

PESTO

1.0

Generated by Doxygen 1.8.11

## Contents

<b>1</b>	<b>PESTO Documentation</b>	<b>2</b>
1.1	Introduction . . . . .	2
1.2	Availability . . . . .	2
1.3	Installation . . . . .	2
1.4	Licensing . . . . .	3
1.5	How to cite . . . . .	3
1.6	Code organization . . . . .	3
1.7	Features . . . . .	3
1.7.1	Global optimization . . . . .	3
1.7.2	Uncertainty analysis . . . . .	4
1.7.3	Parameter sampling . . . . .	4
1.7.4	Plotting . . . . .	5
1.7.5	Properties . . . . .	7
<b>2</b>	<b>Examples</b>	<b>8</b>
2.1	Examples . . . . .	8
2.1.1	Overview . . . . .	8
<b>3</b>	<b>Hierarchical Index</b>	<b>9</b>
3.1	Class Hierarchy . . . . .	9
<b>4</b>	<b>Class Index</b>	<b>10</b>
4.1	Class List . . . . .	10
<b>5</b>	<b>File Index</b>	<b>10</b>
5.1	File List . . . . .	10

<b>6</b>	<b>Class Documentation</b>	<b>12</b>
6.1	DRAMOptions Class Reference . . . . .	12
6.1.1	Detailed Description . . . . .	14
6.1.2	Constructor & Destructor Documentation . . . . .	14
6.1.3	Member Data Documentation . . . . .	14
6.2	MALAOptions Class Reference . . . . .	15
6.2.1	Detailed Description . . . . .	16
6.2.2	Constructor & Destructor Documentation . . . . .	16
6.2.3	Member Data Documentation . . . . .	17
6.3	PestoOptions Class Reference . . . . .	17
6.3.1	Detailed Description . . . . .	20
6.3.2	Constructor & Destructor Documentation . . . . .	20
6.3.3	Member Data Documentation . . . . .	20
6.4	PestoPlottingOptions Class Reference . . . . .	31
6.4.1	Detailed Description . . . . .	33
6.4.2	Constructor & Destructor Documentation . . . . .	33
6.4.3	Member Data Documentation . . . . .	33
6.5	PestoSamplingOptions Class Reference . . . . .	44
6.5.1	Detailed Description . . . . .	45
6.5.2	Constructor & Destructor Documentation . . . . .	46
6.5.3	Member Function Documentation . . . . .	46
6.5.4	Member Data Documentation . . . . .	46
6.6	PHSOptions Class Reference . . . . .	50
6.6.1	Detailed Description . . . . .	51
6.6.2	Constructor & Destructor Documentation . . . . .	51
6.6.3	Member Data Documentation . . . . .	51
6.7	PTOptions Class Reference . . . . .	53
6.7.1	Detailed Description . . . . .	54
6.7.2	Constructor & Destructor Documentation . . . . .	54
6.7.3	Member Data Documentation . . . . .	54

<b>7 File Documentation</b>	<b>56</b>
7.1 collectResults.m File Reference . . . . .	56
7.1.1 Detailed Description . . . . .	56
7.1.2 Function Documentation . . . . .	56
7.2 getMultiStarts.m File Reference . . . . .	57
7.2.1 Detailed Description . . . . .	58
7.2.2 Function Documentation . . . . .	58
7.3 getParameterConfidenceIntervals.m File Reference . . . . .	60
7.3.1 Detailed Description . . . . .	60
7.3.2 Function Documentation . . . . .	61
7.4 getParameterProfiles.m File Reference . . . . .	62
7.4.1 Detailed Description . . . . .	62
7.4.2 Function Documentation . . . . .	62
7.5 getParameterSamples.m File Reference . . . . .	64
7.5.1 Detailed Description . . . . .	65
7.5.2 Function Documentation . . . . .	65
7.6 getPropertyConfidenceIntervals.m File Reference . . . . .	66
7.6.1 Detailed Description . . . . .	66
7.6.2 Function Documentation . . . . .	66
7.7 getPropertyMultiStarts.m File Reference . . . . .	68
7.7.1 Detailed Description . . . . .	68
7.7.2 Function Documentation . . . . .	68
7.8 getPropertyProfiles.m File Reference . . . . .	70
7.8.1 Detailed Description . . . . .	70
7.8.2 Function Documentation . . . . .	71
7.9 getPropertySamples.m File Reference . . . . .	72
7.9.1 Detailed Description . . . . .	73
7.9.2 Function Documentation . . . . .	73
7.10 meigoDummy.m File Reference . . . . .	74
7.10.1 Detailed Description . . . . .	75

7.10.2	Function Documentation	75
7.11	plotConfidenceIntervals.m File Reference	75
7.11.1	Detailed Description	75
7.11.2	Function Documentation	76
7.12	plotMCMCdiagnosis.m File Reference	77
7.12.1	Detailed Description	77
7.12.2	Function Documentation	77
7.13	plotMultiStarts.m File Reference	78
7.13.1	Detailed Description	78
7.13.2	Function Documentation	79
7.14	plotParameterProfiles.m File Reference	80
7.14.1	Detailed Description	80
7.14.2	Function Documentation	80
7.15	plotParameterSamples.m File Reference	82
7.15.1	Detailed Description	82
7.15.2	Function Documentation	82
7.16	plotParameterUncertainty.m File Reference	83
7.16.1	Detailed Description	83
7.16.2	Function Documentation	84
7.17	plotPropertyMultiStarts.m File Reference	85
7.17.1	Detailed Description	85
7.17.2	Function Documentation	85
7.18	plotPropertyProfiles.m File Reference	86
7.18.1	Detailed Description	87
7.18.2	Function Documentation	87
7.19	plotPropertySamples.m File Reference	88
7.19.1	Detailed Description	88
7.19.2	Function Documentation	89
7.20	plotPropertyUncertainty.m File Reference	90
7.20.1	Detailed Description	90
7.20.2	Function Documentation	90
7.21	runPestoTests.m File Reference	91
7.21.1	Detailed Description	92
7.22	testGradient.m File Reference	92
7.22.1	Detailed Description	92
7.22.2	Function Documentation	92

## 1 PESTO Documentation

### 1.1 Introduction

Computational models are commonly used in diverse disciplines such as computational biology, engineering, or meteorology. The parameterization of these models is usually based on measurements or observations. The process of inferring model parameters from such data is called model calibration or parameter estimation. This parameter estimation is often not straightforward due to non-linearities in the model equations or due to the mere size and the resulting computational challenges. Therefore, efficient algorithms are required to provide robust results within acceptable time.

PESTO is a freely available Parameter EStimation TOolbox for MATLAB (MathWorks) implementing a number of state-of-the-art algorithms for parameter estimation. It provides the following features, which are explained in more detail [below](#):

- Parameter estimation from measurement data by global optimization based on multi-start local optimization (some algorithms require the [MATLAB Optimization Toolbox](#))
- Parameter sampling using Markov Chain Monte Carlo (MCMC) algorithms
- Uncertainty analysis based on local approximations, parameter samples and profile-likelihood analysis (some algorithms require the [MATLAB Optimization Toolbox](#))
- Visualization routines for all analyses
- Parallel processing (requires [MATLAB Parallel Computing Toolbox](#))
- ...

PESTO functions can be applied to any user-provided formulation of an optimization problem with an objective function that can be evaluated in MATLAB. Besides the objective function, upper and lower bounds for the function parameters need to be specified.

### 1.2 Availability

PESTO can be freely obtained from <https://github.com/ICB-DCM/PESTO/> by downloading the zip archive at <https://github.com/ICB-DCM/PESTO/archive/master.zip> or cloning the `git` repository via

```
1 git clone git@github.com:ICB-DCM/PESTO.git
```

### 1.3 Installation

If the zip archive was downloaded, it needs to be unzipped and the main folder has to be added to the MATLAB search path (non-recursively).

If the repository was cloned, the main folder needs to be added to the MATLAB search path (non-recursively).

*Note:* Detailed instructions on how to modify your MATLAB search path are provided on the [web](#).

### Third-party packages

PESTO provides an interfaces to several other toolboxes which are not included in the PESTO archive:

- PSwarm (optimizer): <http://www.norg.uminho.pt/aivaz/pswarm/>
- MEIGO (optimizer): <http://gingproc.iim.csic.es/meigo.html>
- DRAM (MCMC): <http://helios.fmi.fi/~lainema/dram/>

To use their functionality, these toolboxes have to be installed separately. Please consult the respective user manuals for details.

## 1.4 Licensing

See `LICENSE` file in the PESTO source directory.

## 1.5 How to cite

This section will be updated upon publication of PESTO.

## 1.6 Code organization

The end-user interface is provided by the MATLAB functions and classes in the top-level directory. PESTO [example](#) applications are provided in `/examples/`. All other folders only contain files used internally in PESTO.

## 1.7 Features

PESTO implements a number of state-of-the-art algorithms related to parameter estimation. The main features are described below. Various [examples](#) demonstrate their application.

### Notations and Terminology

Since most of the examples use analytical approaches for computing the gradient of the respective objective function, which quantifies the deviation of the fit for the current model parameters from the actual measurement data, the usage of the term ‚sensitivity analysis‘ may be misleading. In our context, ‚sensitivity analysis‘ is used in the context of ODE or PDE models and describes the sensitivity of the ODE/PDE state with respect to the model parameters. Those state sensitivities can be implemented in the ODE/PDE system and then used for an analytical calculation of the sensitivity of the objective function. This objective functions sensitivity will always be called the objective function gradient in our context. Finally, the behavior of the objective function by the variation of single parameters in order to find possible (non-)identifiabilities will always be referred to as ‚uncertainty analysis‘.

### 1.7.1 Global optimization

Non-linear optimization problems like those in parameter estimation problems tend to have multiple optima. Usually, nothing is known beforehand about their number or their location, but the user is interested in finding the global optimum. There are different techniques for this kind of problem. PESTO provides a multi-start local optimization framework and provides an interface to two global optimizers.

## Multi-start local optimization

Multi-start local optimization has turned out to be a very efficient method for "global optimization": Here, random points from across the parameter space are chosen as starting points for local optimization. If an adequate number of starting points spanning the domain of interest of the parameter space is selected, the lowest/highest minimum/maximum is accepted to be the global minimum/maximum. By default, `fmincon` from the MATLAB optimization toolbox is used as a local solver.

Multi-start local optimization functionality is provided by `getMultiStarts.m`, `getPropertyMultiStarts.m` and the respective plotting routines `plotMultiStarts.m` and `plotPropertyMultiStarts.m`. See `examples/conversion_reaction/mainConversionReaction.m` for an example.

## Global optimizers

PESTO provides an interface to `PSwarm` and `MEIGO`. Once these toolboxes have been installed - they are not included in the PESTO archive - they can be used for parameter estimation. These optimizers are also accessed via `getMultiStarts.m` by setting `PestoOptions::localOptimizer` and `PestoOptions::localOptimizerOptions` accordingly. In principle, a single optimizer run (`PestoOptions::n_starts = 1`) should be enough for these global optimizers.

An example is included in `mainConversionReaction.m`.

### 1.7.2 Uncertainty analysis

When parameters are inferred from measurement data, the deviation of the data from the fit for the best parameter guess is usually assumed to be of stochastic nature. This means that the estimated parameters themselves are stochastic and underly an uncertainty. This can be quantified by performing uncertainty analysis and computing confidence intervals.

The easiest way to do this is using local approximations (based on the Hessian matrix of the objective function) at the best parameter guess. From those approximations, either threshold-based or mass-based methods can be used to compute confidence intervals for the inferred parameters. Another approach uses sampling based methods in combination with local approximations.

The most reliable way to compute confidence intervals is a third approach, based on profile likelihoods. Here, each model parameter is varied separately while the others are constantly reoptimized. In this way one finds profiles for every parameter. By fixing a confidence level using the inverse chi-squared-distribution, one gets a threshold which, together with the profile likelihood, gives reliable confidence intervals for each parameter. In this way, non-identifiable parameter can be detected.

Those functionalities are provided in `getParameterProfiles.m`, `getPropertyProfiles.m` (for the profile likelihoods), `getParameterConfidenceIntervals.m` and `getPropertyConfidenceIntervals.m` (for the confidence intervals). In order to get confidence intervals based on local approximations or sampling methods, one needs to run the routines `getMultiStarts.m` / `getPropertyMultiStarts.m` or `getParameterSamples.m` / `getPropertySamples.m` first. The respective visualization routines are `plotParameterProfiles.m` and `plotPropertyProfiles.m`. See `mainConversionReaction.m` for an example.

### 1.7.3 Parameter sampling

PESTO provides Markov Chain Monte Carlo (MCMC) algorithms for sampling the posterior distribution. Sampling methods such as the `Metropolis-Hastings (MH)`, adaptive Metropolis (AM) and `Metropolis-adjusted Langevin algorithm (MALA)` are currently implemented. Additionally, PESTO provides an interface to the `Delayed Rejection Adaptive Metropolis (DRAM)` toolbox.

See `getParameterSamples()` for details and `mainConversionReaction.m` for examples.



## 1.7.4 Plotting

An integral part of PESTO are its highly customizable plotting functions for each type of analysis.

Details are provided in the documentation of the specific plotting functions:

- [plotMultiStarts.m](#)
- [plotParameterProfiles.m](#)
- [plotParameterSamples.m](#)
- [plotParameterUncertainty.m](#)
- [plotPropertyMultiStarts.m](#)
- [plotPropertyProfiles.m](#)
- [plotPropertySamples.m](#)
- [plotPropertyUncertainty.m](#)

Here some examples:

Plot of model fit using [plotMultiStarts.m](#):

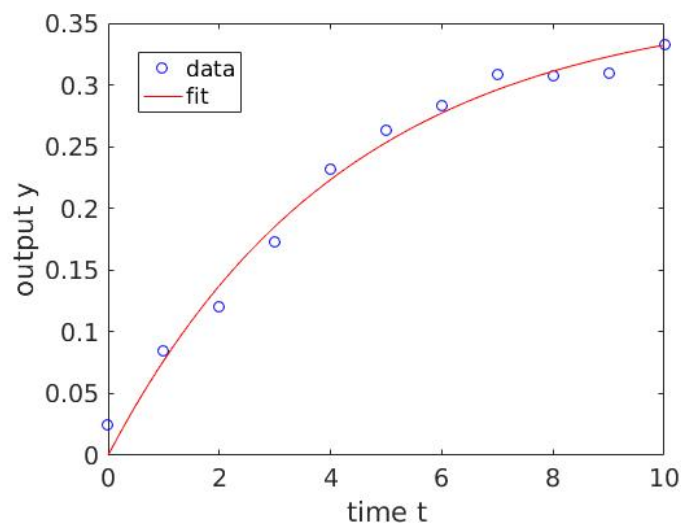


Figure 1 Plot of model fit using [plotMultiStarts.m](#)

Plot of different variants of parameter confidence intervals using [plotParameterUncertainty.m](#):

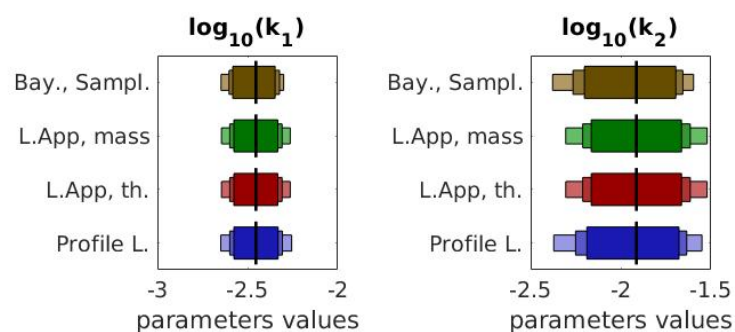
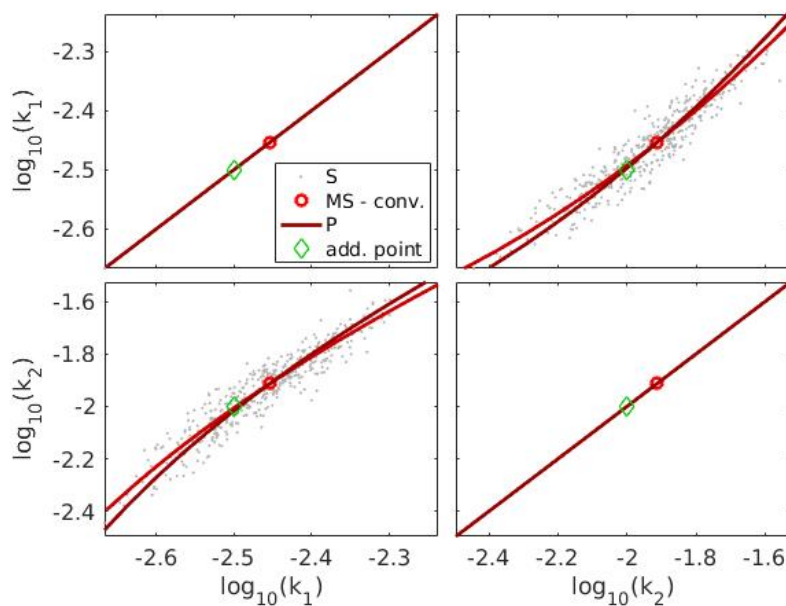


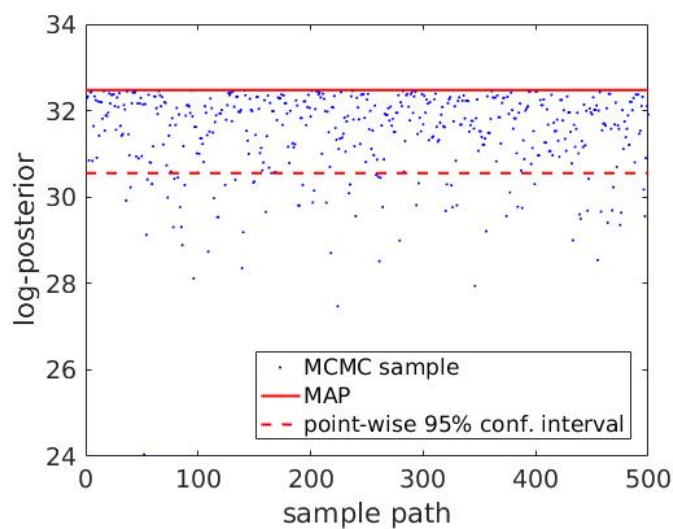
Figure 2 Plot of different variants of parameter confidence intervals using [plotParameterUncertainty.m](#)

2D plot of parameter samples using [plotParameterSamples.m](#):



**Figure 3** 2D plot of parameter samples using [plotParameterSamples.m](#)

Plot of parameter samples using [plotParameterSamples.m](#):



**Figure 4** Plot of parameter samples using [plotParameterSamples.m](#)

Plot of property samples using [plotPropertySamples.m](#):

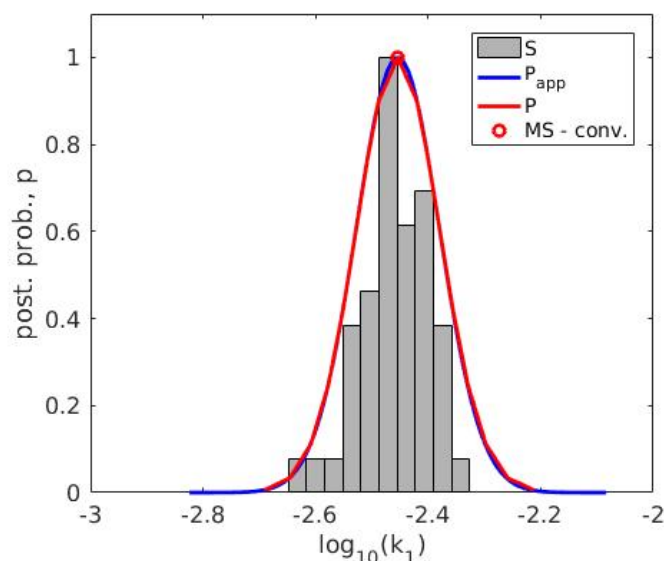


Figure 5 Plot of property samples using `plotPropertySamples.m`

See `mainConversionReaction.m` for live examples.

### 1.7.5 Properties

The above-mentioned methods for parameter estimation, confidence intervals, parameter profiles and parameter samples (`getMultiStarts.m`, `getParameterConfidenceIntervals.m`, `getParameterProfiles.m`, `getParameterSamples.m`) all operate on the objective function parameters directly. However, sometimes not the parameters themselves, but some function thereof is of interest. To this end, PESTO provides a simple interface to achieve this without having to change the objective function. Arbitrary user-provided functions which take the objective function parameter vector as an argument are referred to as 'properties'. After having used any of the `getParameter*.m` functions, the respective `getProperty*.m` function can be called, to evaluate a user-provided property function with the parameters values/samples/confidences obtained from the `getParameter*.m` functions.

The following functions are available to analyze and plot properties:

- `getPropertyConfidenceIntervals.m`
- `getPropertyMultiStarts.m`
- `getPropertyProfiles.m`
- `getPropertySamples.m`
- `plotPropertyMultiStarts.m`
- `plotPropertyProfiles.m`
- `plotPropertySamples.m`
- `plotPropertyUncertainty.m`

See `mainConversionReaction.m` for examples.

