



REGION-BASED MAXIMUM LIKELIHOOD RECONSTRUCTION IN POSITRON EMISSION TOMOGRAPHY FOR QUANTITATIVE ONCOLOGICAL ANALYSIS

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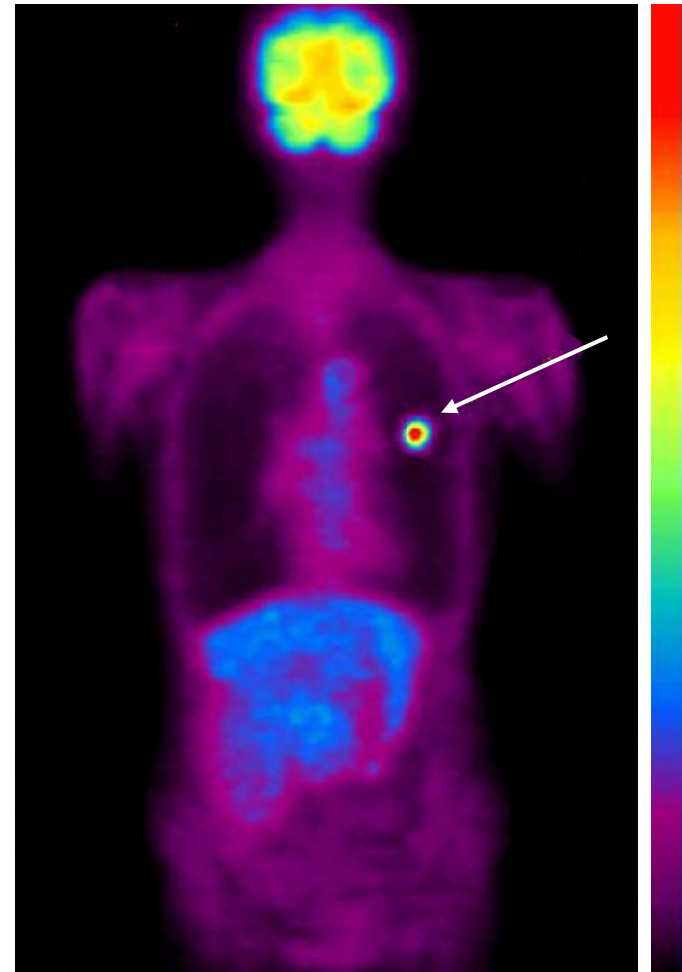
^{18}F -FDG uptake in tissues: index of cellular metabolic activity

^{18}F -FDG PET in clinical oncology:

- cancer diagnosis
- cancer staging
- response to therapy assessment

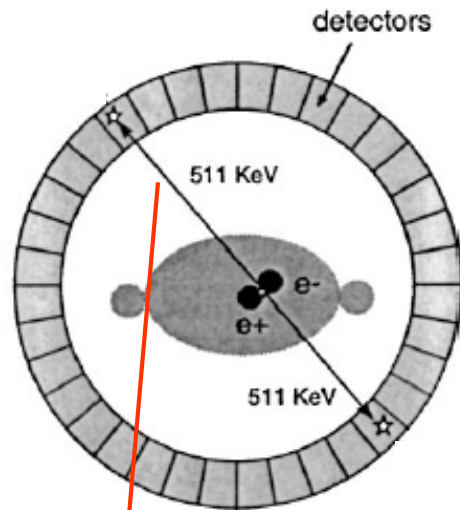
PET is an intrinsically quantitative technique because of:

- radiotracer nature
- attenuation correction

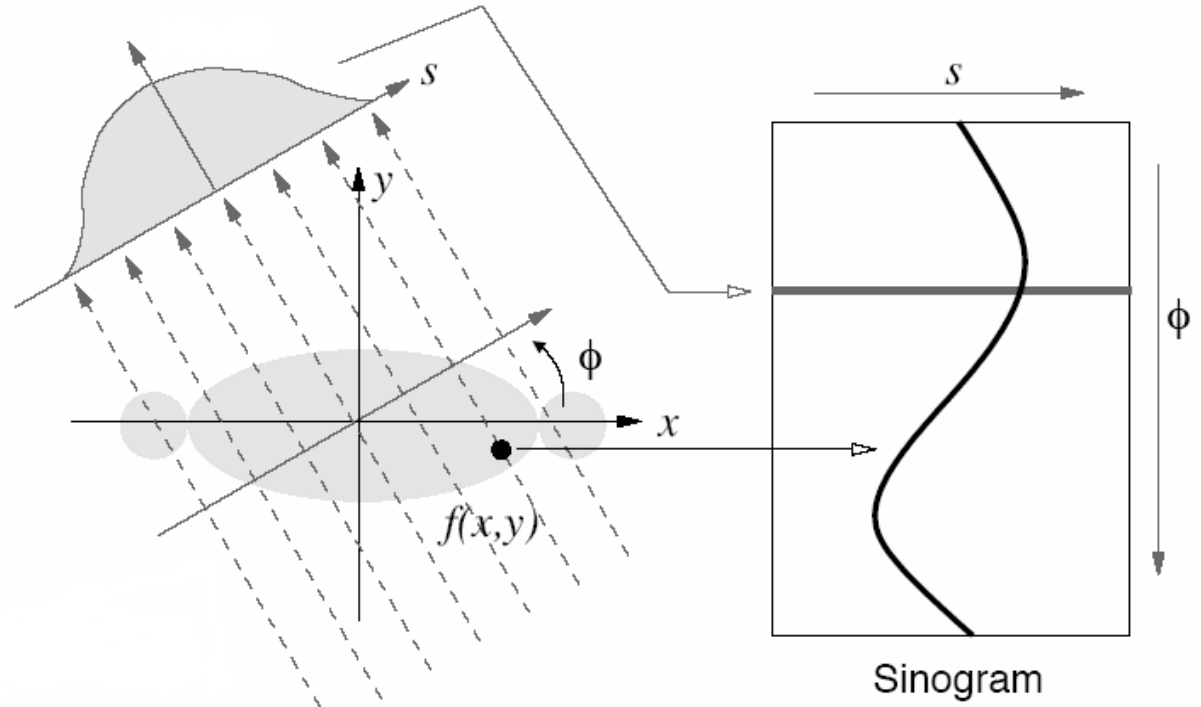




Reconstruction from projections: problem formulation



Line Of Response (LOR)



Sinogram

Acquisition process LSV modeling:

$$\bar{Y}_i = E[Y_i] = \iiint_{FOV} f(x, y, z) h_i(x, y, z) dx dy dz \quad i = 1, \dots, I$$

Poisson variable

integration kernel



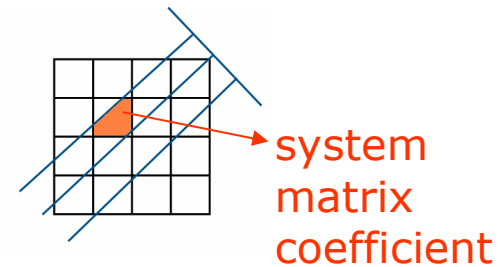
Reconstruction from projections: D-D approach

- Approximation of $f(x,y,z)$ as a linear combination of J basis functions $b_j(x,y,z)$ with unknown weights λ_j :

$$\tilde{f}(x, y, z) = \sum_{j=1}^J \lambda_j b_j(x, y, z)$$

- System matrix coefficients computation:

$$a_{ij} = \iiint_{FOV} h_i(x, y, z) b_j(x, y, z) dx dy dz$$



$$\bar{Y}_i = \sum_{j=1}^J a_{ij} \lambda_j$$

Reconstruction process: iterative search of λ values optimizing an objective function of the acquired data



Maximum Likelihood iterative reconstruction (MLEM)

- maximization of measured data likelihood function
- expectation maximization algorithm

n^{th} iteration:

$$\lambda_j^{n+1} = \lambda_j^n \cdot \frac{\left(\sum_i a_{ij} \frac{y_i}{\sum_k a_{ik} \lambda_k^n} \right)}{\sum_i a_{ij}}$$

i^{th} LOR error

i^{th} LOR estimated counts

weighted average of the errors



Main factors affecting PET accuracy in clinical practice

- Scanner spatial resolution

Point Spread Function (PSF)

