Advanced TOF MLEM reconstruction of a human patient scanned by the modular J-PET

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## Outline

Positron emission tomography List-mode TOF MLEM with a realistic J-PET system matrix Single scatter simulation (SSS) STIR implementation of SSS SSS for the modular J-PET and normalisation Scan setup

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Results

Ways to improve

Conclusions

### Positron emission tomography

 $\beta^+ \rightarrow e^+ e^-$  annihilation  $\rightarrow$  511-keV back-to-back pair





Fig taken from [S. Cherry, S. Gambhir ILAR Journ 2001]

[Shopa R et al IEEE TRPMS 2023]

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### Positron emission tomography

*Radiotracers*: <sup>18</sup>F (FDG) – uptake of glucose (metabolism),
<sup>15</sup>O – blood flow, <sup>64</sup>Cu-ATSM – tissue hypoxia,
<sup>68</sup>Ga DOTATATE – somatostatin-receptor (SSTR),
neuroendocrine tumors

Less used: <sup>44</sup>Sc (preclinical cancer), <sup>22</sup>Na (long lifetime)

<sup>22</sup>Na, <sup>44</sup>Sc, <sup>68</sup>Ga – emit *prompt* photon (**positronium imaging**)





#### Maximum likelihood expectation maximisation

The expected (average) data  $\langle \mathbf{y} \rangle$  vs an unknown radiotracer distribution  $\lambda$ [Lange K. et al. (1984)]:  $\langle \mathbf{y} \rangle = \mathbf{M} \lambda + \hat{\mathbf{b}}$ 

 $\mathbf{M}$  – system matrix,  $\mathbf{\hat{b}}$  – *observation error* (randoms and scatter). Decomposition of  $\mathbf{M}$ :

$$\mathbf{M} = \mathbf{M}_{norm} \mathbf{M}_{det.res} \mathbf{M}_{att} \mathbf{M}_{Radon} \mathbf{M}_{ps.rng}$$

The measured photon counts  $\hat{\mathbf{y}}$  undergo Poisson distribution – a log-likelihood cost function is utilised for MLEM to solve iteratively:

$$oldsymbol{\lambda}^{(k+1)} = rac{oldsymbol{\lambda}^{(k)}}{\mathbf{M}^{\mathrm{T}} \mathbf{1}} \mathbf{M}^{\mathrm{T}} rac{\hat{\mathbf{y}}}{\mathbf{M} oldsymbol{\lambda}^{(k)} + \hat{\mathbf{b}}}$$



 $\langle \mathbf{y} \rangle$ 



5

λ

#### Maximum likelihood expectation maximisation

List-mode MLEM replaces back-projection  $\mathbf{M}^{T}\hat{\mathbf{y}}$  of sinograms with individual events  $\boldsymbol{\varepsilon}$ [Barrett H.H. et al. (1997)]

$$\lambda_{j}^{(k+1)} = \frac{\lambda_{j}^{(k)}}{\sum\limits_{i \in \mathcal{I}} m_{ij}} \sum_{\epsilon \in \mathcal{E}} \frac{m_{i_{\epsilon},j}}{\sum\limits_{j' \in \mathcal{J}} m_{i_{\epsilon},j'} \lambda_{j'}^{(k)} + \hat{b}_{i_{\epsilon}}}$$
$$\lambda_{j} \in \boldsymbol{\lambda}, \ \hat{b}_{i_{\epsilon}} = \hat{r}_{i_{\epsilon}} + \hat{s}_{i}$$

The shift-variant part of M:  $m_{ij} = n_i^{\text{cal}} a_i \chi_{ij}$  $a_i \in \mathbf{M}_{\text{att}}, \chi_{ij}$  jointly represents the geometrical part of  $\mathbf{M}_{\text{norm}}, \mathbf{M}_{\text{det.res}}$  and  $\mathbf{M}_{\text{radon}}$  [Shopa R et al IEEE TRPMS 2023]



[Martinez-Möller, Nekolla ZMP 2012]

## List-mode TOF MLEM for Jagiellonian PET

**J-PET:** plastic scintillators as detectors (Compton scatt.) Allows *time-of-flight* (TOF)! Requires kernels ( $H_{TOF}$ ) and *axial smearing* ( $H_Z$ ).

