

# SlicerRT

## Image-guided radiation therapy research toolkit for 3D Slicer

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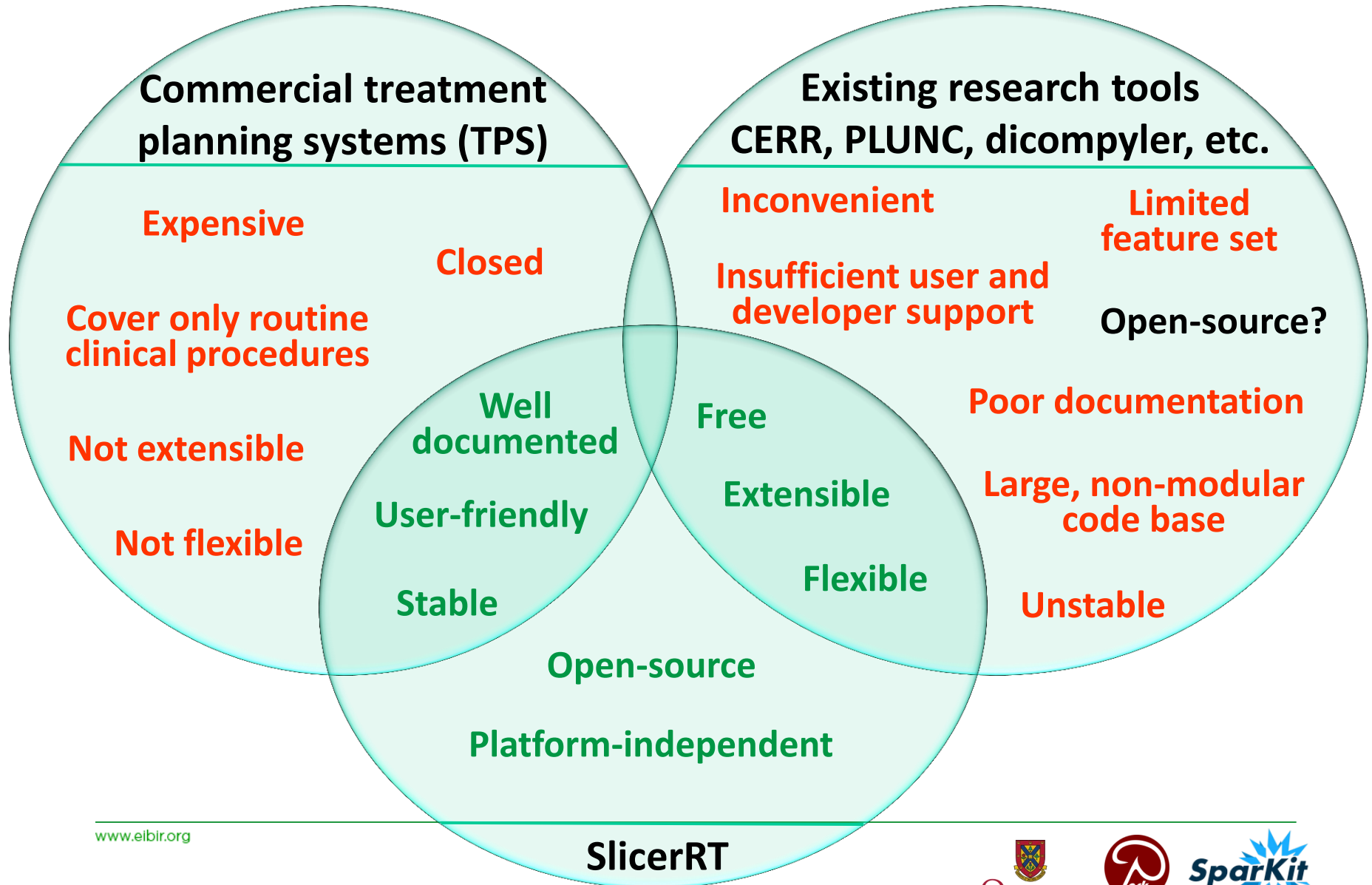


# SparKit project overview

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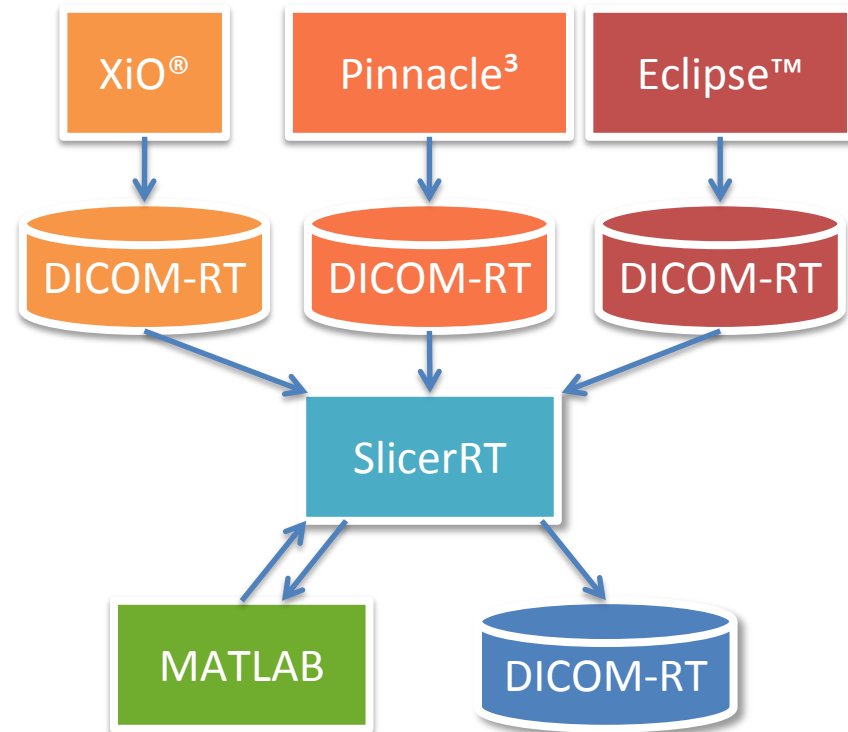
- Goal: provide open-source platform for translational clinical research, mainly cancer
- Themes:
  - **SlicerRT: radiotherapy toolkit for 3D Slicer**
  - SlicerIGT: Image-guided therapy with 3D Slicer
- Funding by Cancer Care Ontario till 2016
- PI & co-PIs: Gabor Fichtinger (Queen's), David Jaffray (Toronto UHN), Terry Peters (Robarts)