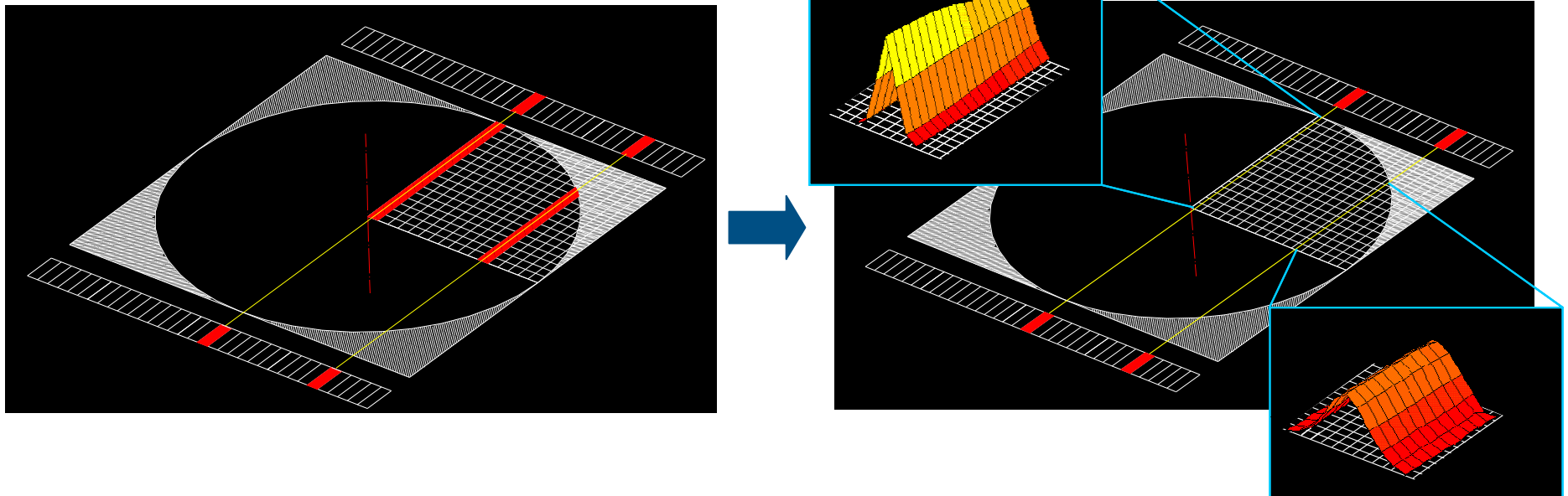


The PSF of a PET scanner is **anisotropic** and **spatially variant** (FWHM typically ranges from 4 to 9 mm)

- Noise component
- Reconstruction algorithm characteristics



Scanner PSF modeling and inclusion in MLEM reconstruction



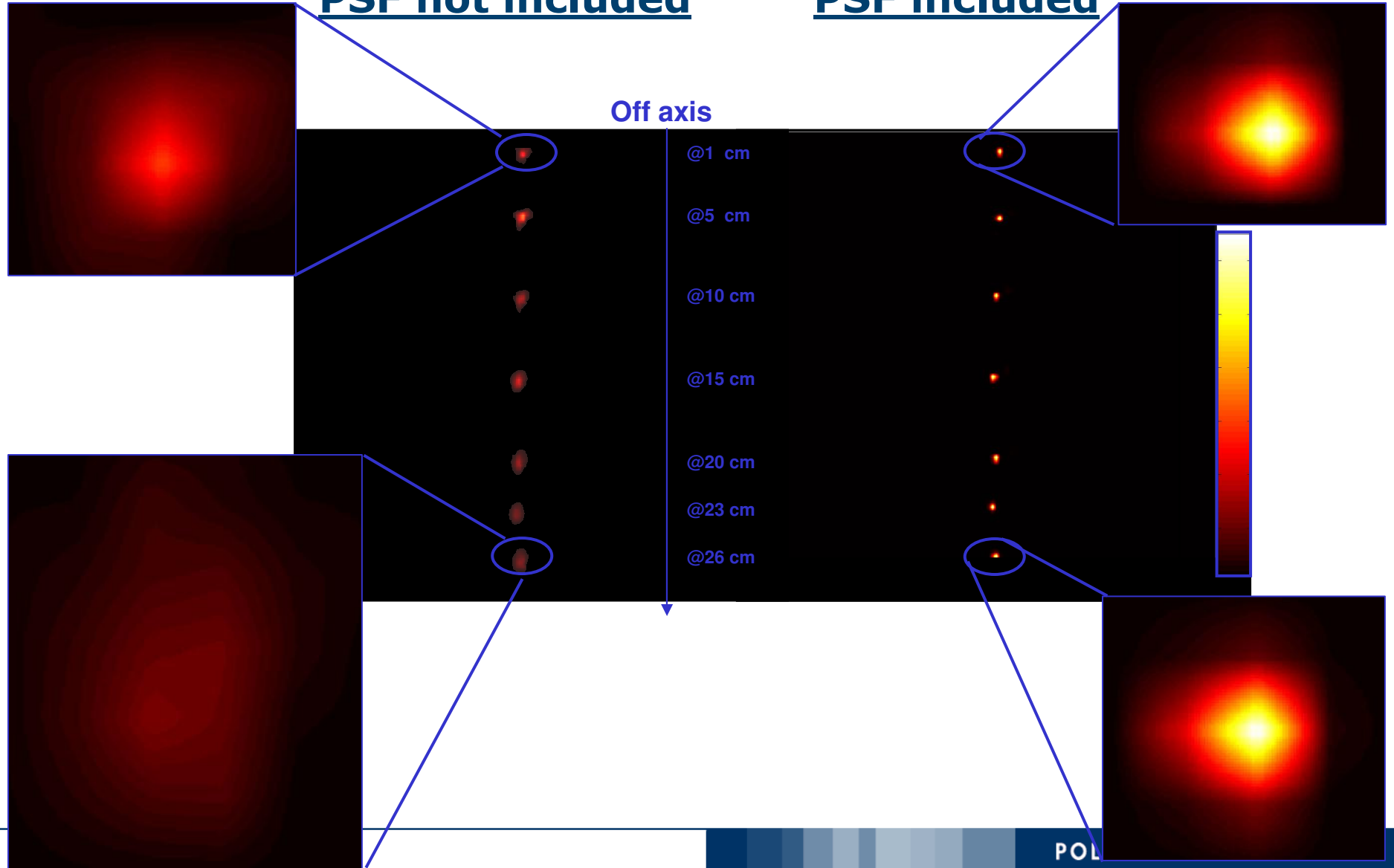


Scanner PSF modeling and inclusion in MLEM reconstruction

Line sources

PSF not included

PSF included





Scanner PSF modeling and inclusion in MLEM reconstruction: results

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Maximum likelihood algorithm drawbacks:

- maximum likelihood solution noisy due the problem ill-conditioning: stopping rules and/or post filtering required
- convergence slow and object-dependent

➡ object dependent bias/noise compromise

Effects of PSF inclusion:

- improvement of contrast and quantitative accuracy
- noise component reduction
- Gibbs-like artifacts
- worsening of the object dependent convergence

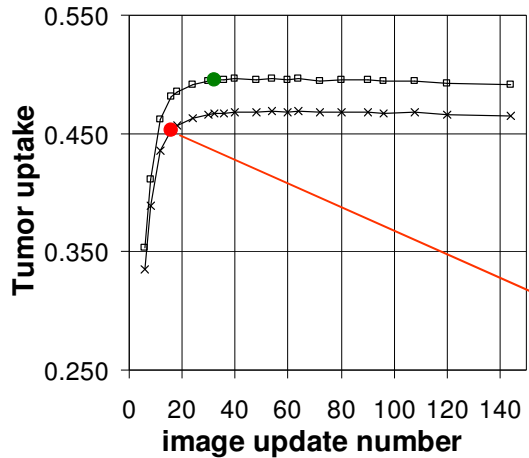
➡ detectability/quantification trade-off: a more correct quantification of small objects is necessarily associated with larger noise level and larger computation time.



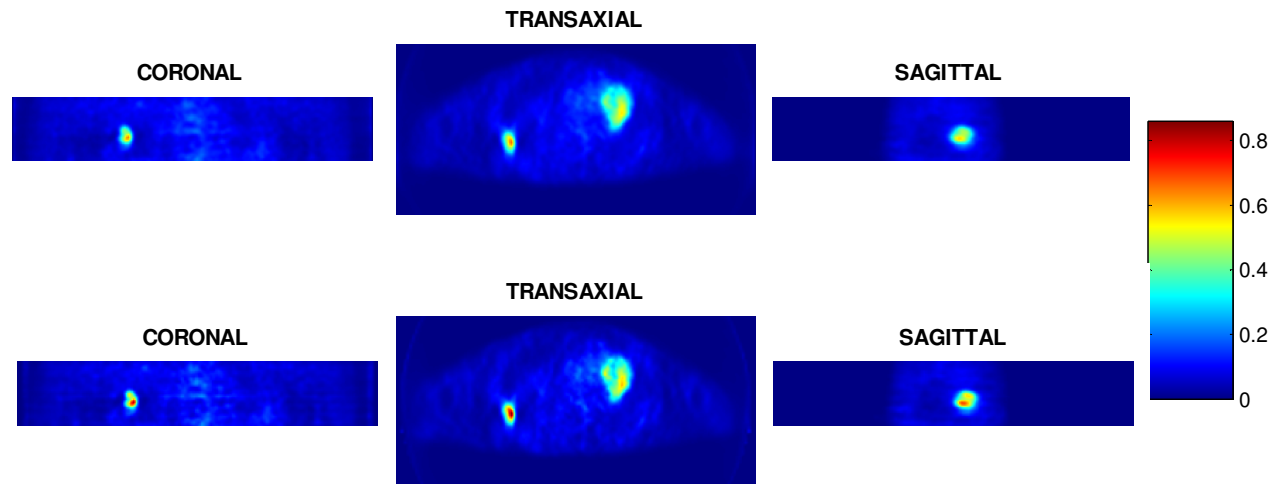
Scanner PSF modeling and inclusion in MLEM reconstruction: results

