

MOLECULAR IMAGING AND NUCLEAR MEDICINE

Molecular Radiotherapy Dosimetry

Practical Dosimetry Achieved

WHY DOSIMETRY?

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New Molecular Therapies Expand the Oncologist's Toolkit

Molecular Radiotherapies provide a distinct advantage over other systemic therapies: their distribution can be seen and measured, enabling personalization. Absorbed dose has shown to be a useful biomarker of response to radiation therapies. As a result, it has significant potential use in clinical decision-making to achieve the best possible patient outcomes. But is access to patient-specific absorbed dose practical?

IMPLEMENTATION

Accurate Dosimetry Performed at Scale

MIM SurePlan™ MRT has been helping institutions drastically reduce the clinical requirements for dosimetry by including AI-based auto-segmentation tools and support for quantitative SPECT reconstruction with existing SPECT/CT cameras, in addition to integrating automation into every facet of its design. This remains a primary focus for continued development.

Hospital administrators can take comfort in knowing that resources are being used efficiently. Likewise, patients and physicians alike will gain insight into these therapies and begin to characterize dose-response relationships for future treatment planning in specific patient populations.

Explore the Dosimetry Workflow

The history of dosimetry for these therapies is fraught with tedious computations and over-simplifications. Let's examine how MIM SurePlan MRT maximizes existing resources and reduces the manual effort to make accurate dosimetry practical for a clinical theranostics center.

PERSONALIZED DOSIMETRY WORKFLOW



MIM SurePlan MRT automates the entire dosimetry process from reconstruction to dose calculation.

IMAGE ACQUISITION AND PROCESSING **SPECT Image Processing**

Capture Quantitative SPECT Imaging with **Existing Hardware**

Standardize and Automate Reconstructions Quantitative reconstruction is supported for many SPECT cameras, bypassing the need to upgrade to a new camera or pay for an expensive quantitative SPECT package. Additionally, having a vendor-neutral platform can help standardize reconstructions across your department.

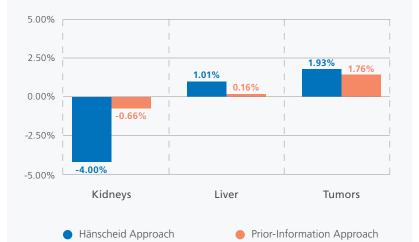
Eliminate the manual SPECT reconstruction process through intelligent programming that automatically performs this task as soon as the images arrive in MIM.

AVERAGE DEVIATION FROM 4 TP RESULTS*

Region of Interest	Hänscheid Approach	Prior- Information Approach
Kidney	-4.0 ± 10.7%	-0.7 ± 9.2%
Liver	1.0 ± 7.8%	0.2 ± 6.5%
Tumors	1.9 ± 11.6%	1.8 ± 12.7%

*Data from ¹⁷⁷Lu-DOTATATE patients. See full details in the MIM Software White Paper "Dosimetry for Targeted Molecular Radiotherapy Using a Single Measurement Timepoint."

SINGLE-TIME-POINT COMPARISON TO MULTIPLE-TIME-POINT DOSIMETRY*



Implement Practical Imaging Protocols

Single-time-point dosimetry is a cutting-edge dosimetry process that allows estimation of the absorbed dose with only a single SPECT/CT, further reducing the requirements needed to perform dosimetry.

Multiple-Time-Point Dosimetry

Model activity over time with two or more post-therapy image acquisitions. Generate patient-specific absorbed dose calculations on the voxel level.

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Prior-Information Approach to Single-Time-Point Dosimetry

Determine activity models with as few as two time points in the first cycle of therapy, then use those models with single-time-point acquisitions in subsequent therapy cycles. Achieve accurate, patient-specific dosimetry with flexible image acquisition times.





Day 1

Single-Time-Point Acquisition

Single-time-point methods reduce the imaging burden without sacrificing the benefits of patient-specific dosimetry.



Hybrid SPECT/Planar Dosimetry

Achieve 3D dosimetry without multiple SPECT/CTs. Voxel-level dosimetry is performed with planar time points for activity modeling, leveraging 3D information from a single SPECT/CT.