## 2 Examples

### 2.1 Examples

#### 2.1.1 Overview

PESTO comes with a number of ready-to-run examples to demonstrate its usage and functionality. The examples mostly stem from computational biology and comprise ordinary and partial differential equation (ODE / PDE), and Gaussian mixture models. More background information is provided in the respective example folder in the `main*.m` script.

The following examples are included:

- *Conversion reaction*, the simplest example and demonstrating all PESTO features (`examples/conversion_reaction/mainConversionReaction.m`, ODE)
- *Enzymatic catalysis* (`examples/enzymatic_catalysis/mainEnzymaticCatalysis.m`, ODE)
- *Gaussian mixture* (`examples/GaussExample/mainExampleGauss.m`)
- *Hyperring* (`examples/RingExample/mainExampleRing.m`)
- *mRNA transfection* § (`examples/mRNA_transfection/mainTransfection.m`, ODE)
- *Pom1p gradient formation* § (`examples/Pom1p_gradient_formation/mainPom1.m`, PDE)
- *Jak-Stat-signaling* § (`examples/jakstat_signaling/mainJakstatSignaling.m`, ODE)
- *ERBB signaling*, largest model among the examples § (`examples/erbb_signaling/mainErbBSignaling.m`, ODE)

§ These models require the freely available AMICI toolbox to run (<http://icb-dcm.github.io/AMICI/>).

The following table provides an overview of which of the PESTO functions are demonstrated in the different examples:

	conversion reaction	enzymatic catalysis	Gaussian mixture	Hyperring	transfection	Pom1p	JakStat	ErbB
<code>getMultiStarts()</code>	X	X		X	X	X	X	X
<code>plotMultiStarts()</code>	X	X		X	X	X	X	X
<code>getParameterConfidenceIntervals()</code>	X	X		X	X			
<code>plotConfidenceIntervals()</code>	X	X		X	X			
<code>getParameterProfiles()</code>	X	X		X	X			
<code>plotParameterProfiles()</code>	X	X	X	X	X	X		

	conversion reaction	enzymatic catalysis	Gaussian mixture	Hyperring	transfection	Pom1p	JakStat	ErbB
<a href="#">get↔ Parameter↔ Samples()</a>	X	X	X	X	X			
<a href="#">plot↔ Parameter↔ Samples()</a>	X	X	X	X	X			
<a href="#">get↔ Property↔ Multi↔ Starts()</a>	X				X			
<a href="#">plot↔ Property↔ Multi↔ Starts()</a>	X				X			
<a href="#">get↔ Property↔ Confidence↔ Intervals()</a>	X				X			
<a href="#">get↔ Property↔ Profiles()</a>	X				X			
<a href="#">plot↔ Property↔ Profiles()</a>	X				X			
<a href="#">get↔ Property↔ Samples()</a>	X				X			
<a href="#">plot↔ Property↔ Samples()</a>	X				X			
<a href="#">test↔ Gradient()</a>								
<a href="#">collect↔ Results()</a>								

### 3 Hierarchical Index

#### 3.1 Class Hierarchy

This inheritance list is sorted roughly, but not completely, alphabetically:

SetGet

<b>DRAMOptions</b>	<b>12</b>
<b>MALAOptions</b>	<b>15</b>
<b>PestoOptions</b>	<b>17</b>
<b>PestoPlottingOptions</b>	<b>31</b>
<b>PestoSamplingOptions</b>	<b>44</b>

<b>PHSOptions</b>	<b>50</b>
<b>PTOptions</b>	<b>53</b>

## 4 Class Index

### 4.1 Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

<b>DRAMOptions</b>	
<b>DRAMOptions</b> provides an option container to pass options as subclass into the <b>PestoSamplingOptions</b> class	<b>12</b>
<b>MALAOptions</b>	
<b>MALAOptions</b> provides an option container to pass options as subclass into the <b>PestoSamplingOptions</b> class	<b>15</b>
<b>PestoOptions</b>	
<b>PestoOptions</b> provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details	<b>17</b>
<b>PestoPlottingOptions</b>	
<b>PestoPlottingOptions</b> is a class for checking and holding information on optimization parameters	<b>31</b>
<b>PestoSamplingOptions</b>	
<b>PestoSamplingOptions</b> provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details	<b>44</b>
<b>PHSOptions</b>	
<b>PHSOptions</b> provides an option container to pass options as subclass into the <b>PestoSamplingOptions</b> class	<b>50</b>
<b>PTOptions</b>	
<b>PTOptions</b> provides an option container to pass options as subclass into the <b>PestoSamplingOptions</b> class	<b>53</b>

## 5 File Index

### 5.1 File List

Here is a list of all documented files with brief descriptions:

<b>collectResults.m</b>	
<b>CollectResults()</b> collects and plots the results stored in a common folder	<b>56</b>
<b>getMultiStarts.m</b>	
<b>GetMultiStarts()</b> computes the maximum a posterior estimate of the parameters of a user-supplied posterior function. Therefore, a multi-start local optimization is used. The parameters from the best value of the posterior function are then used as the global optimum. To ensure that the found maximum is a global one, a sufficiently high number of multistarts must be done. Those starts can be initialized with either randomly sampled parameter values, following either a uniform distribution or a latin hypercube, or they can be sampled by a user provided initial function (provided as option.init_fun)	<b>57</b>

**getParameterConfidenceIntervals.m**

GetParameterConfidenceIntervals() calculates the confidence intervals for the model parameters. This is done by four approaches: The values of CI.local\_PL and CI.PL are determined by the point on which a threshold according to the confidence level alpha (calculated by a chi2-distribution) is reached. local\_PL computes this point by a local approximation around the MAP estimate using the Hessian matrix, PL uses the profile likelihoods instead. The value of CI.local\_B is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate. The value of CI.S is calculated using samples for the model parameters and the according percentiles based on the confidence levels alpha

60

**getParameterProfiles.m**

GetParameterProfiles.m calculates the profiles likelihoods for the model parameters, starting from the maximum a posteriori estimate. This calculation is done by fixing the i-th parameter and repeatedly reoptimizing the likelihood/posterior estimate (for all i). The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization

62

**getParameterSamples.m**

GetParameterSamples.m performs MCMC sampling of the posterior distribution. Note, the D←RAM library routine tooparameters.minox is used internally. This function is capable of sampling with MH, AM, DRAM, MALA, PT and PHS. The sampling plotting routines should no longer be contained in here but as standalone scripts capable of using the resulting par.S

64

**getPropertyConfidenceIntervals.m**

GetPropertyConfidenceIntervals.m calculates the confidence intervals for the model properties. This is done by three approaches: The values of CI.local\_PL and CI.PL are determined by the point on which a threshold according to the confidence level alpha (calculated by a chi2-distribution) is reached. local\_PL computes this point by a local approximation around the MAP estimate using the Hessian matrix, PL uses the profile likelihoods instead. The value of CI.local\_B is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate

66

**getPropertyMultiStarts.m**

GetPropertyMultiStarts.m evaluates the properties for the different mutli-start results

68

**getPropertyProfiles.m**

GetPropertyProfiles.m calculates the profiles of user-supplied property functions, starting from the maximum a posteriori estimate. This calculation is done by varying the value of each property function respectively, starting from the value of this function at the global optimum and by reoptimizing the likelihood/posterior estimate in each variational step of the property. The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization

70

**getPropertySamples.m**

GetPropertySamples.m evaluates the properties for the sampled parameters

72

**meigoDummy.m**

Objective function wrapper for MEIGO / PSwarm / ... which need objective function *file\*name* and cannot use function handles directly

74

**plotConfidenceIntervals.m**

PlotConfidenceIntervals.m visualizes confidence itervals stored in either the parameters or properties struct .CI

75

**plotMCMCdiagnosis.m**

PlotMCMCdiagnosis.m visualizes the Markov chains generated by getSamples.m

77

**plotMultiStartDiagnosis.m**

??

<a href="#">plotMultiStarts.m</a>	
PlotMultiStarts plots the result of the multi-start optimization stored in parameters	78
<a href="#">plotParameterProfiles.m</a>	
PlotParameterProfiles.m visualizes profile likelihood. Note: This routine provides an interface for <a href="#">plotUncertainty.m</a>	80
<a href="#">plotParameterSamples.m</a>	
PlotParameterSamples.m visualizes MCMC samples. Note: This routine provides an interface for <a href="#">plotUncertainty.m</a>	82
<a href="#">plotParameterUncertainty.m</a>	
PlotParameterUncertainty.m visualizes profile likelihood and MCMC samples stored in parameters	83
<a href="#">plotPropertyMultiStarts.m</a>	
PlotPropertyMultiStarts plots the result of the multi-start optimization stored in properties	85
<a href="#">plotPropertyProfiles.m</a>	
PlotPropertyProfiles.m visualizes profile likelihood of model properties. Note: This routine provides an interface for <a href="#">plotPropertyUncertainty.m</a>	86
<a href="#">plotPropertySamples.m</a>	
PlotPropertySamples.m visualizes samples of model properties. Note: This routine provides an interface for <a href="#">plotPropertyUncertainty.m</a>	88
<a href="#">plotPropertyUncertainty.m</a>	
PlotPropertyUncertainty.m visualizes profile likelihood and MCMC samples stored in properties	90
<a href="#">runPestoTests.m</a>	
RunPestoTests Run a set of PESTO unit tests	91
<a href="#">testGradient.m</a>	
TestGradient.m calculates finite difference approximations to the gradient to check an analytical version	92
<a href="#">@DRAMOptions/DRAMOptions.m</a>	??
<a href="#">@MALAOptions/MALAOptions.m</a>	??
<a href="#">@PestoOptions/PestoOptions.m</a>	??
<a href="#">@PestoPlottingOptions/PestoPlottingOptions.m</a>	??
<a href="#">@PestoSamplingOptions/PestoSamplingOptions.m</a>	??
<a href="#">@PHSOptions/PHSOptions.m</a>	??
<a href="#">@PTOptions/PTOptions.m</a>	??

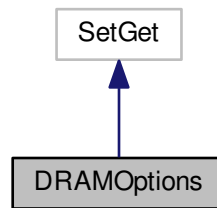
## 6 Class Documentation

### 6.1 DRAMOptions Class Reference

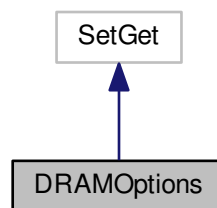
[DRAMOptions](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.



Inheritance diagram for DRAMOptions:



Collaboration diagram for DRAMOptions:



## Public Member Functions

- [DRAMOptions](#) (matlabtypesubstitute varargin)  
[DRAMOptions](#) Construct a new [DRAMOptions](#) object.
- `mlhsInnerSubst < matlabtypesubstitute, new > copy ()`

## Public Attributes

- matlabtypesubstitute [regFactor](#) = 1e-6  
*% Delayed Rejection Adaption Metropolis options Note: This algorithm uses a delayed rejection scheme for better mixing. — Delayed Rejection Adaptive Metropolis — opt.DRAM.regFactor : This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. Larger values equal stronger regularization. opt.DRAM.nTry : The number of tries in the delayed rejection scheme opt.DRAM.verbosityMode : Defines the level of verbosity *silent*, *visual*, *debug* or *text* opt.DRAM.adaptionInterval : Updates the proposal density only every opt.DRAM.adaptionInterval time*
- matlabtypesubstitute [nTry](#) = 1
- matlabtypesubstitute [verbosityMode](#) = "text"
- matlabtypesubstitute [adaptionInterval](#) = 1

### 6.1.1 Detailed Description

[DRAMOptions](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.

This file is based on AMICI amioptions.m (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file `DRAMOptions.m`.

### 6.1.2 Constructor & Destructor Documentation

#### 6.1.2.1 `DRAMOptions::DRAMOptions ( matlabtypesubstitute varargin )`

[DRAMOptions](#) Construct a new [DRAMOptions](#) object.

`OPTS = DRAMOptions()` creates a set of options with each option set to its default value.

`OPTS = DRAMOptions(PARAM, VAL, ...)` creates a set of options with the named parameters altered with the specified values.

`OPTS = DRAMOptions(OLDOPTS, PARAM, VAL, ...)` creates a copy of OLDOPTS with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for [DRAMOptions](#)

Definition at line 68 of file `DRAMOptions.m`.

References `adaptionInterval`, `nTry`, `regFactor`, and `verbosityMode`.

### 6.1.3 Member Data Documentation

#### 6.1.3.1 `DRAMOptions::adaptionInterval = 1`

##### Note

This property has custom functionality when its value is changed.

Definition at line 57 of file `DRAMOptions.m`.

Referenced by `DRAMOptions()`.

#### 6.1.3.2 `DRAMOptions::nTry = 1`

##### Note

This property has custom functionality when its value is changed.

Definition at line 53 of file `DRAMOptions.m`.

Referenced by `DRAMOptions()`.

## 6.1.3.3 DRAMOptions::regFactor = 1e-6

% Delayed Rejection Adaption Metropolis options Note: This algorithm uses a delayed rejection scheme for better mixing. — Delayed Rejection Adaptive Metropolis — opt.DRAM.regFactor : This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. Larger values equal stronger regularization. opt.DRAM.nTry : The number of tries in the delayed rejection scheme opt.DRAM.verbosityMode : Defines the level of verbosity *silent*, *visual*, *debug* or *text* opt.DRAM.adaptionInterval : Updates the proposal density only every opt.DRAM.adaptionInterval time

% Part for checking the correct setting of options

**Default:** 1e-6

**Note**

This property has custom functionality when its value is changed.

Definition at line 32 of file DRAMOptions.m.

Referenced by DRAMOptions().

## 6.1.3.4 DRAMOptions::verbosityMode = "text"

**Note**

This property has custom functionality when its value is changed.

Definition at line 55 of file DRAMOptions.m.

Referenced by DRAMOptions().

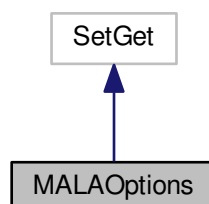
The documentation for this class was generated from the following file:

- @DRAMOptions/DRAMOptions.m

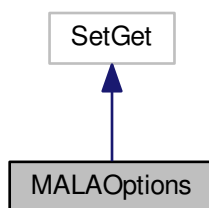
## 6.2 MALAOptions Class Reference

[MALAOptions](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.

Inheritance diagram for MALAOptions:



Collaboration diagram for MALAOptions:



### Public Member Functions

- [MALAOptions](#) (matlabtypesubstitute varargin)  
*[MALAOptions](#) Construct a new [MALAOptions](#) object.*
- mlhsInnerSubst< matlabtypesubstitute, new > **copy** ()

### Public Attributes

- matlabtypesubstitute [regFactor](#) = 1e-6  
*% Metropolis Adaptive Langevin Algorithm options Note: This algorithm uses gradients & Hessians either provided by the user or computed by finite differences. MALA, struct containing the fields .regFactor: This factor is used for regularization in cases where the proposal covariance matrices are ill conditioned. Larger values equal stronger regularization.*

### 6.2.1 Detailed Description

[MALAOptions](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.

This file is based on AMICI amioptions.m (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file MALAOptions.m.

### 6.2.2 Constructor & Destructor Documentation

#### 6.2.2.1 MALAOptions::MALAOptions ( matlabtypesubstitute varargin )

[MALAOptions](#) Construct a new [MALAOptions](#) object.

OPTS = [MALAOptions](#)() creates a set of options with each option set to its default value.

OPTS = [MALAOptions](#)(PARAM, VAL, ...) creates a set of options with the named parameters altered with the specified values.

OPTS = [MALAOptions](#)(OLDOPTS, PARAM, VAL, ...) creates a copy of OLDOPTS with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for [MALAOptions](#)

Definition at line 54 of file MALAOptions.m.

References [regFactor](#).

### 6.2.3 Member Data Documentation

#### 6.2.3.1 MALAOptions::regFactor = 1e-6

% Metropolis Adaptive Langevin Algorithm options Note: This algorithm uses gradients & Hessians either provided by the user or computed by finite differences. MALA, struct containing the fields .regFactor: This factor is used for regularization in cases where the proposal covariance matrices are ill conditioned. Larger values equal stronger regularization.

% Part for checking the correct setting of options

**Default:** 1e-6

#### Note

This property has custom functionality when its value is changed.

Definition at line 31 of file MALAOptions.m.

Referenced by MALAOptions().

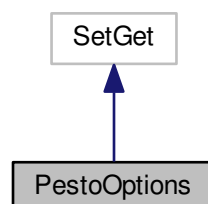
The documentation for this class was generated from the following file:

- @MALAOptions/MALAOptions.m

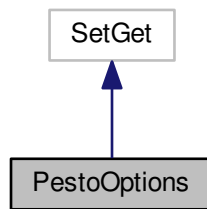
## 6.3 PestoOptions Class Reference

[PestoOptions](#) provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details.

Inheritance diagram for PestoOptions:



Collaboration diagram for PestoOptions:



### Public Member Functions

- **PestoOptions** (matlabtypesubstitute varargin)  
*PestoOptions* Construct a new *PestoOptions* object.
- mlhsInnerSubst< matlabtypesubstitute, new > **copy** ()  
*Creates a copy of the passed PestoOptions instance.*

### Public Attributes

- matlabtypesubstitute **comp\_type** = "sequential"  
*Perform calculations sequentially ('sequential', default), or in parallel ('parallel'). Parallel mode will speed-up the calculations on multi-core systems, but requires the MATLAB Parallel Computing Toolbox to be installed.*
- matlabtypesubstitute **rng** = 0  
*Initialization of random number generator.*
- matlabtypesubstitute **save** = false  
*Determine whether results are saved or not.*
- matlabtypesubstitute **foldername** = strep("datestr(now,31),'\_'\_\_")  
*Name of the folder in which results are stored. If no folder is provided, a random foldername is generated.*
- matlabtypesubstitute **obj\_type** = "log-posterior"  
*Type of objective function provided: log-posterior or negative log-posterior Tells the algorithm that log-posterior or log-likelihood are provided so it performs a maximization of the objective function or that the negative log-posterior or negative log-likelihood are provided so that a minimization of the objective function is performed.*
- matlabtypesubstitute **objOutNumber** = 3  
*Maximum number of outputs, the objective function can provide:*
- matlabtypesubstitute **mode** = "visual"  
*Output mode of algorithm:*
- matlabtypesubstitute **fh** = "[ ]"  
*Figure handle in which results are printed. If no handle is provided, a new figure is used. TODO: move to plot options.*
- matlabtypesubstitute **plot\_options** = **PestoPlottingOptions**("")  
*Plotting options of class PestoPlottingOptions.m.*
- matlabtypesubstitute **n\_starts** = 20  
*Number of local optimizations.*
- matlabtypesubstitute **start\_index** = "[ ]"  
*vector of indices which starts should be performed. default is 1:n\_starts*
- matlabtypesubstitute **init\_threshold** = -inf

- log-likelihood / log-posterior threshold for initialization of optimization.*
- matlabtypesubstitute `localOptimizer` = "fmincon"
  - Which optimizer to use? Current options: [fmincon, meigo-ess, meigo-vns, pswarm].*
- matlabtypesubstitute `localOptimizerOptions`
  - Options for the chosen local optimizer. Setting fmincon options as default local optimizer. See help(fmincon)*
- matlabtypesubstitute `proposal` = "latin hypercube"
  - Method used to propose starting points for fmincon. Can be.*
- matlabtypesubstitute `trace` = false
  - determine whether objective function, parameter values and computation time are stored over iterations*
- matlabtypesubstitute `tempsave` = false
  - determine whether intermediate results are stored every 10 iterations*
- matlabtypesubstitute `resetobjective` = false
  - clears the objective function before every multi-start.*
- matlabtypesubstitute `calc_profiles` = true
  - flag for profile calculation*
- matlabtypesubstitute `P` = {""}
  - parameter bounds (can be removed in a future release, since parameters.min/max can be used instead)*
- matlabtypesubstitute `MAP_index` = 1
  - index MAP - parameter vector starting from which the profile is calculated. This option is helpful if local optima are present.*
- matlabtypesubstitute `R_min` = 0.03
  - Minimal ratio down to which the profile is calculated.*
- matlabtypesubstitute `parameter_index` = "[]"
  - Indices of the parameters for which the profile is calculated.*
- matlabtypesubstitute `profile_optim_index` = "[]"
  - Indices of the parameters for which the profile is calculated by reoptimization.*
- matlabtypesubstitute `profile_integ_index` = "[]"
  - Indices of the parameters for which the profile is calculated by integration.*
- matlabtypesubstitute `profile_method` = "default"
  - How should profiles be computed (if no more precise options are set like profile\_optim\_index or profile\_integ\_index)? Possibilities: {optimization (default), integration, mixed}.*
- matlabtypesubstitute `profileReoptimizationOptions`
  - Optimizer options for profile likelihood.*
- matlabtypesubstitute `dR_max` = 0.10
  - Maximal relative decrease of ratio allowed for two adjacent points in the profile (default = 0.10) if options.dJ = 0;.*
- matlabtypesubstitute `dJ` = 0.5
  - influences step size at small likelihood ratio values*
- matlabtypesubstitute `options_getNextPoint`
  - options for the generation to the next profile point*
- matlabtypesubstitute `solver`
  - Options for profile integration.*
- matlabtypesubstitute `boundary_tol` = 1e-5
  - Tolance for the maximal distance of the list point the lower and upper bounds for the properties.*
- matlabtypesubstitute `property_index` = "[]"
  - Indices of the properties for which the profile is to be calculated (default = 1:properties.number, reoptimization only).*
- matlabtypesubstitute `MCMC`
  - Set MCMC options by calling an `PestoSamplingOptions` Class object.*

### 6.3.1 Detailed Description

[PestoOptions](#) provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details.