Lung cancer patient data: 35 mm tumor, large TBR, 3D mode

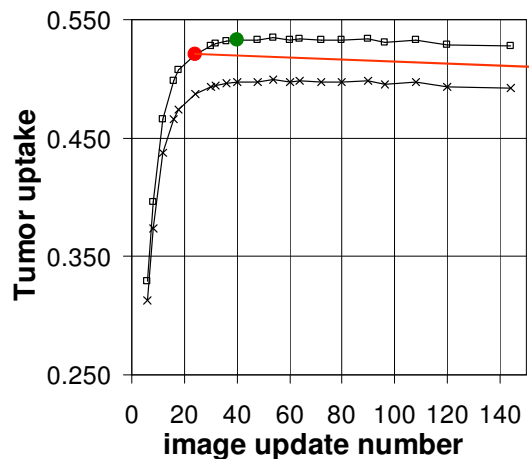
Without PSF



Clinical noise level



With PSF



Uptake +28%

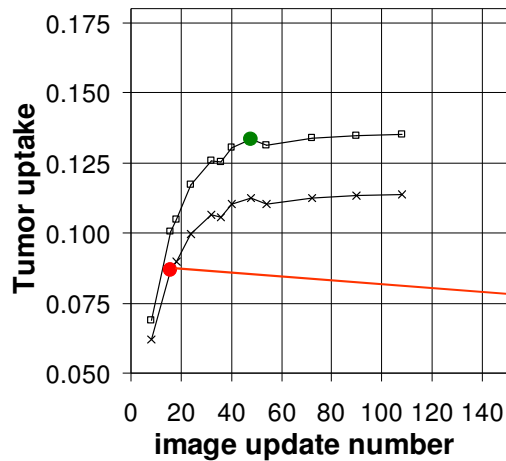
Metabolic volume -39%



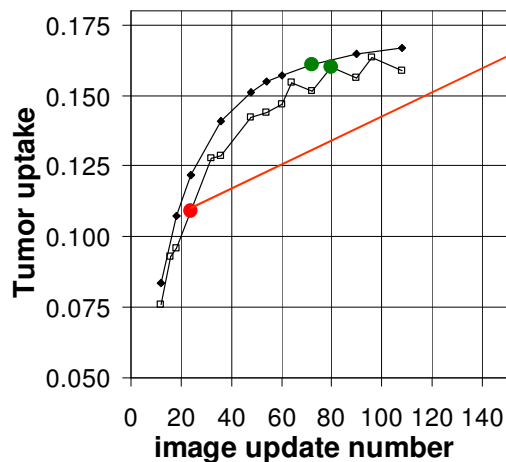
Scanner PSF modeling and inclusion in MLEM reconstruction: results

Lung cancer patient data: 15 mm tumor, low TBR, 2D mode

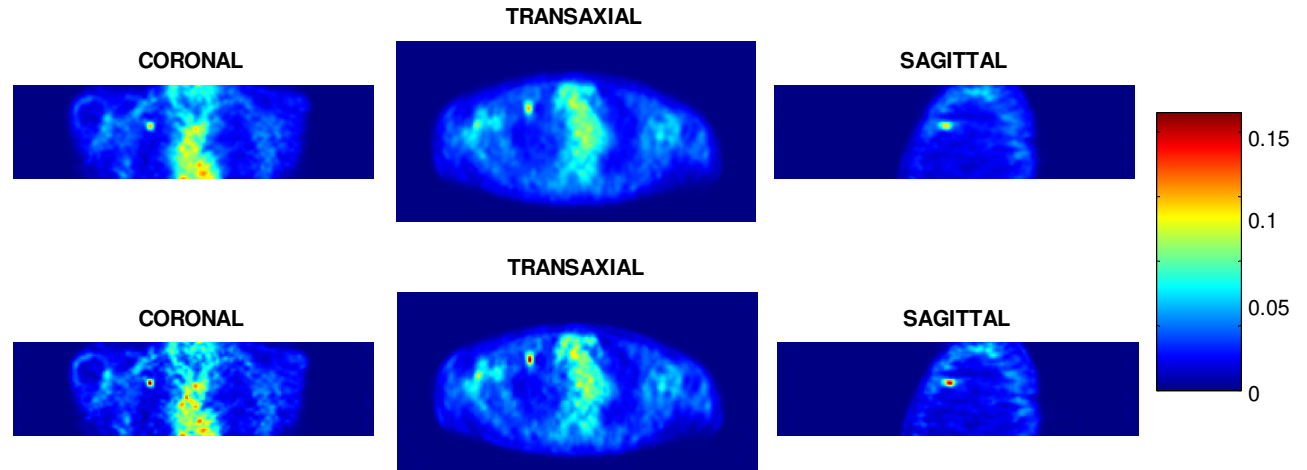
Without PSF



With PSF



Clinical noise level



Uptake +39%

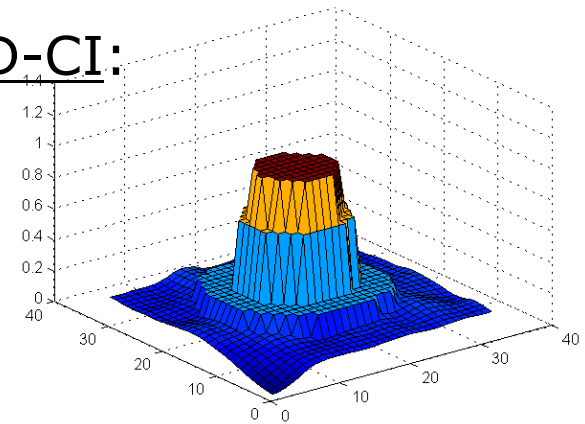
Metabolic volume -33%



Region-based MLEM with PSF inclusion for lesion quantification: the idea

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- Local quantification tool based on a standard smooth clinical image (STD-CI)
- Definition of regional basis functions on STD-CI:
 - the lesion of interest is properly segmented into regional basis functions
 - the image outside the area to be quantified is “frozen” and therefore handled as an unique basis function.
- The ML convergence can be locally achieved without noise increasing (reduced unknown variables; a priori information)





- STD-CI obtained with a standard voxel-based reconstruction process:

$$I^{STD-CI}(x, y, z) = \sum_{j=1}^J \lambda_j^{STD-CI} b_j(x, y, z)$$

- Approximation of $f(x, y, z)$ as linear combination of R regional basis functions $\beta_r(x, y, z)$ with unknown weights μ_r :

$$\tilde{f}(x, y, z) = \sum_{r=1}^R \mu_r \beta_r(x, y, z)$$

- Basis functions definition:

lesion of interest

$$\beta_r(x, y, z) = \begin{cases} T_r(x, y, z) E_{T_r} \left[I^{STD-CI}(x, y, z) \right] & r = 1..N \\ T_r(x, y, z) I^{STD-CI}(x, y, z) & r = bkg \end{cases}$$



- Model of the data collection process:

$$\bar{Y}_i = \sum_{j=1}^J a_{ij} \lambda_j = \sum_{r=1}^R \tilde{a}_{ir} \mu_r$$

- System matrix coefficients of the region-based problem:

$$\tilde{a}_{ir} = \sum_{j \in T_r} a_{ij} \bar{\lambda}_j$$

with $\bar{\lambda}_j$ = value assumed by the basis function $\beta_r(x, y, z)$ in the j^{th} voxel.



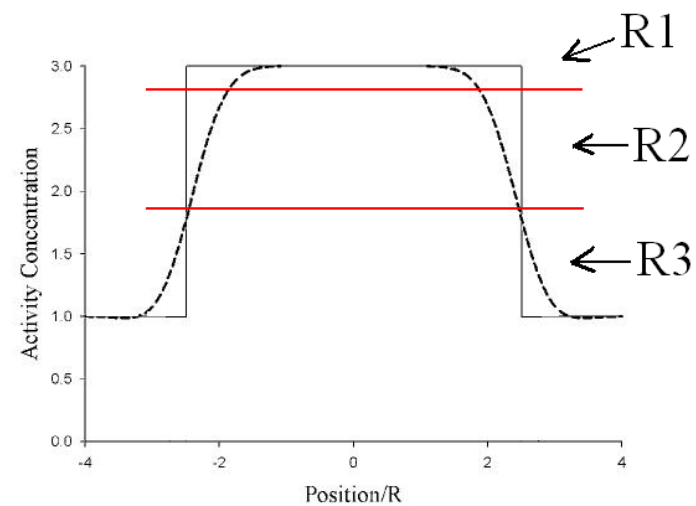
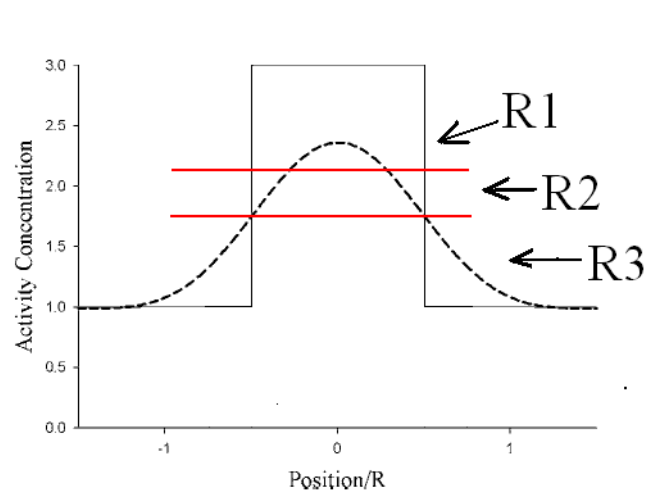
$$\mu_r^{n+1} = \mu_r^n \cdot \frac{\left(\sum_i \tilde{a}_{ir} \frac{y_i}{\sum_k \tilde{a}_{ik} \mu_k^n} \right)}{\sum_i \tilde{a}_{ir}}$$

n^{th} iteration of
region-based
MLEM
reconstruction



Choice of the number of regional basis functions for “uniform” lesions:

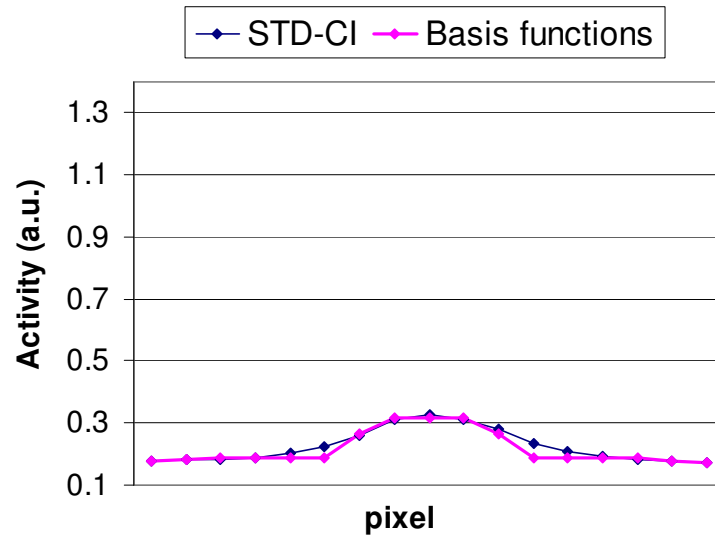
- R1: internal region;
- R2: transition region; activity underestimated for partial volume effect;
- R3: external region; activity overestimated for spillout effect



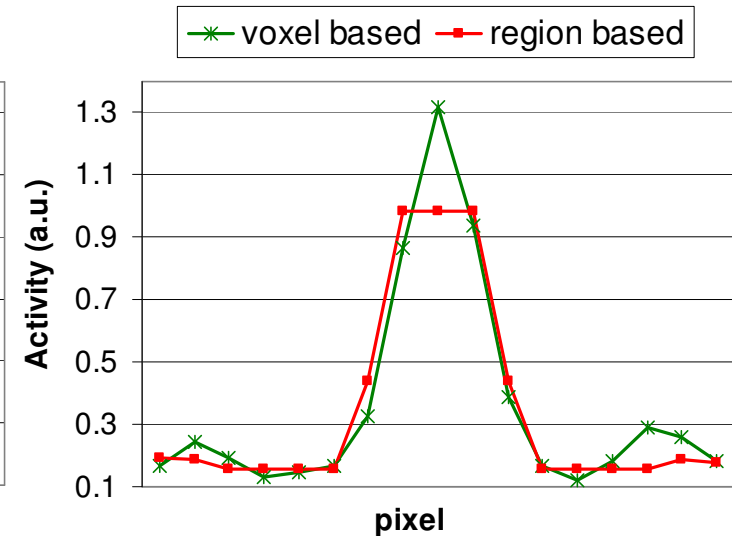


Experimentally acquired sphere phantom

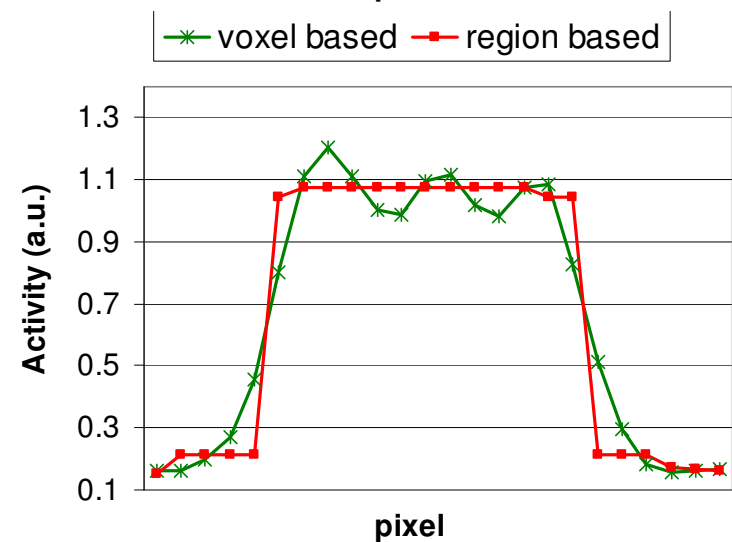
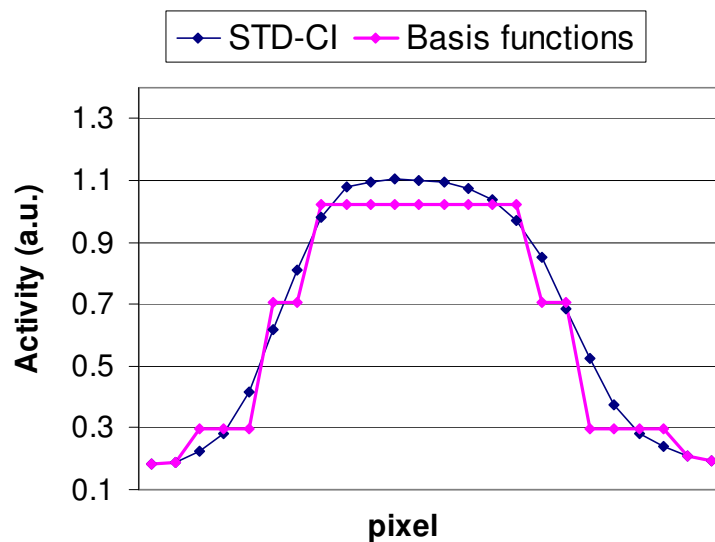
D=9.5mm
TBR=7.5