# Motivation behind SlicerRT



# Development principles

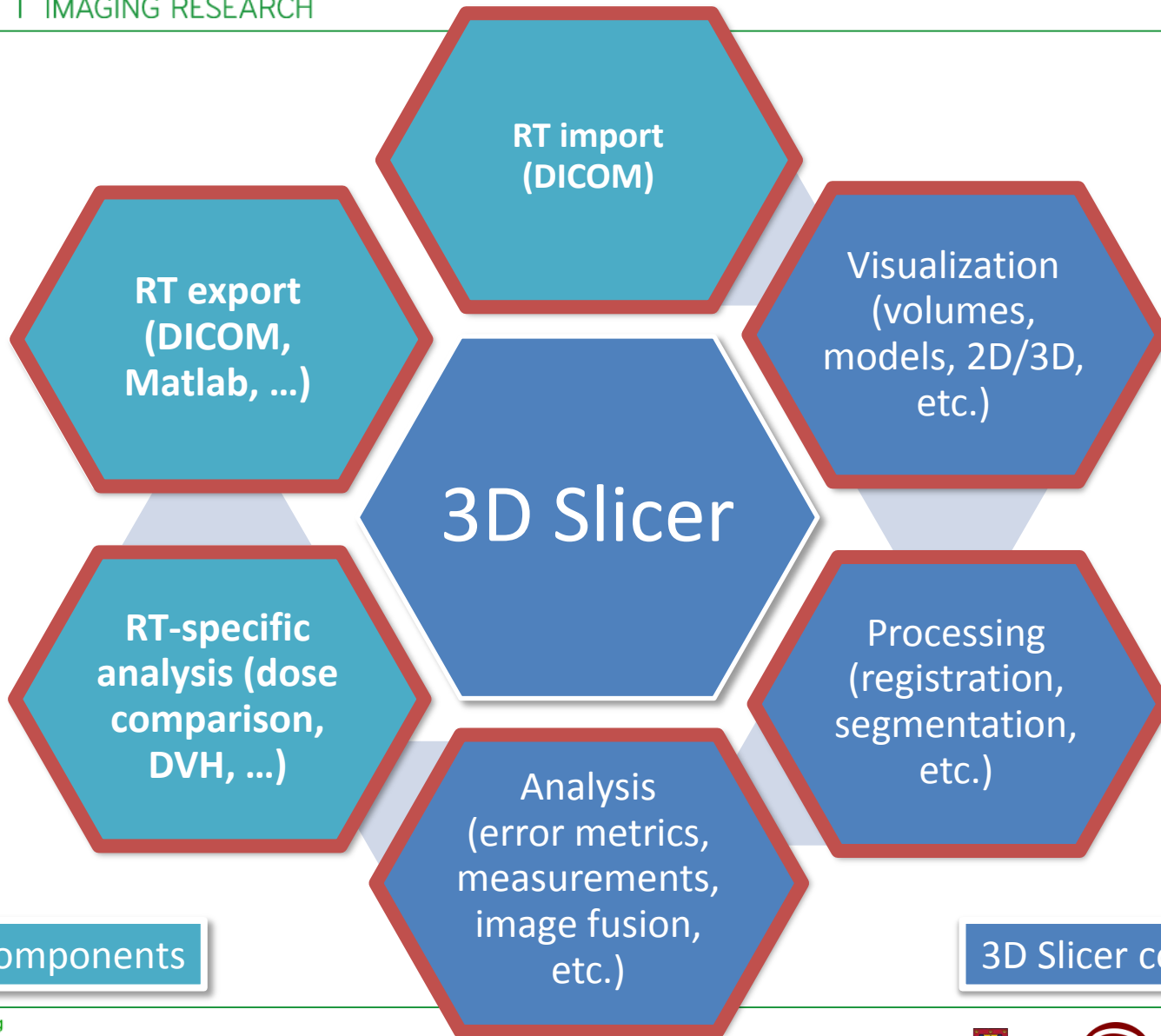
- Leverage existing tools and parallel efforts: 3D Slicer<sup>1</sup>, Plastimatch<sup>2</sup>
- “Hub” for RT data analysis and comparison
- Cover most common RT research workflows
- Free, open-source license (BSD)
- Open to integrate algorithms
- Extensive documentation



<sup>1</sup> S. Pieper, M. Halle, and R. Kikinis, 3D SLICER. *Proceedings of the 1st IEEE International Symposium on Biomedical Imaging: From Nano to Macro* (Brigham and Women’s Hospital, Boston, MA, 2004), pp. 632–635.