Hänscheid Approach to Single-Time-Point Dosimetry

Direct calculation of time-integrated activity using a single time point. The Hänscheid approach is available for post-therapy acquisitions of ¹⁷⁷Lu-DOTATATE and ¹⁷⁷Lu-PSMA.







Day 4

Day 7

NORMAL ORGAN AND TUMOR SEGMENTATION Normal Organ Segmentation

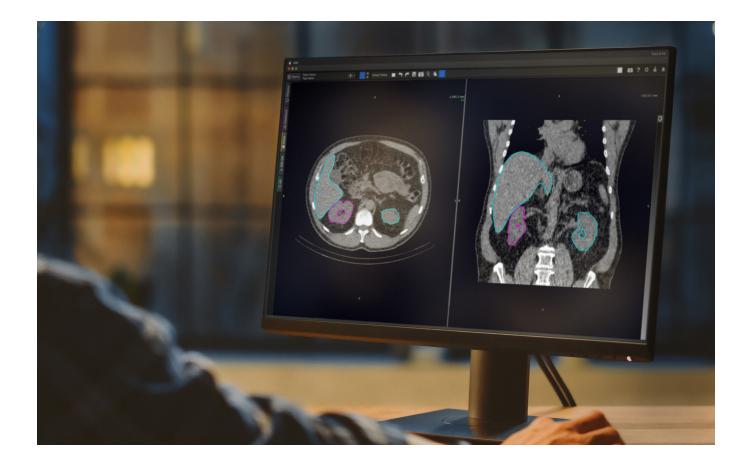
Segment Normal Structures with Artifical Intelligence

Manual segmentation is time-consuming, burdensome, and introduces inter-user variability.

Triggered automatically upon image arrival, Contour ProtégéAl+[™] for MIM SurePlan MRT uses a neural network for segmenting normal structures on CT images. Results are more accurate than atlas-based approaches so users spend less time editing.

Deliver AI on Premise or in the Cloud

Contour ProtégéAI+ for MIM SurePlan MRT has flexible deployment models that allow for easier adoption across any institution.



Tumor Segmentation

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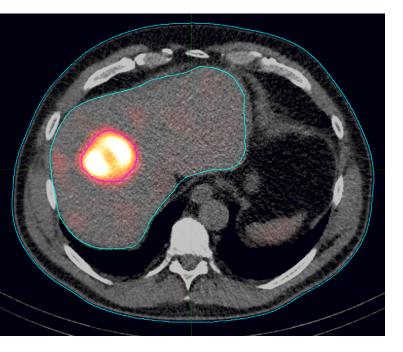
PET Edge®+

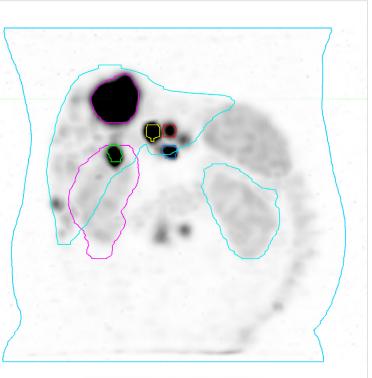
MIM Software's proprietary, gradientbased technique segments lesions with a single click.

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Speed and Consistency

Experience accurate and reproducible tumor segmentation regardless of who is doing the work.





PREPARATION

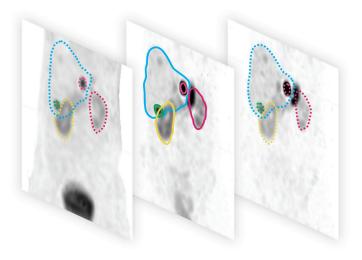
Image Alignment, Curve-Fitting, and Time-Integration

Automatic SPECT Alignment for Accurate Activity **Quantification Across Time**

Automatic Curve-Fitting

and Time-Integration

Optimized contour-based SPECT-to-SPECT activity alignment results in ROIs only needing to be defined on one SPECT/CT. This technique was specially designed for voxel-based dosimetry.



