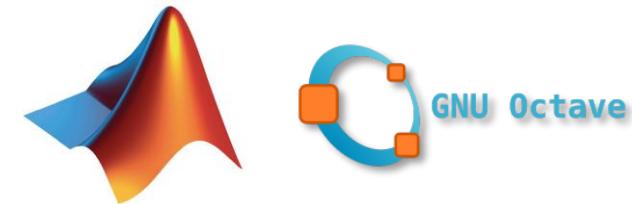


What is matRad?

- 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 (prototyping) & education

Properties:

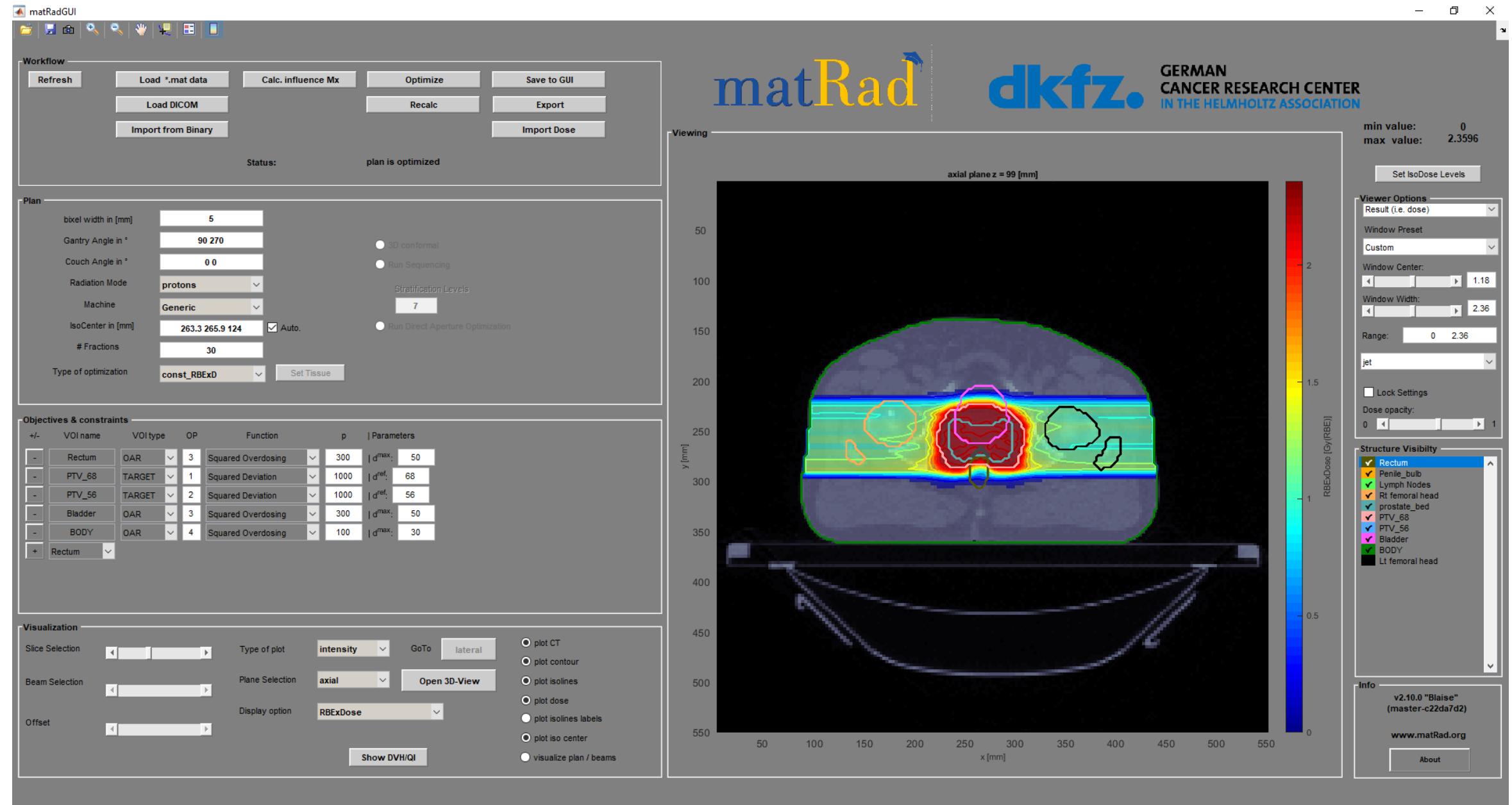
- **open-source code, patients and machine files on GitHub**
- **graphical user interface**
- **Non-linear constrained dose optimization (IPOPT)**
- **Import & export functionalities (DICOM, binary formats)**
- **No Matlab? → Octave compatibility & downloadable standalone**



Why? Supporting open science, reproducibility and education

www.matrad.org

dkfz.



Who is matRad ?



DKFZ Development team

Niklas Wahl

Amit Ben Antony Bennan

Jennifer Hardt

Remo Cristoforetti

Simona Facchiano

Tobias Becher

Noa Homolka

Oliver Jäkel

Alumni

Mark Bangert

Hans-Peter Wieser

Eduardo Cisternas

Ahmad Neishabouri

Cindy Herman

Thomas Klinge

Verena Böswald

Henning Mescher

Alexander Stadler

Guiseppe Pezzano

Lucas-Raphael Müller

Hubert Gabrys

Silke Ulrich

Oliver Schrenk

Paul Meder

Lucas Burigo

HIT cooperation

Benjamin Ackermann

Swantje Ecker

Malte Ellerbrock

Andrea Mairani

Thomas Tessonnier

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Advisors

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Peter Ziegenhein

Other Contributors

Eric Christiansen (Carleton University)

Steven van de Water (PSI)

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).

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



GERMAN
CANCER RESEARCH CENTER
IN THE HELMHOLTZ ASSOCIATION



Deutsche
Forschungsgemeinschaft

Project 265744405

Project 443188743



Detailed core features of the first matRad release

- Data I/O:

DICOM: CT, RTStruct, RTDose (RTPlan import only)
Binary: *.mat & *.nrrd import & export, *.mha & *.vtk export

- 3D dose calculation (validated)

Photons: SVD pencil-beam algorithm + sequencing
Protons: Pencil-Beam algorithm + const. RBE
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 (variable) or effect optimization

- Scripting & Graphical User Interface

- Standalone executable (GUI only)

MEDICAL PHYSICS

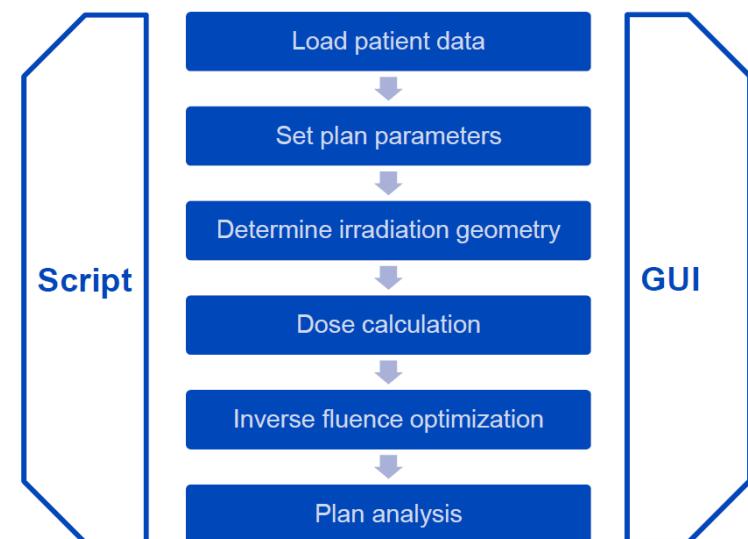
The International Journal of Medical Physics Research and Practice

Research Article | Open Access | CC BY SA

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

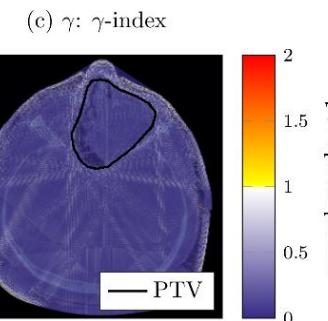
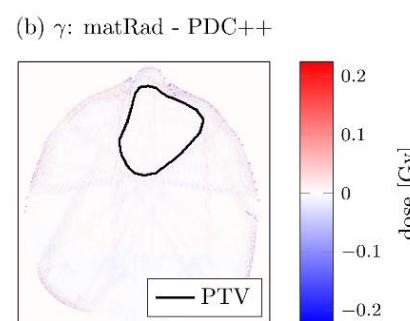
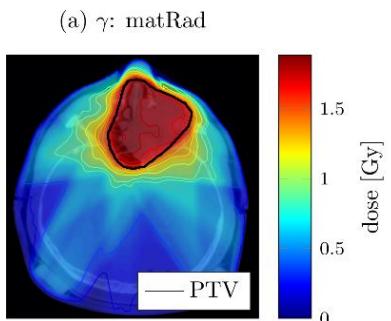
Hans-Peter Wieser, Eduardo Cisternas, Niklas Wahl, Silke Ulrich, Alexander Stadler, Henning Mescher, Lucas-Raphael Müller, Thomas Klinge, Hubert Gabrys, Lucas Burigo, Andrea Mairani, Swantje Ecker, Benjamin Ackermann, Malte Ellerbrock, Katia Parodi, Oliver Jäkel, Mark Bangert