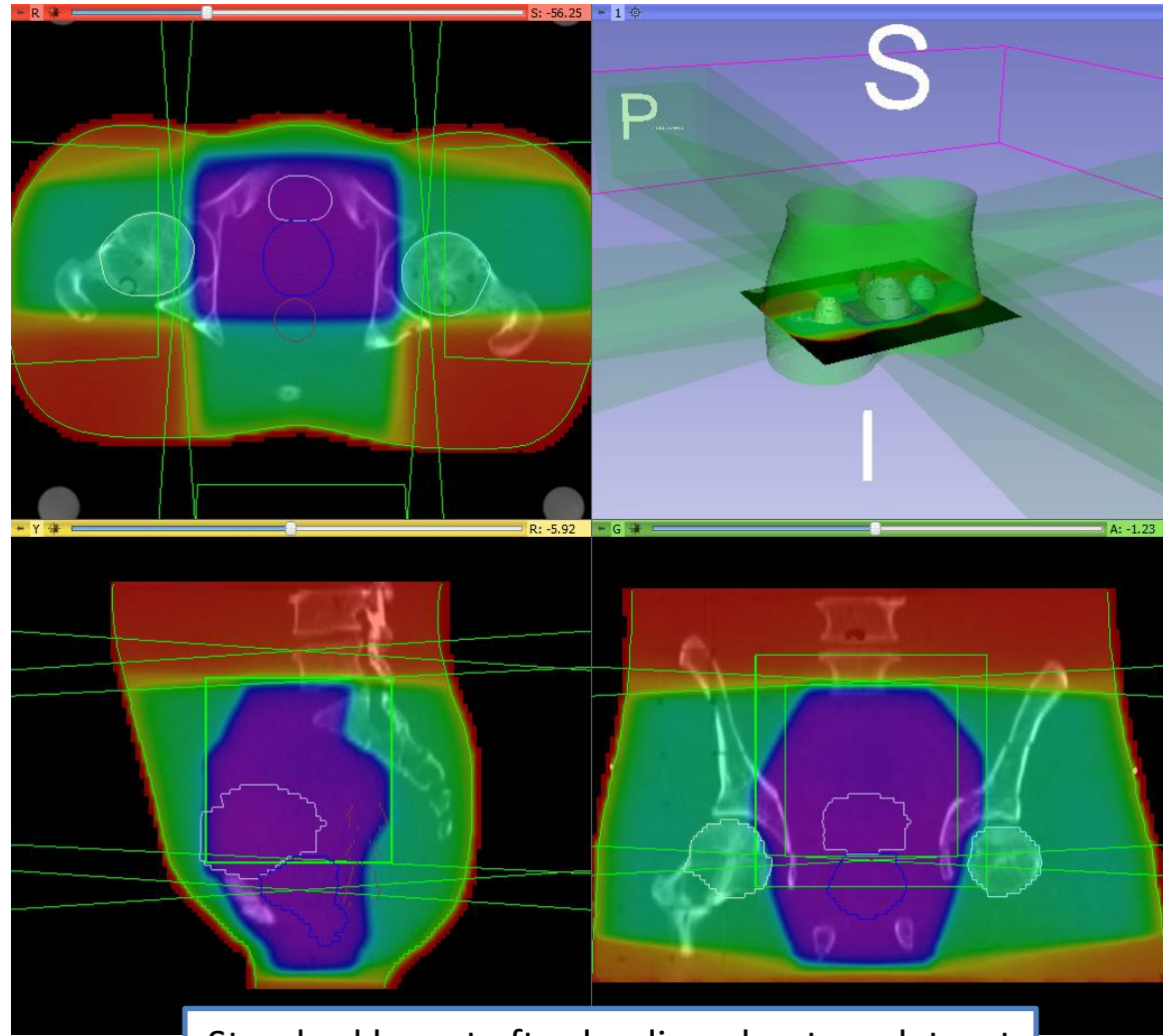
<sup>2</sup> G. C. Sharp, R. Li, J. Wolfgang, G. Chen, M. Peroni, M. F. Spadea, S. Mori, J. Zhang, J. Shackelford, and N. Kandasamy, “Plastimatch: An open source software suite for radiotherapy image processing,” in *Proceedings of the XVIth International Conference on the Use of Computers in Radiotherapy (ICCR)* (Amsterdam, the Netherlands, 2010).