20 iterations

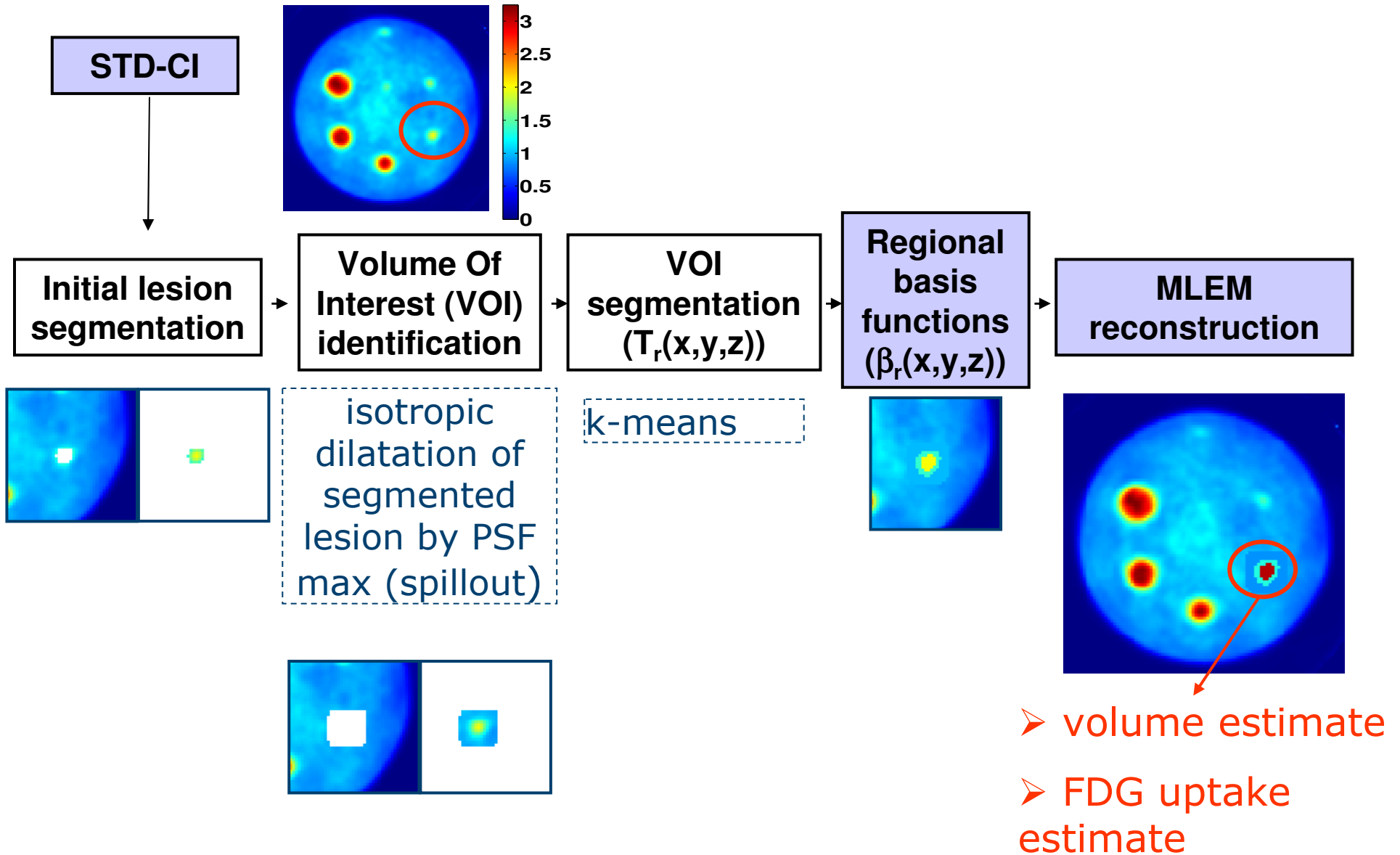


D=28mm
TBR=7.5





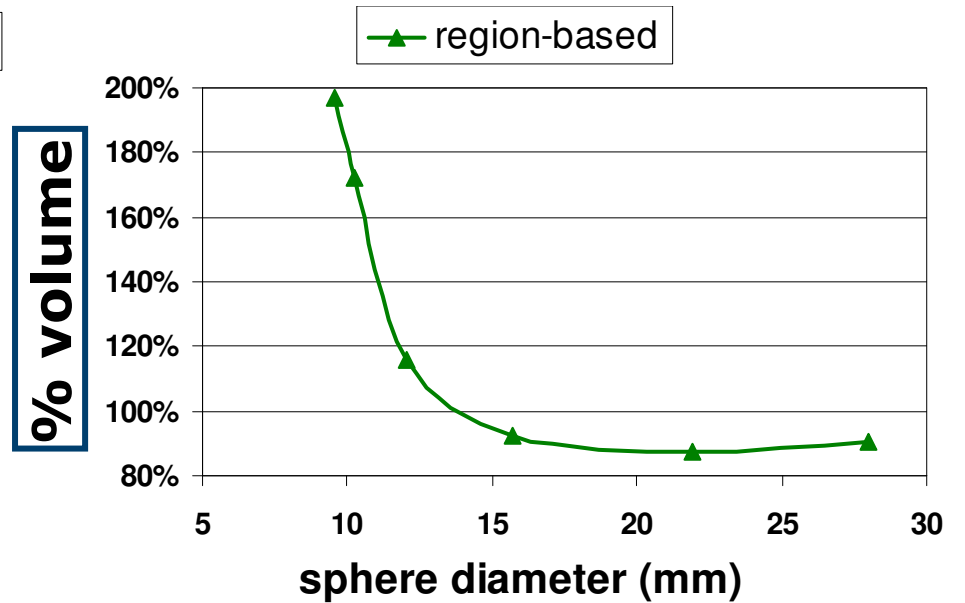
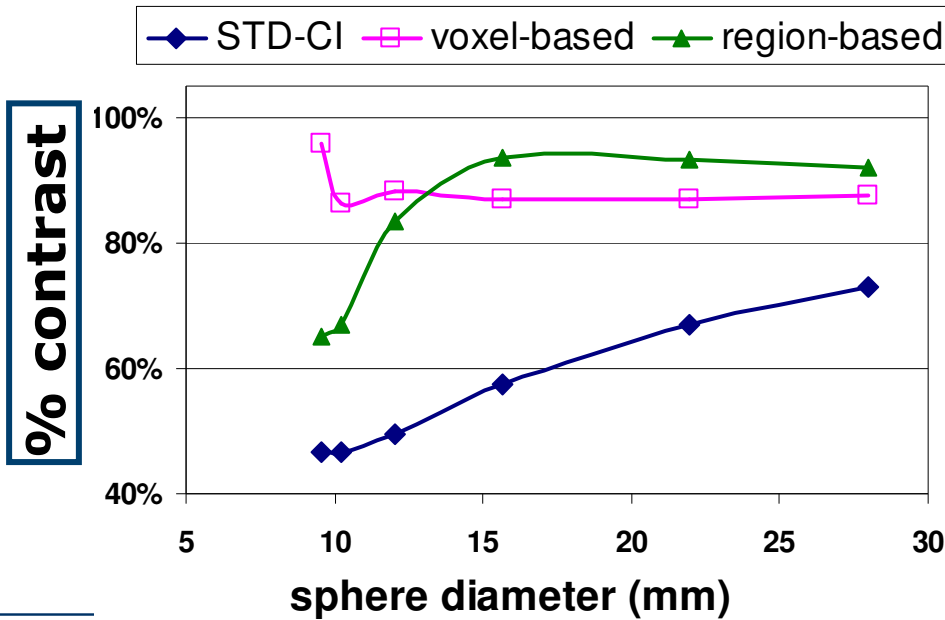
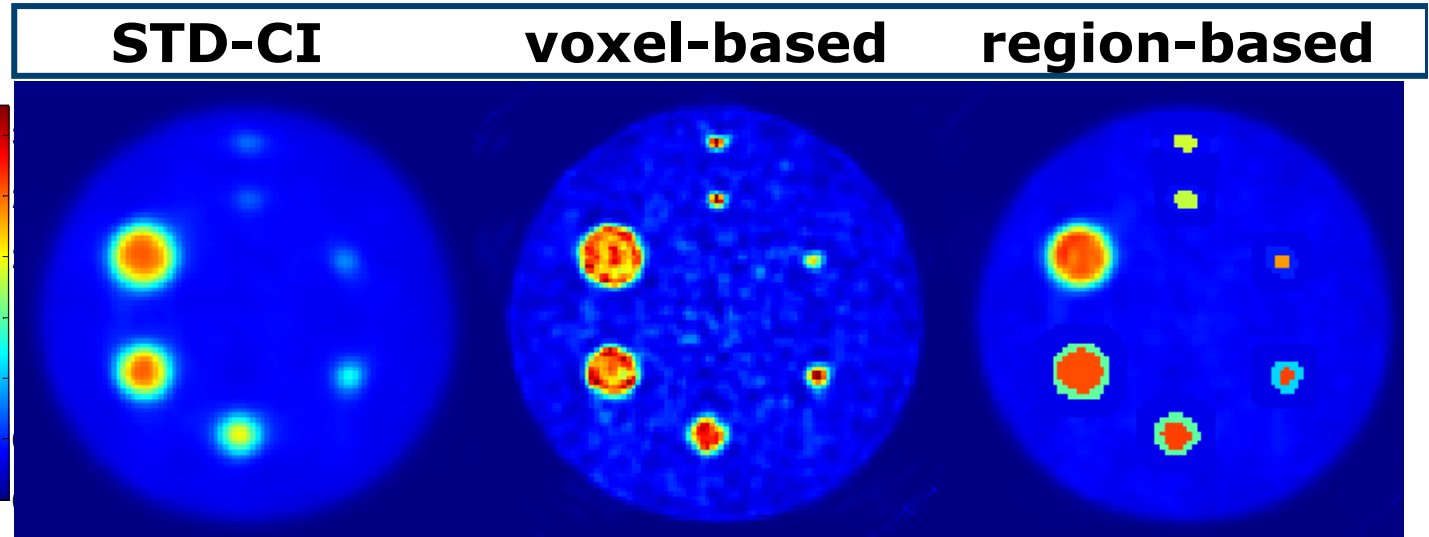
The algorithm