Wieser et al., 2017, Med Phys 44(6)



Validation of dose calculation

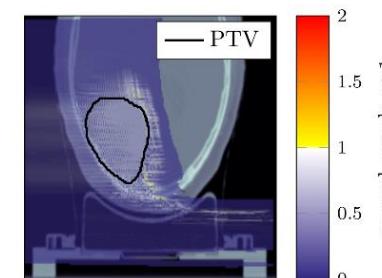
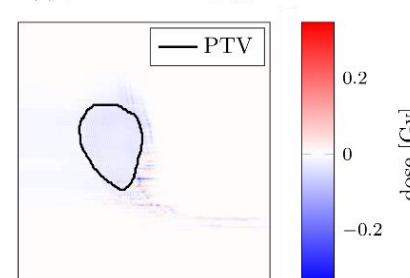
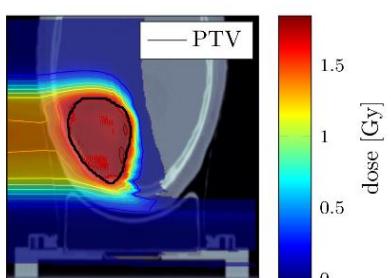
Photons



γ -pass rate
(2%/2mm)

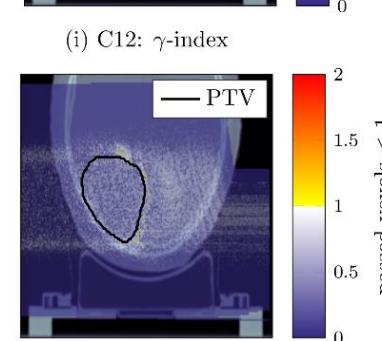
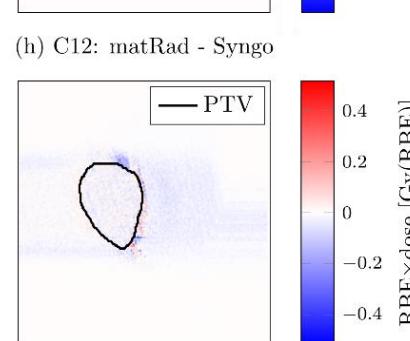
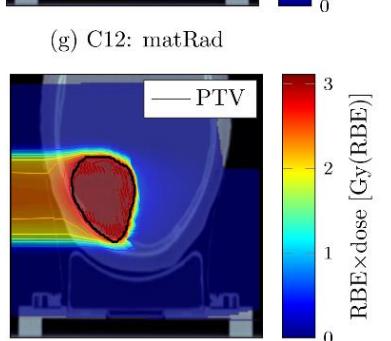
99.82%

Protons



99.84%

Carbon ions



99.67%

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

Inverse dose optimization

Non-linear constrained (Quasi-)Newton methods

- fmincon (Matlab's Optimization Toolbox)
- IPOPT (Interior-point optimizer for non-linear constrained optimization): <https://github.com/coin-or/Ipopt> - EPLv2

$$\begin{aligned} \min_{w \in \mathbb{R}^B} f(w) &= \sum_n p_n f_n(w) \\ \text{s.t. } c_k^l \leq c_k(w) \leq c_k^u \\ 0 \leq w \end{aligned}$$

| | objectives | constraints |
|-------------------------|--------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------|
| $f_{sq\ deviation}$ | $= \frac{1}{N_S} \sum_{i \in S} (d_i - \hat{d})^2$ | |
| $f_{sq\ under\ dosage}$ | $= \frac{1}{N_S} \sum_{i \in S} \Theta(\hat{d} - d_i)(d_i - \hat{d})^2$ | $c_{min\ dose} = d_{min} - \kappa \log \left(\sum_{i \in S} e^{\frac{d_{min} - d_i}{\kappa}} \right)$ |
| $f_{sq\ over\ dosage}$ | $= \frac{1}{N_S} \sum_{i \in S} \Theta(d_i - \hat{d})(d_i - \hat{d})^2$ | $c_{max\ dose} = d_{max} + \kappa \log \left(\sum_{i \in S} e^{\frac{d_i - d_{max}}{\kappa}} \right)$ |
| f_{mean} | $= \frac{1}{N_S} \sum_{i \in S} d_i$ | $c_{mean} = \frac{1}{N_S} \sum_{i \in S} d_i$ |
| f_{EUD} | $= \left(\frac{1}{N_S} \sum_{i \in S} d_i^a \right)^{\frac{1}{a}}$ | $c_{EUD} = \left(\frac{1}{N_S} \sum_{i \in S} d_i^a \right)^{\frac{1}{a}}$ |
| $f_{min\ DVH}$ | $= \frac{1}{N_S} \sum_{i \in S} \Theta(\hat{d} - d_i) \Theta(d_i - \tilde{d}) (d_i - \hat{d})^2$ | $c_{min\ DVH} = \frac{1}{N_S} \sum_{i \in S} \Theta(\hat{d} - d_i)$ |
| $f_{max\ DVH}$ | $= \frac{1}{N_S} \sum_{i \in S} \Theta(d_i - \hat{d}) \Theta(\tilde{d} - d_i) (d_i - \hat{d})^2$ | $c_{max\ DVH} = \frac{1}{N_S} \sum_{i \in S} \Theta(d_i - \hat{d})$ |

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

Performance of matRad - Intel Core i7 2.8 GHz, 32 GB RAM

| modality | setting | #beams | #bixel | D_{ij} elem. [1e6] | D_{ij} size [GB] | t_{dose} [s] | #iter. | t_{opt} [s] |
|----------|----------------------|--------|--------|-------------------------|-----------------------|-------------------|--------|------------------|
| - | - | - | - | - | - | - | - | - |
| photons | 82 mm D_{ij} samp. | 4 | 2608 | 172 | 2.75 | 295 | 145 | 82 |
| photons | 40 mm no samp. | 4 | 2608 | 99 | 1.59 | 101 | 143 | 44 |
| photons | 82 mm D_{ij} samp. | 8 | 3877 | 426 | 6.81 | 741 | 51 | 140 |
| photons | 40 mm no samp. | 8 | 3877 | 236 | 3.77 | 226 | 51 | 66 |
| photons | 40 mm D_{ij} samp. | 72 | 13597 | 567 | 9.07 | 853 | 147 | 407 |
| protons | 99.75 % SG | 1 | 7797 | 19 | 0.29 | 22 | 123 | 41 |
| protons | 99.75 % DG | 1 | 5955 | 87 | 1.38 | 46 | 171 | 109 |
| protons | 99.75 % SG | 3 | 28097 | 56 | 0.89 | 68 | 67 | 187 |
| protons | 99.75 % DG | 3 | 24137 | 269 | 4.30 | 160 | 262 | 330 |
| protons | 99.75 % SG | 2 | 45574 | 116 | 1.86 | 97 | 218 | 137 |
| protons | 99.75 % DG | 2 | 27683 | 520 | 8.33 | 299 | 197 | 486 |
| carbon | 99.75 % SG | 1 | 11780 | 160 | 2.55 | 67 | 72 | 92 |
| carbon | 99.75 % DG | 1 | 9963 | 537 | 8.61 | 203 | 79 | 225 |
| carbon | 99.75 % SG | 3 | 42810 | 411 | 6.68 | 310 | 117 | 193 |
| carbon | 99.75 % DG | 3 | 31205 | 756 | 12.1 | 560 | 107 | 365 |
| carbon | 99.75 % SG | 2 | 24612 | 336 | 5.88 | 137 | 177 | 273 |
| carbon | 99.50 % DG | 2 | 16889 | 855 | 17.94 | 472 | 134 | 521 |

Throughput Optimization: 6 GB/s

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

More than 25 confirmed institutes somehow working with matRad

- Github-
- Citation



A screenshot of a GitHub pull requests page for the "matRad" repository. The repository name is "e0404 / matRad". The "Pull requests" tab is selected, showing 14 open pull requests and 336 closed ones. A filter bar at the top shows "is:pr is:open". The list of pull requests includes:

- #640: Implementation of the analytical dose engine (opened 4 days ago by SimonaFa)
- #639: Dev mix modality effect prescription fix (opened 4 days ago by remocristoforetti)
- #624: Small changes to where penalties are applied (opened on May 26 by tobiasbecher • Changes requested)
- #623: Dev mix mod effect prescription bug fix (opened on May 24 by remocristoforetti)
- #610: Phantom builder added (opened on Apr 3 by tobiasbecher • Changes requested)
- #600: Incorporate multiple weighted lateral Gaussian Profiles (opened on Feb 24 by JenHardt • Changes requested)
- #561: Image Registration / Robustness Evaluation / Cheap Minimax (opened on Jun 10, 2022 by acsevillam • Draft)

The GitHub interface includes various filters, search bars, and navigation buttons at the top.