# General workflow overview



# DICOM-RT import

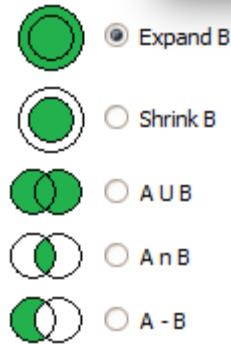
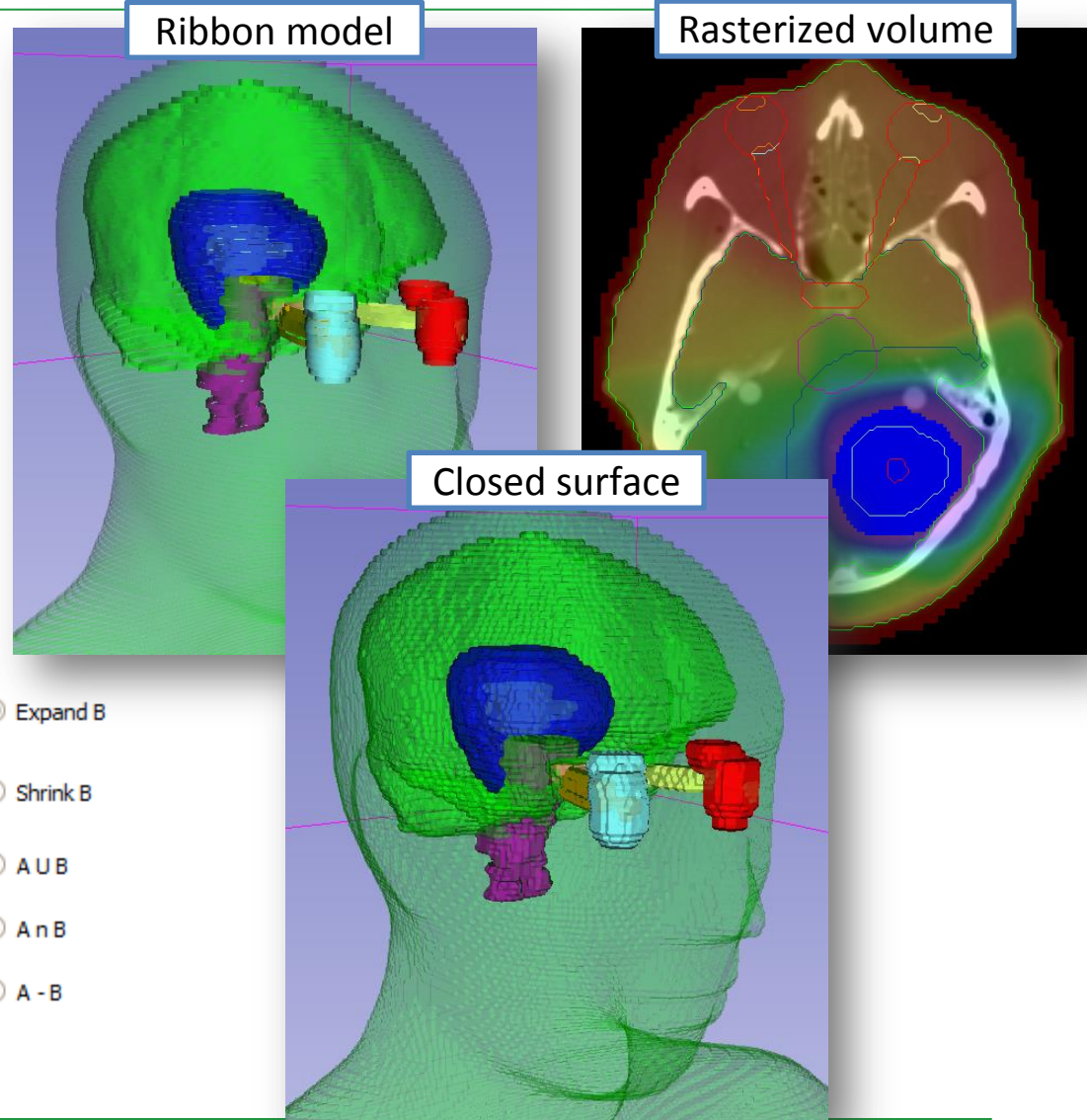
- Integrated into core DICOM import plugin mechanism
- Data is organized in a smart hierarchy
- Supported data types:
  - RT structure sets
    - Contours
    - Fiducial point
  - RT dose map
  - RT image
  - RT plan: isocenter, beams
  - Planning CT, MR, etc.



