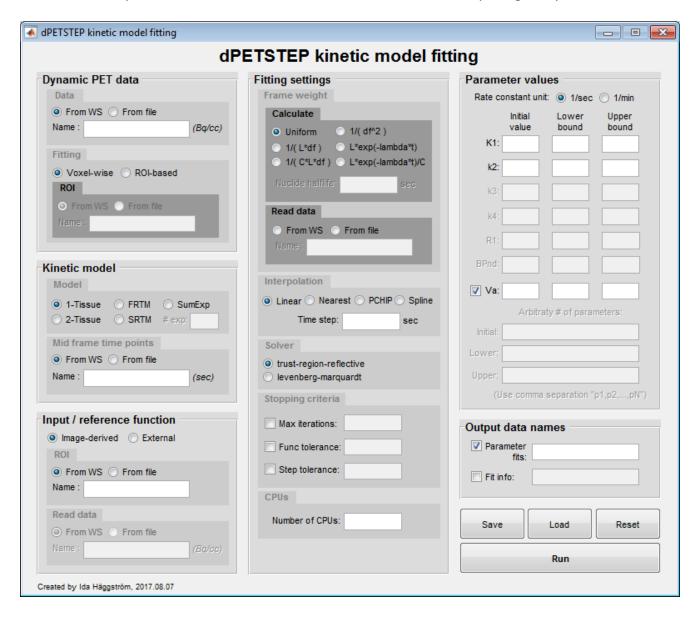
User's guide dPETSTEP kinetic model fitting GUI

This document shows the usage of the dPETSTEP GUI for kinetic model fitting, and explains the different inputs to the GUI.

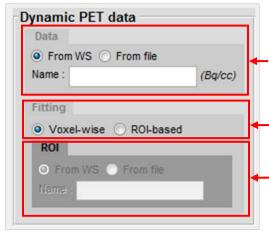
The GUI can be run by typing in the MATLAB command prompt:

>> dPETSTEPgui_fit

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



Dynamic PET data



Read dynamic (4D) PET uptake image matrix 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 dynamic image matrix, no other data.

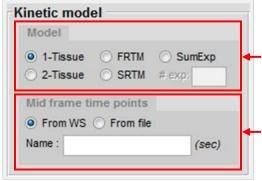
The 4D data matrix [nx,ny,nz,nt] should be in units (Bq/cc).

Fit each voxel of the 4D image to get parametric image, or fit only single ROI time-activity curve.

For ROI-based fitting, enter the name (string) of the WS variable/*.mat file. The data file has to contain ONLY the ROI mask matrix, no other data.

The ROI is a mask matrix [nx,ny,nz] of only values 0 and 1.

Kinetic model

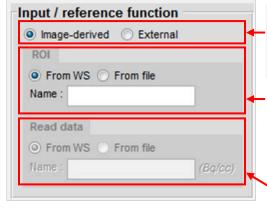


Choose which kinetic model to fit to. For general sum of exponentials model (SumExp), also specify number of exponentials (integer number).

Read mid 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 mid frame vector, no other data.

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

Input/reference function



Choose image-derived (from ROI) input/reference function, or an external one. Used as arterial input function for kinetic models 1-tissue, 2-tissue and sum of exponentials, and as reference function for FRTM and SRTM.

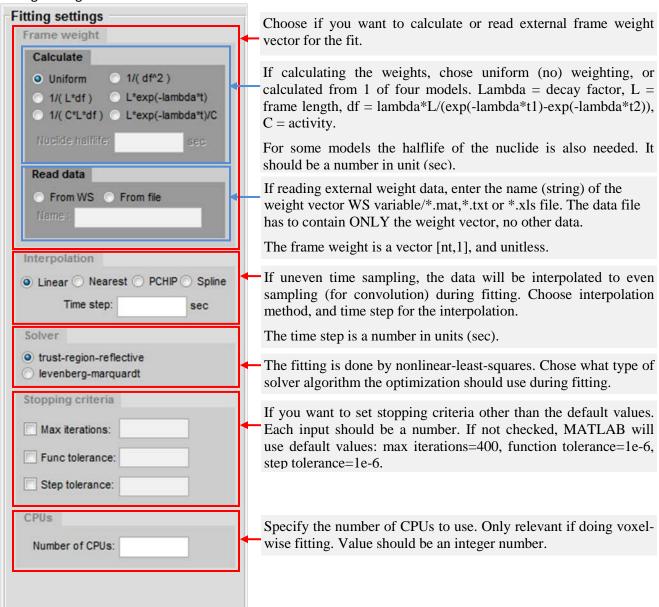
For image-derived input/reference function, enter the name (string) of the ROI mask WS variable/*.mat file. The data file has to contain ONLY the ROI mask matrix, no other data.

The ROI is a mask matrix [nx,ny,nz] of only values 0 and 1.

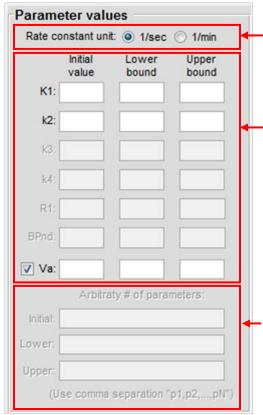
For external input/reference function, enter the name (string) of the WS variable/*.mat,*.txt or *.xls file. The data file has to contain ONLY the input/ref function vector, no other data.

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

Fitting settings



Parameter values



Set unit of parameter initial values, upper and lower bounds. Only relevant for parameters K1, k2, k3, k4, and sum of exponentials parameters.

Set initial values of the parameters relevant to your chosen kinetic model. If using the trust-region-reflective algorithm, also set lower and upper bounds of the parameter estimates.

You can choose to include the blood fraction (Va) in the fitting or not by checking the Va box.

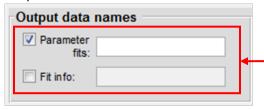
All parameter values should be numbers. K1 in unit (cc/g/sec) or (cc/g/min), k2, k3 and k4 in (1/sec) or (1/min), R1 and BPnd are unitless, and Va in (cc/g).

If chosen kinetic model is an arbitrary sum of exponentials, you specify initial values, lower and upper bounds here.

The number of values specified should be 2*number of exp. You can still add the blood fraction by checking the Va box above.

The inputs should be numbers, and values should be comma separated, like "0.1,0.01,0.01,0.01" (w/o quotation marks).

Output data names

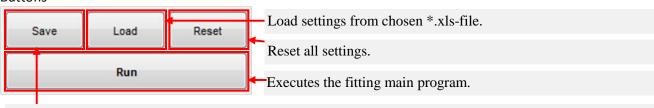


Choose what data you want as output from the fit, and set the name (string) of that data. Will end up in your WS after run.

You can get the parameter fits (parametric image matrix for voxel-wise fit, vector for ROI fit), and the fitting information.

The fitted parameters will be (depending on kinetic model) K1 in unit (cc/g/sec) or (cc/g/min), k2, k3 and k4 in (1/sec) or (1/min), R1 and BPnd unitless, and Va in (ml/g). The fitting information is a structure with resnorm, residual, exitflag, output, lambda, and jacobian. Note that for voxel-wise fitting, there will be one of each structure field per voxel, leading to a potentially very large data structure.

Buttons



Save all current settings to *.csv-file.