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AND HEALTH SERVICES

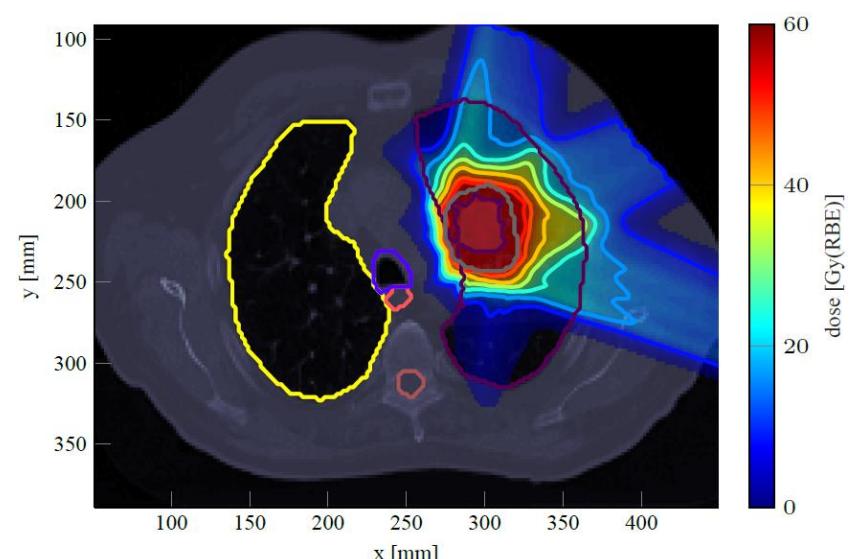
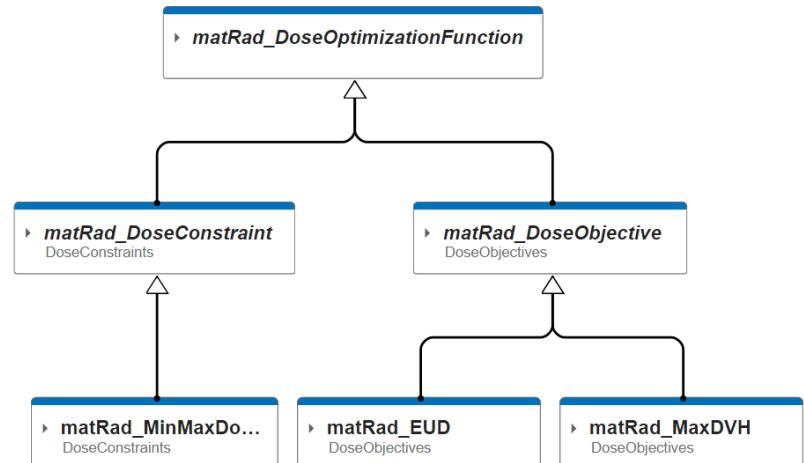


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dkfz.

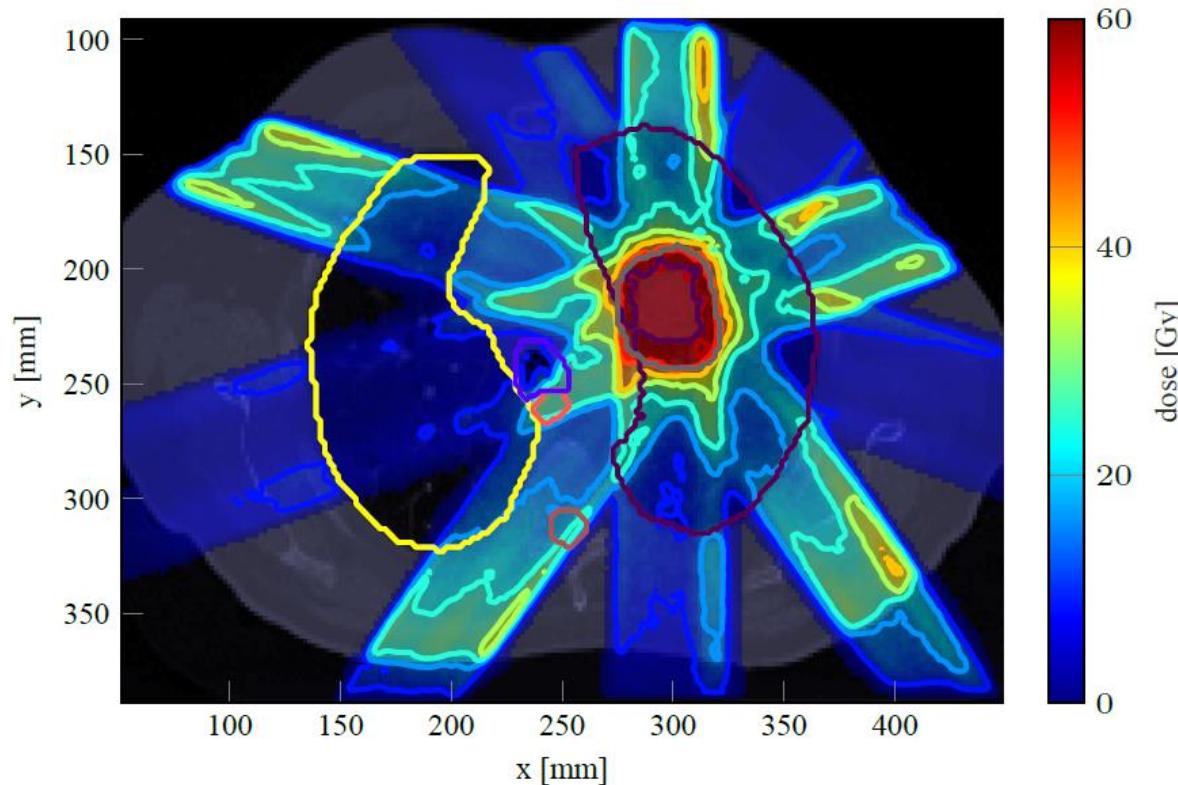
Developments for the last release: 2.10.1 “Blaise” (shown at the Workshop in 2021)

- Bug fixes, additional configuration parameters, etc.
- Independent grids (dose, CT, optimization)
- Extendable object-oriented optimization interface
 - Simple addition of optimizers and objectives / constraints
- Preliminary interfaces to Monte Carlo engines
 - Photons: ompMC - github.com/edoerner/ompMC
 - Protons: MCsquare - openmcsquare.org
- Automated testing (now GitHub Actions)