This file is based on AMICI amioptions.m (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file PestoOptions.m.

### 6.3.2 Constructor & Destructor Documentation

#### 6.3.2.1 PestoOptions::PestoOptions ( matlabtypesubstitute *varargin* )

[PestoOptions](#) Construct a new [PestoOptions](#) object.

OPTS = [PestoOptions](#)() creates a set of options with each option set to its default value.

OPTS = [PestoOptions](#)(PARAM, VAL, ...) creates a set of options with the named parameters altered with the specified values.

OPTS = [PestoOptions](#)(OLDOPTS, PARAM, VAL, ...) creates a copy of OLDOPTS with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for [PestoOptions](#)

#### Parameters

<i>varargin</i>	
-----------------	--

Definition at line 557 of file PestoOptions.m.

### 6.3.3 Member Data Documentation

#### 6.3.3.1 PestoOptions::boundary\_tol = 1e-5

Tolance for the maximal distance of the list point the lower and upper bounds for the properties.

**Default:** 1e-5

Definition at line 522 of file PestoOptions.m.

#### 6.3.3.2 PestoOptions::calc\_profiles = true

flag for profile calculation

- true: profiles are calculated
- false: profiles are not calculated

**Default:** true



**Note**

This property has custom functionality when its value is changed.

Definition at line 278 of file PestoOptions.m.

Referenced by copy().

**6.3.3.3 PestoOptions::comp\_type = "sequential"**

Perform calculations sequentially ('sequential', default), or in parallel ('parallel'). Parallel mode will speed-up the calculations on multi-core systems, but requires the [MATLAB Parallel Computing Toolbox](#) to be installed.

**Default:** "sequential"

**Note**

This property has custom functionality when its value is changed.

Definition at line 31 of file PestoOptions.m.

Referenced by copy().

**6.3.3.4 PestoOptions::dJ = 0.5**

influences step size at small likelihood ratio values

**Default:** 0.5

Definition at line 415 of file PestoOptions.m.

**6.3.3.5 PestoOptions::dR\_max = 0.10**

Maximal relative decrease of ratio allowed for two adjacent points in the profile (default = 0.10) if options.dJ = 0;.

**Default:** 0.10

**Note**

This property has custom functionality when its value is changed.

Definition at line 405 of file PestoOptions.m.

Referenced by copy().

**6.3.3.6 PestoOptions::fh = ""**

Figure handle in which results are printed. If no handle is provided, a new figure is used. TODO: move to plot options.

**Default:** ""

Definition at line 133 of file PestoOptions.m.

### 6.3.3.7 PestoOptions::foldername = strrep("datestr(now,31),'','\_\_'")

Name of the folder in which results are stored. If no folder is provided, a random foldername is generated.

**Default:** strrep("datestr(now,31),'','\_\_'")

Definition at line 69 of file PestoOptions.m.

### 6.3.3.8 PestoOptions::init\_threshold = -inf

log-likelihood / log-posterior threshold for initialization of optimization.

**Default:** -inf

Definition at line 175 of file PestoOptions.m.

### 6.3.3.9 PestoOptions::localOptimizer = "fmincon"

Which optimizer to use? Current options: [fmincon, meigo-ess, meigo-vns, pswarm].

For meigo-ess or meigo-vns, MEIGO (<http://gingproc.iim.csic.es/meigom.html>) has to be installed separately.

For pswarm PSwarm (<http://www.norg.uminho.pt/aivaz/pswarm/>) has to be installed separately

**Default:** "fmincon"

#### Note

This property has custom functionality when its value is changed.

Definition at line 185 of file PestoOptions.m.

Referenced by copy().

### 6.3.3.10 PestoOptions::localOptimizerOptions

#### Initial value:

```
= optimset(" \
    'algorithm', 'interior-point', \
    'display', 'off', \
    'GradObj', 'on', \
    'MaxIter', 2000, \
    'PrecondBandWidth', inf")
```

Options for the chosen local optimizer. Setting fmincon options as default local optimizer. See *help(fmincon)*

MaxIter: fmincon default, necessary to be set for tracing

Options for meigo-ess are described in ess\_kernel.m in the MEIGO folder.

**Default:** optimset(" \ 'algorithm', 'interior-point', \ 'display', 'off', \ 'GradObj', 'on', \ 'MaxIter', 2000, \ 'PrecondBandWidth', inf")

Definition at line 202 of file PestoOptions.m.

#### 6.3.3.11 PestoOptions::MAP\_index = 1

index MAP - parameter vector starting from which the profile is calculated. This option is helpful if local optima are present.

**Default:** 1

##### Note

This property has custom functionality when its value is changed.

Definition at line 304 of file PestoOptions.m.

Referenced by copy().

#### 6.3.3.12 PestoOptions::mode = "visual"

Output mode of algorithm:

- `visual`: plots showing the progress are generated
- `text`: optimization results for multi-start are printed on screen
- `silent`: no output during the multi-start local optimization
- `debug`: print extra debug information (only available in certain functions)

**Default:** "visual"

##### Note

This property has custom functionality when its value is changed.

Definition at line 118 of file PestoOptions.m.

Referenced by copy().

#### 6.3.3.13 PestoOptions::n\_starts = 20

Number of local optimizations.

**Default:** 20

##### Note

This property has custom functionality when its value is changed.

Definition at line 155 of file PestoOptions.m.

Referenced by copy().

#### 6.3.3.14 PestoOptions::obj\_type = "log-posterior"

Type of objective function provided: `log-posterior` or `negative log-posterior` Tells the algorithm that `log-posterior` or `log-likelihood` are provided so it performs a maximization of the objective function or that the negative `log-posterior` or `negative log-likelihood` are provided so that a minimization of the objective function is performed.

**Default:** "log-posterior"

#### Note

This property has custom functionality when its value is changed.

Definition at line 82 of file PestoOptions.m.

Referenced by `copy()`.

#### 6.3.3.15 PestoOptions::objOutNumber = 3

Maximum number of outputs, the objective function can provide:

- 1 ... only objective value
- 2 ... objective value with gradient
- 3 ... objective value, gradient and Hessian

(Missing values will be approximated by finite differences.) Don't confuse this with the number of outputs from the objective function that Pesto really calls! `objOutNumber` just tells Pesto, with how many outputs it can call the objective function. Also, optimization is still possible to be done without gradient information and without the Pesto FD routine.

**Default:** 3

Definition at line 96 of file PestoOptions.m.

#### 6.3.3.16 PestoOptions::options\_getNextPoint

**Initial value:**

```
= struct('mode', 'multi-dimensional', \
        'guess', 1e-2, \
        'min', 1e-6, \
        'max', 1, \
        'update', 1.25)
```

options for the generation fo the next profile point

- `.mode` ... choice of proposal direction
- `= multi-dimensional` (default) ... all parameters are updated.
- The direct is the same as between the last two points.
- `= one-dimensional` ... only parameter for which profile is

- currently calculated is updated.
- `.guess = 1e-2` ... guess for initial update stepsize
- `.min = 1e-6` ... lower bound for update stepsize
- `.min = 1e2` ... upper bound for update stepsize
- `.update = 1.25` ... incremental change if stepsize is too large or
- too small, must be  $> 1$ .

**Default:** `struct('mode', 'multi-dimensional', \ 'guess', 1e-2, \ 'min', 1e-6, \ 'max', 1, \ 'update', 1.25)`

Definition at line 424 of file `PestoOptions.m`.

#### 6.3.3.17 PestoOptions::P = {}

parameter bounds (can be removed in a future release, since `parameters.min/max` can be used instead)

- `.P.min` ... lower bound for profiling parameters, having same dimension as the parameter vector (default = `parameters.min`).
- `.P.max` ... lower bound for profiling parameters, having same dimension as the parameter vector (default = `parameters.max`).

**Default:** `{}`

Definition at line 290 of file `PestoOptions.m`.

#### 6.3.3.18 PestoOptions::parameter\_index = []

Indices of the parameters for which the profile is calculated.

Default: `profile_optim_index` will be set to `1:parameters.number` if left empty

**Default:** `[]`

#### Note

This property has custom functionality when its value is changed.

Definition at line 324 of file `PestoOptions.m`.

Referenced by `copy()`.

#### 6.3.3.19 PestoOptions::plot\_options = PestoPlottingOptions()

Plotting options of class [PestoPlottingOptions.m](#).

**Default:** [PestoPlottingOptions\(\)](#)

Definition at line 143 of file `PestoOptions.m`.

### 6.3.3.20 PestoOptions::profile\_integ\_index = "[]"

Indices of the parameters for which the profile is calculated by integration.

**Default:** "[]"

Definition at line 346 of file PestoOptions.m.

### 6.3.3.21 PestoOptions::profile\_method = "default"

How should profiles be computed (if no more precise options are set like profile\_optim\_index or profile\_integ\_index)? Possibilities: {optimization (default), integration, mixed}.

**Default:** "default"

Definition at line 356 of file PestoOptions.m.

### 6.3.3.22 PestoOptions::profile\_optim\_index = "[]"

Indices of the parameters for which the profile is calculated by reoptimization.

**Default:** "[]"

Definition at line 336 of file PestoOptions.m.

### 6.3.3.23 PestoOptions::profileReoptimizationOptions

**Initial value:**

```
= optimset(" \
    'algorithm', 'interior-point', \
    'display', 'off', \
    'MaxIter', 400, \
    'GradObj', 'on', \
    'GradConstr', 'on', \
    'TolCon', 1e-6 \
")
```

Optimizer options for profile likelihood.

- .algorithm ... choice of algorithm
- = interior-point (default)
- = trust-region-reflective
- = active-set
- .display ... output of optimization process
- = off (default) ... no output
- = iter ... output for every optimization step
- .MaxIter ... maximum of optimization steps
- .GradObj ... are gradients provided?
- .GradConstr ... do we have constraints?
- .TolCon ... Tolerance for constraints

**Default:** optimset(" \ 'algorithm', 'interior-point', \ 'display', 'off', \ 'MaxIter', 400, \ 'GradObj', 'on', \ 'GradConstr', 'on', \ 'TolCon', 1e-6 \ ")

Definition at line 370 of file PestoOptions.m.

#### 6.3.3.24 PestoOptions::property\_index = "[ ]"

Indices of the properties for which the profile is to be calculated (default = 1:properties.number, reoptimization only).

**Default:** "[ ]"

##### Note

This property has custom functionality when its value is changed.

Definition at line 533 of file PestoOptions.m.

Referenced by copy().

#### 6.3.3.25 PestoOptions::proposal = "latin hypercube"

Method used to propose starting points for fmincon. Can be.

- `latin hypercube`: latin hypercube sampling
- `uniform`: uniform random sampling
- `user-supplied`: user supplied function PestoOptions.init\_fun

**Default:** "latin hypercube"

##### Note

This property has custom functionality when its value is changed.

Definition at line 226 of file PestoOptions.m.

Referenced by copy().

#### 6.3.3.26 PestoOptions::R\_min = 0.03

Minimal ratio down to which the profile is calculated.

**Default:** 0.03

##### Note

This property has custom functionality when its value is changed.

Definition at line 315 of file PestoOptions.m.

Referenced by copy().

### 6.3.3.27 PestoOptions::resetobjective = false

clears the objective function before every multi-start.

- false: persistent variables are preserved.
- true: remove all temporary/persistent variables.

WHEN TRUE THIS OPTION REMOVES ALL OBJECTIVE FUNCTION BREAK POINTS

**Default:** false

#### Note

This property has custom functionality when its value is changed.

Definition at line 262 of file PestoOptions.m.

Referenced by copy().

### 6.3.3.28 PestoOptions::rng = 0

Initialization of random number generator.

- Any real number r: random generator is initialized with r.
- []: random number generator is not initialized. (Initializing the random number generator with a specific seed can be helpful to reproduce problems.)

**Default:** 0

Definition at line 45 of file PestoOptions.m.

### 6.3.3.29 PestoOptions::save = false

Determine whether results are saved or not.

- false: results are not saved
- true: results are stored to an extra folder

**Default:** false

#### Note

This property has custom functionality when its value is changed.

Definition at line 58 of file PestoOptions.m.

Referenced by copy().



## 6.3.3.30 PestoOptions::solver

## Initial value:

```
= struct('type', 'ode113', \
        'algorithm', 'Adams', \
        'nonlinSolver', 'Newton', \
        'linSolver', 'Dense', \
        'gamma', 0, \
        'eps', 1e-8, \
        'minCond', 1e-10, \
        'hessian', 'user-supplied', \
        'gradient', true, \
        'MaxStep', 0.1, \
        'MinStep', 1e-4, \
        'MaxNumSteps', 1e5, \
        'RelTol', 1e-5, \
        'AbsTol', 1e-8
    )
```

Options for profile integration.

- .type ... choice of ODE integrator
- = ode113 (default) ... Adams-Bashf.-Solver by Matlab
- = ode15s ... BDF-Solver by Matlab
- = ode45 ... Runge-Kutta-Solver by Matlab
- = CVODE ... BDF and AB-Solver by sundials (to be implemented!)
- .algorithm ... choice of algorithm (CVODE only)
- = Adams (default) ... Adams-Bashford solver
- = BDF ... BDF solver
- .nonlinSolver ... choice of nonlinear solver (CVODE only)
- = Newton (default)
- = Functional
- .linSolver ... choice of linear solver (CVODE only)
- = Dense (default) ... solver for small problems
- = Band ... solver for bigger problems
- .gamma ... Retraction factor
- .eps ... regularization for poorly conditioned Hessian
- .minCond ... Minimum condition number, when regularization is to be used
- .hessian ... how is the Hessian Matrix provided?
- = user-supplied (default) ... Hessian is 3rd output of ObjFun
- = bfgs ... BFGS approximation to Hessian
- = srl ... symmetric-rank 1 approximation to Hessian
- .gradient ... is a gradient provided?
- .MaxStep ... minimum step size of the solver
- .MaxStep ... maximum step size of the solver

- `.MaxNumSteps` ... maximum steps to be taken
- `.RelTol` ... maximum relative integration error
- `.AbsTol` ... maximum absolute integration error

**Default:** `struct('type', 'ode113', \ 'algorithm', 'Adams', \ 'nonlinSolver', 'Newton', \ 'linSolver', 'Dense', \ 'gamma', 0, \ 'eps', 1e-8, \ 'minCond', 1e-10, \ 'hessian', 'user-supplied', \ 'gradient', true, \ 'MaxStep', 0.1, \ 'MinStep', 1e-4, \ 'MaxNumSteps', 1e5, \ 'RelTol', 1e-5, \ 'AbsTol', 1e-8 )`

Definition at line 454 of file PestoOptions.m.

#### 6.3.3.31 `PestoOptions::start_index = []`

vector of indices which starts should be performed. default is `1:n_starts`

**Default:** `[]`

#### Note

This property has custom functionality when its value is changed.

Definition at line 165 of file PestoOptions.m.

Referenced by `copy()`.

#### 6.3.3.32 `PestoOptions::tempsave = false`

determine whether intermediate results are stored every 10 iterations

- `false`: not saved
- `true`: results are stored to an extra folder

**Default:** `false`

#### Note

This property has custom functionality when its value is changed.

Definition at line 250 of file PestoOptions.m.

Referenced by `copy()`.

### 6.3.3.33 PestoOptions::trace = false

determine whether objective function, parameter values and computation time are stored over iterations

- false: not saved
- true: stored in fields par\_trace, fval\_trace and time\_trace

**Default:** false

#### Note

This property has custom functionality when its value is changed.

Definition at line 238 of file PestoOptions.m.

Referenced by copy().

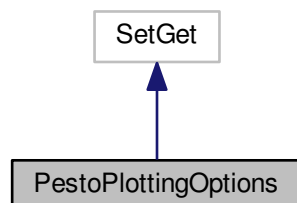
The documentation for this class was generated from the following file:

- @PestoOptions/PestoOptions.m

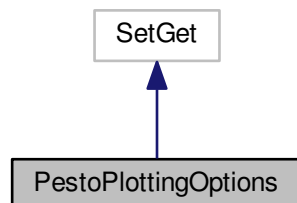
## 6.4 PestoPlottingOptions Class Reference

[PestoPlottingOptions](#) is a class for checking and holding information on optimization parameters.

Inheritance diagram for PestoPlottingOptions:



Collaboration diagram for PestoPlottingOptions:



## Public Member Functions

- [PestoPlottingOptions](#) (matlabtypesubstitute varargin)  
*PestoPlottingOptions* Construct a new *PestoPlottingOptions* object.
- `mlhsInnerSubst<` matlabtypesubstitute, new `>` [copy](#) ()  
*Creates a copy of the passed [PestoPlottingOptions](#) instance.*

## Public Attributes

- matlabtypesubstitute [title](#) = true  
*Title of PESTO-generated plots.*
- matlabtypesubstitute [add\\_points](#)  
*Additional points to include in the plots, e.g. true parameter in the case of test examples.*
- matlabtypesubstitute [mark\\_constraint](#) = false  
*TODO: from plotmultistarts.*
- matlabtypesubstitute [subplot\\_size\\_1D](#) = "[]"  
*TODO from plotparameteruncertainty.*
- matlabtypesubstitute [subplot\\_indexing\\_1D](#) = "[]"  
*TODO.*
- matlabtypesubstitute [labels](#)  
*TODO.*
- matlabtypesubstitute [hold\\_on](#) = false  
*Indicates whether plots are redrawn or whether something is added to the plot.*
- matlabtypesubstitute [interval](#) = "dynamic"  
*Way of choosing x limits for plotting.*
- matlabtypesubstitute [draw\\_bounds](#) = true  
*Draw bounds.*
- matlabtypesubstitute [bounds](#) = {""}  
*Bounds used for visualization if options.interval = static*
- matlabtypesubstitute [P](#)  
*Options for profile plots.*
- matlabtypesubstitute [S](#)  
*Options for sample plots.*
- matlabtypesubstitute [MS](#)  
*Options for multi-start optimization plots.*
- matlabtypesubstitute [A](#)  
*Options for distribution approximation plots.*
- matlabtypesubstitute [MCMC](#) = "multistart"  
*Option if a user provided sampling initialization should be used for plotting an approximation of the distribution.*
- matlabtypesubstitute [boundary](#)  
*Options for boundary visualization.*
- matlabtypesubstitute [CL](#)  
*Options for confidence level plots.*
- matlabtypesubstitute [group\\_CI\\_by](#) = "parprop"  
*Options for the way to plot confidence intervals.*
- matlabtypesubstitute [op2D](#) = struct("b1", 0.15, 'b2', 0.02, 'r', 0.95")  
*Settings for 2D plot to position subplot axes.*
- matlabtypesubstitute [legend](#)  
*Legend options.*
- matlabtypesubstitute [fontsize](#) = struct ("tick", 12")

- Fontsize for labels.*
  - matlabtypesubstitute `fh_logPost_trace` = "[ ]"  
*figure handle for log-posterior trace plot*
  - matlabtypesubstitute `fh_par_trace` = "[ ]"  
*figure handle for parameter trace plots.*
  - matlabtypesubstitute `fh_par_dis_1D` = "[ ]"  
*figure handle for the 1D parameter distribution plot.*
  - matlabtypesubstitute `fh_par_dis_2D` = "[ ]"  
*figure handle for the 2D parameter distribution plot.*
  - matlabtypesubstitute `plot_type` = {"parameter", "posterior"}  
*plot type*
  - matlabtypesubstitute `n_max` = 1e4  
*max*

#### 6.4.1 Detailed Description

`PestoPlottingOptions` is a class for checking and holding information on optimization parameters.

This file is based on AMICI amioptions.m (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file `PestoPlottingOptions.m`.

#### 6.4.2 Constructor & Destructor Documentation

##### 6.4.2.1 `PestoPlottingOptions::PestoPlottingOptions ( matlabtypesubstitute varargin )`

`PestoPlottingOptions` Construct a new `PestoPlottingOptions` object.

`OPTS = PestoPlottingOptions()` creates a set of options with each option set to its default value.

`OPTS = PestoPlottingOptions(PARAM, VAL, ...)` creates a set of options with the named parameters altered with the specified values.

`OPTS = PestoPlottingOptions(OLDOPTS, PARAM, VAL, ...)` creates a copy of OLDOPTS with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for `PestoPlottingOptions`

#### Parameters

<code>varargin</code>	
-----------------------	--

Definition at line 489 of file `PestoPlottingOptions.m`.

#### 6.4.3 Member Data Documentation

##### 6.4.3.1 `PestoPlottingOptions::A`

**Initial value:**

```
= struct("'plot_type', 1, \
        'col', [0,0,1], \
        'lw', 2, \
        'sigma_level', 2, \
        'name', 'P_{app}')
```

Options for distribution approximation plots.