$$\chi_{(it)_{\epsilon},j} \to \chi_{i_{\epsilon}} \left( l_{j}^{\parallel}, l_{j}^{\perp}, \theta_{\epsilon} \right) \cdot H_{\text{TOF}} \left[ l_{j}^{\parallel}, \left( l_{0}^{\parallel} \right)_{\epsilon} \right] \\ \times \left[ H_{Z}^{\text{CRT}} \left( z_{j}, l_{j}^{\parallel} \right) * H_{Z}^{\text{prlx}} \left( z_{j}, l_{j}^{\parallel}, i_{\epsilon}, \theta_{\epsilon} \right) \right].$$

$$\lambda_j^{(k+1)} = \frac{\lambda_j^{(n)}}{\sum\limits_{i \in \mathcal{I}} n_i^{\text{cal}} a_i \chi_{ij}} \sum_{\epsilon \in \mathcal{E}} \frac{\chi_{(it)_{\epsilon},j}}{\sum\limits_{j' \in \mathcal{J}_{\epsilon}} \chi_{(it)_{\epsilon},j'} \lambda_{j'}^{(k)} + \hat{b}_{i_{\epsilon}}^*}.$$
$$\hat{b}_{i_{\epsilon}}^* = (\hat{r}_{i_{\epsilon}} + \hat{s}_{i_{\epsilon}}) / (n_{i_{\epsilon}}^{\text{cal}} a_{i_{\epsilon}}) \qquad \forall i \in \mathcal{I} \ n_i^{\text{cal}} = 1$$



Monte Carlo simulation ( $\theta = \pi/4$ )



fitted by  $\chi_i(l^{||}, l^{\perp}, \theta)$ 



Simulated 2-layer 2-m long J-PET:  $\chi_{ij}$  defined via Monte Carlo, then fitted by polynomial [Shopa et al IEEE TRPMS 2023]

# List-mode TOF MLEM

**Simpler solution (c++):** Monte Carlo for a grid  $r \times l \times \vartheta_{b2b}$ , do not fit, calc only for a half-module (symmetry), i.e. ~300 × 7 bins for a *single-layer modular J-PET* (can be extended to  $2\gamma$  + prompt)





### List-mode TOF MLEM

**Add TOF**: post-MC ( $2\gamma$ ) or for each event ( $2\gamma$  + prompt). Interpolate rough grid to match voxel size.



### **Types of coincidence data**

[Kowalski P. et al. (2018) PMB]



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#### Single scatter simulation (SSS)

Introduced by Watson C.C. et al. (1996) – expected single Compton scattered events for a detector pair (*A*,*B*), no TOF



$$S^{AB} = \int_{V_s} dV_s \left( \frac{\sigma_{AS} \sigma_{BS}}{4\pi R_{AS}^2 R_{BS}^2} \right) \underbrace{\mu}_{\sigma_c} \frac{d\sigma_c}{d\Omega} \left[ I^A + I^B \right]$$
$$I^A = \varepsilon_{AS} \varepsilon'_{BS} e^{-\left( \int_s^A \mu ds + \int_s^B \mu' ds \right)} \int_s^A \lambda(s) ds$$
activity
$$I^B = \varepsilon_{BS} \varepsilon'_{AS} e^{-\left( \int_s^A \mu' ds + \int_s^B \mu ds \right)} \int_s^B \lambda(s) ds.$$

Referring to Fig. 1,  $V_s$  is the total scatter volume (that is, the sentire volume of finite density), S is the scatter point,  $dV_s$  is an incremental volume element at S, s is the distance from the scatter point along a ray such as AS,  $\sigma_{AS}$  is the geometrical cross section of detector A for  $\gamma$ -rays incident along AS,  $R_{AS}$  is the distance from S to A,  $\varepsilon_{AS}$  is the efficiency of detector A for  $\gamma$ -rays incident along AS,  $\mu$  is the linear attenuation coefficient,  $\lambda$  is the emitter density in the object,  $\sigma_c$  is the total Compton interaction cross section, and  $\Omega$  is the scattering solid angle.

# **STIR implementation of SSS**

STIR – Software for Tomographic Image Reconstruction, current version 6.0. Two options:

 $> \texttt{simulate\_scatter simulate\_scatter.par}$ 

> estimate\_scatter scatter\_estimate.par

We use non-TOF **simulate\_scatter**: easier for J-PET Computationally expensive – requires downsampling and low-res activity & attenuation images.