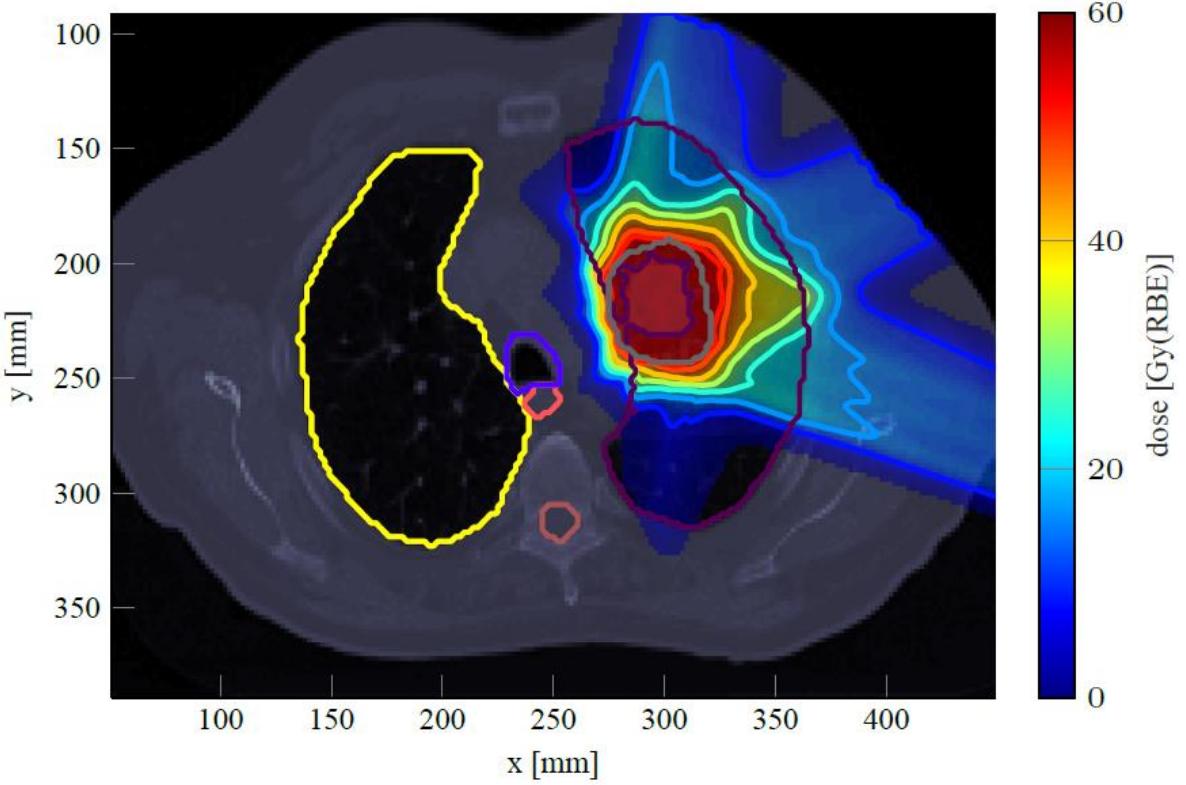


Proton plan with MCsquare beamlet computation
2e4 histories/beamlet, 4689 beamlets
120 min at (2.5mm)³ resolution

matRad 2.10.1: Novel developments – Basic open-source Monte Carlo Interfaces



- Photon plan with ompMC
- 1e5 histories/beamlet, 654 beamlets
- 25 min at $(2.5\text{mm})^3$ resolution

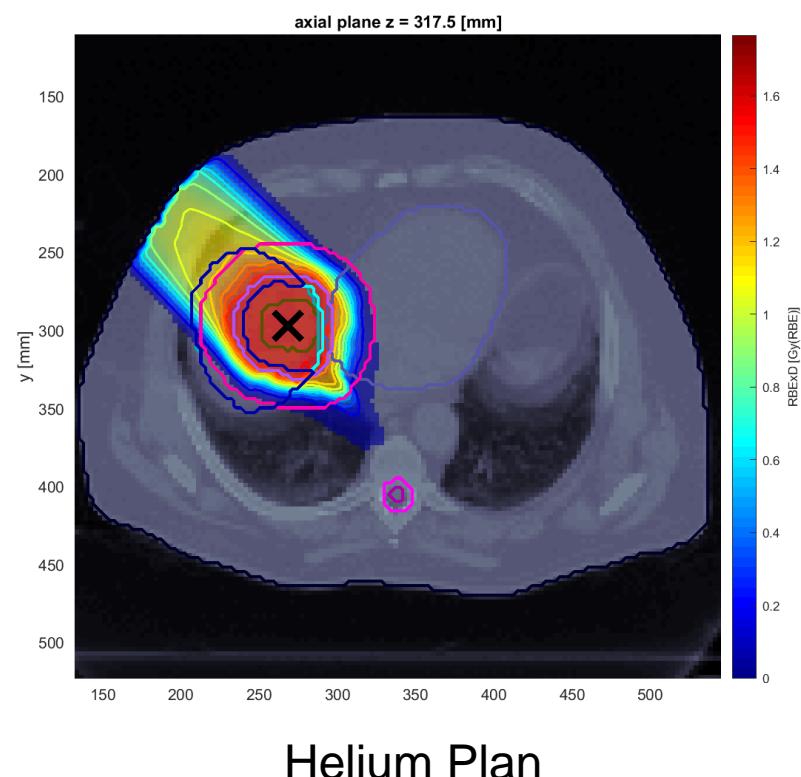


- Proton plan with MCsquare
- 2e4 histories/beamlet, 4689 beamlets
- 120 min at $(2.5\text{mm})^3$ resolution

Evaluated on Desktop PC, i7-6700 @ 3.4 GHz (4 cores +HT)

Post 2020 Development & Research Branches (shown at the Workshop in 2021)

- Helium base data (physical & biological)*
-dev
- 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 (e.g., TOPAS)*
-dev_MonteCarlo
- External contributions:
 - VMAT -dev_VMAT (E. Christiansen)
 - optimization -dev_exactOpt (S. van de Water)



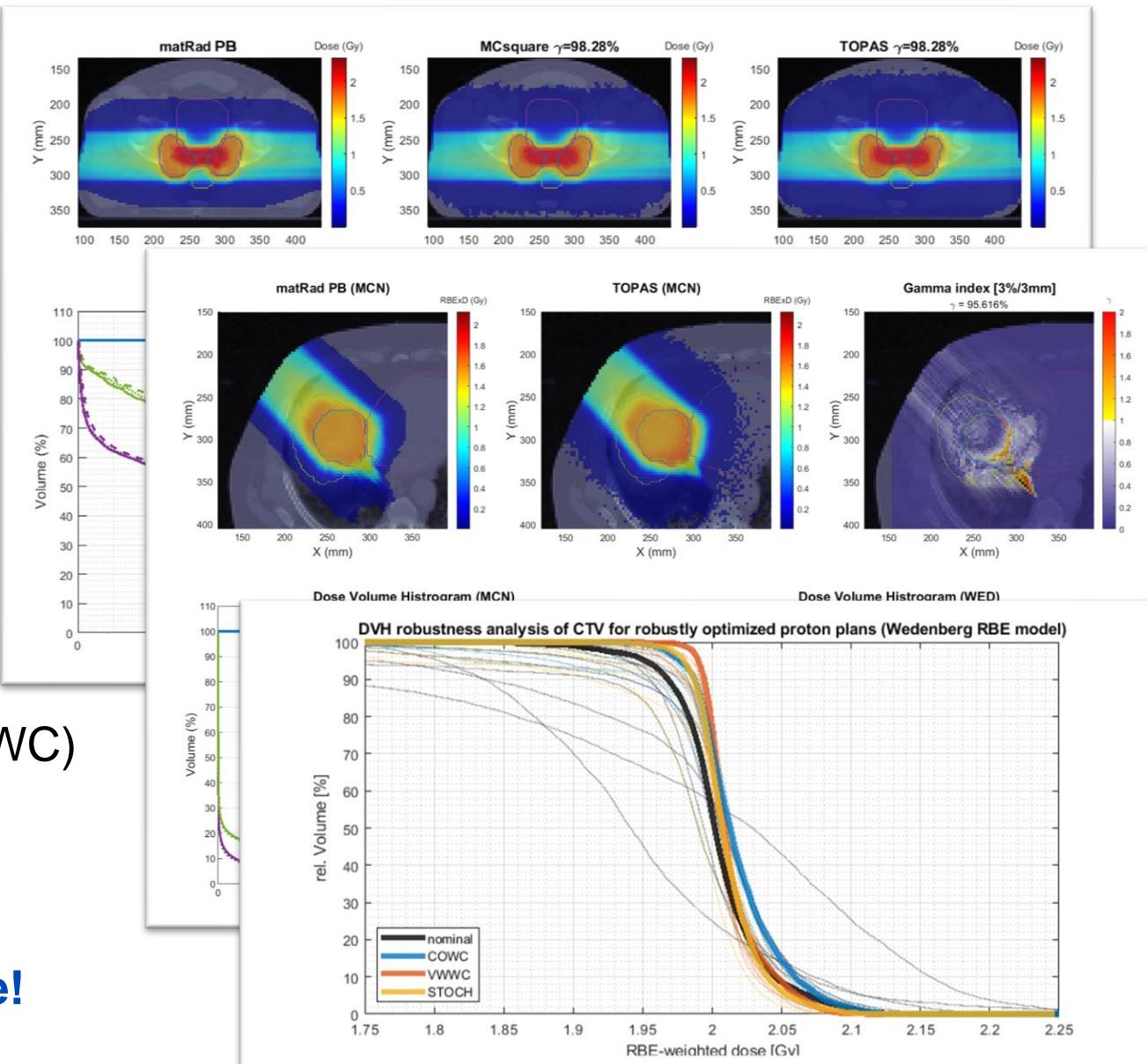
→ Sustainable research software call by (German Research Foundation, DFG)

→ Funding beginning 2022 enabled us to work towards release of v3 (*)

Next Release matRad v3.0.0 – Main features

- Improved Monte Carlo dose calculation interface
 - TOPAS + MCsquare (extendable)
 - Protons & Ions (He + C)
- Variable (Proton) RBE-models
 - LET-dependent models
 - Tabulated RBE models
- Robust Optimization methods
 - Worst-case optimization (COWC, OWC, VWWC)
 - Probabilistic optimization
 - Robustness Analysis

→ Best Feature: Extend & Combine all of those!