Struct with

- `.plot_type`: plot type
  - = 0 (default if no MS are provided) ... no plot
  - = 1 (default if MS are provided) ... likelihood ratio
  - = 2 ... negative log-likelihood
- `.col`: color of approximation lines (default: [0,0,1])
- `.lw`: line width of approximation lines (default: 1.5)
- `.sigma_level`: sigma-level which is visualized (default = 2)
- `.name`: name of legend entry (default = `P_{app}`)

**Default:** `struct("'plot_type', 1, \ 'col', [0,0,1], \ 'lw', 2, \ 'sigma_level', 2, \ 'name', 'P_{app}')`

Definition at line 277 of file `PestoPlottingOptions.m`.

#### 6.4.3.2 `PestoPlottingOptions::add_points`

**Initial value:**

```
= struct("'par', [], \
        'logPost', [], \
        'col', [0,0.8,0], \
        'ls', '-', \
        'lw', 1, \
        'm', 'd', \
        'ms', 8, \
        'name', 'add. point')
```

Additional points to include in the plots, e.g. true parameter in the case of test examples.

Struct with the following fields

- `.par`:  $n \times m$  matrix of  $m$  additional points
- `.col`: color used for additional points (default = [0,0,0]). This can also be a  $m \times 3$  matrix of colors.
- `.ls`: line style (default = `-`)
- `.lw`: line width (default = 2)
- `.m`: marker style (default = `s`)
- `.ms`: line width (default = 8)
- `.name`: name of legend entry (default = `add. point`)
- `.property_MS`: line width (default = 8).
- `.logPost`

**Default:** `struct("'par', [], \ 'logPost', [], \ 'col', [0,0.8,0], \ 'ls', '-', \ 'lw', 1, \ 'm', 'd', \ 'ms', 8, \ 'name', 'add. point')`

Definition at line 42 of file `PestoPlottingOptions.m`.

## 6.4.3.3 PestoPlottingOptions::boundary

**Initial value:**

```
= struct('mark', true, \
        'eps', 1e-4)
```

Options for boundary visualization.

Struct with

- `.mark`: marking of profile points which are on the boundary
  - `= 0` ... no visualization
  - `= 1` (default) ... indicates points which are close to the boundaries in one or more dimensions.
- `.eps`: minimal distance from boundary for which points are considered to be close to the boundary (default = `1e-4`). Note that a one-norm is used.

**Default:** `struct('mark', true, 'eps', 1e-4)`

Definition at line 318 of file `PestoPlottingOptions.m`.

## 6.4.3.4 PestoPlottingOptions::bounds = {}

Bounds used for visualization if `options.interval = static`

struct with

- `.min`: lower bound
- `.max`: upper bound

**Default:** `{}`

Definition at line 152 of file `PestoPlottingOptions.m`.

#### 6.4.3.5 PestoPlottingOptions::CL

**Initial value:**

```
= struct("plot_type", 0, \
        'alpha', 0.95, \
        'type', 'point-wise', \
        'col', [0,0,0], \
        'lw', 2, \
        'name', 'cut-off')
```

Options for confidence level plots.

Struct with

- `.plot_type`: plot type
  - = 0 (default) ... no plot
  - = 1 ... likelihood ratio
  - = 2 ... negative log-likelihood
- `.alpha`: visualized confidence level (default = 0.95)
- `.type`: type of confidence interval
  - = `point-wise` (default) ... point-wise confidence interval
  - = `simultaneous` ... point-wise confidence interval
  - = `{point-wise,simultaneous}` ... both
- `.col`: color of profile lines (default: [0,0,0])
- `.lw`: line width of profile lines (default: 1.5)
- `.name`: name of legend entry (default = `cut-off`)

**Default:** `struct("plot_type", 0, \ 'alpha', 0.95, \ 'type', 'point-wise', \ 'col', [0,0,0], \ 'lw', 2, \ 'name', 'cut-off')`

Definition at line 339 of file `PestoPlottingOptions.m`.

#### 6.4.3.6 PestoPlottingOptions::draw\_bounds = true

Draw bounds.

- `true`: yes
- `false`: no

**Default:** `true`

**Note**

This property has custom functionality when its value is changed.

Definition at line 141 of file `PestoPlottingOptions.m`.

Referenced by `copy()`.



#### 6.4.3.7 PestoPlottingOptions::fh\_logPost\_trace = "[ ]"

figure handle for log-posterior trace plot

**Default:** "[ ]"

Definition at line 430 of file PestoPlottingOptions.m.

#### 6.4.3.8 PestoPlottingOptions::fh\_par\_dis\_1D = "[ ]"

figure handle for the 1D parameter distribution plot.

**Default:** "[ ]"

Definition at line 448 of file PestoPlottingOptions.m.

#### 6.4.3.9 PestoPlottingOptions::fh\_par\_dis\_2D = "[ ]"

figure handle for the 2D parameter distribution plot.

**Default:** "[ ]"

Definition at line 457 of file PestoPlottingOptions.m.

#### 6.4.3.10 PestoPlottingOptions::fh\_par\_trace = "[ ]"

figure handle for parameter trace plots.

**Default:** "[ ]"

Definition at line 439 of file PestoPlottingOptions.m.

#### 6.4.3.11 PestoPlottingOptions::fontsize = struct ('tick', 12)

Fontsize for labels.

- .tick: fontsize for ticklabels (default = 12)

**Default:** struct ('tick', 12)

Definition at line 420 of file PestoPlottingOptions.m.

#### 6.4.3.12 PestoPlottingOptions::group\_CI\_by = "parprop"

Options for the way to plot confidence intervals.

Either all confidence intervals of one method are plotted to one window `params`, or the confidence intervals for one parameter from all methods are plotted to one window `methods`, or everthing is grouped together `all`.

**Default:** "parprop"

##### Note

This property has custom functionality when its value is changed.

Definition at line 373 of file PestoPlottingOptions.m.

Referenced by `copy()`.

#### 6.4.3.13 PestoPlottingOptions::hold\_on = false

Indicates whether plots are redrawn or whether something is added to the plot.

- `true`: extension of plot
- `false`: new plot

**Default:** false

##### Note

This property has custom functionality when its value is changed.

Definition at line 116 of file PestoPlottingOptions.m.

Referenced by `copy()`.

#### 6.4.3.14 PestoPlottingOptions::interval = "dynamic"

Way of choosing x limits for plotting.

- `dynamic`: x limits depending on analysis results
- `static`: x limits depending on `parameters.min` and `.max` or on user-defined bound `options.bounds.min` and `.max`. The later are used if provided.

**Default:** "dynamic"

##### Note

This property has custom functionality when its value is changed.

Definition at line 128 of file PestoPlottingOptions.m.

Referenced by `copy()`.

## 6.4.3.15 PestoPlottingOptions::labels

**Initial value:**

```
= struct('y_always', true, \
        'y_name', [])
```

TODO.

**Default:** struct("y\_always", true, \ 'y\_name', [])

Definition at line 105 of file PestoPlottingOptions.m.

## 6.4.3.16 PestoPlottingOptions::legend

**Initial value:**

```
= struct('color', 'none', \
        'box', 'on', \
        'orientation', 'vertical', \
        'position', [])
```

Legend options.

- .color: background color (default = none).
- .box: legend outline (default = on).
- .orientation: orientation of list (default = vertical)

**Default:** struct("color", 'none', \ 'box', 'on', \ 'orientation', 'vertical', \ 'position', [])

Definition at line 402 of file PestoPlottingOptions.m.

## 6.4.3.17 PestoPlottingOptions::mark\_constraint = false

TODO: from plotmultistarts.

**Default:** false

**Note**

This property has custom functionality when its value is changed.

Definition at line 78 of file PestoPlottingOptions.m.

Referenced by copy().

#### 6.4.3.18 PestoPlottingOptions::MCMC = "multistart"

Option if a user provided sampling initialization should be used for plotting an approximation of the distribution.

- user-provided
- multistart (default)

**Default:** "multistart"

#### Note

This property has custom functionality when its value is changed.

Definition at line 305 of file PestoPlottingOptions.m.

Referenced by copy().

#### 6.4.3.19 PestoPlottingOptions::MS

**Initial value:**

```
= struct("plot_type", 1, \
        'col', [1,0,0], \
        'lw', 2, \
        'name_conv', 'MS - conv.', \
        'name_nconv', 'MS - not conv.', \
        'only_optimum', false")
```

Options for multi-start optimization plots.

Struct with

- .plot\_type: plot type
  - = 0 (default if no MS are provided) ... no plot
  - = 1 (default if MS are provided) ... likelihood ratio and position of optima above threshold
  - = 2 ... negative log-likelihood and position of optima above threshold
- .col: color of local optima (default: [1,0,0])
- .lw: line width of local optima (default: 1.5)
- .name\_conv: name of legend entry (default = MS - conv.)
- .name\_nconv: name of legend entry (default = MS - not conv.)
- .only\_optimum: only optimum is plotted

**Default:** struct("plot\_type", 1, \ 'col', [1,0,0], \ 'lw', 2, \ 'name\_conv', 'MS - conv.', \ 'name\_nconv', 'MS - not conv.', \ 'only\_optimum', false")

Definition at line 244 of file PestoPlottingOptions.m.

## 6.4.3.20 PestoPlottingOptions::n\_max = 1e4

max

**Default:** 1e4

Note

This property has custom functionality when its value is changed.

Definition at line 475 of file PestoPlottingOptions.m.

Referenced by copy().

## 6.4.3.21 PestoPlottingOptions::op2D = struct("b1", 0.15, 'b2', 0.02, 'r', 0.95")

Settings for 2D plot to position subplot axes.

Struct with

- .b1 ... offset from left and bottom border (default = 0.15)
- .b2 ... offset from left and bottom border (default = 0.02)
- .r ... relative width of subplots (default = 0.95)

**Default:** struct("b1", 0.15, 'b2', 0.02, 'r', 0.95")

Definition at line 387 of file PestoPlottingOptions.m.

## 6.4.3.22 PestoPlottingOptions::P

**Initial value:**

```
= struct('plot_type', 1, \
        'col', [1,0,0], \
        'lw', 2, \
        'name', 'P')
```

Options for profile plots.

Struct with

- .plot\_type: plot type
  - = 0 (default if no profiles are provided) ... no plot
  - = 1 (default if profiles are provided) ... likelihood ratio
  - = 2 ... negative log-likelihood
- .col: color of profile lines (default: [1,0,0])
- .lw: line width of profile lines (default: 1.5)

**Default:** struct('plot\_type', 1, \ 'col', [1,0,0], \ 'lw', 2, \ 'name', 'P')

Definition at line 165 of file PestoPlottingOptions.m.

#### 6.4.3.23 PestoPlottingOptions::plot\_type = {"parameter','posterior'"}

plot type

**Default:** {"parameter','posterior'"}

Definition at line 466 of file PestoPlottingOptions.m.

#### 6.4.3.24 PestoPlottingOptions::S

**Initial value:**

```
= struct('plot_type', 0, \
        'bins', 'optimal', \
        'scaling', [], \
        'hist_col', [0.7,0.7,0.7], \
        'sp_col', [0.7,0.7,0.7], \
        'lin_col', [1,0,0], \
        'lin_lw', 2, \
        'sp_m', '.', \
        'sp_ms', 5, \
        'col', [1,0,0], 'lw', 2, \
        'PT', struct('sp_m', '.', \
                    'sp_ms', 5, \
                    'lw', 1.5, \
                    'ind', [], \
                    'col', [], \
                    'plot_type', 0), \
        'name', 'S')
```

Options for sample plots.

- .plot\_type: plot type
  - = 0 (default if no samples are provided) ... no plot
  - = 1 (default if samples are provided) ... histogram
  - = 2 ... kernel-density estimates
- .col ... color of profile lines (default: [0.7,0.7,0.7])
- .hist\_col ... color of histogram (default = [0.7,0.7,0.7])
- .bins ... number of histogram bins (default: 30)
  - = optimal ... selection using Scott's rule
  - = conservative ... selection using Scott's rule / 2
  - = N (with N being an integer) ... N bins
- .sp\_col: color of scatter plot (default = [0.7,0.7,0.7])
- .sp\_m: marker for scatter plot (default = .)
- .sp\_ms: marker size for scatter plot (default = 5)
- .name: name of legend entry (default = S)

**Default:** struct('plot\_type', 0, \ 'bins', 'optimal', \ 'scaling', [], \ 'hist\_col', [0.7,0.7,0.7], \ 'sp\_col', [0.7,0.7,0.7], \ 'lin\_col', [1,0,0], \ 'lin\_lw', 2, \ 'sp\_m', '.', \ 'sp\_ms', 5, \ 'col', [1,0,0], 'lw', 2, \ 'PT', struct('sp\_m', '.', \ 'sp\_ms', 5, \ 'lw', 1.5, \ 'ind', [], \ 'col', [], \ 'plot\_type', 0), \ 'name', 'S')

Definition at line 188 of file PestoPlottingOptions.m.

#### 6.4.3.25 PestoPlottingOptions::subplot\_indexing\_1D = "[ ]"

TODO.

**Default:** "[ ]"

Definition at line 96 of file PestoPlottingOptions.m.

#### 6.4.3.26 PestoPlottingOptions::subplot\_size\_1D = "[ ]"

TODO from plotparameteruncertainty.

**Default:** "[ ]"

Definition at line 87 of file PestoPlottingOptions.m.

#### 6.4.3.27 PestoPlottingOptions::title = true

Title of PESTO-generated plots.

- true: show
- false: don't show

**Default:** true

#### Note

This property has custom functionality when its value is changed.

Definition at line 30 of file PestoPlottingOptions.m.

Referenced by copy().

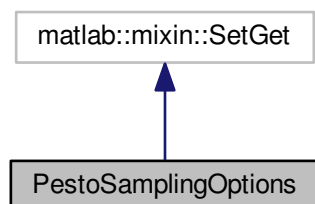
The documentation for this class was generated from the following file:

- @PestoPlottingOptions/PestoPlottingOptions.m

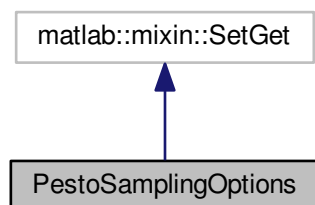
## 6.5 PestoSamplingOptions Class Reference

[PestoSamplingOptions](#) provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details.

Inheritance diagram for PestoSamplingOptions:



Collaboration diagram for PestoSamplingOptions:



### Public Member Functions

- [PestoSamplingOptions](#) (matlabtypesubstitute varargin)  
*[PestoSamplingOptions](#) Construct a new [PestoSamplingOptions](#) object.*
- mlhsInnerSubst< matlabtypesubstitute, new > **copy** ()
- mlhsInnerSubst< matlabtypesubstitute, this > [checkDependentDefaults](#) (matlabtypesubstitute par)  
*Should be called providing the parameter struct for dependent defaults as theta0. Does both, checking already set dependent options and defaulting based on par if not set yet.*

### Public Attributes

- matlabtypesubstitute [obj\\_type](#) = "log-posterior"  
*% General options Type of objective function provided: log-posterior (default) or negative log-posterior*
- matlabtypesubstitute [rndSeed](#) = "shuffle"



- Random seed, either a number or `shuffle` (default)
- matlabtypesubstitute `samplingAlgorithm` = "PT"  
Sampling algorithm, can be `PT` (parallel tempering), `PHS`, (parallel hierarchical sampling), `MALA` (Metropolis adjusted Langevin algorithm), or `DRAM` (delayed rejection adapted Metropolis algorithm, only if the `DRAM` toolbox is installed). Default value is `PT`
  - matlabtypesubstitute `nIterations` = 1e5  
Number of iterations, integer (1e5 is not too high, e.g. 1e6 is a more reasonable value for reliable results, but computationally more intensive).
  - matlabtypesubstitute `theta0` = "[]"  
Initialization points for all chains. If the algorithm uses multiple chains (as `PT` and `PHS`), one can specify multiple `theta0`, e.g.: `opt.theta0 = repmat([0.1, 1.5, -2.5, -0.5, 0.4], opt.nTemps, 1)`; If there is just one chain, please specify as `opt.theta0 = [1; 2; 3; 4]`; It is recommended to set `theta0` by taking into account the results from a preceding optimization (see Pesto examples).
  - matlabtypesubstitute `sigma0` = "[]"  
Initial covariance matrices for all chains. Example for single-chain algorithms: `opt.sigma0 = 1e5 * diag(ones(1,5))`; Example for multi-chain algorithms: `opt.sigma0 = repmat(1e5*diag(ones(1,5)), opt.nTemps, 1)`; It is recommended to set `sigma0` by taking into account the results from a preceding optimization.
  - matlabtypesubstitute `mode` = "visual"  
Output mode for sampling algorithms (except `DRAM`, which has its own format), can be chosen as `visual`, `text`, `silent`, or `debug`. Default: `visual`.
  - matlabtypesubstitute `objOutNumber` = 1  
Maximum number of outputs, the objective function can provide: 1 ... only objective value 2 ... objective value with gradient 3 ... objective value, gradient and Hessian (Default)
  - matlabtypesubstitute `PT`  
% Parallel Tempering Options `PT`, struct containing the fields `.nTemps`: Initial number of temperatures (default 10) `.exponentT`: The initial temperatures are set by a power law to  $^{\text{opt.exponentT}}$  (default 4) `.alpha`: Parameter which controls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.51. No adaption (classical Metropolis-Hastings) for 0. `.temperatureAlpha`: Parameter which controls the adaption degeneration velocity of the temperature adaption. Value between 0 and 1. Default 0.51. No effect for value = 0. `.memoryLength`: The higher the value the more it lowers the impact of early adaption steps. Default 1. `.regFactor`: Regularization factor for ill conditioned covariance matrices of the adapted proposal density. Regularization might happen if the eigenvalues of the covariance matrix strongly differ in order of magnitude. In this case, the algorithm adds a small diag-matrix to the covariance matrix with elements `regFactor`. `.temperatureAdaptionScheme`: Follows the temperature adaption scheme from Vousden16 or Lack15. Can be set to `none` for no temperature adaption.
  - matlabtypesubstitute `PHS` = struct  
% Parallel Hierarchical Sampling options `PHS`, struct containing the fields `.nChains`: Number of chains (1 mother-chain and `nChains-1` auxillary chains) `.alpha`: Control parameter for adaption decay. Needs values between 0 and 1. Higher values lead to faster decays, meaning that new iterations influence the single-chain proposal adaption only very weakly very quickly. `.memoryLength`: Control parameter for adaption. Higher values suppress strong early adaption. `.regFactor`: This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. `nChains` larger values equal stronger regularization. `.trainingTime`: The iterations before the first chain swap is invoked
  - matlabtypesubstitute `MALA` = struct  
% Metropolis Adaptive Langevin Algorithm options Note: This algorithm uses gradients & Hessians either provided by the user or computed by finite differences. `MALA`, struct containing the fields `.regFactor`: This factor is used for regularization in cases where the proposal covariance matrices are ill conditioned. Larger values equal stronger regularization.
  - matlabtypesubstitute `DRAM` = struct  
% Delayed Rejection Adaptive Metropolis options `DRAM`, struct containing the fields `.adaptionInterval`: Updates the proposal density only every `adaptionInterval`-th time `.nTry`: The number of tries in the delayed rejection scheme `.regFactor`: This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. Larger values equal stronger regularization. `.verbosityMode`: Defines the level of verbosity `silent`, `visual`, `debug`, or `text`

### 6.5.1 Detailed Description

`PestoSamplingOptions` provides an option container to pass options to various PESTO functions. Not all options are used by all functions, consult the respective function documentation for details.

This file is based on AMICI amioptions.m (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file PestoSamplingOptions.m.

## 6.5.2 Constructor & Destructor Documentation

### 6.5.2.1 PestoSamplingOptions::PestoSamplingOptions ( matlabtypesubstitute *varargin* )

**PestoSamplingOptions** Construct a new **PestoSamplingOptions** object.

`OPTS = PestoSamplingOptions()` creates a set of options with each option set to its default value.

`OPTS = PestoSamplingOptions(PARAM, VAL, ...)` creates a set of options with the named parameters altered with the specified values.

`OPTS = PestoSamplingOptions(OLDOPTS, PARAM, VAL, ...)` creates a copy of OLDOPTS with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for **PestoSamplingOptions**

Definition at line 246 of file PestoSamplingOptions.m.

References mode, nIterations, obj\_type, objOutNumber, rndSeed, and samplingAlgorithm.

## 6.5.3 Member Function Documentation

### 6.5.3.1 mlhsInnerSubst< matlabtypesubstitute, this > PestoSamplingOptions::checkDependentDefaults ( matlabtypesubstitute *par* )

Should be called providing the parameter struct for dependent defaults as theta0. Does both, checking already set dependent options and defaulting based on par if not set yet.

Required fields of par:

Generated fields of this:

Definition at line 487 of file PestoSamplingOptions.m.

## 6.5.4 Member Data Documentation

### 6.5.4.1 PestoSamplingOptions::DRAM = struct

% Delayed Rejection Adaptive Metropolis options DRAM, struct containing the fields .adaptionInterval: Updates the proposal density only every adaptionInterval-th time .nTry: The number of tries in the delayed rejection scheme .regFactor: This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. Larger values equal stronger regularization. .verbosityMode: Defines the level of verbosity *silent*, *visual*, *debug*, or *text*

**Default:** struct

Definition at line 219 of file PestoSamplingOptions.m.