Standard layout after loading phantom dataset

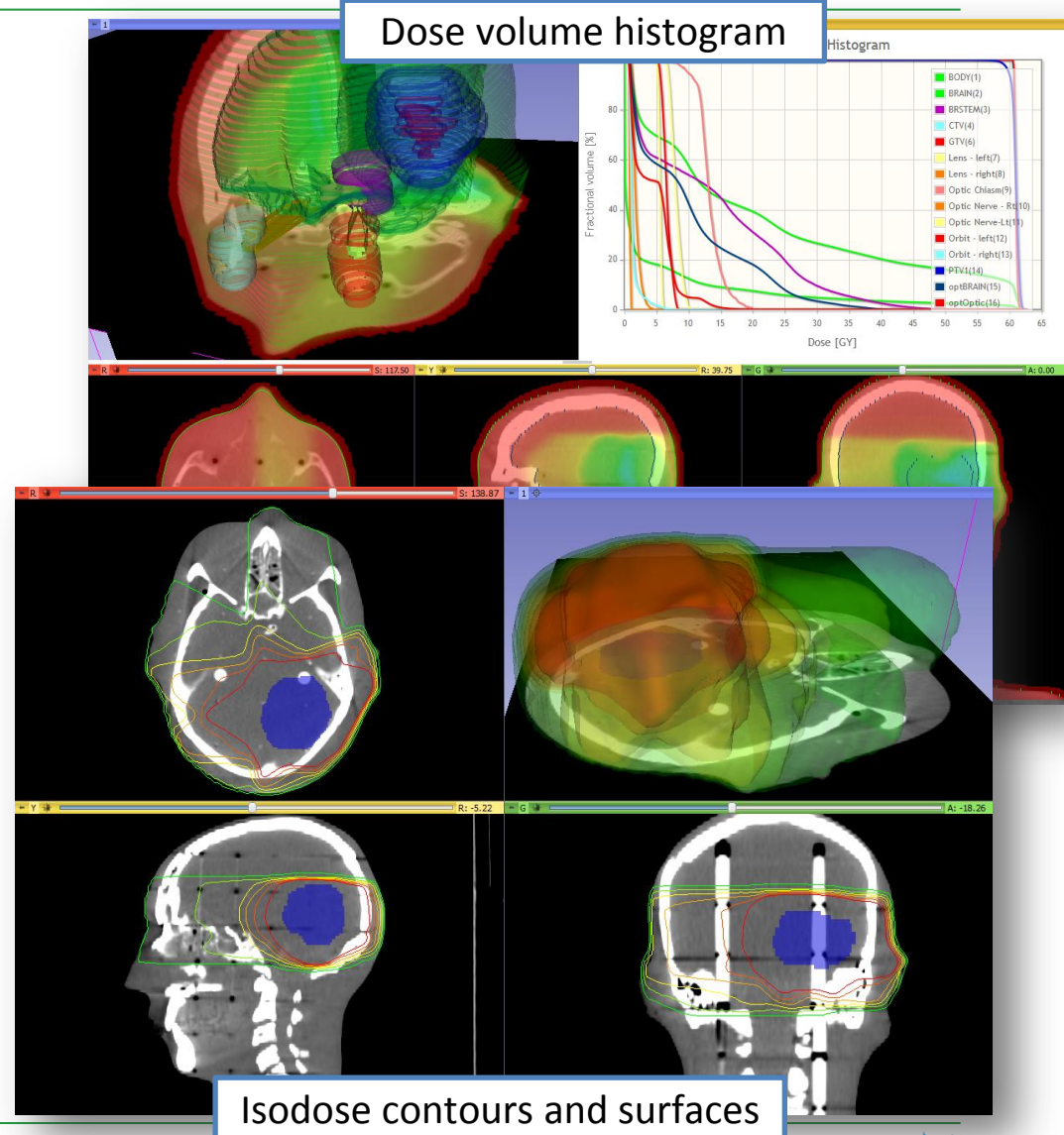
# Contour analysis

- Multiple representations (automatic conversion)
  - Ribbon model
  - Rasterized volume
  - Closed surface model
- Contour comparison
  - Dice coefficient
  - Hausdorff distance
- Contour morphology
  - Expand, shrink
  - Combine using logical operators



# Dose analysis

- Dose volume histogram (plot visualization + metrics)
- Dose accumulation
- Dose comparison (gamma)
- Isodose contours / surfaces
- Visualize deformation fields
- Proton dose computation
- Registration
  - BSpline registration
  - Landwarp registration





# Example use case

Use case: Evaluate the effectiveness of RT plan adaptation

- Import, load and visualize data
- Register day 1 CT with day 2 CT
  - Rigid
  - Bspline
- Resample day 2 dose using results
- Accumulate doses
- Compute and display DVH for all methods
- Compare DVH curves and metrics for target volume and organs at risk