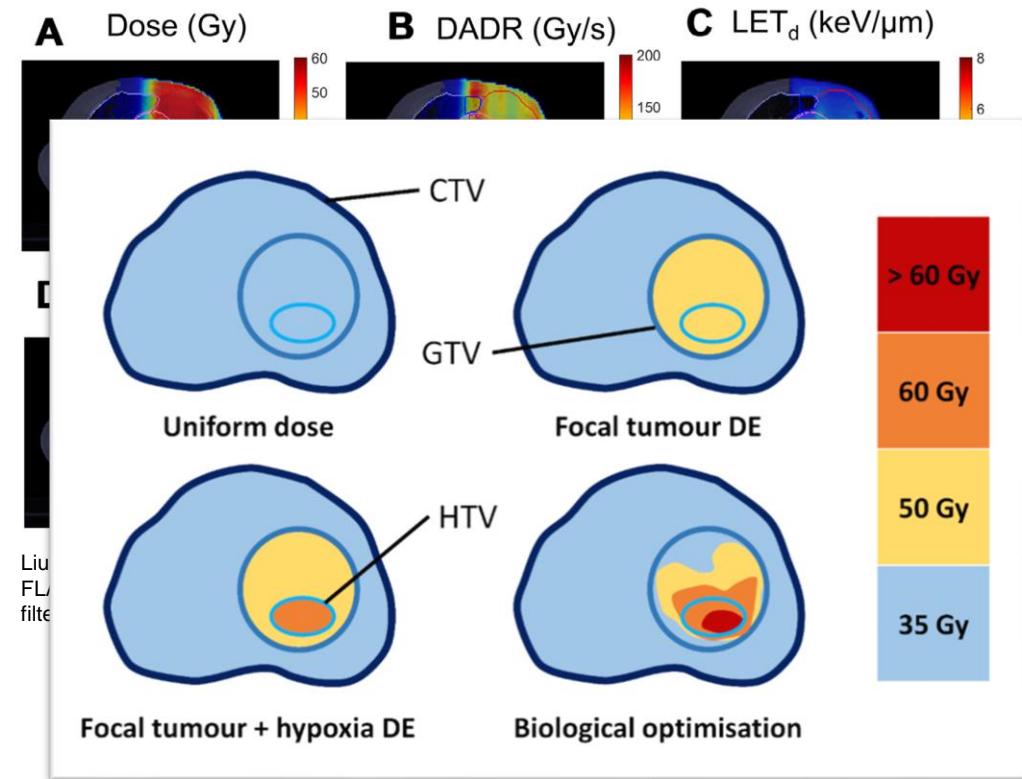


Release Candidate: <https://github.com/e0404/matRad/tree/rc/v3.0.0>

Current / Future Plans: Going further beyond dose?

Current research on planning with alternative quantities

- FLASH – different dose-rate concepts (DADR, IDR, PBSDR, etc.)
- Reduce sensitiveness to RBE (models)
 - LET-based optimization (LET minimization, dirty dose, etc.)
 - Micro- and nanodosimetric quantities
- Biology-guided planning
 - “dose painting” based on functional imaging
 - (voxel-based) NTCP optimization



Her EJ, Haworth A, Sun Y, et al. Biologically Targeted Radiation Therapy: Incorporating Patient-Specific Hypoxia Data Derived from Quantitative Magnetic Resonance Imaging. *Cancers*. 2021;13(19):4897. doi:10.3390/cancers13194897

→ matRad facilitates the addition of new quantities q for dose calculation & optimization

- data-structure for dose-like containers/matrices
- Framework for defining new objectives $f(q)$ and constraints $c(q)$
- Framework chaining fluence to the respective quantity & objectives: $\frac{df}{dw} = \frac{df}{dq} \frac{dq}{dw}$

Current / Future Plans: What about Python? – pyRadPlan

- Bidirectional Python interface to be published with next subrelease of matRad (probably v3.1)

Call matRad algorithms from Python



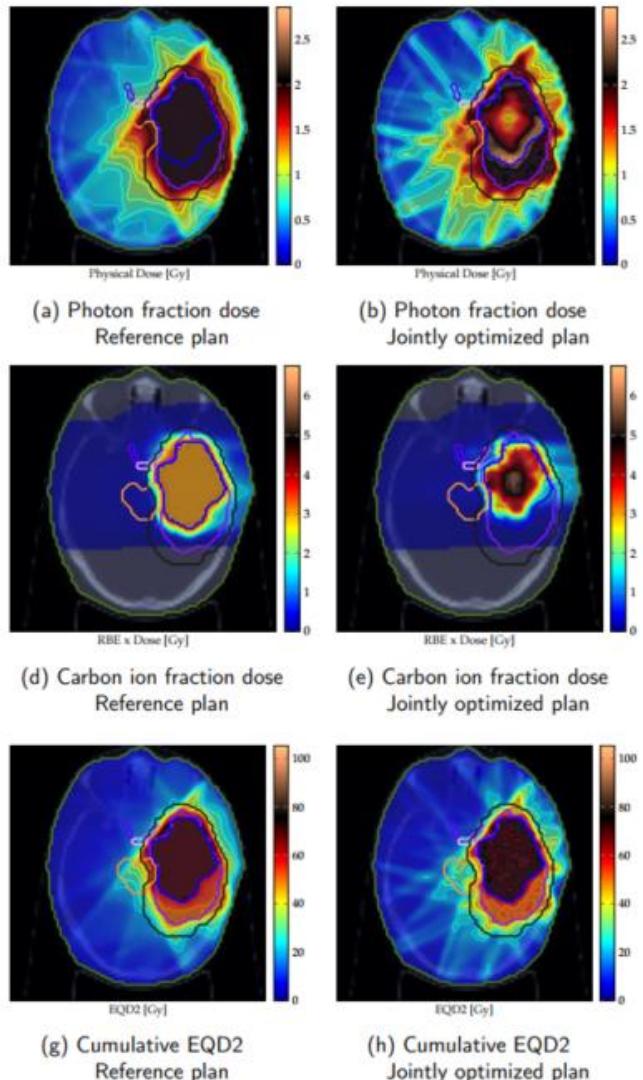
Call Python algorithms from matRad

- matRad run via MATLAB Engine API for Python, oct2py (Octave), or as compiled Python package
- Designed to work on the high-level API of matRad (geometry generation, dose calculation, optimization/superiorization, analysis)
- pyRadPlan will also support Python native planning
 - basic geometry generation possible
 - optimization reimplemented for physical dose & constant RBE using scipy solvers or cyipopt
 - Already used in an internal research project for ML-NTCP-based optimization
 - Exploit Python's flexibility for more performance



Example Research Projects

Example Project: Joint mixed-modality planning (A. Bennan, R. Cristoforetti)



- Conventional combined modality treatments are individually optimized and summed up based on RBE weighted dose
→ Not fully exploiting the biological potential
- Simultaneous optimization of the *total treatment effect* based on the LQ-Model:

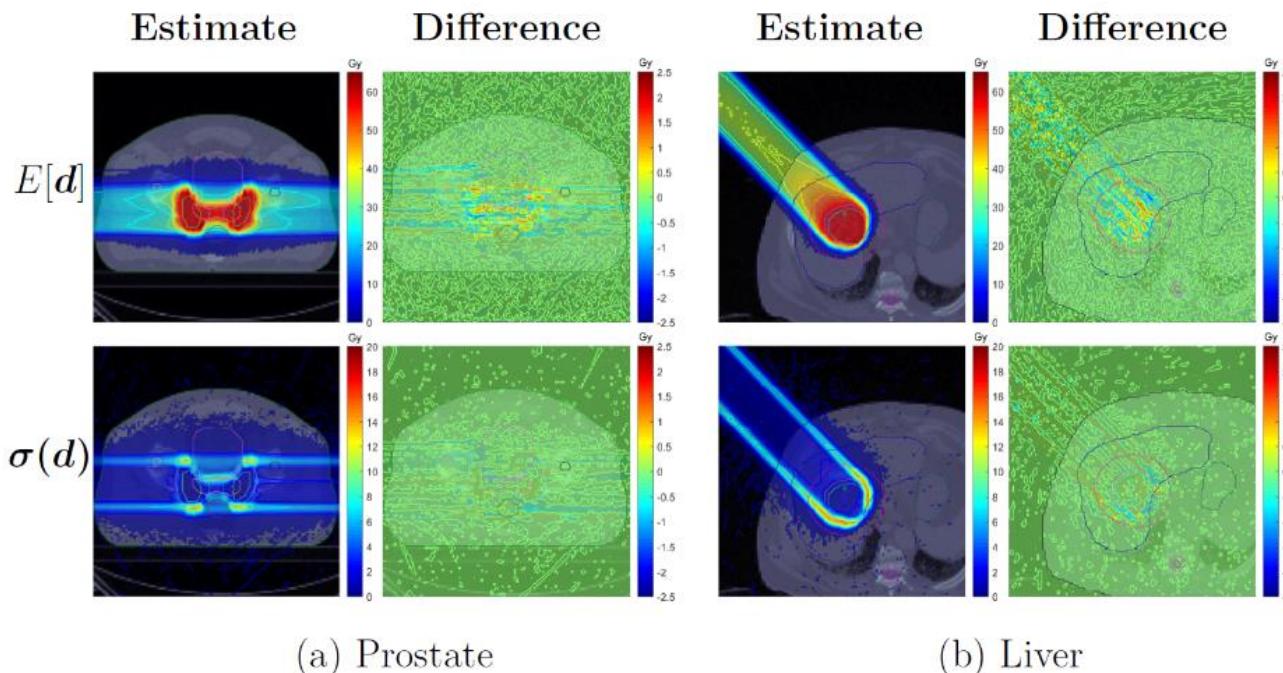
$$\varepsilon = -\ln(S) = \underbrace{n_\gamma(\alpha_\gamma d_\gamma + \beta_\gamma d_\gamma^2)}_{\text{photons}} + \underbrace{n_c(\alpha_c d_c + \beta_c d_c^2)}_{\text{carbon ions}}$$

- **Possibilities:**
 - Exploiting radiobiological properties of the modalities on a voxel-basis
 - improving conformality of biological effect in the adjacent critical structures
 - Exploiting fractionation in mixed (e.g. infiltrated) tissues

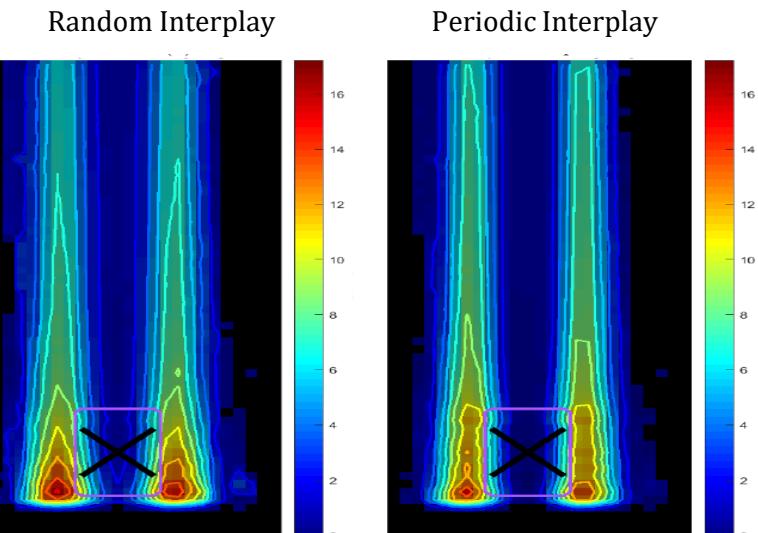
Bennan, A.B.A., Unkelbach, J., Wahl, N., Salome, P., Bangert, M., Joint optimization of photon – carbon ion treatments for Glioblastoma. *Int. J. Radiat. Oncol.*, 2021, <https://doi.org/10.1016/j.ijrobp.2021.05.126>

Example Project: MC Dose Uncertainty Quantification (P. Stammer, KIT / DKFZ)

- Avoid computation of explicit error / sample scenarios for robustness analysis and robust optimization with Monte Carlo codes
 - Re-weighting of histories of a single simulation for uncertainty estimation
 - Input Uncertainty modeling of static & time-dependent beam application/movement patterns using pencil beam correlations



Estimates of expected dose and variance using the reweighting approach and their difference to a reference [1]



Dose standard deviation in a waterbox for random vs. periodic movement pattern / interplay during treatment

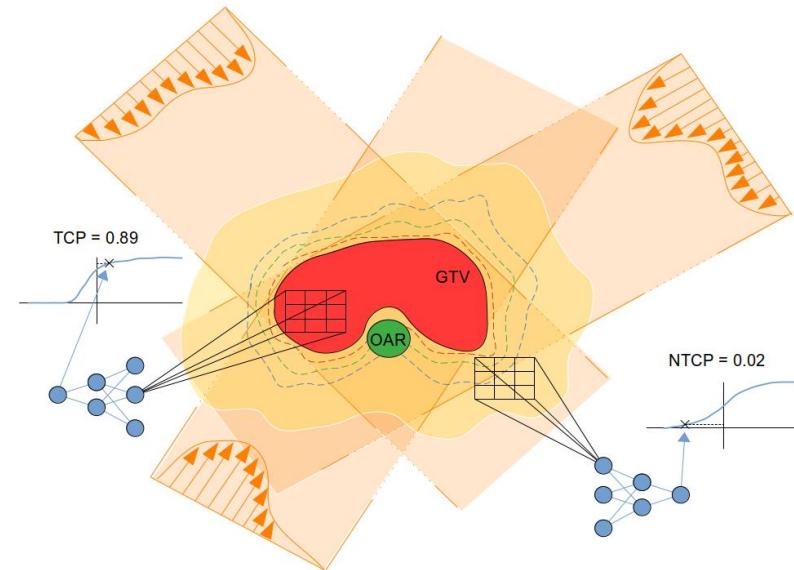
Other Involvements at KIT:

- Development of KiT-RT: A Kinetic Transport Solver for Radiation Therapy (<https://github.com/CSMMLab/KiT-RT>)
 - (Dynamical) low rank methods for more time and space efficient UQ in radiative transport

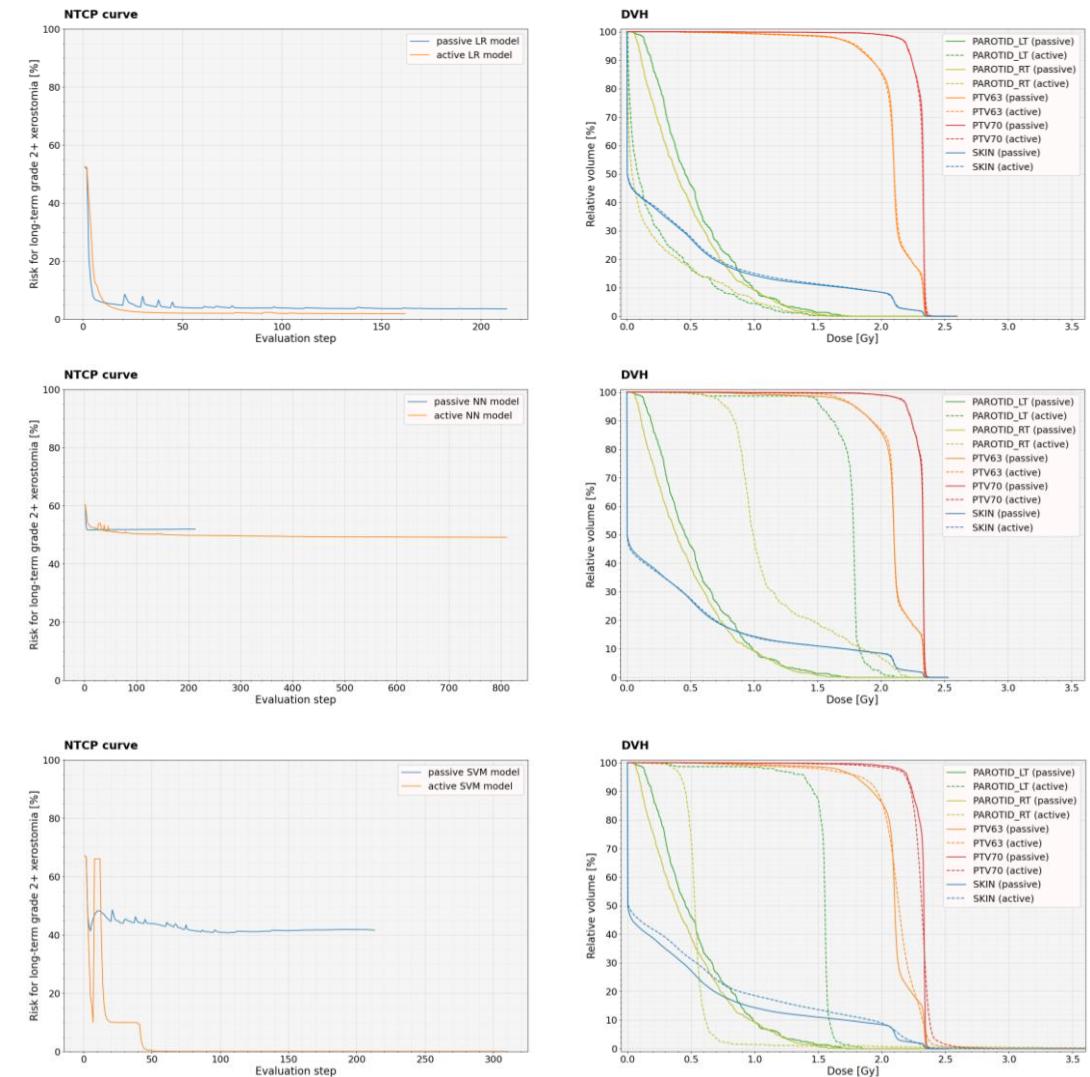
P. Stammer, L. Burigo, O. Jäkel, M. Frank, and N. Wahl, "Multivariate error modeling and uncertainty quantification using importance (re-)weighting for Monte Carlo simulations in particle transport," *JCP* 2023;473:111725, doi: [10.1016/j.jcp.2022.111725](https://doi.org/10.1016/j.jcp.2022.111725).