#### 6.5.4.2 PestoSamplingOptions::MALA = struct

% Metropolis Adaptive Langevin Algorithm options Note: This algorithm uses gradients & Hessians either provided by the user or computed by finite differences. MALA, struct containing the fields .regFactor: This factor is used for regularization in cases where the proposal covariance matrices are ill conditioned. Larger values equal stronger regularization.

**Default:** struct

Definition at line 200 of file PestoSamplingOptions.m.

#### 6.5.4.3 PestoSamplingOptions::mode = "visual"

Output mode for sampling algorithms (except DRAM, which has its own format), can be chosen as `visual`, `text`, `silent`, or `debug`. Default: `visual`.

**Default:** "visual"

##### Note

This property has custom functionality when its value is changed.

Definition at line 109 of file PestoSamplingOptions.m.

Referenced by PestoSamplingOptions().

#### 6.5.4.4 PestoSamplingOptions::nIterations = 1e5

Number of iterations, integer (1e5 is not too high, e.g. 1e6 is a more reasonable value for reliable results, but computationally more intensive).

**Default:** 1e5

##### Note

This property has custom functionality when its value is changed.

Definition at line 67 of file PestoSamplingOptions.m.

Referenced by PestoSamplingOptions().

#### 6.5.4.5 PestoSamplingOptions::obj\_type = "log-posterior"

% General options Type of objective function provided: `log-posterior` (default) or `negative log-posterior`

% Part for checking the correct setting of options

Tells the algorithm that log-posterior or log-likelihood are provided so it takes into account the correct sign for performing all algorithms correctly.

**Default:** "log-posterior"

##### Note

This property has custom functionality when its value is changed.

Definition at line 32 of file PestoSamplingOptions.m.

Referenced by PestoSamplingOptions().

#### 6.5.4.6 PestoSamplingOptions::objOutNumber = 1

Maximum number of outputs, the objective function can provide: 1 ... only objective value 2 ... objective value with gradient 3 ... objective value, gradient and Hessian (Default)

Missing values will be approximated by finite differences.

**Default:** 1

#### Note

This property has custom functionality when its value is changed.

Definition at line 120 of file PestoSamplingOptions.m.

Referenced by PestoSamplingOptions().

#### 6.5.4.7 PestoSamplingOptions::PHS = struct

% Parallel Hierarchical Sampling options PHS, struct containing the fields .nChains: Number of chains (1 mother-chain and nChains-1 auxiliary chains) .alpha: Control parameter for adaption decay. Needs values between 0 and 1. Higher values lead to faster decays, meaning that new iterations influence the single-chain proposal adaption only very weakly very quickly. .memoryLength: Control parameter for adaption. Higher values suppress strong early adaption. .regFactor: This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. nChainsarger values equal stronger regularization. .trainingTime: The iterations before the first chain swap is invoked

**Default:** struct

Definition at line 172 of file PestoSamplingOptions.m.

#### 6.5.4.8 PestoSamplingOptions::rndSeed = "shuffle"

Random seed, either a number or `shuffle` (default)

**Default:** "shuffle"

#### Note

This property has custom functionality when its value is changed.

Definition at line 47 of file PestoSamplingOptions.m.

Referenced by PestoSamplingOptions().

#### 6.5.4.9 PestoSamplingOptions::samplingAlgorithm = "PT"

Sampling algorithm, can be `PT` (parallel tempering), `PHS`, (parallel hierarchical sampling), `MALA` (Metropolis adjusted Langevin algorithm), or `DRAM` (delayed rejection adapted Metropolis algorithm, only if the `DRAM` toolbox is installed). Default value is `PT`

**Default:** "PT"

##### Note

This property has custom functionality when its value is changed.

Definition at line 55 of file `PestoSamplingOptions.m`.

Referenced by `PestoSamplingOptions()`.

#### 6.5.4.10 PestoSamplingOptions::sigma0 = "[]"

Initial covariance matrices for all chains. Example for single-chain algorithms: `opt.sigma0 = 1e5 * diag(ones(1,5));` Example for multi-chain algorithms: `opt.sigma0 = repmat(1e5*diag(ones(1,5)),opt.nTemps,1);` It is recommended to set `sigma0` by taking into account the results from a preceeding optimization.

**Default:** "[]"

Definition at line 94 of file `PestoSamplingOptions.m`.

#### 6.5.4.11 PestoSamplingOptions::theta0 = "[]"

Initialization points for all chains. If the algorithm uses multiple chains (as `PT` and `PHS`), one can specify multiple `theta0`, e.g.: `opt.theta0 = repmat([0.1,1.5,-2.5,-0.5,0.4],opt.nTemps,1);` If there is just one chain, please specify as `opt.theta0 = [1;2;3;4];` It is recommended to set `theta0` by taking into account the results from a preceeding optimization (see `Pesto` examples).

**Default:** "[]"

Definition at line 78 of file `PestoSamplingOptions.m`.

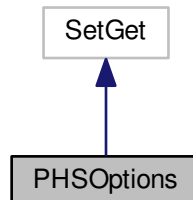
The documentation for this class was generated from the following file:

- `@PestoSamplingOptions/PestoSamplingOptions.m`

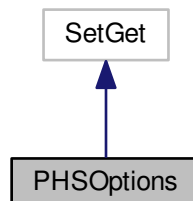
## 6.6 PHSOPTIONS Class Reference

[PHSOPTIONS](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.

Inheritance diagram for PHSOPTIONS:



Collaboration diagram for PHSOPTIONS:



### Public Member Functions

- [PHSOPTIONS](#) (matlabtypesubstitute varargin)  
[PHSOPTIONS](#) Construct a new [PHSOPTIONS](#) object.
- mlhsInnerSubst< matlabtypesubstitute, new > **copy** ()

### Public Attributes

- matlabtypesubstitute [nChains](#) = 10  
*% Parallel Hierarchical Sampling Options PHS, struct containing the fields .nChains: Initial number of temperatures (default 10) .alpha: Parameter which controls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.51. No adaption (classical Metropolis-Hastings) for 0. .memoryLength: The higher the value the more it lowers the impact of early adaption steps. Default 1. .regFactor: Regularization factor for ill conditioned covariance matrices of the adapted proposal density. Regularization might happen if the eigenvalues of the covariance matrix strongly differ in order of magnitude. In this case, the algorithm adds a small diag-matrix to the covariance matrix with elements regFactor. .trainingTime: The number of iterations before the chains start to swap. Might get used to ensure a proper local adaption of the single chains before exchanging them.*

- matlabtypesubstitute `alpha` = 0.51
  - matlabtypesubstitute `memoryLength` = 1
  - matlabtypesubstitute `regFactor` = 1e-6
- % Part for checking the correct setting of options*
- matlabtypesubstitute `trainingTime` = 1

### 6.6.1 Detailed Description

`PHSOPTIONS` provides an option container to pass options as subclass into the `PestoSamplingOptions` class.

This file is based on AMICI `amioptions.m` (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file `PHSOPTIONS.m`.

### 6.6.2 Constructor & Destructor Documentation

#### 6.6.2.1 `PHSOPTIONS::PHSOPTIONS ( matlabtypesubstitute varargin )`

`PHSOPTIONS` Construct a new `PHSOPTIONS` object.

`OPHSS = PHSOPTIONS()` creates a set of options with each option set to its default value.

`OPHSS = PHSOPTIONS(PARAM, VAL, ...)` creates a set of options with the named parameters altered with the specified values.

`OPHSS = PHSOPTIONS(OLDOPHSS, PARAM, VAL, ...)` creates a copy of `OLDOPHSS` with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for `PHSOPTIONS`

Definition at line 72 of file `PHSOPTIONS.m`.

References `alpha`, `memoryLength`, `nChains`, `regFactor`, and `trainingTime`.

### 6.6.3 Member Data Documentation

#### 6.6.3.1 `PHSOPTIONS::alpha = 0.51`

##### Note

This property has custom functionality when its value is changed.

Definition at line 56 of file `PHSOPTIONS.m`.

Referenced by `PHSOPTIONS()`.

#### 6.6.3.2 `PHSOPTIONS::memoryLength = 1`

##### Note

This property has custom functionality when its value is changed.

Definition at line 58 of file `PHSOPTIONS.m`.

Referenced by `PHSOPTIONS()`.

### 6.6.3.3 PHSOptions::nChains = 10

% Parallel Hierarchical Sampling Options PHS, struct containing the fields .nChains: Initial number of temperatures (default 10) .alpha: Parameter which controls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.51. No adaption (classical Metropolis-Hastings) for 0. .memoryLength: The higher the value the more it lowers the impact of early adaption steps. Default 1. .regFactor: Regularization factor for ill conditioned covariance matrices of the adapted proposal density. Regularization might happen if the eigenvalues of the covariance matrix strongly differ in order of magnitude. In this case, the algorithm adds a small diag-matrix to the covariance matrix with elements regFactor. .trainingTime: The number of iterations before the chains start to swap. Might get used to ensure a proper local adaption of the single chains before exchanging them.

**Default:** 10

#### Note

This property has custom functionality when its value is changed.

Definition at line 30 of file PHSOptions.m.

Referenced by PHSOptions().

### 6.6.3.4 PHSOptions::regFactor = 1e-6

% Part for checking the correct setting of options

#### Note

This property has custom functionality when its value is changed.

Definition at line 60 of file PHSOptions.m.

Referenced by PHSOptions().

### 6.6.3.5 PHSOptions::trainingTime = 1

#### Note

This property has custom functionality when its value is changed.

Definition at line 62 of file PHSOptions.m.

Referenced by PHSOptions().

The documentation for this class was generated from the following file:

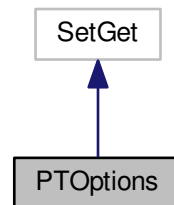
- @PHSOptions/PHSOptions.m



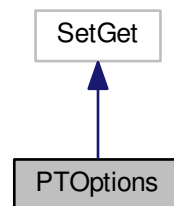
## 6.7 PTOptions Class Reference

[PTOptions](#) provides an option container to pass options as subclass into the [PestoSamplingOptions](#) class.

Inheritance diagram for PTOptions:



Collaboration diagram for PTOptions:



### Public Member Functions

- [PTOptions](#) (matlabtypesubstitute varargin)  
[PTOptions](#) Construct a new [PTOptions](#) object.
- `mlhsInnerSubst< matlabtypesubstitute, new > copy ()`

### Public Attributes

- matlabtypesubstitute `nTemps` = 10  
*% Parallel Tempering Options PT, struct containing the fields .nTemps: Initial number of temperatures (default 10) .exponentT: The initial temperatures are set by a power law to  $\wedge^{opt.exponentT}$ . (default 4) .alpha: Parameter which controls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.↔ 51. No adaption (classical Metropolis-Hastings) for 0. .temperatureAlpha: Parameter which controls the adaption degeneration velocity of the temperature adaption. Value between 0 and 1. Default 0.51. No effect for value = 0. .memoryLength: The higher the value the more it lowers the impact of early adaption steps. Default 1. .regFactor: Regularization factor for ill conditioned covariance matrices of the adapted proposal density. Regularization might happen if the eigenvalues of the covariance matrix strongly differ in order of magnitude. In this case, the algorithm adds a small diag-matrix to the covariance matrix with elements regFactor. .temperatureAdaptionScheme: Follows the temperature adaption scheme from `Vousden16` or `Lack15`. Can be set to `none` for no temperature adaption.*

- matlabtypesubstitute `exponentT` = 4
- matlabtypesubstitute `alpha` = 0.51
- matlabtypesubstitute `temperatureAlpha` = 0.51
- matlabtypesubstitute `memoryLength` = 1
- matlabtypesubstitute `regFactor` = 1e-6
- *% Part for checking the correct setting of options*
- matlabtypesubstitute `temperatureAdaptionScheme` = "Vousden16"

### 6.7.1 Detailed Description

`PTOptions` provides an option container to pass options as subclass into the `PestoSamplingOptions` class.

This file is based on AMICI `amioptions.m` (<http://icb-dcm.github.io/AMICI/>)

Definition at line 17 of file `PTOptions.m`.

### 6.7.2 Constructor & Destructor Documentation

#### 6.7.2.1 `PTOptions::PTOptions ( matlabtypesubstitute varargin )`

`PTOptions` Construct a new `PTOptions` object.

`OPTS = PTOptions()` creates a set of options with each option set to its default value.

`OPTS = PTOptions(PARAM, VAL, ...)` creates a set of options with the named parameters altered with the specified values.

`OPTS = PTOptions(OLDOPTS, PARAM, VAL, ...)` creates a copy of `OLDOPTS` with the named parameters altered with the specified value

Note to see the parameters, check the documentation page for `PTOptions`

Definition at line 83 of file `PTOptions.m`.

References `alpha`, `exponentT`, `memoryLength`, `nTemps`, `regFactor`, `temperatureAdaptionScheme`, and `temperatureAlpha`.

### 6.7.3 Member Data Documentation

#### 6.7.3.1 `PTOptions::alpha = 0.51`

##### Note

This property has custom functionality when its value is changed.

Definition at line 65 of file `PTOptions.m`.

Referenced by `PTOptions()`.

#### 6.7.3.2 PTOptions::exponentT = 4

##### Note

This property has custom functionality when its value is changed.

Definition at line 63 of file PTOptions.m.

Referenced by PTOptions().

#### 6.7.3.3 PTOptions::memoryLength = 1

##### Note

This property has custom functionality when its value is changed.

Definition at line 69 of file PTOptions.m.

Referenced by PTOptions().

#### 6.7.3.4 PTOptions::nTemps = 10

% Parallel Tempering Options PT, struct containing the fields .nTemps: Initial number of temperatures (default 10) .exponentT: The initial temperatures are set by a power law to  $\wedge^{\text{opt.exponentT}}$ . (default 4) .alpha: Parameter which controls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.51. No adaption (classical Metropolis-Hastings) for 0. .temperatureAlpha: Parameter which controls the adaption degeneration velocity of the temperature adaption. Value between 0 and 1. Default 0.51. No effect for value = 0. .memoryLength: The higher the value the more it lowers the impact of early adaption steps. Default 1. .regFactor: Regularization factor for ill conditioned covariance matrices of the adapted proposal density. Regularization might happen if the eigenvalues of the covariance matrix strongly differ in order of magnitude. In this case, the algorithm adds a small diag-matrix to the covariance matrix with elements regFactor. .temperatureAdaptionScheme: Follows the temperature adaption scheme from Vousden16 or Lacki15. Can be set to none for no temperature adaption.

**Default:** 10

##### Note

This property has custom functionality when its value is changed.

Definition at line 31 of file PTOptions.m.

Referenced by PTOptions().

#### 6.7.3.5 PTOptions::regFactor = 1e-6

% Part for checking the correct setting of options

##### Note

This property has custom functionality when its value is changed.

Definition at line 71 of file PTOptions.m.

Referenced by PTOptions().

### 6.7.3.6 PTOptions::temperatureAdaptionScheme = "Vousden16"

#### Note

This property has custom functionality when its value is changed.

Definition at line 73 of file PTOptions.m.

Referenced by PTOptions().

### 6.7.3.7 PTOptions::temperatureAlpha = 0.51

#### Note

This property has custom functionality when its value is changed.

Definition at line 67 of file PTOptions.m.

Referenced by PTOptions().

The documentation for this class was generated from the following file:

- @PTOptions/PTOptions.m

## 7 File Documentation

### 7.1 collectResults.m File Reference

[collectResults\(\)](#) collects and plots the results stored in a common folder

#### Functions

- `mlhsInnerSubst< matlabtypesubstitute, obj > collectResults (matlabtypesubstitute foldername)`  
*[collectResults\(\)](#) collects and plots the results stored in a common folder*

#### 7.1.1 Detailed Description

[collectResults\(\)](#) collects and plots the results stored in a common folder

#### 7.1.2 Function Documentation

7.1.2.1 `mlhsInnerSubst< matlabtypesubstitute, obj > collectResults ( matlabtypesubstitute foldername )`

[collectResults\(\)](#) collects and plots the results stored in a common folder

#### USAGE

`[parameters] = collectResults(foldername)`

#### History

2014/06/12 Jan Hasenauer % Initialization

## Parameters

<i>foldername</i>	Name of folder from which results are collected.
-------------------	--

## Return values

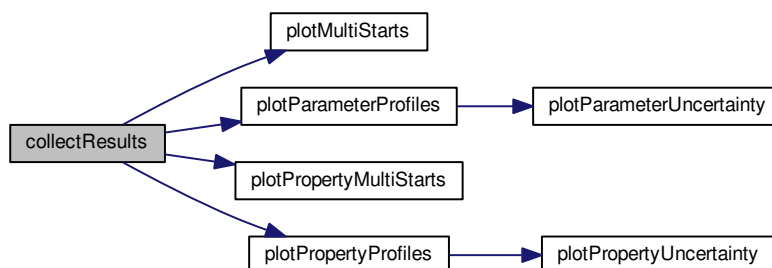
<i>parameters</i>	parameter struct.
-------------------	-------------------

Generated fields of obj:

Definition at line 17 of file collectResults.m.

References `plotMultiStarts()`, `plotParameterProfiles()`, `plotPropertyMultiStarts()`, and `plotPropertyProfiles()`.

Here is the call graph for this function:



## 7.2 getMultiStarts.m File Reference

`getMultiStarts()` computes the maximum a posterior estimate of the parameters of a user-supplied posterior function. Therefore, a multi-start local optimization is used. The parameters from the best value of the posterior function are then used as the global optimum. To ensure that the found maximum is a global one, a sufficiently high number of multistarts must be done. Those starts can be initialized with either randomly sampled parameter values, following either a uniform distribution or a latin hypercube, or they can be sampled by a user provided initial function (provided as `option.init_fun`).

## Functions

- `mlhsSubst < mlhsInnerSubst < matlabtypesubstitute, parameters >, mlhsInnerSubst < matlabtypesubstitute, fh >` `> getMultiStarts` (matlabtypesubstitute parameters, matlabtypesubstitute objective\_function, matlabtypesubstitute varargin)

*`getMultiStarts()` computes the maximum a posterior estimate of the parameters of a user-supplied posterior function. Therefore, a multi-start local optimization is used. The parameters from the best value of the posterior function are then used as the global optimum. To ensure that the found maximum is a global one, a sufficiently high number of multistarts must be done. Those starts can be initialized with either randomly sampled parameter values, following either a uniform distribution or a latin hypercube, or they can be sampled by a user provided initial function (provided as `option.init_fun`).*

- `mlhsInnerSubst < matlabtypesubstitute, stringTimePrediction >` `mtoc_subst_getMultiStarts_m_tsbust` `← cotm_updateWaitBar` (matlabtypesubstitute timePredicted)
- `noret::substitute mtoc_subst_getMultiStarts_m_tsbust cotm_saveResults` (matlabtypesubstitute parameters, matlabtypesubstitute options, matlabtypesubstitute i)

### 7.2.1 Detailed Description

`getMultiStarts()` computes the maximum a posterior estimate of the parameters of a user-supplied posterior function. Therefore, a multi-start local optimization is used. The parameters from the best value of the posterior function are then used as the global optimum. To ensure that the found maximum is a global one, a sufficiently high number of multistarts must be done. Those starts can be initialized with either randomly sampled parameter values, following either a uniform distribution or a latin hypercube, or they can be sampled by a user provided initial function (provided as `option.init_fun`).

### 7.2.2 Function Documentation

**7.2.2.1** `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, parameters >,mlhsInnerSubst< matlabtypesubstitute, fh > >`  
`getMultiStarts ( matlabtypesubstitute parameters, matlabtypesubstitute objective_function, matlabtypesubstitute varargin )`

`getMultiStarts()` computes the maximum a posterior estimate of the parameters of a user-supplied posterior function. Therefore, a multi-start local optimization is used. The parameters from the best value of the posterior function are then used as the global optimum. To ensure that the found maximum is a global one, a sufficiently high number of multistarts must be done. Those starts can be initialized with either randomly sampled parameter values, following either a uniform distribution or a latin hypercube, or they can be sampled by a user provided initial function (provided as `option.init_fun`).

Note: This function can exploit up to  $(n\_start + 1)$  workers when running in `parallel` mode.