Example: use sampled low-res bins and interpolate [Werling A. et al. (2002)]



### **Template for SSS sinogram in STIR**

Ideal cylinder, D = 73.8 cm, 24 axial 2-cm rings, 96 detectors per ring Energy cuts: 425 keV – 650 keV



### Workflow



Factors  $s_{ie}^{(SSS)}$  and  $a_{ie}$  are calculated *for each event* (using *interpolation* for SSS sinogram)

#### Normalisation problem

#### Important: ignore randoms!

$$\lambda_j^{(k+1)} = \frac{\lambda_j^{(k)}}{\sum\limits_{i \in \mathcal{I}} n_i^{\text{cal}} a_i \chi_{ij}} \sum_{\epsilon \in \mathcal{E}} \frac{\chi_{(it)_{\epsilon},j}}{\sum\limits_{j' \in \mathcal{J}_{\epsilon}} \chi_{(it)_{\epsilon},j'} \lambda_{j'}^{(k)} + \hat{b}_{i_{\epsilon}}^*}.$$
$$\hat{b}_{i_{\epsilon}}^* = (i + \hat{s}_{i_{\epsilon}}) / (n_{i_{\epsilon}}^{\text{cal}} a_{i_{\epsilon}})$$

STIR uses *tail-fitting* for normalisation of SSS-sinograms (as projection data). Difficult to apply to list-mode TOF data of J-PET.

 $\hat{s}_{i_{\epsilon}} \to n^{(\mathrm{SSS})} \cdot s_{i_{\epsilon}}^{(\mathrm{SSS})}$ 

Current solution – normalise by each event by an *adjustable regularisation factor*  $\beta$ .

$$n_{\epsilon}^{(\mathrm{SSS})} = \beta \cdot n_{\chi_{\epsilon}} \equiv \beta \cdot \sum_{j \in \mathcal{J}_{\epsilon}} \chi_{it,j}$$
$$\lambda_{j}^{(k+1)} = \frac{\lambda_{j}^{(k)}}{\sum_{i \in \mathcal{I}} a_{i}\chi_{ij}} \sum_{\epsilon \in \mathcal{E}} \frac{\chi_{(it)_{\epsilon},j}}{\sum_{j' \in \mathcal{J}_{\epsilon}} \chi_{(it)_{\epsilon},j'}(\lambda_{j'}^{(k)} + \beta \cdot s_{i_{\epsilon}}^{(\mathrm{SSS})}/a_{i_{\epsilon}})}$$

### Scan setup

#### "Patient 10":

F, 49 y.o., multifocal gastric neuroendocrine neoplasm tumour (NET G1, Ki67 2 percent)

#### Stats:

Ga-68-DOTATATE 158 MBq, 59 min wait, 20 min PET/CT scan (Biograph 64), *10 min scan (J-PET)* 

**2γ data**: 5.36M events;

 $2\gamma$  + prompt: 17.1k events (that's why <sup>68</sup>Ga)

#### (too few for Ps imaging!!!)

After geometrical pre-filtering (2 $\gamma$ ): ~ 3.7M coincidences

CT to the attenuation map – based on the work [Jonathan P. J. Carney, Med Phys 2006] + isocentres adjusted to match PET scan (*some ML techniques by Manish Das, J-PET collaboration*)

MIP along Z

#### Attenuation map (511 keV, cm<sup>-1</sup>)



MIP along Y

#### Results -22 Ш SSS sinograms segID Max value = 3.231525 -12 segID 4 segID = Ļ Ш segID zID = -12 zID = -7 zID = -3 zID = -1 zID = +1 zID = +3 zID = +7 zID = +12

#### J-PET group photo at the Medical University of Warsaw

# transverse

coronal



Iteration 1









20-

40

-8



Iteration 3



Results

#### Uncorrected image, maximum intensity projection (MIP) Iteration 3 Iteration 7

20-

-6

-8

-20

-9-

-0

Iteration 10



Iteration 10



#### Uncorrected images

It. 7, Y = -12 cm

-6

-8

-20

-9

0

coronal



No correction,

Cross-sections And MIPs

Axial smearing in J-PET is seen (FWHM ~ 1.5–2 cm)



lt. 7, MIPz





kBq/ml 10 zo zo do 40

It. 7, X = 8 cm



#### With SSS applied



Iteration 7 SSS correction,

Optimal norm:  $\beta = 0.0025$ 







-0



#### With SSS and regularisation (AMD-FMH)

#### It. 7, Y = -12 cm lt. 7, Z = -20.25 cm It. 7, X = 8 cm**Results** -8 -09 = 2.5 mm) 60 transverse sagittal coronal 50 20 20-40 -6 -6 one slice (d -8 8 -8 20 50 50kBq/ml -9-9 9 0 0 It. 7, MIPz It. 7, MIP<sub>Y</sub> It. 7, MIP<sub>X</sub> -9 -09 -99 Full MIP projections 20-20 -02 40 -4 -6 -8 30 30-20 20 20 kBq/ml -9--9-9

0

Iteration 7 SSS correction + Anisotropic Median-Diffusion [Jian Ling et al. IEEE TMI (2002)] + Finite-impluseresponse median hybrid filter (AMD-FMH) [Shopa R, RAP proc, 2021] **MLEM converges!** 