P. Stammer, L. Burigo, O. Jäkel, M. Frank, and N. Wahl, "Efficient uncertainty quantification for Monte Carlo dose calculations using importance (re-)weighting," *Phys Med Biol*, 2021;66(20):205003, doi: [10.1088/1361-6560/ac287f](https://doi.org/10.1088/1361-6560/ac287f)

Example Project: Inverse Planning using Machine Learning Outcome Prediction Models (T. Ortkamp, KIT / DKFZ)



- Explicit integration of machine learning (N)TCP models into the inverse planning problem of radiotherapy optimization
 - Investigation of model stability, i.e. generalization error and uncertainty
 - Efficient fast forward and gradient computations/approximations
- Development of strategies to enable custom model integration and to facilitate the generation of clinically acceptable treatment plans
 - Python-based advanced numerical nonlinear optimization for radiotherapy (pyanno4rt)
 - Software prototype for machine learning model-based photon & proton treatment plan optimization



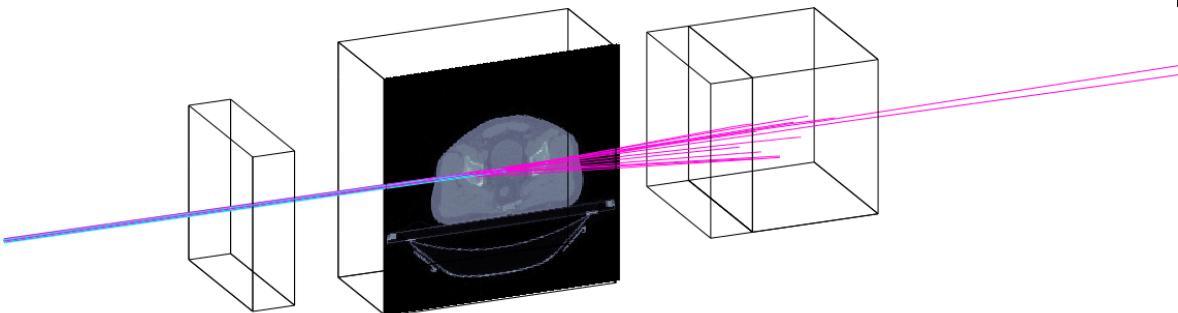
Logistic regression

Neural network

Support vector machine

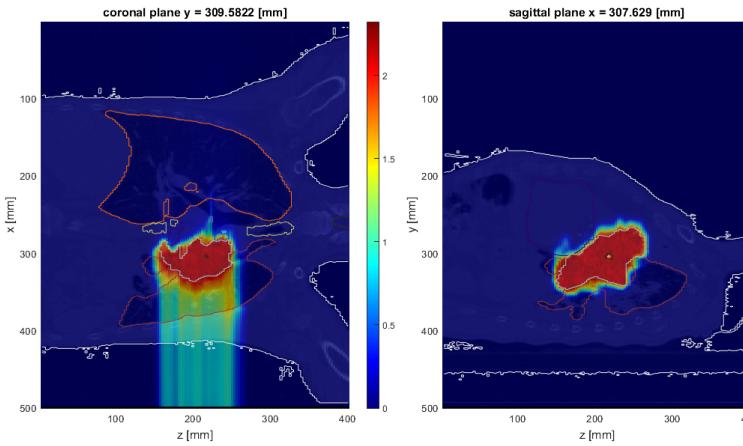
Example Project (Following Talk by Jennifer): Developing a HELium Imaging Oncology Scanner for Range Guided Radiotherapy (RGRT)

- (Theoretically) possible to accelerate a mixed beam of carbon / helium ions
- Helium v. carbon range: $R_{He} = 3R_C$
→ He-monitoring?
- How to use in treatment planning / decision making / intervention?

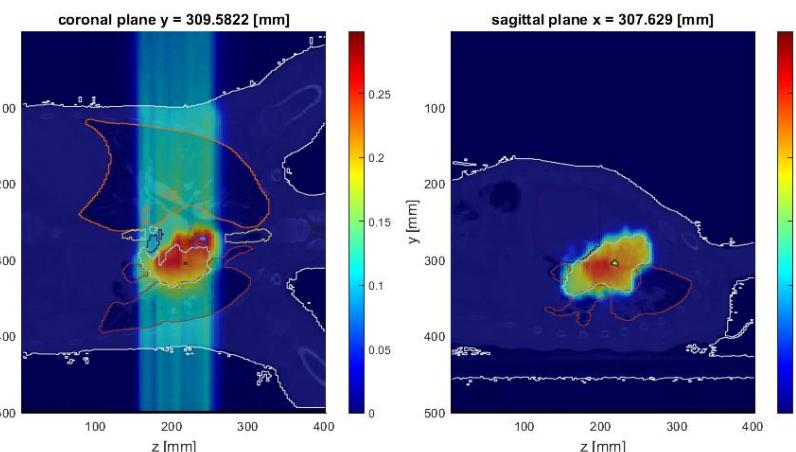


DFG
Deutsche
Forschungsgemeinschaft

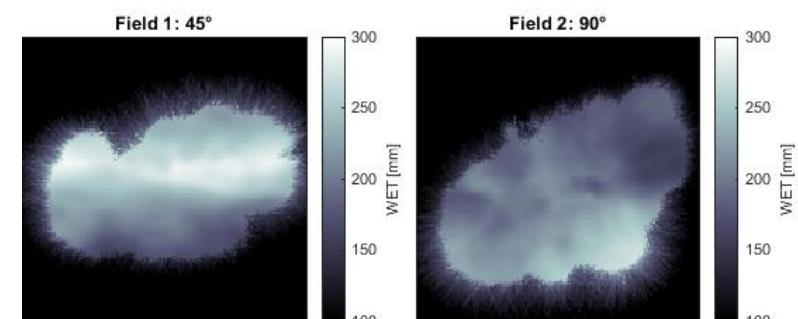
Full dose



He dose → 3x carbon range



WEPL image



Example Part I

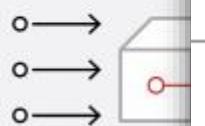
Nanodosimetry

Collaborators

- Nanodosimetry
→ improved

PARTICLE BEAM

Low-LET



High-LET



Goal: Facilitate

```
%> %% Cluster Dose Setup
pln.propDoseCalc.calcClusterDose = true;
pln.propDoseCalc.clusterDoseIP = 'F4';

%> %% generate steering file
stf = matRad_generateStf(ct,cst,pln);

%> %% dose calculation
dij = matRad_calcDose(ct,cst,stf,pln);

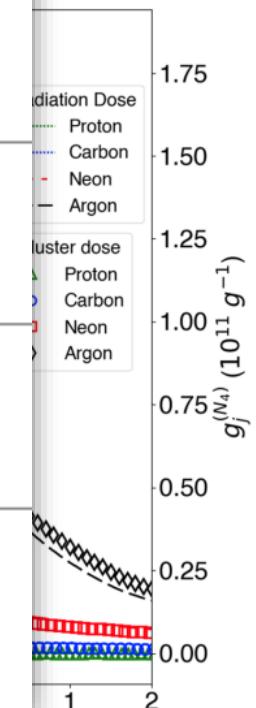
%> %% standard optimization without cluster dose
resultGUI = matRad_fluenceOptimization(dij,cst,pln);

%> %% start gui for visualization of result
matRadGUI

%> %% cluster dose optimization
cst{16,6}{2} = ClusterDoseObjectives.matRad_ClusterDoseVariance(1e6);
resultGUI2 = matRad_fluenceOptimization(dij,cst,pln);
```