#### USAGE

- `[...] = getMultiStarts(parameters,objective_function)`
- `[...] = getMultiStarts(parameters,objective_function,options)`
- `[parameters,fh] = getMultiStarts(...)`

`getMultiStarts()` uses the following `PestoOptions` members

- `PestoOptions::start_index`
- `PestoOptions::n_starts`
- `PestoOptions::mode`
- `PestoOptions::fh`
- `PestoOptions::fmincon`
- `PestoOptions::rng`
- `PestoOptions::proposal`
- `PestoOptions::save`
- `PestoOptions::foldername`
- `PestoOptions::trace`
- `PestoOptions::comp_type`
- `PestoOptions::tempsave`
- `PestoOptions::resetobjective`
- `PestoOptions::obj_type`
- `PestoOptions::init_threshold`
- `PestoOptions::plot_options`

## History

- 2012/05/31 Jan Hasenauer
- 2012/07/11 Jan Hasenauer
- 2014/06/11 Jan Hasenauer
- 2015/07/28 Fabian Froehlich
- 2015/11/10 Fabian Froehlich
- 2016/06/07 Paul Stapor
- 2016/10/04 Daniel Weindl
- 2016/12/04 Paul Stapor

## Parameters

<i>parameters</i>	parameter struct
<i>objective_function</i>	objective function to be optimized. This function should accept one input, the parameter vector.
<i>varargin</i>	<pre>1 getMultiStarts ( ..., options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• options A <a href="#">PestoOptions</a> object holding various options for the algorithm.</li> </ul>

## Return values

<i>parameters</i>	updated parameter object
<i>fh</i>	figure handle

## Required fields of parameters:

- `number` -- Number of parameters
- `min` -- Lower bound for each parameter
- `max` -- upper bound for each parameter name = {name1, ...}: names of the parameters
- `guess` -- initial guess for the parameters (Optional, will be initialized empty if not provided)
- `init_fun` -- function to draw starting points for local optimization, must have the structure `init_fun(theta_0, theta_min, theta_max)`. (Only required if `proposal == user-supplied`)

## Generated fields of parameters:

- `MS` -- information about multi-start optimization
  - `par0(:,i)`: starting point yielding ith MAP
  - `par(:,i)`: ith MAP
  - `logPost(i)`: log-posterior for ith MAP
  - `logPost0(i)`: log-posterior for starting point yielding ith MAP
  - `gradient(_,i)`: gradient of log-posterior at ith MAP
  - `hessian(:,i)`: hessian of log-posterior at ith MAP
  - `n_objfun(i)`: # objective evaluations used to calculate ith MAP
  - `n_iter(i)`: # iterations used to calculate ith MAP
  - `t_cpu(i)`: CPU time for calculation of ith MAP

- `exitflag(i)`: exitflag the optimizer returned for ith MAP
- `par_trace(:,i)`: parameter trace for ith MAP (if `options.trace == true`)
- `fval_trace(:,i)`: objective function value trace for ith MAP (if `options.trace == true`)
- `time_trace(:,i)`: computation time trace for ith MAP (if `options.trace == true`)

Definition at line 17 of file `getMultiStarts.m`.

References `plotMultiStarts()`.

Here is the call graph for this function:



### 7.3 `getParameterConfidenceIntervals.m` File Reference

`getParameterConfidenceIntervals()` calculates the confidence intervals for the model parameters. This is done by four approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level `alpha` (calculated by a `chi2`-distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate. The value of `CI.S` is calculated using samples for the model parameters and the according percentiles based on the confidence levels `alpha`.

#### Functions

- `mlhsInnerSubst < matlabtypesubstitute, parameters > getParameterConfidenceIntervals (matlabtypesubstitute parameters, matlabtypesubstitute alpha, matlabtypesubstitute varargin)`

*`getParameterConfidenceIntervals()` calculates the confidence intervals for the model parameters. This is done by four approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level `alpha` (calculated by a `chi2`-distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate. The value of `CI.S` is calculated using samples for the model parameters and the according percentiles based on the confidence levels `alpha`.*

#### 7.3.1 Detailed Description

`getParameterConfidenceIntervals()` calculates the confidence intervals for the model parameters. This is done by four approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level `alpha` (calculated by a `chi2`-distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate. The value of `CI.S` is calculated using samples for the model parameters and the according percentiles based on the confidence levels `alpha`.



## 7.3.2 Function Documentation

7.3.2.1 `mlhsInnerSubst < matlabtypesubstitute, parameters > getParameterConfidenceIntervals ( matlabtypesubstitute parameters, matlabtypesubstitute alpha, matlabtypesubstitute varargin )`

`getParameterConfidenceIntervals()` calculates the confidence intervals for the model parameters. This is done by four approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level `alpha` (calculated by a chi2-distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate. The value of `CI.S` is calculated using samples for the model parameters and the according percentiles based on the confidence levels `alpha`.

## USAGE

- `parameters = getParameterConfidenceIntervals(parameters, alpha)`

## History

- 2013/11/29 Jan Hasenauer
- 2016/12/01 Paul Stapor

## Parameters

<i>parameters</i>	parameter struct
<i>alpha</i>	vector with desired confidence levels for the intervals
<i>varargin</i>	<pre>1 getParameterConfidenceIntervals ( ..., options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• options A <a href="#">PestoOptions</a> instance</li> </ul>

## Return values

<i>parameters</i>	updated parameter struct
-------------------	--------------------------

## Required fields of parameters:

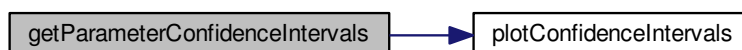
## Generated fields of parameters:

- `CI` -- Information about confidence levels
  - `local_PL`: Threshold based approach, uses a local approximation by the Hessian matrix at the MAP estimate (requires `parameters.MS`, e.g. from `getMultiStarts`)
  - `PL`: Threshold based approach, uses profile likelihoods (requires `parameters.P`, e.g. from `getParameterProfiles`)
  - `local_B`: Mass based approach, uses a local approximation by the Hessian matrix at the MAP estimate (requires `parameters.MS`, e.g. from `getMultiStarts`)
  - `S`: Bayesian approach, uses percentiles based on samples (requires `parameters.S`, e.g. from `getParameterSamples`)

Definition at line 17 of file `getParameterConfidenceIntervals.m`.

References `plotConfidenceIntervals()`.

Here is the call graph for this function:



## 7.4 `getParameterProfiles.m` File Reference

[getParameterProfiles.m](#) calculates the profiles likelihoods for the model parameters, starting from the maximum a posteriori estimate. This calculation is done by fixing the  $i$ -th parameter and repeatedly reoptimizing the likelihood/posterior estimate (for all  $i$ ). The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

### Functions

- `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, parameters >,mlhsInnerSubst< matlabtypesubstitute, fh > > getParameterProfiles (matlabtypesubstitute parameters, matlabtypesubstitute objective_function, matlabtypesubstitute varargin)`

*[getParameterProfiles.m](#) calculates the profiles likelihoods for the model parameters, starting from the maximum a posteriori estimate. This calculation is done by fixing the  $i$ -th parameter and repeatedly reoptimizing the likelihood/posterior estimate (for all  $i$ ). The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.*

#### 7.4.1 Detailed Description

[getParameterProfiles.m](#) calculates the profiles likelihoods for the model parameters, starting from the maximum a posteriori estimate. This calculation is done by fixing the  $i$ -th parameter and repeatedly reoptimizing the likelihood/posterior estimate (for all  $i$ ). The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

#### 7.4.2 Function Documentation

- 7.4.2.1 `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, parameters >,mlhsInnerSubst< matlabtypesubstitute, fh > > getParameterProfiles ( matlabtypesubstitute parameters, matlabtypesubstitute objective_function, matlabtypesubstitute varargin )`

[getParameterProfiles.m](#) calculates the profiles likelihoods for the model parameters, starting from the maximum a posteriori estimate. This calculation is done by fixing the  $i$ -th parameter and repeatedly reoptimizing the likelihood/posterior estimate (for all  $i$ ). The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

Note: This function can exploit up to  $(n\_theta + 1)$  workers when running in `parallel` mode.

## USAGE

```
[...] = getParameterProfiles(parameters, objective_function) [...] = getParameterProfiles(parameters,
objective_function, options) [parameters, fh] = getParameterProfiles(...)
```

`getParameterProfiles()` uses the following `PestoOptions` members

- `PestoOptions::calc_profiles`
- `PestoOptions::comp_type`
- `PestoOptions::dJ`
- `PestoOptions::dR_max`
- `PestoOptions::fh`
- `PestoOptions::MAP_index`
- `PestoOptions::mode`
- `PestoOptions::obj_type`
- `PestoOptions::options_getNextPoint` .guess .min .max .update .mode
- `PestoOptions::parameter_index`
- `PestoOptions::parameter_method_index`
- `PestoOptions::profile_method`
- `PestoOptions::profileReoptimizationOptions`
- `PestoOptions::plot_options`
- `PestoOptions::R_min`
- `PestoOptions::save`

## History

- 2012/05/16 Jan Hasenauer
- 2014/06/12 Jan Hasenauer
- 2016/10/04 Daniel Weindl
- 2016/10/12 Paul Stapor

## Parameters

<i>parameters</i>	parameter struct
<i>objective_function</i>	objective function to be optimized. This function should accept one input, the parameter vector.
<i>varargin</i>	<pre>1 getParameterProfiles ( ..., options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• options A <code>PestoOptions</code> object holding various options for the algorithm.</li> </ul>

## Return values

<i>parameters</i>	updated parameter struct
<i>fh</i>	figure handle

**Required fields of parameters:**

- `number` -- Number of parameters
- `min` -- Lower bound for each parameter
- `max` -- upper bound for each parameter name = {name1, ...}: names of the parameters
- `MS` -- results of global optimization, obtained using for instance the routine [getMultiStarts.m](#). MS has to contain at least
  - `par`: sorted list `n_theta` x `n_starts` of parameter estimates. The first entry is assumed to be the best one.
  - `logPost`: sorted list `n_starts` x 1 of log-posterior values corresponding to the parameters listed in `.par`.
  - `hessian`: Hessian matrix (or approximation) at the optimal point

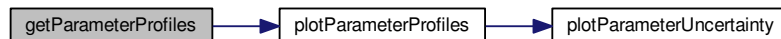
**Generated fields of parameters:**

- `P(i)` -- profile for i-th parameter
  - `par`: MAPs along profile
  - `logPost`: maximum log-posterior along profile
  - `R`: ratio

Definition at line 17 of file `getParameterProfiles.m`.

References `plotParameterProfiles()`.

Here is the call graph for this function:



## 7.5 `getParameterSamples.m` File Reference

[getParameterSamples.m](#) performs MCMC sampling of the posterior distribution. Note, the DRAM library routine `tooparameters.minox` is used internally. This function is capable of sampling with MH, AM, DRAM, MALA, PT and PHS. The sampling plotting routines should no longer be contained in here but as standalone scripts capable of using the resulting `par.S`.

**Functions**

- `mlhsInnerSubst < matlabtypesubstitute, parameters > getParameterSamples` (matlabtypesubstitute parameters, matlabtypesubstitute objFkt, matlabtypesubstitute opt)

*[getParameterSamples.m](#) performs MCMC sampling of the posterior distribution. Note, the DRAM library routine `tooparameters.minox` is used internally. This function is capable of sampling with MH, AM, DRAM, MALA, PT and PHS. The sampling plotting routines should no longer be contained in here but as standalone scripts capable of using the resulting `par.S`.*

## 7.5.1 Detailed Description

`getParameterSamples.m` performs MCMC sampling of the posterior distribution. Note, the DRAM library routine `tooparameters.minox` is used internally. This function is capable of sampling with MH, AM, DRAM, MALA, PT and PHS. The sampling plotting routines should no longer be contained in here but as standalone scripts capable of using the resulting `par.S`.

## 7.5.2 Function Documentation

7.5.2.1 `mlhsInnerSubst < matlabtypesubstitute, parameters > getParameterSamples ( matlabtypesubstitute parameters, matlabtypesubstitute objFkt, matlabtypesubstitute opt )`

`getParameterSamples.m` performs MCMC sampling of the posterior distribution. Note, the DRAM library routine `tooparameters.minox` is used internally. This function is capable of sampling with MH, AM, DRAM, MALA, PT and PHS. The sampling plotting routines should no longer be contained in here but as standalone scripts capable of using the resulting `par.S`.

`parameters`: parameter struct covering model options and results obtained by optimization, profiles and sampling. Optimization results can be used for initialization. The parameter struct should at least contain

- `par.min`: Lower parameter bounds
- `par.max`: Upper parameter bounds
- `par.number`: Number of parameters
- `par.obj_type`: Type of objective function, e.g. `log-posterior` `objFkt`: Objective function which measures the difference of model output and data `opt` : An options object holding various options for the sampling. Depending on the algorithm and particular flavor, different options must be set: For details, please visit [PE↔STOSamplingOptions.m](#)

## History

- 2012/07/11 Jan Hasenauer
- 2015/04/29 Jan Hasenauer
- 2016/10/17 Benjamin Ballnus
- 2016/10/19 Daniel Weindl
- 2016/11/04 Paul Stapor
- 2017/02/01 Benjamin Ballnus

## Return values

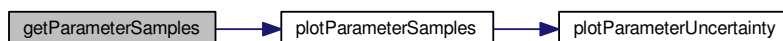
<code>parameters</code>	The provided parameters struct with the obtained sampling results added.
-------------------------	--

Required fields of `opt`:Generated fields of `parameters`:

Definition at line 17 of file `getParameterSamples.m`.

References `plotParameterSamples()`.

Here is the call graph for this function:



## 7.6 `getPropertyConfidenceIntervals.m` File Reference

[getPropertyConfidenceIntervals.m](#) calculates the confidence intervals for the model properties. This is done by three approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level  $\alpha$  (calculated by a  $\chi^2$ -distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate.

### Functions

- `mlhsInnerSubst < matlabtypesubstitute, properties > getPropertyConfidenceIntervals (matlabtypesubstitute properties, matlabtypesubstitute alpha, matlabtypesubstitute varargin)`

*[getPropertyConfidenceIntervals.m](#) calculates the confidence intervals for the model properties. This is done by three approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level  $\alpha$  (calculated by a  $\chi^2$ -distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate.*

### 7.6.1 Detailed Description

[getPropertyConfidenceIntervals.m](#) calculates the confidence intervals for the model properties. This is done by three approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level  $\alpha$  (calculated by a  $\chi^2$ -distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate.

### 7.6.2 Function Documentation

- #### 7.6.2.1 `mlhsInnerSubst < matlabtypesubstitute, properties > getPropertyConfidenceIntervals ( matlabtypesubstitute properties, matlabtypesubstitute alpha, matlabtypesubstitute varargin )`

[getPropertyConfidenceIntervals.m](#) calculates the confidence intervals for the model properties. This is done by three approaches: The values of `CI.local_PL` and `CI.PL` are determined by the point on which a threshold according to the confidence level  $\alpha$  (calculated by a  $\chi^2$ -distribution) is reached. `local_PL` computes this point by a local approximation around the MAP estimate using the Hessian matrix, `PL` uses the profile likelihoods instead. The value of `CI.local_B` is computed by using the cumulative distribution function of a local approximation of the profile based on the Hessian matrix at the MAP estimate.

## USAGE

- `properties = getPropertyConfidenceIntervals(properties, alpha)`

## History

- 2013/11/29 Jan Hasenauer
- 2016/12/01 Paul Stapor

## Parameters

<i>properties</i>	property struct
<i>alpha</i>	vector with desired confidence levels for the intervals
<i>varargin</i>	<pre>1 getPropertyConfidenceIntervals ( ..., options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• options A <a href="#">PestoOptions</a> instance</li> </ul>

## Return values

<i>properties</i>	updated properties struct
-------------------	---------------------------

## Required fields of properties:

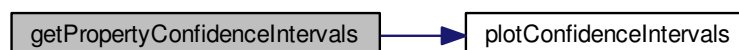
## Generated fields of properties:

- `CI` -- Information about confidence levels
  - `local_PL`: Threshold based approach, uses a local approximation by the Hessian matrix at the MAP estimate (requires `parameters.MS`, e.g. from `getMultiStarts`)
  - `PL`: Threshold based approach, uses profile likelihoods (requires `parameters.P`, e.g. from `getParameterProfiles`)
  - `local_B`: Mass based approach, uses a local approximation by the Hessian matrix at the MAP estimate (requires `parameters.MS`, e.g. from `getMultiStarts`)

Definition at line 17 of file `getPropertyConfidenceIntervals.m`.

References `plotConfidenceIntervals()`.

Here is the call graph for this function:



## 7.7 getPropertyMultiStarts.m File Reference

[getPropertyMultiStarts.m](#) evaluates the properties for the different mutli-start results.

### Functions

- `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh > > getPropertyMultiStarts (matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute varargin)`

*[getPropertyMultiStarts.m](#) evaluates the properties for the different mutli-start results.*

### 7.7.1 Detailed Description

[getPropertyMultiStarts.m](#) evaluates the properties for the different mutli-start results.

### 7.7.2 Function Documentation

**7.7.2.1** `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh > > getPropertyMultiStarts ( matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

[getPropertyMultiStarts.m](#) evaluates the properties for the different mutli-start results.

### USAGE

```
[...] = getPropertyMultiStarts(properties,parameters) [...] = getPropertyMultiStarts(properties,parameters,options)
[parameters,fh] = getPropertyMultiStarts(...)
```

`getPropertyMultiStarts()` uses the following `PestoOptions` members

- `PestoOptions::mode`
- `PestoOptions::fh`
- `PestoOptions::save`
- `PestoOptions::foldername`
- `PestoOptions::comp_type`

### History

- 2015/03/03 Jan Hasenauer
- 2016/04/10 Daniel Weindl

### Parameters

<i>properties</i>	property struct containing at least:
<i>parameters</i>	parameter struct containing at least:



## Parameters

<i>varargin</i>	<pre>1 getPropertyMultiStarts ( ..., MS, number, min, max, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• MS information about multi-start optimization</li> <li>• number Number of properties</li> <li>• min lower bound for property values</li> <li>• max upper bound for property values name = {name1,...}: names of the properties function = {function1,...}: functions to evaluate property values. These functions provide the values of the respective properties and the corresponding 1st and 2nd order derivatives.</li> <li>• options A <a href="#">PestoOptions</a> object holding the options for the algorithm.</li> </ul>
-----------------	---

## Return values

<i>properties</i>	updated parameter object containing:
<i>fh</i>	figure handle
<i>MS</i>	properties for multi-start optimization results <ul style="list-style-type: none"> <li>• par(:,i): ith MAP</li> <li>• logPost(i): log-posterior for ith MAP</li> <li>• exitflag(i): exit flag of ith MAP</li> <li>• prop(j,i): values of jth property for ith MAP</li> <li>• prop_Sigma(:,i): covariance of properties for ith MAP</li> </ul>

Required fields of parameters:

Required fields of properties:

Generated fields of properties:

Definition at line 17 of file getPropertyMultiStarts.m.

References [plotPropertyMultiStarts\(\)](#).

Here is the call graph for this function:



## 7.8 getPropertyProfiles.m File Reference

[getPropertyProfiles.m](#) calculates the profiles of user-supplied property functions, starting from the maximum a posteriori estimate. This calculation is done by varying the value of each property function respectively, starting from the value of this function at the global optimum and by reoptimizing the likelihood/posterior estimate in each variational step of the property. The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

### Functions

- `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh > > getPropertyProfiles (matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute objective_function, matlabtypesubstitute varargin)`

*[getPropertyProfiles.m](#) calculates the profiles of user-supplied property functions, starting from the maximum a posteriori estimate. This calculation is done by varying the value of each property function respectively, starting from the value of this function at the global optimum and by reoptimizing the likelihood/posterior estimate in each variational step of the property. The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.*

- `mlhsInnerSubst< matlabtypesubstitute, varargout > mtoc_subst_getPropertyProfiles_m_tsbust_cotm↔  
_obj` (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute type)
- `mlhsInnerSubst< matlabtypesubstitute, varargout > mtoc_subst_getPropertyProfiles_m_tsbust_cotm↔  
_obj_con` (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute fun\_min, matlabtypesubstitute type)
- `mlhsInnerSubst< matlabtypesubstitute, varargout > mtoc_subst_getPropertyProfiles_m_tsbust_cotm↔  
prop_fun` (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute prop\_min, matlabtypesubstitute prop\_max, matlabtypesubstitute s)
- `mlhsInnerSubst< matlabtypesubstitute, varargout > mtoc_subst_getPropertyProfiles_m_tsbust_cotm↔  
prop_con_fun` (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute prop\_min, matlabtypesubstitute prop\_max, matlabtypesubstitute s)

### 7.8.1 Detailed Description

[getPropertyProfiles.m](#) calculates the profiles of user-supplied property functions, starting from the maximum a posteriori estimate. This calculation is done by varying the value of each property function respectively, starting from the value of this function at the global optimum and by reoptimizing the likelihood/posterior estimate in each variational step of the property. The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

## 7.8.2 Function Documentation

7.8.2.1 `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh >`  
`> getPropertyProfiles ( matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute`  
`objective_function, matlabtypesubstitute varargin )`

`getPropertyProfiles.m` calculates the profiles of user-supplied property functions, starting from the maximum a posteriori estimate. This calculation is done by varying the value of each property function respectively, starting from the value of this function at the global optimum and by reoptimizing the likelihood/posterior estimate in each variational step of the property. The initial guess for the next reoptimization point is computed by extrapolation from the previous points to ensure a quick optimization.

Note: This function can exploit up to  $(n\_theta + 1)$  workers when running in `parallel` mode.

