

Accuracy. Ensured.

Monte Carlo Secondary Dose Check and Plan QA with SciMoCa TM



DISCOVER SCIMOCATM

Discover the power of Monte Carlo accuracy for secondary dose check and plan QA.

What is SciMoCa? How will it improve daily QA work?

SciMoCa establishes a new standard for accuracy in secondary dose check and independent plan QA. SciMoCa is powered by the accuracy of Monte Carlo dose calculations. It identifies TPS dose errors and discrepancies with minimal effort, and provides instantaneous feedback regarding treatment plan quality and achievement of planning objectives. SciMoCa utilizes a best-inclass Monte Carlo dose calculation algorithm with an outstanding academic pedigree to assure the quality of your QA. Enhance the quality of your treatment planning and patient QA with the accuracy of Monte Carlo.

Monte Carlo. Unlimited.

SciMoCa supports all Elekta™, all Siemens, all Varian LINACs [except Halcyon™] and CyberKnife®. TomoTherapy® will be supported soon.*

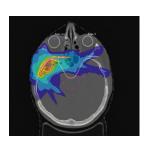
SciMoCa supports all treatment planning systems that provide DICOM export.

Supported treatment techniques are 3D, IMRT, VMAT, SBRT,

Accuracy

SciMoCa achieves an accuracy of < 1% ** and matches the precision of such top-tier algorithms as Varian's Acuros® and Accuray's Precision® Monte Carlo. The Monte Carlo calculation provides dose to medium and dose to water.

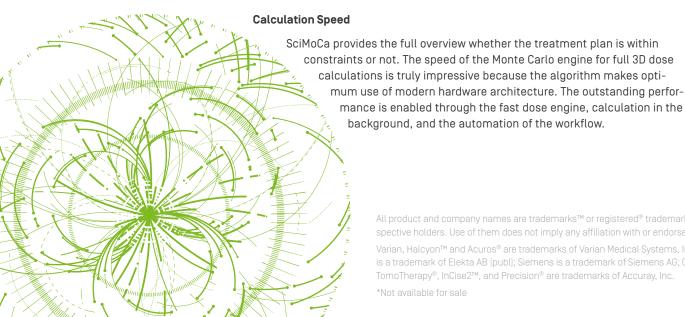
This accuracy is achieved due to the quality of this Monte Carlo implementation in addition to a custom machine-specific Linac beam model. This model is generated for the customer using proprietary processes and dedicated expert tools, backed by extensive expertise.



Accuray InCise2™ CyberKnife® dose distribution by SciMoCa



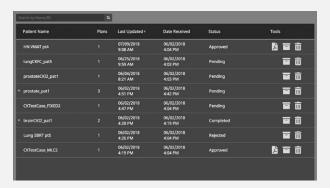
SciMoCa compared against Accuray Precision 1.0 [Monte Carlo], Gamma analysis 1%/1mm



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Varian, Halcyon™ and Acuros® are trademarks of Varian Medical Systems, Inc.; Elekta is a trademark of Elekta AB (publ); Siemens is a trademark of Siemens AG; CyberKnife®, TomoTherapy®, InCise2™, and Precision® are trademarks of Accuray, Inc.

Automated Workflow — Application Simplicity — Quality Outcome



AUTOMATIC PROCESS

Treatment plans are simply exported from the TPS via DICOM to the SciMoCa dose calculation server. Then SciMoCa starts the Monte Carlo calculation automatically.

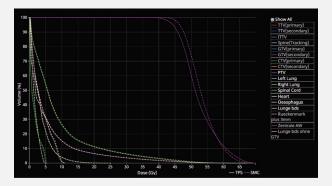
The patient worklist is automatically updated to keep the user informed about the current QA status of each patient plan.



AUTOMATIC ANALYSIS

SciMoCa provides automatic evaluation of dose calculation results based on customizable protocols and plan quality measures for both target volumes and OARs.

This allows users to evaluate plan quality as well as identify dose discrepancies based on user-defined dose-difference thresholds.



COMPARATIVE DVH

DVHs for the treatment plan as well as the independent SciMoCa dose distribution are compared in the comparative DVH section.

The software allows individual selection of structures for the DVH display, thereby enabling a detailed and focused evaluation of relevant ROIs.



3D ANALYSIS VIA SLICE VIEWER

SciMoCa includes a slice viewer to visualize the TPS dose, SciMoCa-recalculated Monte Carlo dose, and Gamma analysis on individual CT slices. The 3D viewer permits detecting dose discrepancies on the patient anatomy level.

^{**} SciMoCa published results: Gamma 2%/2mm pass > 97%, with deviation SciMoCa vs. Acuros® of mean dose in patients < 1%: Hoffmann et al., Validation of the Acuros® XB dose calculation algorithm versus Monte Carlo for clinical treatment plans. Med Phys. 2018 June 16

Discover more about SciMoCa:



Monte Carlo. Unlimited.

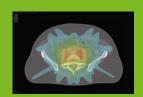
Monte Carlo Calculation Speed and Accuracy

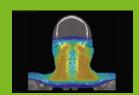
 PTV volume
 193.3 cc
 prostate / LN, dMLC [7 beams, 140 control points]
 head & neck, VMAT [2 arcs, 293 control points]

 voxel size
 3 mm
 3 mm
 2 mm

 calculation time
 15.8 sec
 55.6 sec
 40.9 sec
 118.9 sec







Calculation times on dual 8-core Intel Xeon E5-2690 server with hyperthreading [32 logical cores] for dose calculation with final statistical uncertainty of 1%.

SciMoCa Recommended System Requirements [v. 1.0 +]

Minimum recommended specifications: Recommended specifications:

СРИ	x64-based processors 4C/8T, 1.4GHz or faster	x64-based processors 8C/16T, 1.4GHz or faster
RAM		
Operating System	Windows™ Server 2012 R2 Standard or newer with IIS enabled	
Software		
Database	Microsoft SQL Server 2012 Standard or newer	
Monitor		
Language	English [US] or English [UK]	
Free Hard Disk Space		

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