#### With SSS and regularisation (AMD-FMH)



#### Ways to improve

Test data selection using ML (ongoing studies lead by Konrad Klimaszewski)
Dedicated measurements with <sup>18</sup>F or <sup>22</sup>Na (negligible *positron range*) or account for M<sub>ps.rng</sub>

$$\mathbf{M} = \mathbf{M}_{\mathrm{norm}} \mathbf{M}_{\mathrm{det.res}} \mathbf{M}_{\mathrm{att}} \mathbf{M}_{\mathrm{Radon}} \mathbf{M}_{\mathrm{ps.rng}}$$

– Longer scans, longer FOV, proper Ps imaging







### Conclusions

- The first static clinical PET image acquired for the modular J-PET
- Realistic system matrix modelling that accounts for detector resolution is better than traditional PSF (e.g. in STIR or CASToR)
- Scatter correction introduced for the first time in the J-PET experiments, some issues with the normalisation yet to be resolved
- Regularisation by AMD-FHM further improves PET image as MLEM converges
- Ga-68-DOTATATE radiotracer can provide information on the positronium lifetime
- Still significant positron range to be resolved in the future

### Thank You for Your attention!



#### **Template header for SSS sinogram**

#### **!INTERFILE** := !imaging modality := PT name of data file := sss djpet50 updAct p10.s originating system := unknown !version of keys := STIR6.0 **!GENERAL DATA :=** GENERAL IMAGE DATA := type of data := PET patient orientation := head in patient rotation := supine What if other isotope? imagedata byte order := LITTLEENDIAN number of radionuclides := 1 radionuclide name[1] := ^18^Fluorine radionuclide halflife (sec)[1] := 6584.04 radionuclide branching factor[1] := 0.9686 !PET STUDY (General) := **!PET** data type **:= Emission** applied corrections := {None} !number format := float !number of bytes per pixel := 4 number of dimensions := 4 matrix axis label [4] := segment !matrix size [4] := 47 matrix axis label [3] := view !matrix size [3] := 48 matrix axis label [2] := axial coordinate !matrix size [2] := { 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,23,22,21,20,19,18,17,16,15,14,13,12,11,10,9,8,7,6,5,4,3,2,1} matrix axis label [1] := tangential coordinate !matrix size [1] := 31 minimum ring difference per segment := maximum ring difference per segment := -23,-22,-21,-20,-19,-18,-17,-16,-15,-14,-13,-12,-11,-10,-9,-8,-7,-6,-5,-4,-3,-2,-1,0,1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23} Scanner parameters:= Scanner type := unknown Simple cylindrical geometry Number of rings := 24 Number of detectors per ring := 96 **Inner** ring diameter (cm) := 73.872 Average depth of interaction (cm) := 0.01 **Distance** between rings (cm) **Default** bin size (cm) View offset (degrees) := 0 Maximum number of non-arc-corrected bins := 31 Default number of arc-corrected bins

#### **Template header for SSS sinogram**

Scanner parameters:=		
Scanner type := unknown		
Number of rings	:= 24	
Number of detectors per ring	:= 96	
<b>Inner</b> ring diameter (cm)	:= 73.872	
Average depth of interaction (cm)	:= 0.01	
Distance between rings (cm)	:= 2	
<b>Default</b> bin size (cm)	:= -1	
View offset (degrees)	:= 0	
Maximum number of non-arc-corrected bins	:= 31	
Default number of arc-corrected bins	:= 0	
Energy resolution := 0.16		
<b>Reference</b> energy (in keV) := 511		
Number of blocks per bucket in transaxia	l direction	:= 0
Number of blocks per bucket in axial dire	ection	:= 0
Number of crystals per block in axial di	rection	:= 0
Number of crystals per block in transaxia	al direction	:= 0
Number of detector layers		:= 1
Number of crystals per singles unit in a	xial direction	:= -1
Number of crystals per singles unit in the	ransaxial direction	:= -1
Scanner geometry (BlocksOnCylindrical/Cy	lindrical/Generic)	:= Cylindric
End scanner parameters:=		
number of time frames := 1	Enoro	
<pre>number of energy windows := 1</pre>		y I
energy window lower level[1] := 425	cuto	×
energy window upper level[1] := 650	Cuis	
start vertical bed position (mm) := 0		•
start horizontal bed position (mm) := 0		
!END OF INTERFILE :=		

Tangential: sAxial: ZView:  $\phi$ Segment:  $\theta$ 





# Single scatter simulation (SSS)





\*Primed quantities are evaluated at the scattered photon's energy, while unprimed ones are evaluated at 511 keV.