

User's guide

dPETSTEP simulation GUI

This document shows the usage of the dPETSTEP GUI for PET image simulation, and explains the different inputs to the GUI.

The GUI can be run by typing in the MATLAB command prompt:

```
>> dPETSTEPgui_sim
```

This command opens the full window seen below. All fields have tool tips to guide you.

dPETSTEP simulation

Input data

Data

☒ From WS ☐ From file

Name : (Bq/cc)

Time frame

☒ From WS ☐ From file

Name : (sec)

Scanner settings

Sinogram radial size

☒ Radial bins ☐ Scanner diameter

Radial bins : (near)

Ring diameter: mm

Angular bins:

System PSF: mm

Maximum ring diff:

System sensitivity: cnts per kBq/cc

Acquisition settings

Replicates :

Counts scale factor :

Scatter fraction:

Random fraction:

Nuclide half-life: sec

☒ No decay

Kinetic modelling

Input function

☒ From WS ☐ From file

Name : (Bq/cc)

Scale factor:

Model

☒ 1-Tissue ☐ FRTM ☐ SumExp

☐ 2-Tissue ☐ SRTM # exp:

Interpolation

☒ Linear ☐ Nearest ☐ PCHIP ☐ Spline

Time step: sec

Biologic variability

☒ None ☐ Scale:

Reconstruction settings

Image matrix size:

FOV: mm

Method

☐ FBP ☐ OSEM ☐ OSEM w/ PSF

iterations:

Subsets:

PSF correction: mm

Postfiltering

☐ XY: XY FWHM: mm

☐ Z: ☒ From WS ☐ From file

Name:

Output data names

☒ Data:

☒ Counts:

☒ Counts noisy:

☒ Prompts sino:

☒ Trues sino:

☒ Scatters sino:

☒ Randoms sino:

☒ WCC:

☐ FBP:

☐ OSEM:

☐ OSEM w/ PSF:

Buttons: Save Load Reset Run

Created by Ida Häggström, 2017.07.07

Buttons for type of simulation

Parametric image input **Dynamic image input** → Start with pristine dynamic uptake image to simulate realistic PET scan of it.

Start with parametric image, which calculates to a pristine dynamic uptake image which is used to simulate realistic PET scan.

Input data

Input data

Data

☒ From WS
Name :

Time frame

☒ From WS ☐ From file
Name : (sec)

Read data structure (containing attenuation data, and parametric image or dynamic image) either from workspace (WS), or from *.mat-file. Enter the name (string) of the WS variable/file. The *.mat-file has to contain ONLY the data structure, no other data.

For parametric image input, the structure should contain parametric image.

For dynamic image input, the structure should contain dynamic uptake image.

Read frame time points either from workspace (WS), or from *.mat, *.txt, or *.xls file. Enter the name (string) of the WS variable/file. The data file has to contain ONLY the frame vector, no other data.

The frame data vector [nt+1,1] should be in unit (sec).

Scanner settings

Scanner settings

Sinogram radial size

☒ Radial bins ☐ Scanner diameter

Radial bins : (near)

Ring diameter: mm

Angular bins:

System PSF: mm

Maximum ring diff:

System sensitivity: cnts per kBq/cc

Choose if you want to set the number of radial bins directly, or have it calculated indirectly via the scanner diameter.

Number of radial bins is an integer number, and the program will get to the desired number as closely as possible (uneven number).

The ring diameter is a number in unit (mm).

The number of angular bins is an integer number.

The system PSF is the blurring point spread function FWHM of the imaging system in unit (mm).

The maximum ring difference is an integer number with max separation between scanner rings allowed for coincidences.

The system sensitivity is the number of counts per kBq/cc for the particular scanner.

Acquisition settings

Acquisition settings

Replicates :

Counts scale factor :

Scatter fraction:

Random fraction:

Nuclide half-life: sec

☒ No decay

Number of replicates is an integer with number of desired noise realizations of the simulation.

Counts scale factor is a factor that scales the dynamic image activity, in turn number of counts (noise level).

Scatter fraction is a unitless number, $SF=S/(T+S)$.

Random fraction is a unitless number $RF=R/(T+S+R)$.

Uncheck if you want to simulate physical decay, and specify nuclide half-life in unit (sec).

Kinetic modeling

Kinetic modelling

Input function

☒ From WS ☐ From file

Name : (Bq/cc)

Scale factor:

Model

☒ 1-Tissue ☐ FRTM ☐ SumExp

☐ 2-Tissue ☐ SRTM # exp:

Interpolation

☒ Linear ☐ Nearest ☐ PCHIP ☐ Spline

Time step: sec

Biologic variability

☒ None ☐ Scale:

Read input/reference function vector either from workspace (WS), or from *.mat, *.txt or *.xls-file. Enter the name (string) of the WS variable/file. The data file has to contain ONLY the input/ref function vector, no other data.

The input/ref function vector [nt,1] should be in units (Bq/cc).

The unitless scale factor is a multiplicative scalar that scales the amplitude of the input function. Set to 1 for no scaling.

Choose which kinetic model to simulate. For general sum of exponentials model (SumExp), also specify number of exponentials (integer number).