## USAGE

`[...] = getPropertyProfiles(properties, parameters, objective_function)` `[...] = getPropertyProfiles(properties, parameters, objective_function, options)` `[parameters, fh] = getPropertyProfiles(...)`

%

`getPropertyProfiles()` uses the following `PestoOptions` members

- `PestoOptions::boundary`
- `PestoOptions::calc_profiles`
- `PestoOptions::comp_type`
- `PestoOptions::dJ`
- `PestoOptions::dR_max`
- `PestoOptions::fh`
- `PestoOptions::fmincon`
- `PestoOptions::foldername`
- `PestoOptions::MAP_index`
- `PestoOptions::mode`
- `PestoOptions::obj_type`
- `PestoOptions::options_getNextPoint` `.guess` `.min` `.max` `.update` `.mode`
- `PestoOptions::plot_options`
- `PestoOptions::property_index`
- `PestoOptions::R_min`
- `PestoOptions::save`

## History

- 2012/03/02 Jan Hasenauer
- 2016/04/10 Daniel Weindl
- 2016/10/12 Paul Stapor

## Parameters

<i>properties</i>	property struct
<i>parameters</i>	parameter struct
<i>objective_function</i>	objective function to be optimized. This function should accept one input, the parameter vector.
Generated by Doxygen <i>varargin</i>	<pre>1 getPropertyProfiles ( ..., options )</pre>

**Return values**

<i>properties</i>	updated property struct
<i>fh</i>	figure handle

**Required fields of properties:**

- `number` -- Number of properties
- `min` -- Lower bound for each properties
- `max` -- upper bound for each properties `name = {name1, ...}`: names of the properties `function = {function1, ...}`: functions to evaluate property values. These functions provide the values of the respective properties and the corresponding 1st and 2nd order derivatives.

**Required fields of parameters:**

- `number` -- Number of parameters
- `min` -- Lower bound for each parameter
- `max` -- upper bound for each parameter `name = {name1, ...}`: names of the parameters
- `MS` -- results of global optimization, obtained using for instance the routine [getMultiStarts.m](#). `MS` has to contain at least
  - `par`: sorted list `n_theta x n_starts` of parameter estimates. The first entry is assumed to be the best one.
  - `logPost`: sorted list `n_starts x 1` of log-posterior values corresponding to the parameters listed in `.par`.
  - `hessian`: Hessian matrix (or approximation) at the optimal point

**Generated fields of properties:**

- `P(i)` -- profile for i-th parameter
  - `prop`: MAPs along profile
  - `par`: MAPs along profile
  - `logPost`: maximum log-posterior along profile
  - `R`: ratio

Definition at line 17 of file `getPropertyProfiles.m`.

References `plotPropertyProfiles()`.

Here is the call graph for this function:



## 7.9 `getPropertySamples.m` File Reference

[getPropertySamples.m](#) evaluates the properties for the sampled parameters.

## Functions

- `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh > > getPropertySamples (matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute varargin)`

*[getPropertySamples.m](#) evaluates the properties for the sampled parameters.*

### 7.9.1 Detailed Description

[getPropertySamples.m](#) evaluates the properties for the sampled parameters.

### 7.9.2 Function Documentation

7.9.2.1 `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, properties >,mlhsInnerSubst< matlabtypesubstitute, fh > > getPropertySamples ( matlabtypesubstitute properties, matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

[getPropertySamples.m](#) evaluates the properties for the sampled parameters.

## USAGE

```
[...] = getPropertySamples(properties,parameters) [...] = getPropertySamples(properties,parameters,options)
[parameters,fh] = getPropertySamples(...)
```

`getPropertySamples()` uses the following `PestoOptions` members

- `PestoOptions::property_index`
- `PestoOptions::mode`
- `PestoOptions::fh`
- `PestoOptions::save`
- `PestoOptions::foldername`
- `PestoOptions::comp_type`
- `PestoOptions::plot_options`
- `PestoOptions::MCMC.thinning`

## History

- 2015/04/01 Jan Hasenauer
- 2016/10/04 Daniel Weindl

## Parameters

<i>properties</i>	property struct
<i>parameters</i>	parameter struct
<i>varargin</i>	<pre>1 getPropertySamples ( ..., options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• options A <a href="#">PestoOptions</a> object holding various options for the algorithm.</li> </ul>

## Return values

<i>properties</i>	updated parameter object
<i>fh</i>	figure handle

## Required fields of properties:

- `number` -- number of parameter
- `min` -- lower bound for property values
- `max` -- upper bound for property values
- `name` -- = {`name1`,...} ... names of the parameters
- `function` -- = {`function1`,...} ... functions to evaluate property values. These functions provide the values of the respective properties and the corresponding 1st and 2nd order derivatives.

## Required fields of parameters:

- `S` -- parameter and posterior sample. `logPost` ... log-posterior function along chain `par` ... parameters along chain *Note* This struct is obtained using `getSamples.m`.

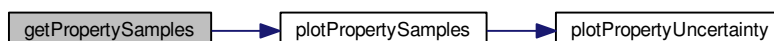
## Generated fields of properties:

- `S` -- properties for sampling results
  - `par(*,i)`: *i*th samples parameter vector
  - `logPost(i)`: log-posterior for *i*th samples parameter vector
  - `prop(j,i)`: values of *j*th property for *i*th samples parameter vector
  - `prop_Sigma(*,*,i)`: covariance of properties for *i*th samples parameter vector

Definition at line 17 of file `getPropertySamples.m`.

References `plotPropertySamples()`.

Here is the call graph for this function:



## 7.10 meigoDummy.m File Reference

Objective function wrapper for MEIGO / PSwarm / ... which need objective function *file\*name* and cannot use function handles directly.

## Functions

- `mlhsInnerSubst< matlabtypesubstitute, f > meigoDummy` (matlabtypesubstitute `theta`, matlabtypesubstitute `varargin`)

*Objective function wrapper for MEIGO / PSwarm / ... which need objective function file\*name and cannot use function handles directly.*

## 7.10.1 Detailed Description

Objective function wrapper for MEIGO / PSwarm / ... which need objective function *file\*name* and cannot use function handles directly.

## 7.10.2 Function Documentation

7.10.2.1 `mlhsInnerSubst< matlabtypesubstitute, f > meigoDummy ( matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute varargin )`

Objective function wrapper for MEIGO / PSwarm / ... which need objective function *file\*name* and cannot use function handles directly.

## Parameters

<i>theta</i>	parameter vector
<i>fun</i>	objective function handle
<i>varargin</i>	

## Return values

<i>f</i>	Objective function value
----------	--------------------------

Definition at line 17 of file `meigoDummy.m`.

7.11 `plotConfidenceIntervals.m` File Reference

[plotConfidenceIntervals.m](#) visualizes confidence intervals stored in either the parameters or properties struct `.CI`

## Functions

- `mlhsInnerSubst< matlabtypesubstitute, fh > plotConfidenceIntervals (matlabtypesubstitute pStruct, matlabtypesubstitute alpha, matlabtypesubstitute varargin)`
- `mlhsInnerSubst< matlabtypesubstitute, methodsOut > mtoc_subst_plotConfidenceIntervals_m_tsbu↔_cotm_checkMeth (matlabtypesubstitute methodsIn, matlabtypesubstitute pStruct, matlabtypesubstitute boolWarning)`

## 7.11.1 Detailed Description

[plotConfidenceIntervals.m](#) visualizes confidence intervals stored in either the parameters or properties struct `.CI`

### 7.11.2 Function Documentation

7.11.2.1 `mlhsInnerSubst< matlabtypesubstitute, fh > plotConfidenceIntervals ( matlabtypesubstitute pStruct,  
matlabtypesubstitute alpha, matlabtypesubstitute varargin )`

[plotConfidenceIntervals.m](#) visualizes confidence intervals stored in either the parameters or properties struct `.CI`

#### USAGE

`fh = plotParameterUncertainty(pStruct)` `fh = plotParameterUncertainty(pStruct, methods)` `fh = plotParameterUncertainty(pStruct, methods, options)`

`plotMultiStarts()` uses the following `PestoPlottingOptions` members

- [PestoPlottingOptions::P](#)
- [PestoPlottingOptions::S](#)
- [PestoPlottingOptions::MS](#)
- [PestoPlottingOptions::boundary](#)
- [PestoPlottingOptions::subplot\\_size\\_1D](#)
- [PestoPlottingOptions::subplot\\_indexing\\_1D](#)
- [PestoPlottingOptions::CL](#)
- [PestoPlottingOptions::hold\\_on](#)
- [PestoPlottingOptions::interval](#)
- [PestoPlottingOptions::bounds](#)
- [PestoPlottingOptions::A](#)
- [PestoPlottingOptions::add\\_points](#)
- [PestoPlottingOptions::labels](#)
- [PestoPlottingOptions::legend](#)
- [PestoPlottingOptions::op2D](#)
- [PestoPlottingOptions::fontsize](#)

#### History

- 2016/11/14 Paul Stapor

#### Parameters

<i>pStruct</i>	either the parameter or the property struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of <a href="#">plotMultiStarts.m</a> , <a href="#">getProfiles.m</a> or <a href="#">plotSamples.m</a> .
<i>alpha</i>	significance levels
<i>varargin</i>	<p><code>1 plotConfidenceIntervals ( ..., method, options )</code></p> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• method integer array, from which method confidence intervals should be plotted:</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>



## Return values

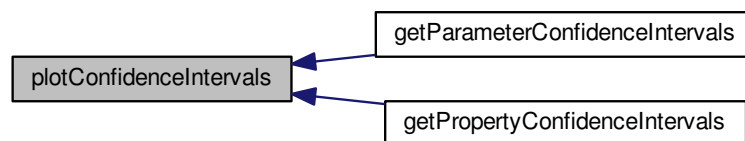
<i>fh</i>	figure handle
-----------	---------------

Required fields of pStruct:

Definition at line 17 of file plotConfidenceIntervals.m.

Referenced by `getParameterConfidenceIntervals()`, and `getPropertyConfidenceIntervals()`.

Here is the caller graph for this function:



## 7.12 plotMCMCdiagnosis.m File Reference

[plotMCMCdiagnosis.m](#) visualizes the Markov chains generated by `getSamples.m`.

## Functions

- `mlhsInnerSubst< matlabtypesubstitute, fh > plotMCMCdiagnosis` (`matlabtypesubstitute parameters`, `matlabtypesubstitute varargin`)  
*[plotMCMCdiagnosis.m](#) visualizes the Markov chains generated by `getSamples.m`.*

## 7.12.1 Detailed Description

[plotMCMCdiagnosis.m](#) visualizes the Markov chains generated by `getSamples.m`.

## 7.12.2 Function Documentation

7.12.2.1 `mlhsInnerSubst< matlabtypesubstitute, fh > plotMCMCdiagnosis` ( `matlabtypesubstitute parameters`, `matlabtypesubstitute varargin` )

[plotMCMCdiagnosis.m](#) visualizes the Markov chains generated by `getSamples.m`.

## USAGE

```
fh = plotMCMCdiagnosis(parameters) fh = plotMCMCdiagnosis(parameters,type) fh = plotMCMCdiagnosis(parameters,type,fh) fh = plotMCMCdiagnosis(parameters,type,fh,l,options)
```

## History

- 2014/06/20 Jan Hasenauer
- 2016/10/10 Daniel Weindl

**Parameters**

<i>parameters</i>	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.S). This structures is the output of <a href="#">plotMultiStarts.m</a> , <a href="#">getProfiles.m</a> or <a href="#">plotSamples.m</a> .
<i>varargin</i>	<pre>1 plotMCMCdiagnosis ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: <code>parameters</code> (default) and <code>log-posterior</code></li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of parameters which are updated. If no index is provided all parameters are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

**Return values**

<i>fh</i>	figure handle
-----------	---------------

**Required fields of parameters:**

Definition at line 17 of file `plotMCMCdiagnosis.m`.

**7.13 plotMultiStarts.m File Reference**

`plotMultiStarts` plots the result of the multi-start optimization stored in `parameters`.

**Functions**

- `mlhsInnerSubst < matlabtypesubstitute, fh > plotMultiStarts` (matlabtypesubstitute `parameters`, matlabtype-substitute `varargin`)

*plotMultiStarts plots the result of the multi-start optimization stored in parameters.*

**7.13.1 Detailed Description**

`plotMultiStarts` plots the result of the multi-start optimization stored in `parameters`.

## 7.13.2 Function Documentation

7.13.2.1 `mlhsInnerSubst < matlabtypesubstitute, fh > plotMultiStarts ( matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

`plotMultiStarts` plots the result of the multi-start optimization stored in `parameters`.

## USAGE

`fh = plotMultiStarts(parameters)` `fh = plotMultiStarts(parameters,fh)` `fh = plotMultiStarts(parameters,fh,options)`

`plotMultiStarts()` uses the following `PestoPlottingOptions` members

- [PestoPlottingOptions::add\\_points](#)
- [PestoPlottingOptions::title](#)
- [PestoPlottingOptions::draw\\_bounds](#)

History:

- 2012/05/31 Jan Hasenauer
- 2016/10/07 Daniel Weindl

## Parameters

<i>parameters</i>	parameter struct containing information about parameters and log-posterior.
<i>varargin</i>	<pre>1 plotMultiStarts ( ..., fh, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• <code>fh</code> handle of figure in which profile likelihood is plotted. If no figure handle is provided, a new figure is opened.</li> <li>• <code>options</code> options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

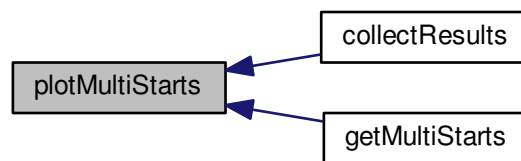
<i>fh</i>	figure handle
-----------	---------------

Required fields of `parameters`:

Definition at line 17 of file `plotMultiStarts.m`.

Referenced by `collectResults()`, and `getMultiStarts()`.

Here is the caller graph for this function:



## 7.14 `plotParameterProfiles.m` File Reference

[plotParameterProfiles.m](#) visualizes profile likelihood. Note: This routine provides an interface for `plotUncertainty.m`.

### Functions

- `mlhsInnerSubst < matlabtypesubstitute, fh > plotParameterProfiles (matlabtypesubstitute parameters, matlabtypesubstitute varargin)`  
*[plotParameterProfiles.m](#) visualizes profile likelihood. Note: This routine provides an interface for `plotUncertainty.m`.*

### 7.14.1 Detailed Description

[plotParameterProfiles.m](#) visualizes profile likelihood. Note: This routine provides an interface for `plotUncertainty.m`.

### 7.14.2 Function Documentation

- 7.14.2.1 `mlhsInnerSubst < matlabtypesubstitute, fh > plotParameterProfiles ( matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

[plotParameterProfiles.m](#) visualizes profile likelihood. Note: This routine provides an interface for `plotUncertainty.m`.

### USAGE

```
fh = plotParameterProfiles(parameters) fh = plotParameterProfiles(parameters,type) fh = plotParameterProfiles(parameters,type,fh) fh = plotParameterProfiles(parameters,type,fh,l) fh = plotParameterProfiles(parameters,type,fh,l,options)
```

### History

- 2012/05/31 Jan Hasenauer
- 2014/06/20 Jan Hasenauer
- 2016/10/10 Daniel Weindl

## Parameters

<i>parameters</i>	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structure is the output of <a href="#">plotMultiStarts.m</a> , <a href="#">getProfiles.m</a> or <a href="#">plotSamples.m</a> .
<i>varargin</i>	<pre>1 plotParameterProfiles ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D or 2D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of parameters which are updated. If no index is provided all parameters are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

<i>fh</i>	figure handle
-----------	---------------

Definition at line 17 of file `plotParameterProfiles.m`.

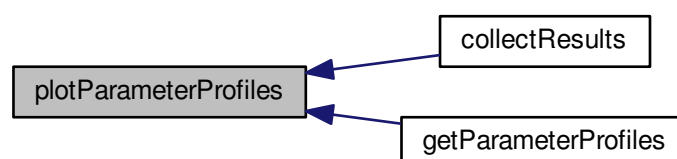
References `plotParameterUncertainty()`.

Referenced by `collectResults()`, and `getParameterProfiles()`.

Here is the call graph for this function:



Here is the caller graph for this function:



## 7.15 plotParameterSamples.m File Reference

[plotParameterSamples.m](#) visualizes MCMC samples. Note: This routine provides an interface for [plotUncertainty.m](#).

### Functions

- `mlhsInnerSubst< matlabtypesubstitute, fh > plotParameterSamples (matlabtypesubstitute parameters, matlabtypesubstitute varargin)`  
*[plotParameterSamples.m](#) visualizes MCMC samples. Note: This routine provides an interface for [plotUncertainty.m](#).*

### 7.15.1 Detailed Description

[plotParameterSamples.m](#) visualizes MCMC samples. Note: This routine provides an interface for [plotUncertainty.m](#).

### 7.15.2 Function Documentation

- 7.15.2.1 `mlhsInnerSubst< matlabtypesubstitute, fh > plotParameterSamples ( matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

[plotParameterSamples.m](#) visualizes MCMC samples. Note: This routine provides an interface for [plotUncertainty.m](#).

### USAGE

```
fh = plotParameterSamples(parameters) fh = plotParameterSamples(parameters,type) fh = plotParameterSamples(parameters,type,fh) fh = plotParameterSamples(parameters,type,fh,l) fh = plotParameterSamples(parameters,type,fh,l,options)
```

### History

- 2012/05/31 Jan Hasenauer
- 2014/06/20 Jan Hasenauer
- 2016/10/10 Daniel Weindl

### Parameters

<i>parameters</i>	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of <a href="#">plotMultiStarts.m</a> , <a href="#">getProfiles.m</a> or <a href="#">plotSamples.m</a> .
<i>varargin</i>	<pre>1 plotParameterSamples ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D or 2D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of parameters which are updated. If no index is provided all parameters are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

<i>fh</i>	figure handle
-----------	---------------

Definition at line 17 of file `plotParameterSamples.m`.

References `plotParameterUncertainty()`.

Referenced by `getParameterSamples()`.

Here is the call graph for this function:



Here is the caller graph for this function:



## 7.16 `plotParameterUncertainty.m` File Reference

[plotParameterUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in parameters.

## Functions

- `mlhsInnerSubst < matlabtypesubstitute, fh > plotParameterUncertainty` (`matlabtypesubstitute` parameters, `matlabtypesubstitute` varargin)  
*[plotParameterUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in parameters.*

### 7.16.1 Detailed Description

[plotParameterUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in parameters.

### 7.16.2 Function Documentation

#### 7.16.2.1 `mlhslInnerSubst< matlabtypesubstitute, fh > plotParameterUncertainty ( matlabtypesubstitute parameters, matlabtypesubstitute varargin )`

[plotParameterUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in parameters.

#### USAGE

```
fh = plotParameterUncertainty(parameters) fh = plotParameterUncertainty(parameters,type) fh = plot↵
ParameterUncertainty(parameters,type,fh) fh = plotParameterUncertainty(parameters,type,fh,l) fh = plot↵
ParameterUncertainty(parameters,type,fh,l,options)
```

`plotMultiStarts()` uses the following `PestoPlottingOptions` members

- [PestoPlottingOptions::P](#)
- [PestoPlottingOptions::S](#)
- [PestoPlottingOptions::MS](#)
- [PestoPlottingOptions::boundary](#)
- [PestoPlottingOptions::subplot\\_size\\_1D](#)
- [PestoPlottingOptions::subplot\\_indexing\\_1D](#)
- [PestoPlottingOptions::CL](#)
- [PestoPlottingOptions::hold\\_on](#)
- [PestoPlottingOptions::interval](#)
- [PestoPlottingOptions::bounds](#)
- [PestoPlottingOptions::A](#)
- [PestoPlottingOptions::add\\_points](#)
- [PestoPlottingOptions::labels](#)
- [PestoPlottingOptions::legend](#)
- [PestoPlottingOptions::op2D](#)
- [PestoPlottingOptions::fontsize](#)

#### History

- 2012/05/31 Jan Hasenauer
- 2014/06/20 Jan Hasenauer
- 2016/10/10 Daniel Weindl

#### Parameters

<i>parameters</i>	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of <a href="#">plotMultiStarts.m</a> , <a href="#">getProfiles.m</a> or <a href="#">plotSamples.m</a> .
<i>varargin</i>	<pre>1 plotParameterUncertainty ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D or 2D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of parameters which are updated. If no index is provided all parameters are updated.</li> </ul>
	<ul style="list-style-type: none"> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>



## Return values

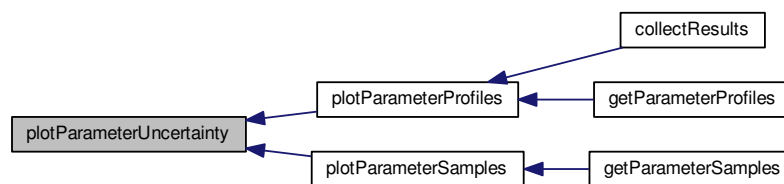
<i>fh</i>	figure handle
-----------	---------------

## Required fields of parameters:

Definition at line 17 of file plotParameterUncertainty.m.

Referenced by plotParameterProfiles(), and plotParameterSamples().

Here is the caller graph for this function:



## 7.17 plotPropertyMultiStarts.m File Reference

`plotPropertyMultiStarts` plots the result of the multi-start optimization stored in properties.

## Functions

- `mlhsInnerSubst < matlabtypesubstitute, fh > plotPropertyMultiStarts` (`matlabtypesubstitute properties`, `matlabtypesubstitute varargin`)  
*plotPropertyMultiStarts plots the result of the multi-start optimization stored in properties.*

## 7.17.1 Detailed Description

`plotPropertyMultiStarts` plots the result of the multi-start optimization stored in properties.