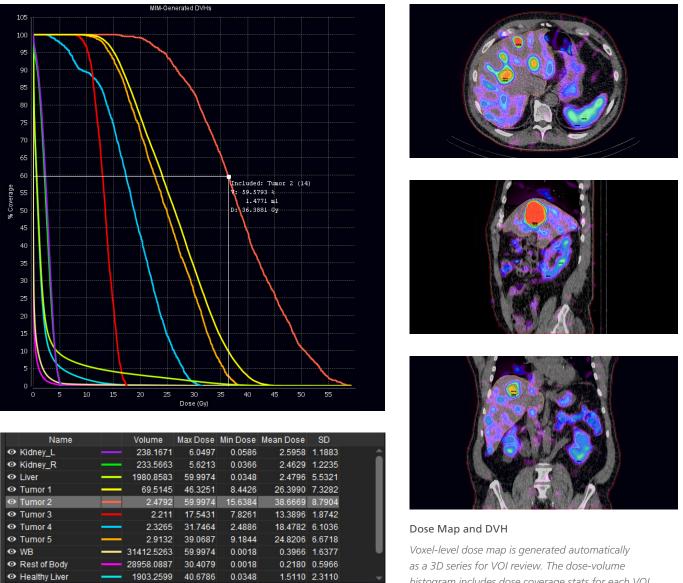
Activity-based alignment optimized per dosimetry structure.

Time-activity curve-fitting is designed to be flexible and smart. It can be performed using several available fit models or using trapezoidal integration, and fitting can be performed per-voxel or per-ROI.

Logical constraints on fit curve functions are in place to mitigate issues with noisy data.

Dose Calculation

Fast, Patient-Specific Dose Calculation on the Voxel Level



	Name	Volume	Max Dose	Min Dose	Mean Dose	
0	Kidney_L	238.1671	6.0497	0.0586	2.5958	1
0	Kidney_R	 233.5663	5.6213	0.0366	2.4629	
0	Liver	 1980.8583	59.9974	0.0348	2.4796	ŧ
0	Tumor 1	 69.5145	46.3251	8.4426	26.3990	7
0	Tumor 2	2.4792	59.9974	15.6384	38.6669	
0	Tumor 3	2.211	17.5431	7.8261	13.3896	
0	Tumor 4	 2.3265	31.7464	2.4886	18.4782	6
0	Tumor 5	 2.9132	39.0687	9.1844	24.8206	6
0	WB	 31412.5263	59.9974	0.0018	0.3966	
0	Rest of Body	28958.0887	30.4079	0.0018	0.2180	(
0	Healthy Liver	 1903.2599	40.6786	0.0348	1.5110	2

Voxel S-Value (VSV) kernels developed with Monte Carlo simulation are included in MIM SurePlan MRT for clinically used isotopes.

as a 3D series for VOI review. The dose-volume histogram includes dose coverage stats for each VOI.

Dosimetry Results Review

Dosimetry Reports Allow for Effective Communication with Patients and Referring Physicians

Track Dose Across Therapy Cycles and Quickly Review Tumor Progression MIM's structured reports can be customized according to your specific needs. For example, structured reports can be generated for referring physicians or for patients seeking to understand the effect of the therapy.

Dose accumulation runs in the background through MIM Assistant[®]. Cumulative doses and volume changes across cycles can be reviewed with a single interface.



49.9 (+21.7

65.8 (+15.9)

83.8 (+18

Dose accumulation statistics provide a therapy summary of absorbed dose and volume changes.

"MIM Software is devoted to providing superior customer service.

For MIM SurePlan MRT, that means around-the-clock support for questions and technical issues and customized setup and training. We design our dosimetry tools to streamline the clinical process and also allow for research-focused analysis.

We offer guidance for those new to dosimetry and access to clinical scientists for in-depth questions and technical understanding. Our goal is to provide both the tools and the support you need to improve patient care."

Aaron S. Nelson, MD Chief Medical Officer MIM SOFTWARE INC.



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