Experimentally scanned sphere phantom: results

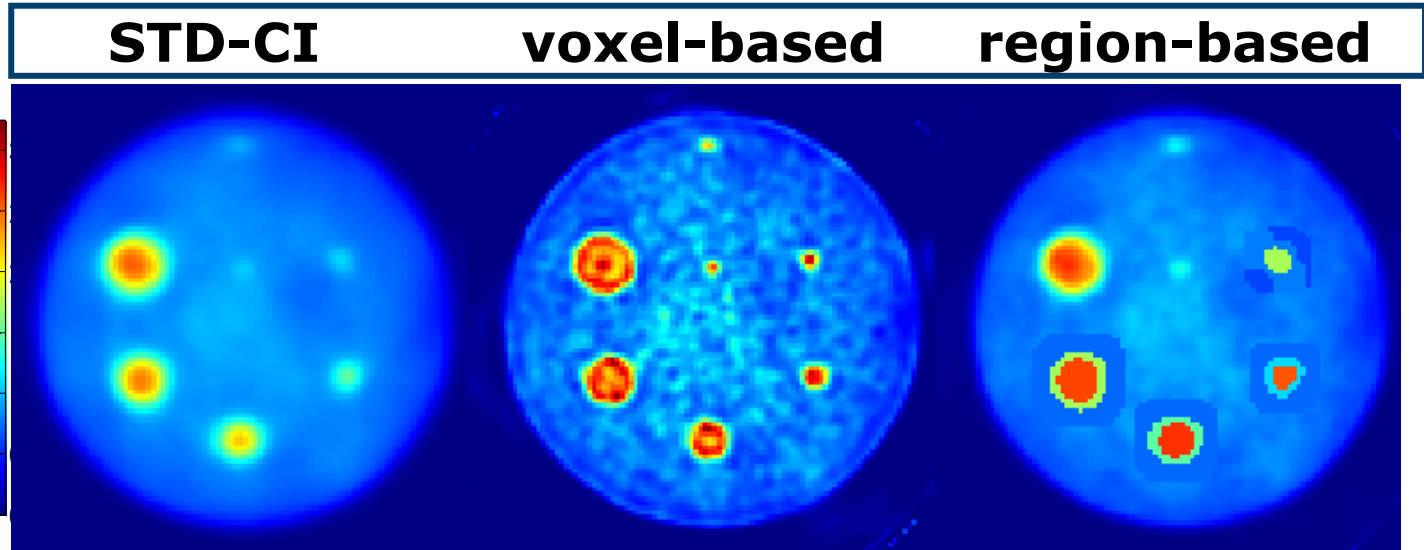
Activity contrast=7.5



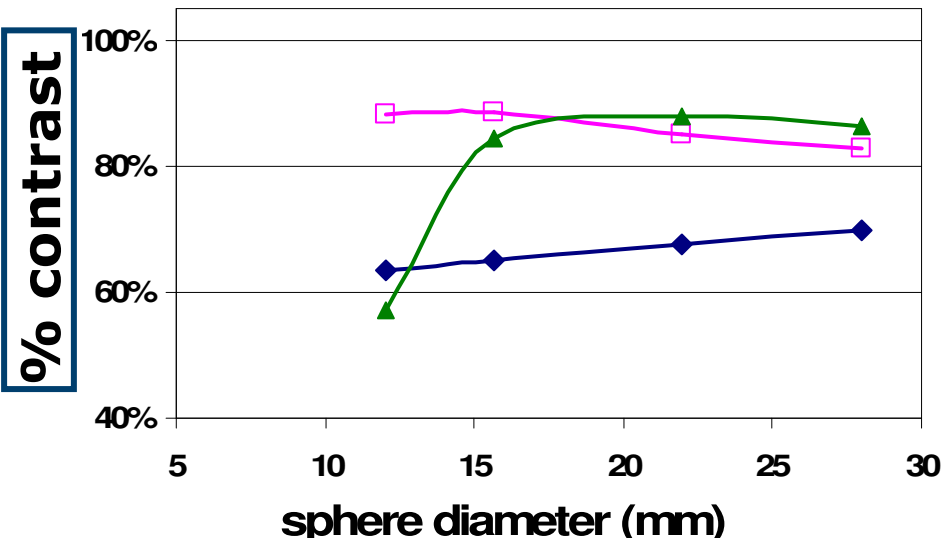


Experimentally scanned sphere phantom: results

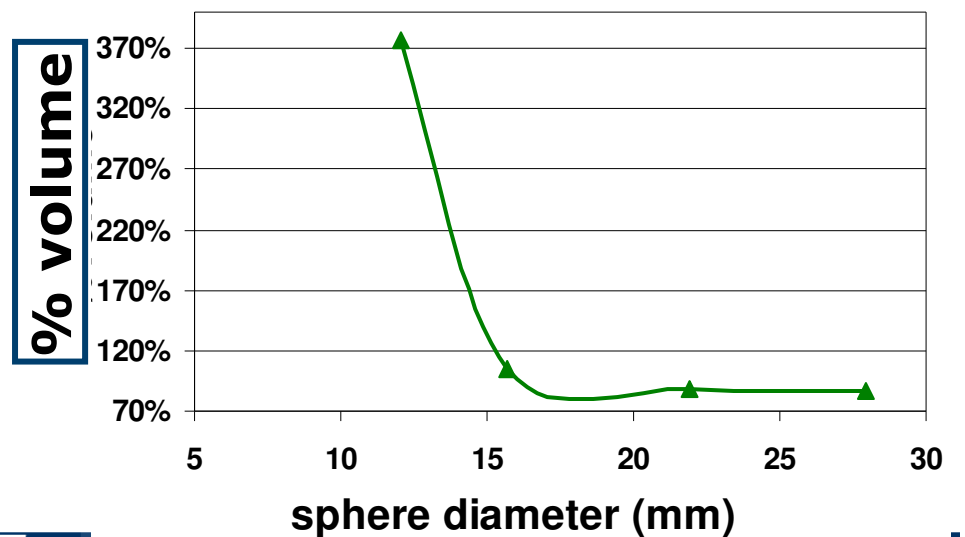
Activity
contrast=4



◆ STD-CI □ voxel-based ▲ region-based



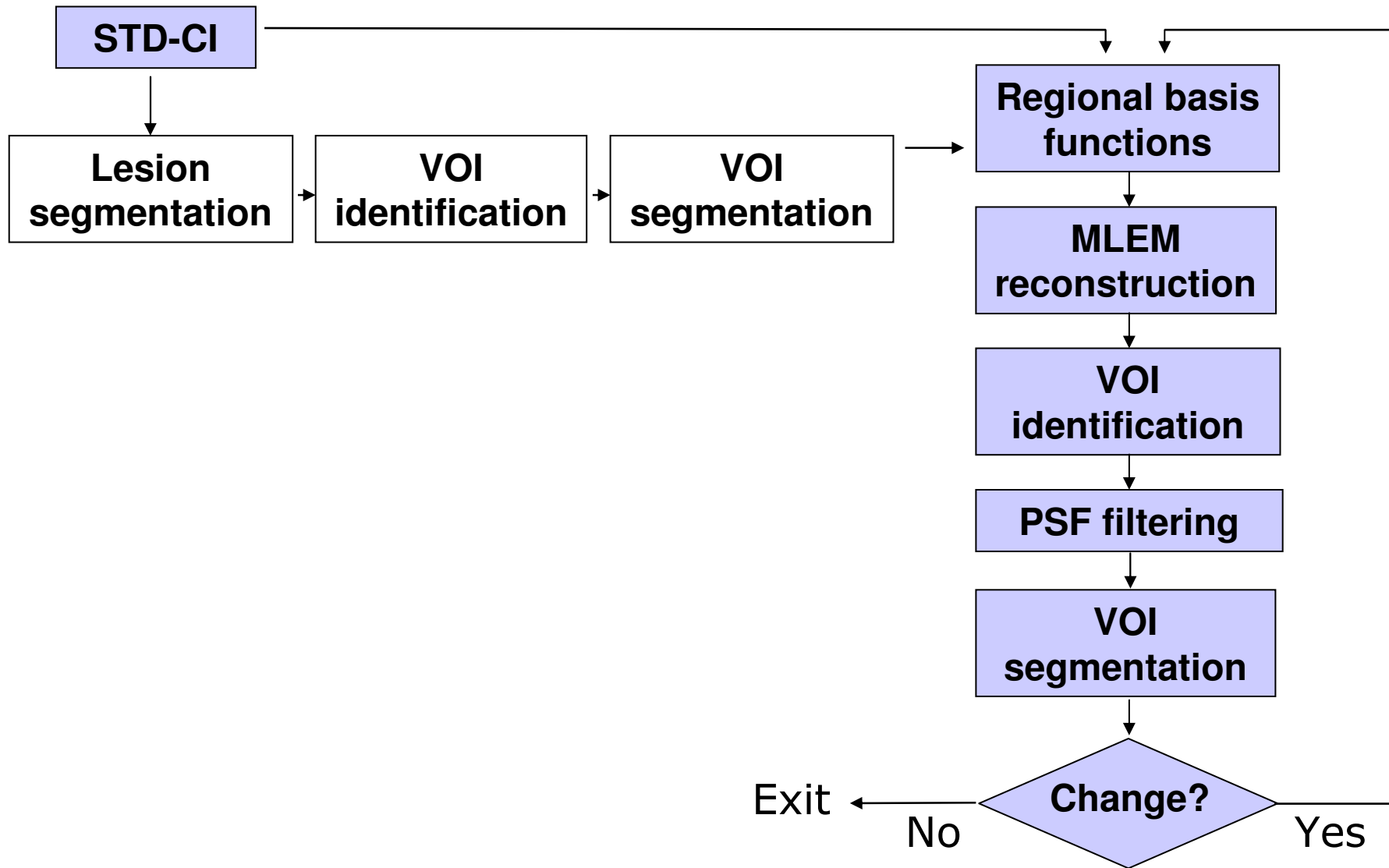
▲ region-based





The algorithm (2)

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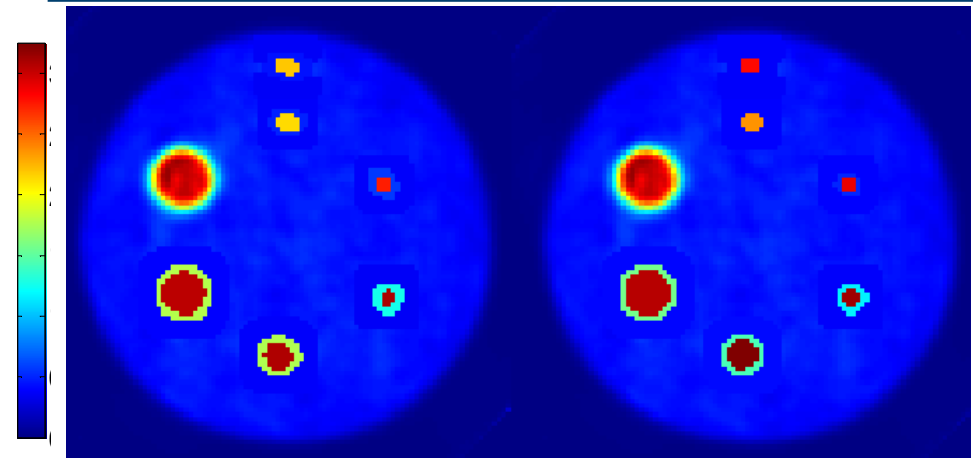




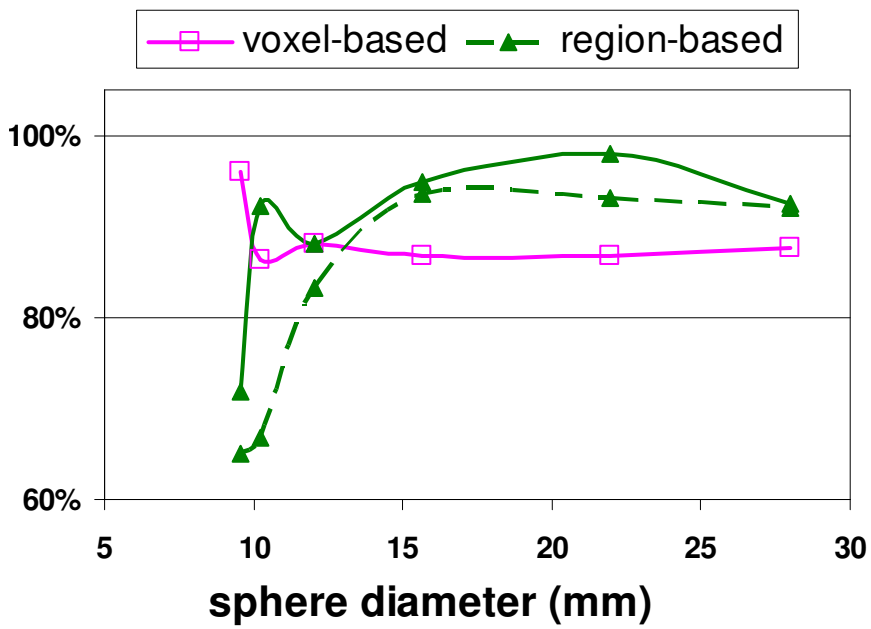
Experimentally scanned sphere phantom: results