## 7.17.2 Function Documentation

- 7.17.2.1 `mlhsInnerSubst < matlabtypesubstitute, fh > plotPropertyMultiStarts` ( `matlabtypesubstitute properties`, `matlabtypesubstitute varargin` )

`plotPropertyMultiStarts` plots the result of the multi-start optimization stored in properties.

## USAGE

```
fh = plotPropertyMultiStarts(properties) fh = plotPropertyMultiStarts(properties,fh) fh = plotPropertyMultiStarts(properties,fh,options)
```

## History

- 2015/03/03 Jan Hasenauer

## Parameters

<i>properties</i>	property struct containing information about properties and log-posterior.
<i>varargin</i>	<pre>1 plotPropertyMultiStarts ( ..., fh, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• fh handle of figure in which profile likelihood is plotted. If no figure handle is provided, a new figure is opened.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

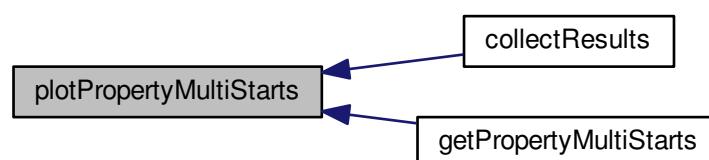
<i>fh</i>	figure handle
-----------	---------------

Required fields of properties:

Definition at line 17 of file plotPropertyMultiStarts.m.

Referenced by `collectResults()`, and `getPropertyMultiStarts()`.

Here is the caller graph for this function:



## 7.18 plotPropertyProfiles.m File Reference

[plotPropertyProfiles.m](#) visualizes profile likelihood of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## Functions

- `mlhsInnerSubst < matlabtypesubstitute, fh > plotPropertyProfiles` (matlabtypesubstitute properties, matlabtypesubstitute varargin)  
[plotPropertyProfiles.m](#) visualizes profile likelihood of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## 7.18.1 Detailed Description

[plotPropertyProfiles.m](#) visualizes profile likelihood of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## 7.18.2 Function Documentation

7.18.2.1 `mlhslInnerSubst< matlabtypesubstitute, fh > plotPropertyProfiles ( matlabtypesubstitute properties,  
matlabtypesubstitute varargin )`

[plotPropertyProfiles.m](#) visualizes profile likelihood of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## USAGE

```
fh = plotPropertyProfiles(properties) fh = plotPropertyProfiles(properties,type) fh = plotPropertyProfiles(properties,type,fh) fh = plotPropertyProfiles(properties,type,fh,I) fh = plotPropertyProfiles(properties,type,fh,I,options)
```

## History

- 2015/03/02 Jan Hasenauer
- 2016/10/10 Daniel Weindl

## Parameters

<i>properties</i>	property struct containing information about properties and results of optimization (.MS) and uncertainty analysis (.P and .S).
<i>varargin</i>	<pre>1 plotPropertyProfiles ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D or 2D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of properties which are updated. If no index is provided all properties are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

<i>fh</i>	figure handle
-----------	---------------

Definition at line 17 of file `plotPropertyProfiles.m`.

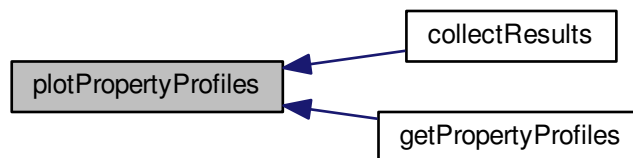
References `plotPropertyUncertainty()`.

Referenced by `collectResults()`, and `getPropertyProfiles()`.

Here is the call graph for this function:



Here is the caller graph for this function:



## 7.19 plotPropertySamples.m File Reference

[plotPropertySamples.m](#) visualizes samples of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

### Functions

- `mlhsInnerSubst< matlabtypesubstitute, fh > plotPropertySamples` (matlabtypesubstitute properties, matlabtypesubstitute varargin)

*[plotPropertySamples.m](#) visualizes samples of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).*

### 7.19.1 Detailed Description

[plotPropertySamples.m](#) visualizes samples of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## 7.19.2 Function Documentation

7.19.2.1 `mlhsInnerSubst< matlabtypesubstitute, fh > plotPropertySamples ( matlabtypesubstitute properties,  
matlabtypesubstitute varargin )`

[plotPropertySamples.m](#) visualizes samples of model properties. Note: This routine provides an interface for [plotPropertyUncertainty.m](#).

## USAGE

```
fh = plotPropertySamples(properties) fh = plotPropertySamples(properties,type) fh = plotPropertySamples(properties,type,fh) fh = plotPropertySamples(properties,type,fh,I) fh = plotPropertySamples(properties,type,fh,I,options)
```

## History

- 2015/04/01 Jan Hasenauer
- 2016/10/10 Daniel Weindl

## Parameters

<i>properties</i>	property struct containing information about properties and results of optimization (.MS) and uncertainty analysis (.P and .S).
<i>varargin</i>	<pre>1 plotPropertySamples ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D or 2D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of properties which are updated. If no index is provided all properties are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

<i>fh</i>	figure handle
-----------	---------------

Definition at line 17 of file `plotPropertySamples.m`.

References `plotPropertyUncertainty()`.

Referenced by `getPropertySamples()`.

Here is the call graph for this function:



Here is the caller graph for this function:



## 7.20 plotPropertyUncertainty.m File Reference

[plotPropertyUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in properties.

### Functions

- `mlhsInnerSubst< matlabtypesubstitute, fh > plotPropertyUncertainty` (`matlabtypesubstitute properties`, `matlabtypesubstitute varargin`)  
*[plotPropertyUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in properties.*

### 7.20.1 Detailed Description

[plotPropertyUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in properties.

### 7.20.2 Function Documentation

7.20.2.1 `mlhsInnerSubst< matlabtypesubstitute, fh > plotPropertyUncertainty` ( `matlabtypesubstitute properties`, `matlabtypesubstitute varargin` )

[plotPropertyUncertainty.m](#) visualizes profile likelihood and MCMC samples stored in properties.

### USAGE

`fh = plotPropertyUncertainty(properties,type)` `fh = plotPropertyUncertainty(properties,type,fh)` `fh = plotPropertyUncertainty(properties,type,fh,l)` `fh = plotPropertyUncertainty(properties,type,fh,l,options)`

### History

- 2012/05/31 Jan Hasenauer
- 2014/06/20 Jan Hasenauer
- 2016/10/10 Daniel Weindl

## Parameters

<i>properties</i>	properties struct.
<i>varargin</i>	<pre>1 plotPropertyUncertainty ( ..., type, fh, I, options )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• type string indicating the type of visualization: 1D</li> <li>• fh handle of figure. If no figure handle is provided, a new figure is opened.</li> <li>• I index of properties which are updated. If no index is provided all parameters are updated.</li> <li>• options options of plotting as instance of <a href="#">PestoPlottingOptions</a></li> </ul>

## Return values

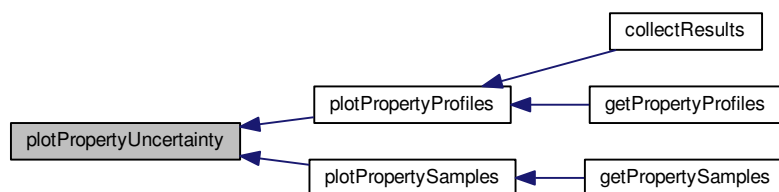
<i>fh</i>	figure handle
-----------	---------------

Required fields of properties:

Definition at line 17 of file plotPropertyUncertainty.m.

Referenced by plotPropertyProfiles(), and plotPropertySamples().

Here is the caller graph for this function:



## 7.21 runPestoTests.m File Reference

runPestoTests Run a set of PESTO unit tests

## Functions

- noret::substitute [runPestoTests](#) ()  
*runPestoTests Run a set of PESTO unit tests*

### 7.21.1 Detailed Description

runPestoTests Run a set of PESTO unit tests

## 7.22 testGradient.m File Reference

[testGradient.m](#) calculates finite difference approximations to the gradient to check an analytical version.

### Functions

- `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, g >,mlhsInnerSubst< matlabtypesubstitute, g_fd_↵  
f >,mlhsInnerSubst< matlabtypesubstitute, g_fd_b >,mlhsInnerSubst< matlabtypesubstitute, g_fd_c > >  
testGradient (matlabtypesubstitute varargin)`
- *[testGradient.m](#) calculates finite difference approximations to the gradient to check an analytical version.*
- `noret::substitute mtoc_subst_testGradient_m_tsbu cotm_error_plot` (matlabtypesubstitute g1, matlabtypesubstitute g2, matlabtypesubstitute ee)
- `noret::substitute mtoc_subst_testGradient_m_tsbu cotm_ratio_plot` (matlabtypesubstitute g1, matlabtypesubstitute g2, matlabtypesubstitute rr, matlabtypesubstitute ee)

### 7.22.1 Detailed Description

[testGradient.m](#) calculates finite difference approximations to the gradient to check an analytical version.

### 7.22.2 Function Documentation

7.22.2.1 `mlhsSubst< mlhsInnerSubst< matlabtypesubstitute, g >,mlhsInnerSubst< matlabtypesubstitute, g_fd_f  
>,mlhsInnerSubst< matlabtypesubstitute, g_fd_b >,mlhsInnerSubst< matlabtypesubstitute, g_fd_c > >  
testGradient ( matlabtypesubstitute varargin )`

[testGradient.m](#) calculates finite difference approximations to the gradient to check an analytical version.

backward differences:  $g\_fd\_f = (f(\theta + \epsilon \cdot e_i) - f(\theta)) / \epsilon$

forward differences:  $g\_fd\_b = (f(\theta) - f(\theta - \epsilon \cdot e_i)) / \epsilon$

central differences:  $g\_fd\_c = (f(\theta + \epsilon \cdot e_i) - f(\theta - \epsilon \cdot e_i)) / (2 \cdot \epsilon)$

in order to work with tensors of order n the gradient must be returned as tensor of order n+1 where the n+1th tensor dimension indexes the parameters with respect to which the differentiation was carried out

### USAGE

`[...] = testGradient(theta,fun,eps,il,ig) [g,g_fd_f,g_fd_b,g_fd_c] = testGradient(...)`

### History

- 2014/06/11 Jan Hasenauer
- 2015/01/16 Fabian Froehlich
- 2015/04/03 Jan Hasenauer
- 2015/07/28 Fabian Froehlich



## Parameters

<i>varargin</i>	<pre>1 testGradient ( theta, fun, eps, il, ig )</pre> <p><i>Required Parameters for varargin:</i></p> <ul style="list-style-type: none"> <li>• theta parameter vector at which gradient is evaluated.</li> <li>• fun function of theta for which gradients are checked.</li> <li>• eps epsilon used for finite difference approximation of gradient (eps = 1e-4).</li> <li>• il argout index/fieldname at which function values are returned (default = 1).</li> <li>• ig argout index/fieldname at which gradient values are returned (default = 2).</li> </ul>
-----------------	--

## Return values

<i>g</i>	gradient computed by f
<i>g_fd↔ _f</i>	backward differences
<i>g_fd↔ _b</i>	forward differences
<i>g_fd↔ _c</i>	central differences

Definition at line 17 of file testGradient.m.



## Index

### A

- PestoPlottingOptions, 33
- adaptionInterval
  - DRAMOptions, 14
- add\_points
  - PestoPlottingOptions, 34
- alpha
  - PHSOptions, 51
  - PTOptions, 54
- boundary
  - PestoPlottingOptions, 34
- boundary\_tol
  - PestoOptions, 20
- bounds
  - PestoPlottingOptions, 35
- calc\_profiles
  - PestoOptions, 20
- checkDependentDefaults
  - PestoSamplingOptions, 46
- CL
  - PestoPlottingOptions, 35
- collectResults
  - collectResults.m, 56
- collectResults.m, 56
  - collectResults, 56
- comp\_type
  - PestoOptions, 21
- dR\_max
  - PestoOptions, 21
- DRAMOptions, 12
  - adaptionInterval, 14
  - DRAMOptions, 14
  - nTry, 14
  - regFactor, 14
  - verbosityMode, 15
- DRAM
  - PestoSamplingOptions, 46
- dJ
  - PestoOptions, 21
- draw\_bounds
  - PestoPlottingOptions, 36
- exponentT
  - PTOptions, 54
- fh
  - PestoOptions, 21
- fh\_logPost\_trace
  - PestoPlottingOptions, 36
- fh\_par\_dis\_1D
  - PestoPlottingOptions, 37
- fh\_par\_dis\_2D
  - PestoPlottingOptions, 37

- fh\_par\_trace
  - PestoPlottingOptions, 37
- foldername
  - PestoOptions, 21
- fontsize
  - PestoPlottingOptions, 37
- getMultiStarts
  - getMultiStarts.m, 58
- getMultiStarts.m, 57
  - getMultiStarts, 58
- getParameterConfidenceIntervals
  - getParameterConfidenceIntervals.m, 61
- getParameterConfidenceIntervals.m, 60
  - getParameterConfidenceIntervals, 61
- getParameterProfiles
  - getParameterProfiles.m, 62
- getParameterProfiles.m, 62
  - getParameterProfiles, 62
- getParameterSamples
  - getParameterSamples.m, 65
- getParameterSamples.m, 64
  - getParameterSamples, 65
- getPropertyConfidenceIntervals
  - getPropertyConfidenceIntervals.m, 66
- getPropertyConfidenceIntervals.m, 66
  - getPropertyConfidenceIntervals, 66
- getPropertyMultiStarts
  - getPropertyMultiStarts.m, 68
- getPropertyMultiStarts.m, 68
  - getPropertyMultiStarts, 68
- getPropertyProfiles
  - getPropertyProfiles.m, 71
- getPropertyProfiles.m, 70
  - getPropertyProfiles, 71
- getPropertySamples
  - getPropertySamples.m, 73
- getPropertySamples.m, 72
  - getPropertySamples, 73
- group\_CI\_by
  - PestoPlottingOptions, 37
- hold\_on
  - PestoPlottingOptions, 38
- init\_threshold
  - PestoOptions, 22
- interval
  - PestoPlottingOptions, 38
- labels
  - PestoPlottingOptions, 38
- legend
  - PestoPlottingOptions, 39
- localOptimizer
  - PestoOptions, 22

- localOptimizerOptions
  - PestoOptions, 22
- MALAOptions, 15
  - MALAOptions, 16
  - regFactor, 17
- MALA
  - PestoSamplingOptions, 46
- MAP\_index
  - PestoOptions, 22
- MCMC
  - PestoPlottingOptions, 39
- mark\_constraint
  - PestoPlottingOptions, 39
- meigoDummy
  - meigoDummy.m, 75
- meigoDummy.m, 74
  - meigoDummy, 75
- memoryLength
  - PHSOOptions, 51
  - PTOptions, 55
- mode
  - PestoOptions, 23
  - PestoSamplingOptions, 47
- MS
  - PestoPlottingOptions, 40
- n\_max
  - PestoPlottingOptions, 40
- n\_starts
  - PestoOptions, 23
- nChains
  - PHSOOptions, 51
- nIterations
  - PestoSamplingOptions, 47
- nTemps
  - PTOptions, 55
- nTry
  - DRAMOptions, 14
- obj\_type
  - PestoOptions, 23
  - PestoSamplingOptions, 47
- objOutNumber
  - PestoOptions, 24
  - PestoSamplingOptions, 47
- op2D
  - PestoPlottingOptions, 41
- options\_getNextPoint
  - PestoOptions, 24
- P
  - PestoOptions, 25
  - PestoPlottingOptions, 41
- PHSOOptions, 50
  - alpha, 51
  - memoryLength, 51
  - nChains, 51
  - PHSOOptions, 51
  - regFactor, 52
  - trainingTime, 52
- PHS
  - PestoSamplingOptions, 48
- PTOptions, 53
  - alpha, 54
  - exponentT, 54
  - memoryLength, 55
  - nTemps, 55
  - PTOptions, 54
  - regFactor, 55
  - temperatureAdaptionScheme, 55
  - temperatureAlpha, 56
- parameter\_index
  - PestoOptions, 25
- PestoOptions, 17
  - boundary\_tol, 20
  - calc\_profiles, 20
  - comp\_type, 21
  - dR\_max, 21
  - dJ, 21
  - fh, 21
  - foldername, 21
  - init\_threshold, 22
  - localOptimizer, 22
  - localOptimizerOptions, 22
  - MAP\_index, 22
  - mode, 23
  - n\_starts, 23
  - obj\_type, 23
  - objOutNumber, 24
  - options\_getNextPoint, 24
  - P, 25
  - parameter\_index, 25
  - PestoOptions, 20
  - plot\_options, 25
  - profile\_integ\_index, 25
  - profile\_method, 26
  - profile\_optim\_index, 26
  - profileReoptimizationOptions, 26
  - property\_index, 26
  - proposal, 27
  - R\_min, 27
  - resetobjective, 27
  - rng, 28
  - save, 28
  - solver, 28
  - start\_index, 30
  - tempsave, 30
  - trace, 30
- PestoPlottingOptions, 31
  - A, 33
  - add\_points, 34
  - boundary, 34
  - bounds, 35
  - CL, 35
  - draw\_bounds, 36
  - fh\_logPost\_trace, 36

- fh\_par\_dis\_1D, 37
- fh\_par\_dis\_2D, 37
- fh\_par\_trace, 37
- fontsize, 37
- group\_CI\_by, 37
- hold\_on, 38
- interval, 38
- labels, 38
- legend, 39
- MCMC, 39
- mark\_constraint, 39
- MS, 40
- n\_max, 40
- op2D, 41
- P, 41
- PestoPlottingOptions, 33
- plot\_type, 41
- S, 42
- subplot\_indexing\_1D, 42
- subplot\_size\_1D, 43
- title, 43
- PestoSamplingOptions, 44
  - checkDependentDefaults, 46
  - DRAM, 46
  - MALA, 46
  - mode, 47
  - nIterations, 47
  - obj\_type, 47
  - objOutNumber, 47
  - PHS, 48
  - PestoSamplingOptions, 46
  - rndSeed, 48
  - samplingAlgorithm, 48
  - sigma0, 49
  - theta0, 49
- plot\_options
  - PestoOptions, 25
- plot\_type
  - PestoPlottingOptions, 41
- plotConfidenceIntervals
  - plotConfidenceIntervals.m, 76
- plotConfidenceIntervals.m, 75
  - plotConfidenceIntervals, 76
- plotMCMCdiagnosis
  - plotMCMCdiagnosis.m, 77
- plotMCMCdiagnosis.m, 77
  - plotMCMCdiagnosis, 77
- plotMultiStarts
  - plotMultiStarts.m, 79
- plotMultiStarts.m, 78
  - plotMultiStarts, 79
- plotParameterProfiles
  - plotParameterProfiles.m, 80
- plotParameterProfiles.m, 80
  - plotParameterProfiles, 80
- plotParameterSamples
  - plotParameterSamples.m, 82
- plotParameterSamples.m, 82
  - plotParameterSamples, 82
- plotParameterUncertainty
  - plotParameterUncertainty.m, 84
- plotParameterUncertainty.m, 83
  - plotParameterUncertainty, 84
- plotPropertyMultiStarts
  - plotPropertyMultiStarts.m, 85
- plotPropertyMultiStarts.m, 85
  - plotPropertyMultiStarts, 85
- plotPropertyProfiles
  - plotPropertyProfiles.m, 87
- plotPropertyProfiles.m, 86
  - plotPropertyProfiles, 87
- plotPropertySamples
  - plotPropertySamples.m, 89
- plotPropertySamples.m, 88
  - plotPropertySamples, 89
- plotPropertyUncertainty
  - plotPropertyUncertainty.m, 90
- plotPropertyUncertainty.m, 90
  - plotPropertyUncertainty, 90
- profile\_integ\_index
  - PestoOptions, 25
- profile\_method
  - PestoOptions, 26
- profile\_optim\_index
  - PestoOptions, 26
- profileReoptimizationOptions
  - PestoOptions, 26
- property\_index
  - PestoOptions, 26
- proposal
  - PestoOptions, 27
- R\_min
  - PestoOptions, 27
- regFactor
  - DRAMOptions, 14
  - MALAOptions, 17
  - PHSOptions, 52
  - PTOptions, 55
- resetobjective
  - PestoOptions, 27
- rndSeed
  - PestoSamplingOptions, 48
- rng
  - PestoOptions, 28
- runPestoTests.m, 91
- S
  - PestoPlottingOptions, 42
- samplingAlgorithm
  - PestoSamplingOptions, 48
- save
  - PestoOptions, 28
- sigma0
  - PestoSamplingOptions, 49
- solver
  - PestoOptions, 28

- start\_index
  - PestoOptions, [30](#)
- subplot\_indexing\_1D
  - PestoPlottingOptions, [42](#)
- subplot\_size\_1D
  - PestoPlottingOptions, [43](#)
- temperatureAdaptionScheme
  - POptions, [55](#)
- temperatureAlpha
  - POptions, [56](#)
- tempsave
  - PestoOptions, [30](#)
- testGradient
  - testGradient.m, [92](#)
- testGradient.m, [92](#)
  - testGradient, [92](#)
- theta0
  - PestoSamplingOptions, [49](#)
- title
  - PestoPlottingOptions, [43](#)
- trace
  - PestoOptions, [30](#)
- trainingTime
  - PHSOptions, [52](#)
- verbosityMode
  - DRAMOptions, [15](#)