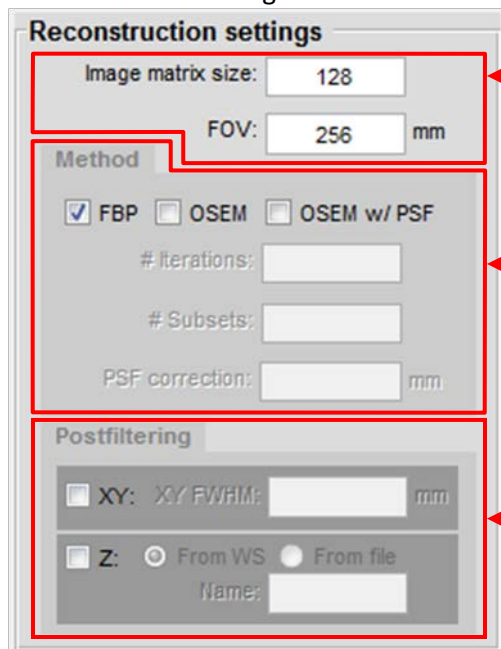
If uneven time sampling, the data will be interpolated to even sampling (for convolution) during fitting. Choose interpolation method, and time step for the interpolation.

The time step is a number in units (sec).

Chose if you want to simulate biologic variability by setting the scale to a unitless number.

Scale of added varaibility: variance of Gaussian noise = image value/scale.

Reconstruction settings



The 'Reconstruction settings' panel contains three main sections. The top section, 'Image matrix size', has a text box for 'Image matrix size' (value 128) and a text box for 'FOV' (value 256) followed by 'mm'. The middle section, 'Method', has three checkboxes: 'FBP' (checked), 'OSEM' (unchecked), and 'OSEM w/ PSF' (unchecked). Below these are text boxes for '# iterations' and '# Subsets'. The bottom section, 'Postfiltering', has a checkbox for 'XY' (unchecked) followed by 'XY FWHM' (text box) and 'mm'. Below that is a checkbox for 'Z' (unchecked) followed by radio buttons for 'From WS' (selected) and 'From file' (unchecked), and a 'Name' text box.

Set desired simulated image matrix XY-size (integer number).

Set image XY field of view (FOV) in (mm). Pixel size will be FOV/matrix size.

Check which type of reconstructed images you want to simulate; filtered back projection (FBP), ordered subset expectation maximization (OSEM), and/or OSEM with point spread function (PSF) correction.

For OSEM and OSEM+PSF, specify number of iterations (integer) and number of subsets (integer).

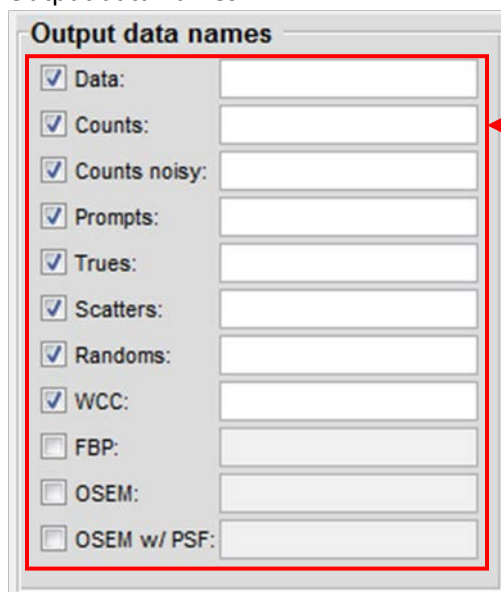
For OSEM+PSF, specify the correction PSF FWHM in (mm).

Check boxes if you want to perform XY and/or Z postfiltering.

Set the XY-postfilter Gaussian FWHM in (mm).

Read the Z-filter either from workspace (WS), or from *.mat, *.txt or *.xls-file. Enter the name (string) of the WS variable/file.

Output data names



The 'Output data names' panel has a list of checkboxes on the left and corresponding text boxes on the right. The checkboxes are: 'Data' (checked), 'Counts' (checked), 'Counts noisy' (checked), 'Prompts' (checked), 'Trues' (checked), 'Scatters' (checked), 'Randoms' (checked), 'WCC' (checked), 'FBP' (unchecked), 'OSEM' (unchecked), and 'OSEM w/ PSF' (unchecked). Each checked checkbox has a text box next to it for naming the output data.

Choose what data you want as output from the simulation, and set the name (string) of that data. Will end up in your WS after run.

Data is the structure used as input, with (for parametric image input) a field of the pristine dynamic uptake image added.

Counts is the noiseless sinogram counts of prompts, trues, scatters and randoms.

Counts noisy is the noisy sinogram counts of prompts, trues, scatters and randoms.

Prompts is the prompts sinogram matrix.

Trues is the trues sinogram matrix.

Scatters is the scatters sinogram matrix.

Randoms is the randoms sinogram matrix.

WCC is the well counter calibration factor to calibrate reconstructed image to (Bq/cc).

FBP is the reconstructed 4D FBP image.

OSEM is the reconstructed 4D OSEM image. Each iteration is saved.

OSEM w/ PSF is the reconstructed 4D OSEM w/ PSF correction image. Each iteration is saved.

Buttons



Load settings from chosen *.xls-file.

Reset all settings.

Executes the simulation main program.

Save all current settings to *.xls-file.