Activity
contrast=7.5

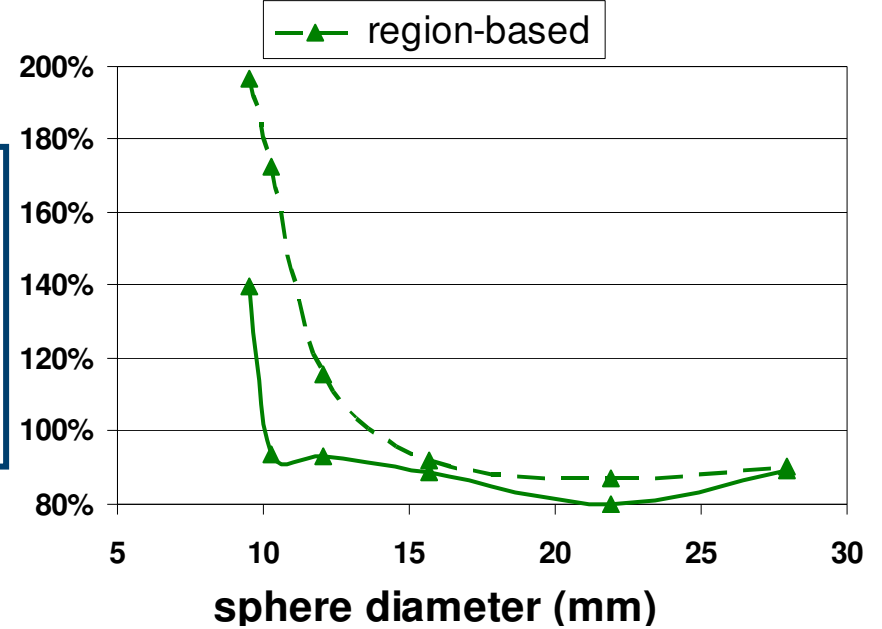
first iteration last iteration



% contrast



% volume

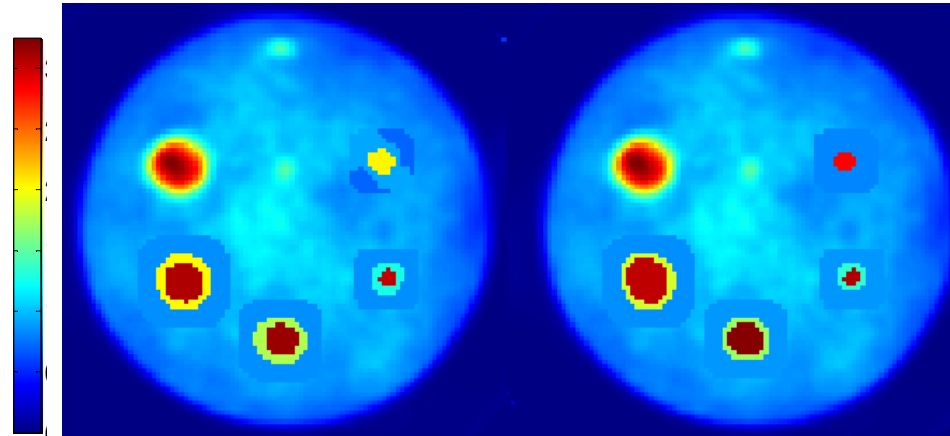




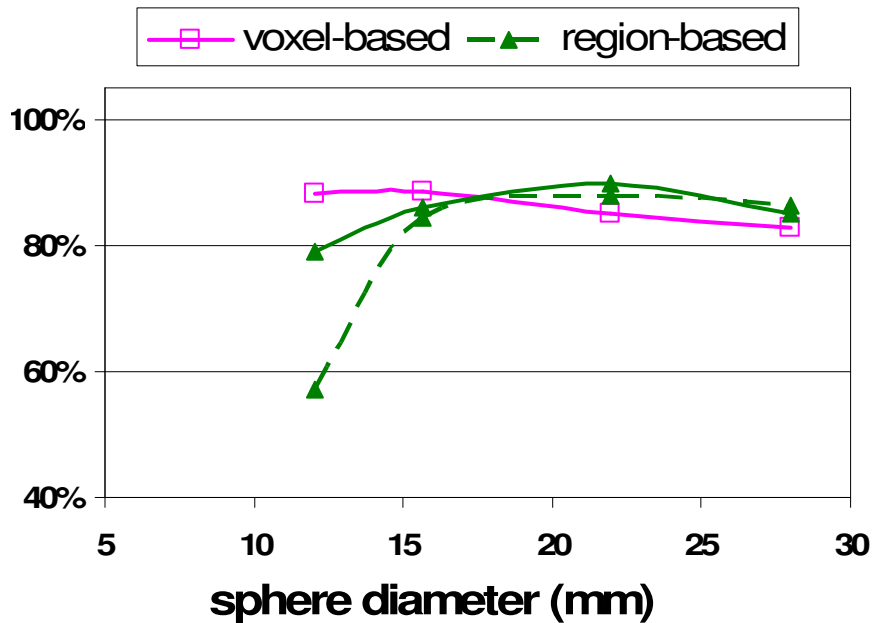
Experimentally scanned sphere phantom: results

Activity
contrast=4

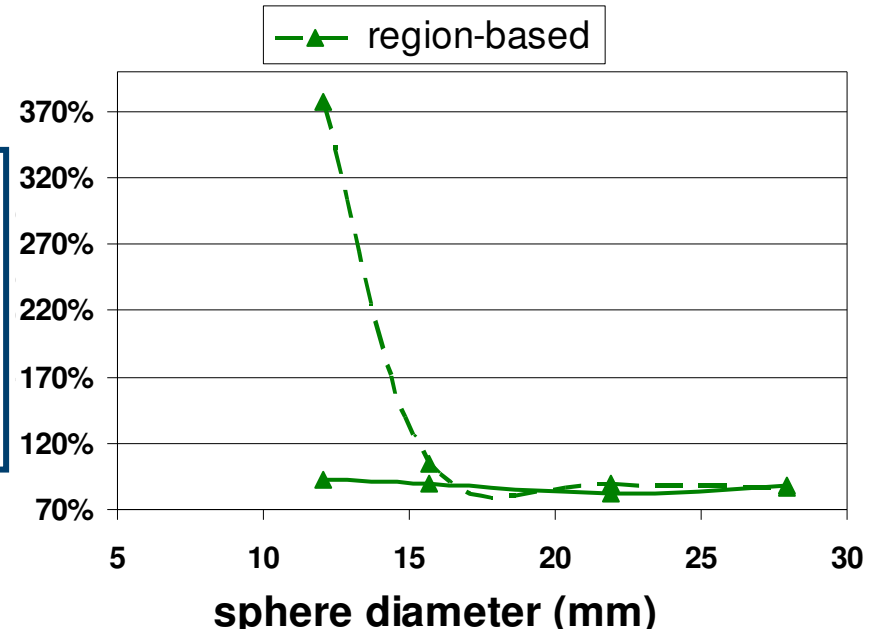
first iteration last iteration



% contrast



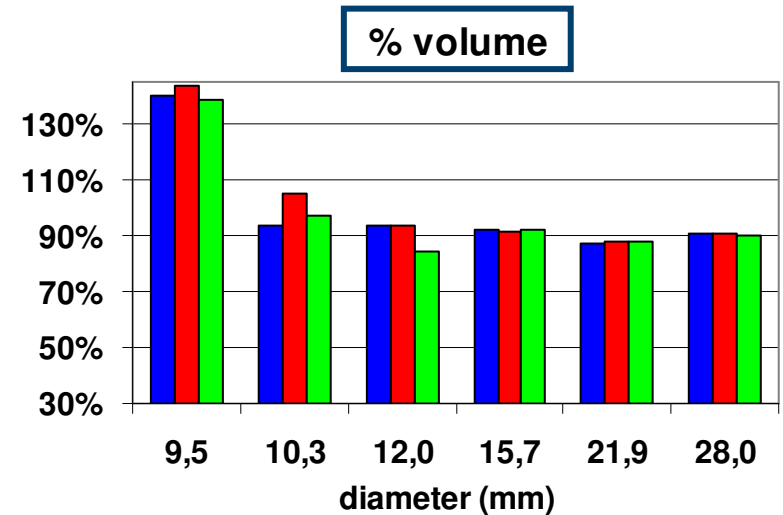
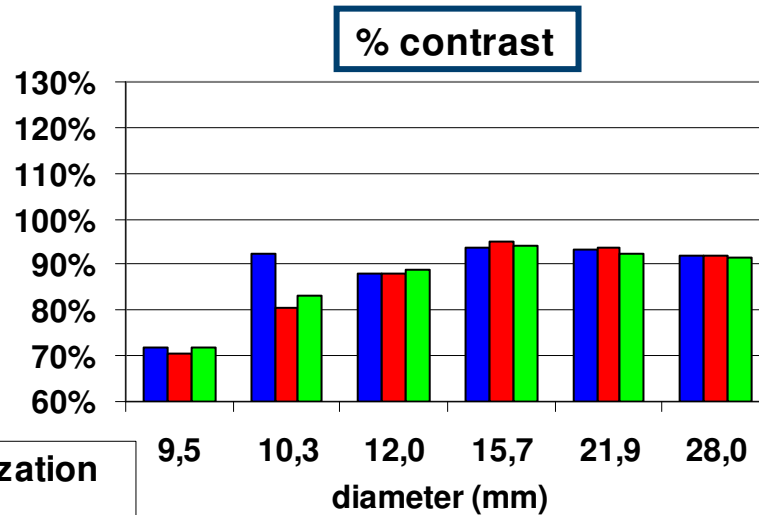
% volume