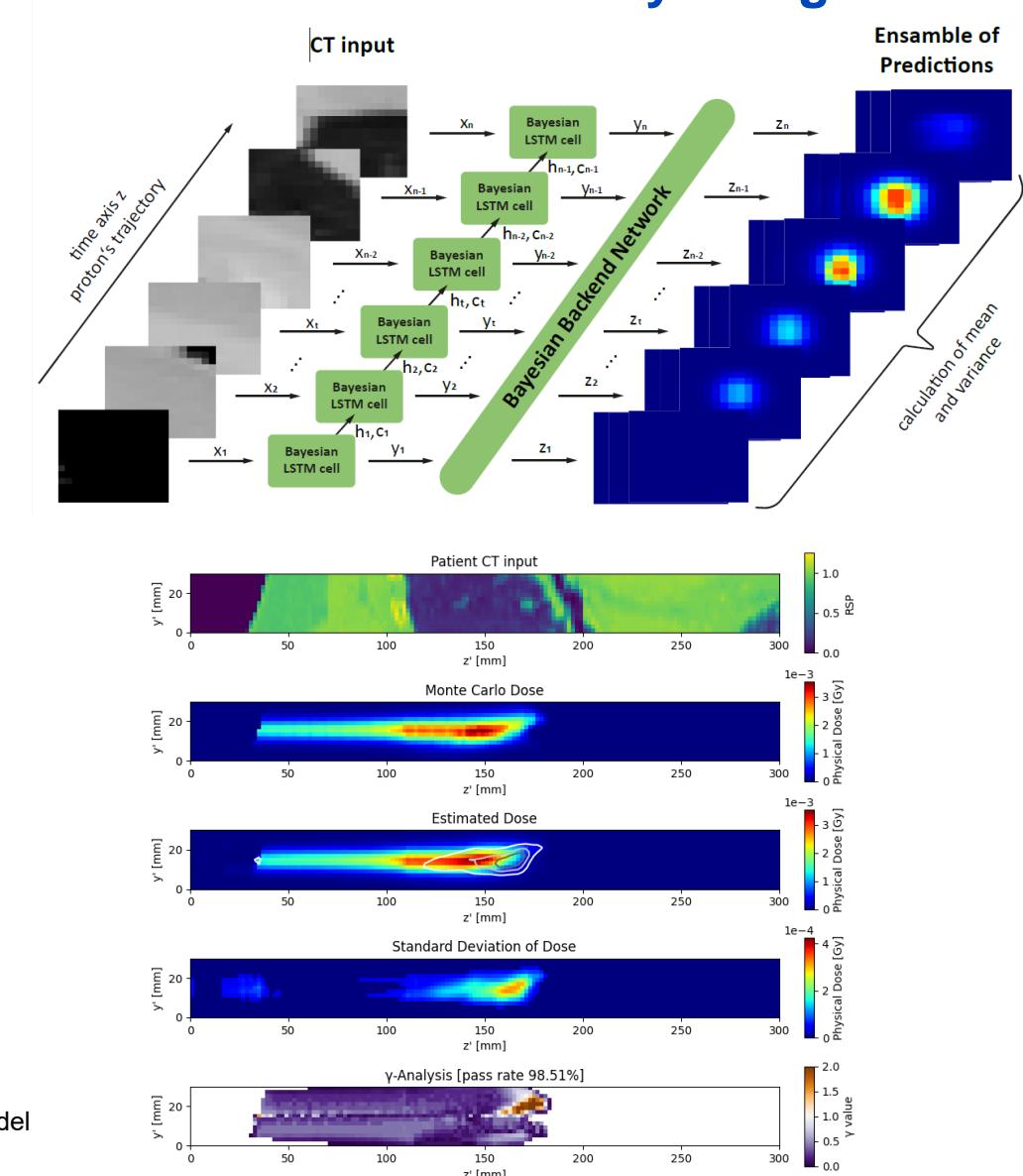
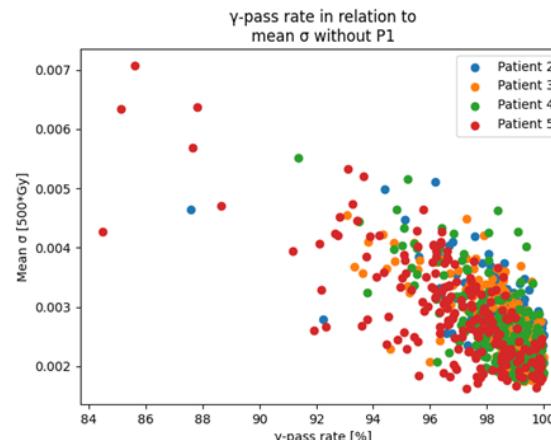
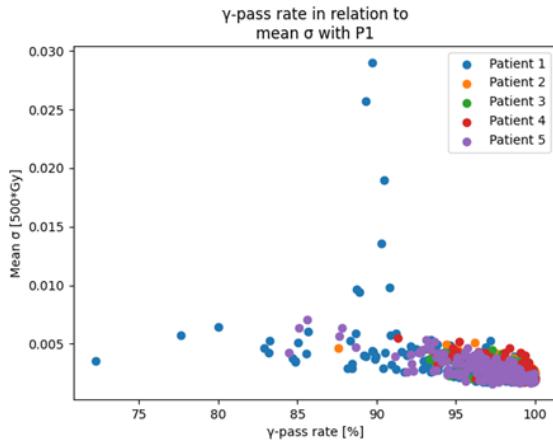


constitutes
health



E040-4: BayesDose: Comprehensive proton dose prediction with model uncertainty using Bayesian LSTMs (L. Voss)

- Need for fast and accurate dose calculation techniques with quantifiably uncertainty estimation in proton therapy
- Using a Bayesian Approach to prior LSTM research, the model achieves fast predictions with equal quality compared to the prior deterministic model
- The obtained prediction's standard deviation correlates well with dosimetric inaccuracy
 - Models like BayesDose could support decision making and quality assurance in clinic



L. Voss, A. Neishabouri, T. Ortakamp, A. Mairani, and N. Wahl, "BayesDose: Comprehensive proton dose prediction with model uncertainty using Bayesian LSTMs." arXiv 2023. doi: [10.48550/arXiv.2307.01151](https://arxiv.org/abs/2307.01151).

How to get going with matRad?

1. Go to our page on GitHub: www.matRad.org
2. Download the Code, or even better: Familiarize with **git** and Clone
3. Checkout the UI & the code
 - `matRadGUI.m` & `matRad.m`
 - many **examples** in the `./examples/` folder
 - Wiki on GitHub: <https://github.com/e0404/matRad/wiki>
4. Profit (and contribute!)
→ Never hesitate to contact us!

How to get going with matRad – Examples

1. How to create your own phantom?
2. How to setup a photon treatment plan?
3. How to perform a direct aperture optimization for photons?
4. How to perform a Monte Carlo dose calculation for photon
5. How to create a treatment plan for protons?
6. How to manipulate stopping powers or to simulate an isocenter shift?
7. How to define carbon ion treatment plan?
8. How to perform robust optimization?

Example: Generate your own phantom geometry

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%%%%%%%

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%%%%%%%

In this example we will show (i) how to create arbitrary ct data (resolution, ct numbers) (ii) how to create a cst structure containing the volume of interests of the phantom (iii) generate a treatment plan for this phantom

`clc, clear, close all`

Create a CT image series

```
xDim = 200;
yDim = 200;
zDim = 50;

ct.cubeDim      = [xDim yDim zDim];
ct.resolution.x = 2;
ct.resolution.y = 2;
ct.resolution.z = 3;
ct.numOfCtScen = 1;

% create an ct image series with zeros - it will be filled later
ct.cubeHU(1) = ones(ct.cubeDim) * -1000; % assign HU of Air
```



GitHub



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/Ipopt>

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 will be available in matRad 3