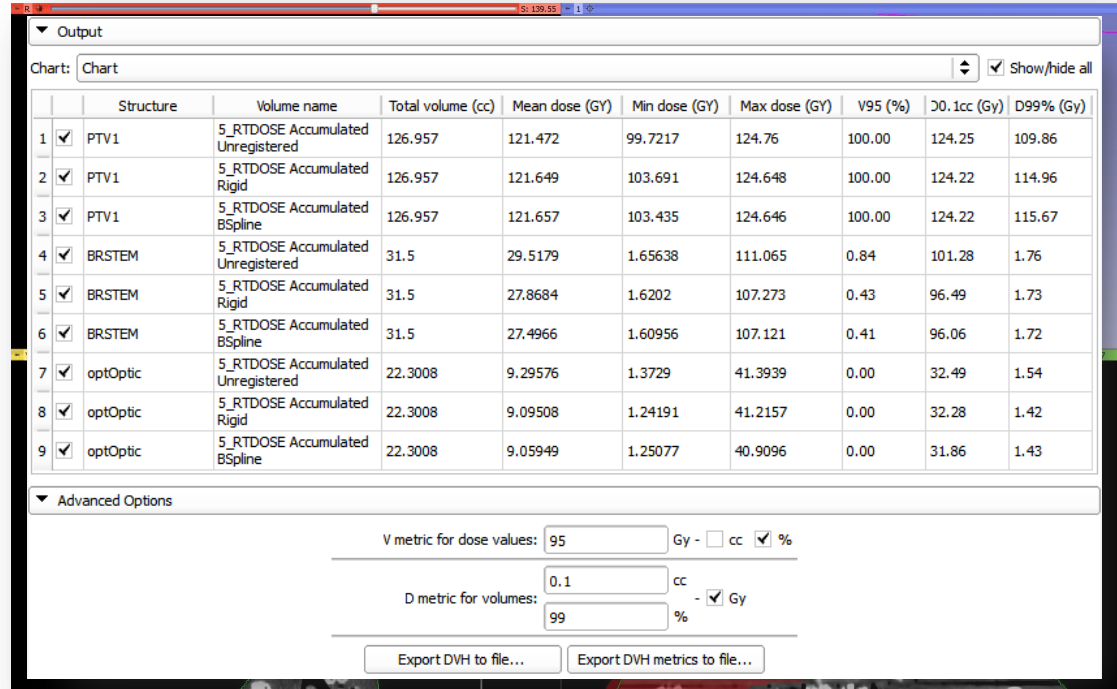


Chart: Chart Show/hide all

	Structure	Volume name	Total volume (cc)	Mean dose (Gy)	Min dose (Gy)	Max dose (Gy)	V95 (%)	D0.1cc (Gy)	D99% (Gy)	
1	<input checked="" type="checkbox"/>	PTV1	5_RTDOSE Accumulated Unregistered	126.957	121.472	99.7217	124.76	100.00	124.25	109.86
2	<input checked="" type="checkbox"/>	PTV1	5_RTDOSE Accumulated Rigid	126.957	121.649	103.691	124.648	100.00	124.22	114.96
3	<input checked="" type="checkbox"/>	PTV1	5_RTDOSE Accumulated BSpline	126.957	121.657	103.435	124.646	100.00	124.22	115.67
4	<input checked="" type="checkbox"/>	BRSTEM	5_RTDOSE Accumulated Unregistered	31.5	29.5179	1.65638	111.065	0.84	101.28	1.76
5	<input checked="" type="checkbox"/>	BRSTEM	5_RTDOSE Accumulated Rigid	31.5	27.8684	1.6202	107.273	0.43	96.49	1.73
6	<input checked="" type="checkbox"/>	BRSTEM	5_RTDOSE Accumulated BSpline	31.5	27.4966	1.60956	107.121	0.41	96.06	1.72
7	<input checked="" type="checkbox"/>	optOptic	5_RTDOSE Accumulated Unregistered	22.3008	9.29576	1.3729	41.3939	0.00	32.49	1.54
8	<input checked="" type="checkbox"/>	optOptic	5_RTDOSE Accumulated Rigid	22.3008	9.09508	1.24191	41.2157	0.00	32.28	1.42
9	<input checked="" type="checkbox"/>	optOptic	5_RTDOSE Accumulated BSpline	22.3008	9.05949	1.25077	40.9096	0.00	31.86	1.43

Advanced Options

V metric for dose values: 95 Gy -  cc  %

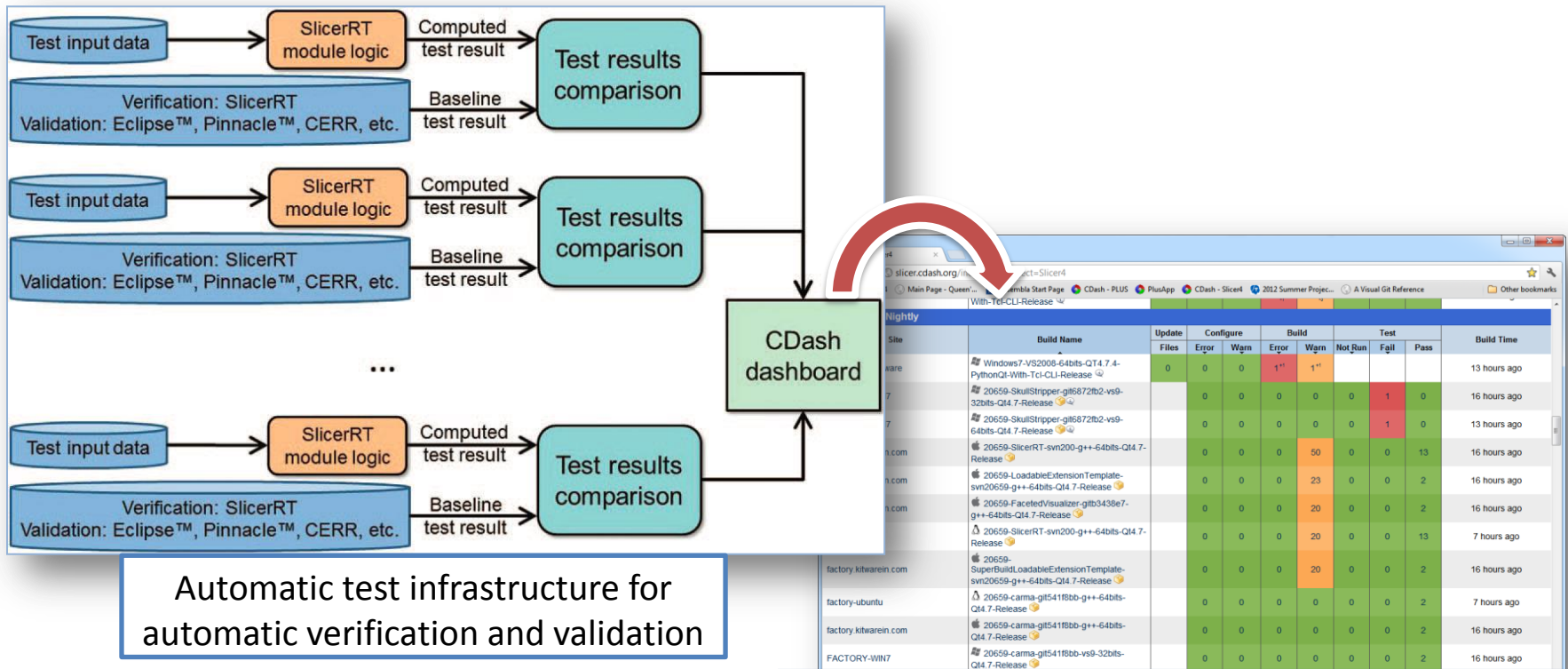
D metric for volumes: 0.1 cc -  Gy

99 %

Export DVH to file... Export DVH metrics to file...