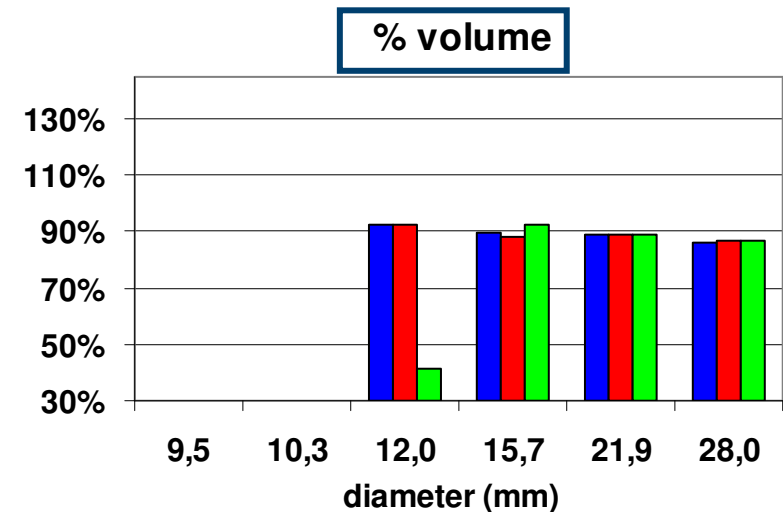
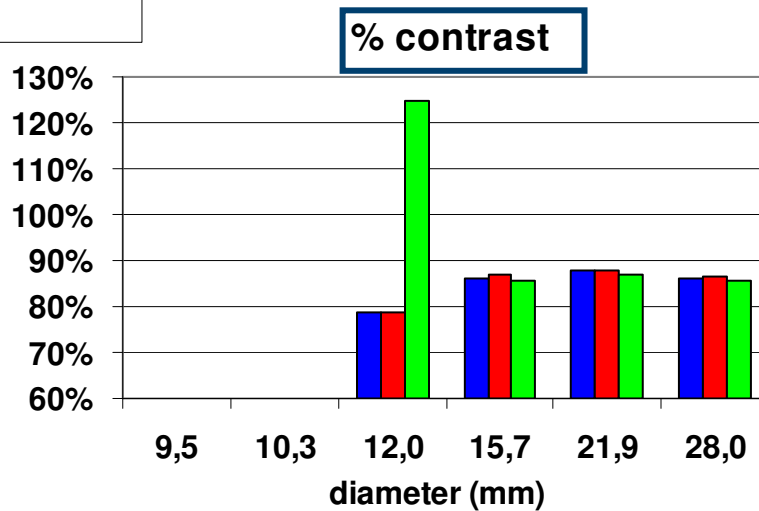
Errors in sphere segmentation: robustness evaluation

TBR 7.5



- correct initialization
- error +30%
- error -30%

TBR 4

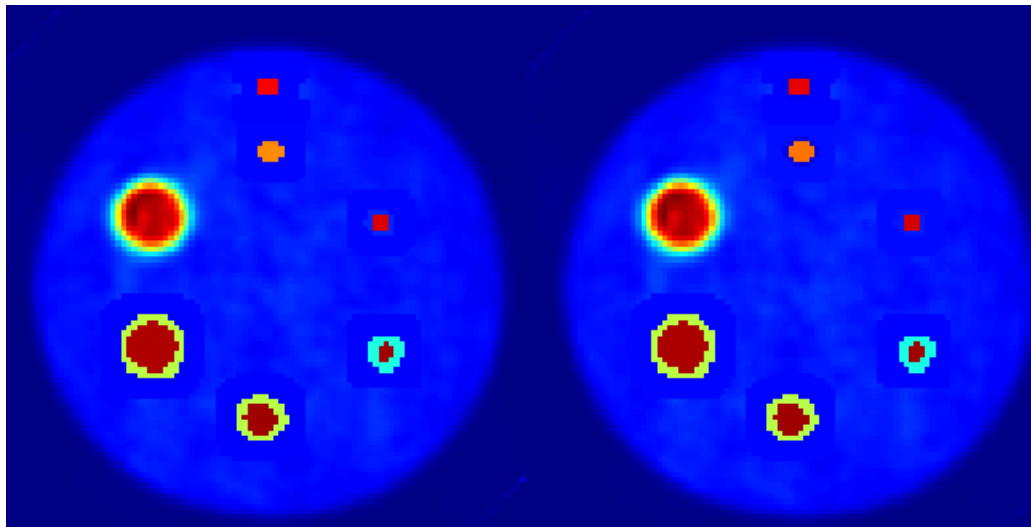
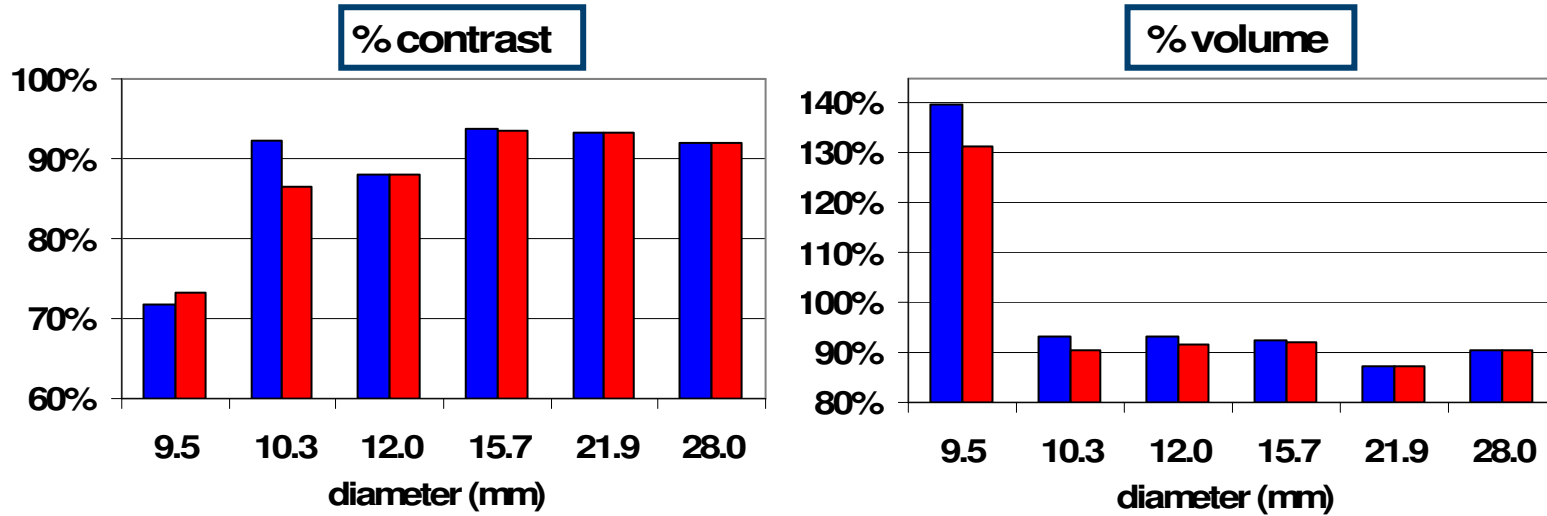


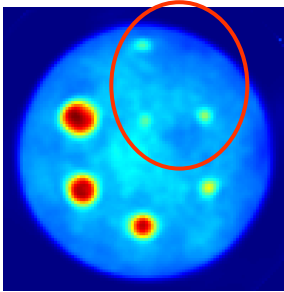


Experimentally acquired sphere phantom: results

TBR 7.5

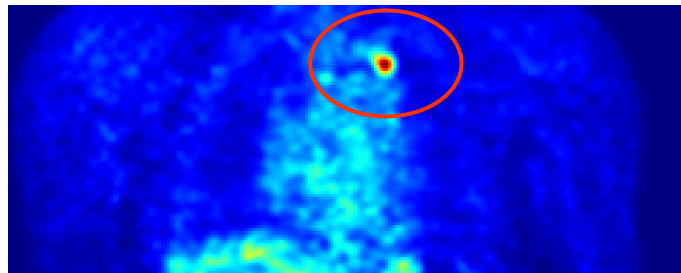
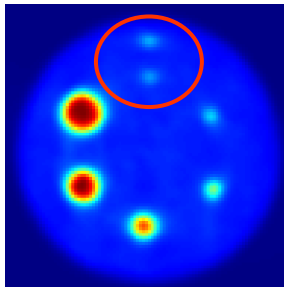
■ one object segmented ■ all objects segmented





1. Uniform, small, low-contrast objects:

- initial lesion segmentation with acceptable error ($\pm 30\%$)
- VOI segmentation in connected $T_r(x,y,z)$



2. Contiguous objects:

- VOI identification
- VOI segmentation in $Tr(x,y,z)$

3. Inhomogeneous objects:

- VOI segmentation in $Tr(x,y,z)$