# Software quality

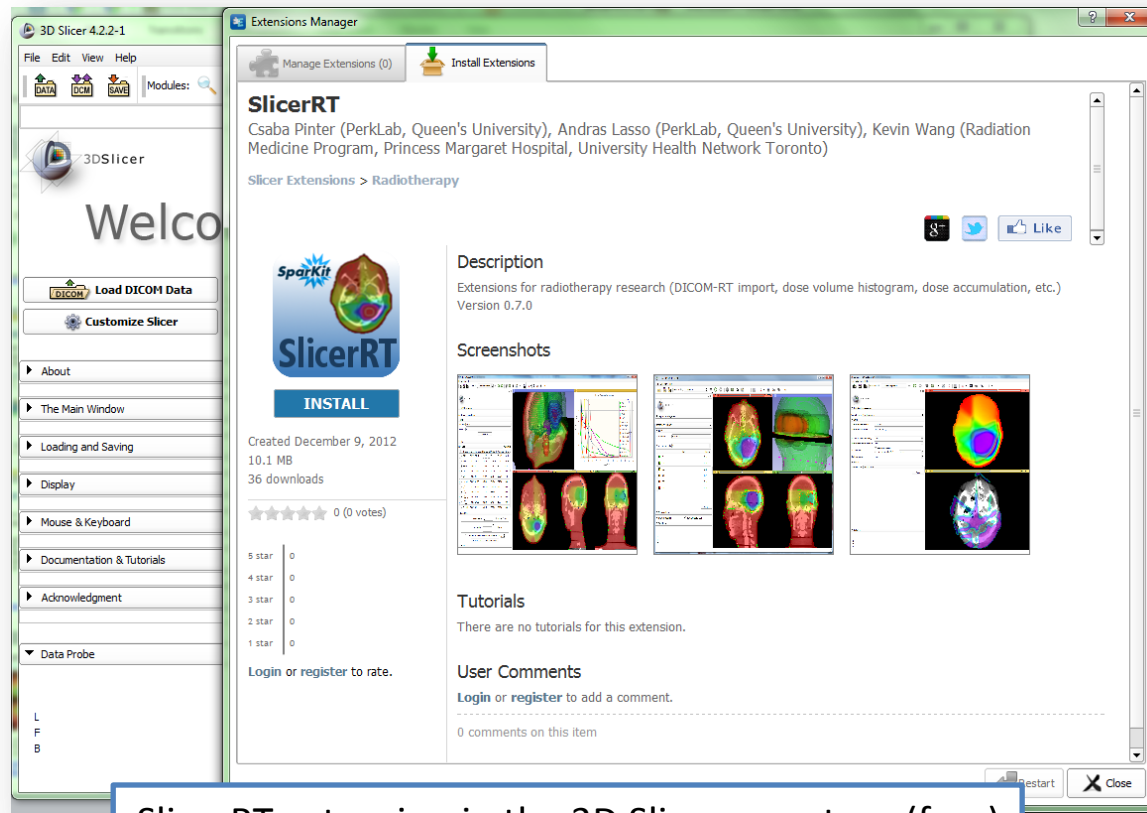
- Extensive automatic testing done on multiple platforms every night
- Validated against other software packages (Pinnacle, CERR, ...)



Test results reported to the web-based dashboard  
<http://slicer.cdash.org/index.php?project=Slicer4>

# Extension for 3D Slicer

- Collection of RT-specific modules, includes *Plaslimatch*
- Distributed as a 3D Slicer extension: can be downloaded, installed, upgraded using the extension manager in Slicer

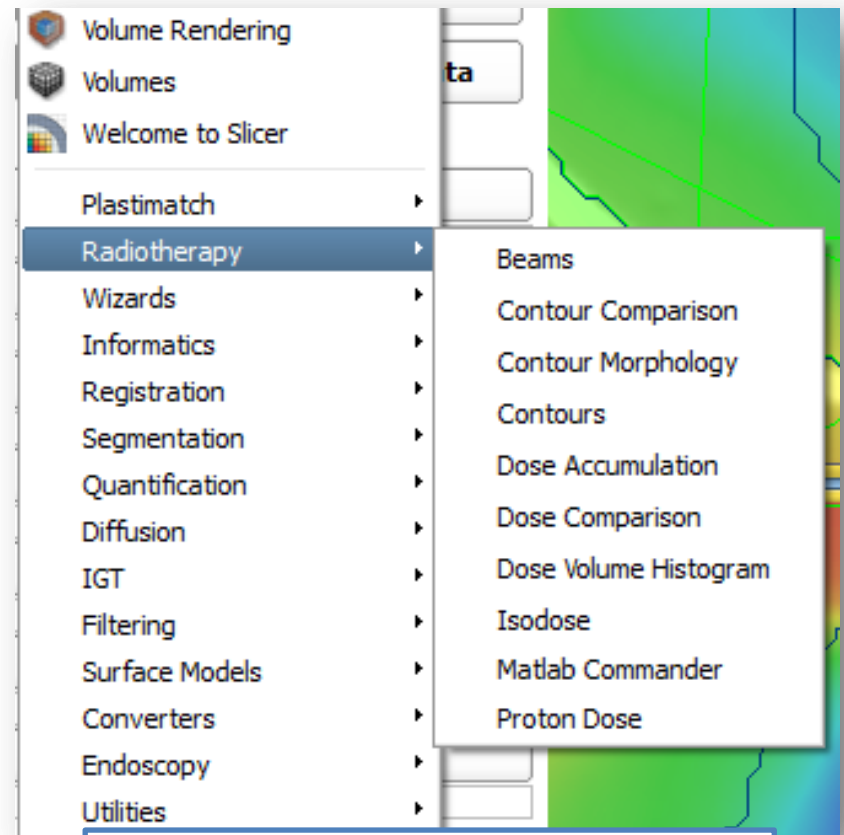


SlicerRT extension in the 3D Slicer app store (free)

# Next steps

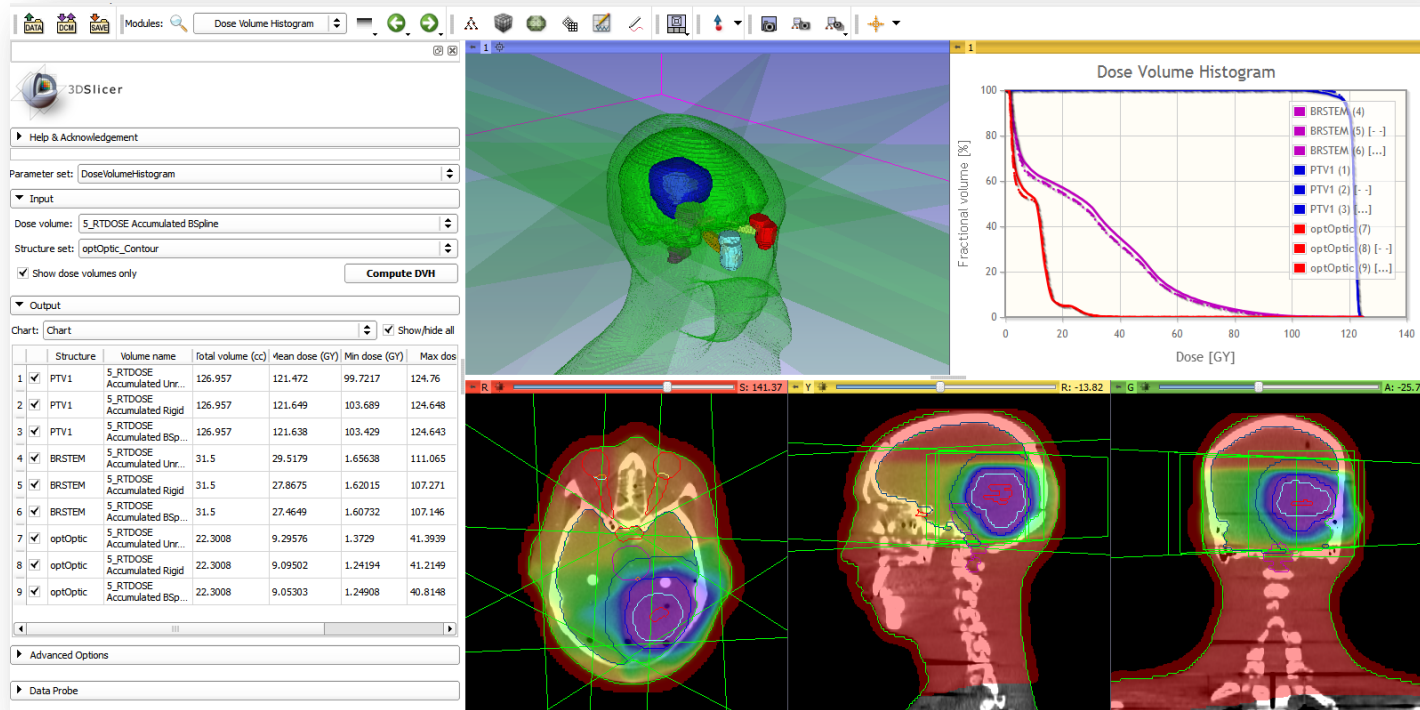
Enhancements planned for the next 6 months:

- DICOM-RT export
- Matlab bridge interface: execute Matlab functions from 3D Slicer
- Digitally reconstructed radiograph
- Support CERR PlanC format
- Scripting examples
- More testing and validation



Current radiotherapy modules  
(including work-in-progress modules)

Detailed plan: <https://www.assembla.com/spaces/slicerrt/tickets>



- Overview paper: Csaba Pinter, Andras Lasso, An Wang, David Jaffray, and Gabor Fichtinger, “SlicerRT: Radiation therapy research toolkit for 3D Slicer”, Med. Phys. 39 (10), October 2012
- Project homepage: <http://www.SlicerRT.org/>
- Contact: Csaba Pinter ([pinter@cs.queensu.ca](mailto:pinter@cs.queensu.ca))