PESTO

1.0

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Contents

1 PESTO Documentation

1.1 Introduction

Computational models are commonly used in diverse disciplines such as computational biology, engineering, or meteorology. The parameterization of the 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/main↩ ConversionReaction.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:

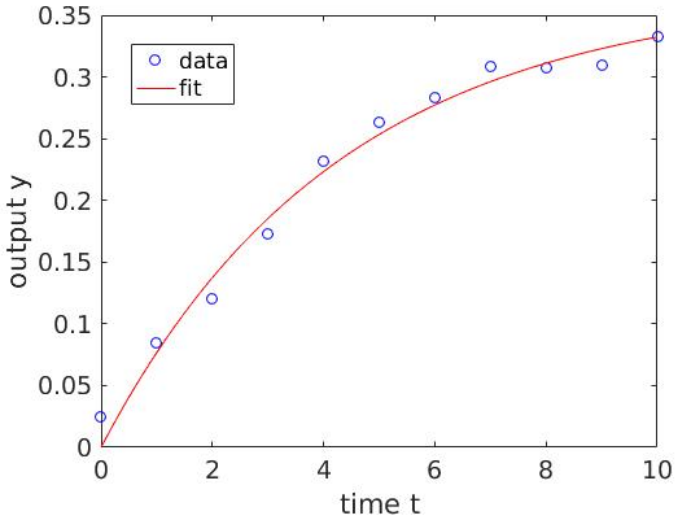


Figure 1 Plot of model fit using plotMultiStarts.m

Plot of different variants of parameter confidence intervals using plotParameterUncertainty.m:

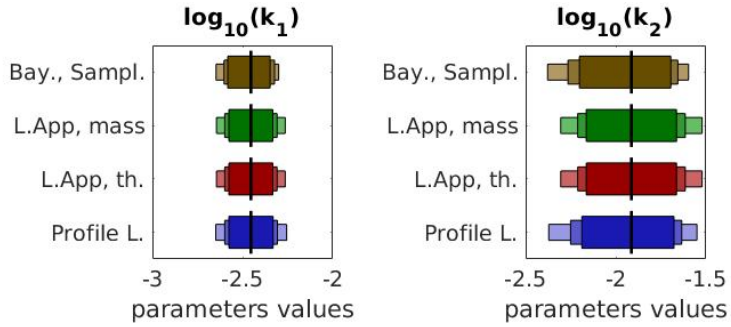


Figure 2 Plot of different variants of parameter confidence intervals using plotParameterUncertainty.m

2D plot of parameter samples using plotParameterSamples.m:

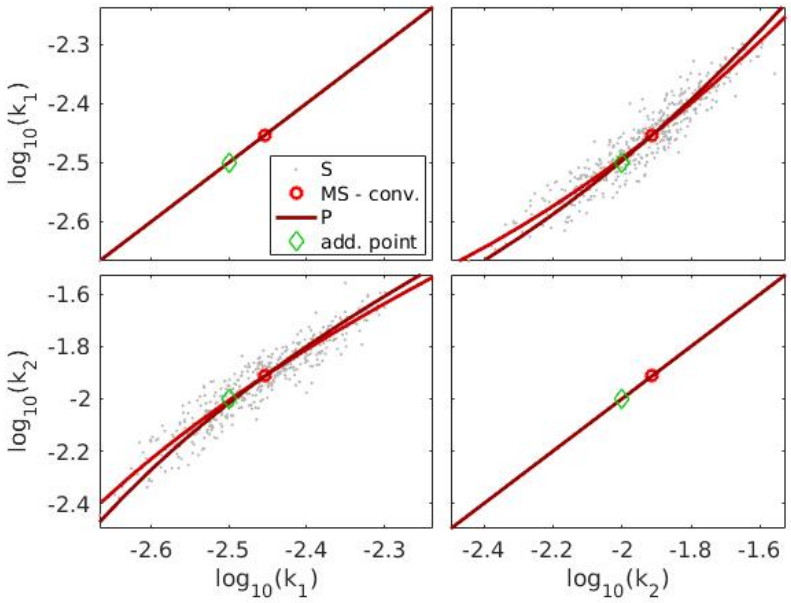


Figure 3 2D plot of parameter samples using plotParameterSamples.m

Plot of parameter samples using plotParameterSamples.m:

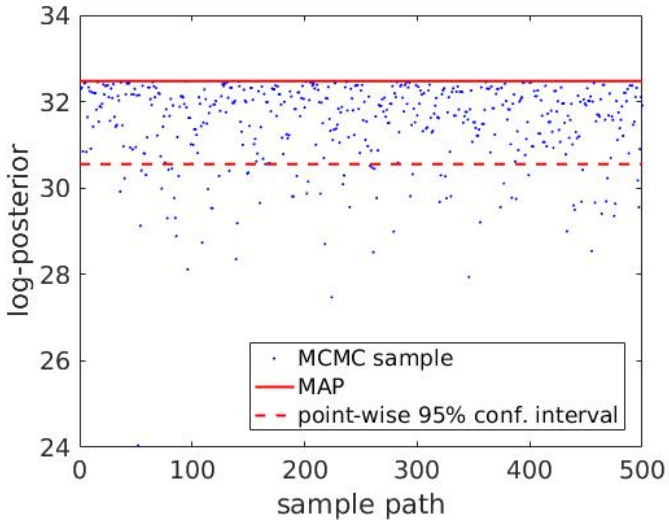


Figure 4 Plot of parameter samples using plotParameterSamples.m

Plot of property samples using plotPropertySamples.m:

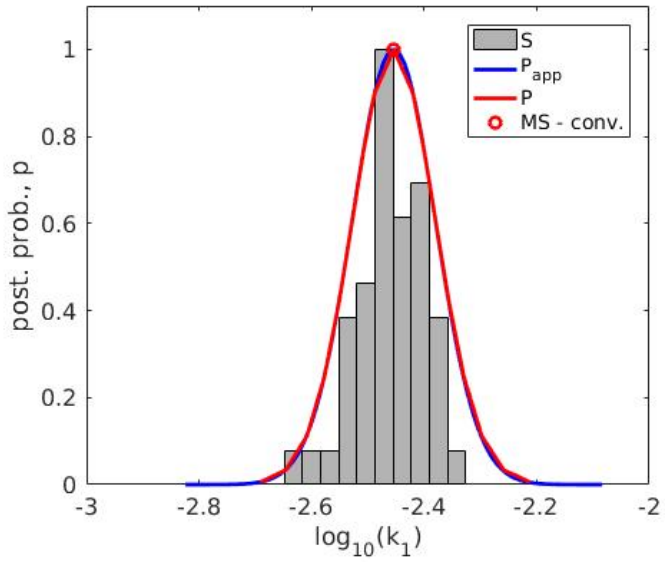


Figure 5 Plot of property samples using plotPropertySamples.m

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
getMultiStarts()	X	X		X	X	X	X	X
plotMultiStarts()	X	X		X	X	X	X	X
getParameterConfidenceIntervals()	X	X		X	X			
plotConfidenceIntervals()	X	X		X	X			
getParameterProfiles()	X	X		X	X			
plotParameterProfiles()	X	X	X	X	X	X		

	conversion reaction	enzymatic catalysis	Gaussian mixture	Hyperring	transfection	Pom1p	JakStat	ErbB
getParameterSamples()	X	X	X	X	X			
plotParameterSamples()	X	X	X	X	X			
getPropertyMultiStarts()	X				X			
plotPropertyMultiStarts()	X				X			
getPropertyConfidenceIntervals()	X				X			
getPropertyProfiles()	X				X			
plotPropertyProfiles()	X				X			
getPropertySamples()	X				X			
plotPropertySamples()	X				X			
testGradient()								
collectResults()								

3 Hierarchical Index

3.1 Class Hierarchy

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

SetGet

PestoPlottingOptions 12

PestoSamplingOptions 23

4 Class Index

4.1 Class List

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

5 File Index

5.1 File List

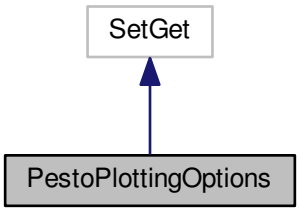
Here is a list of all documented files with brief descriptions:

6 Class Documentation

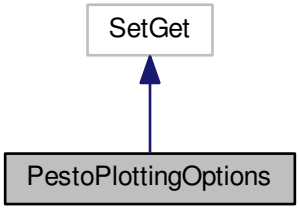
6.1 PestoPlottingOptions Class Reference

PestoPlottingOptions is 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** ()

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 parameter distribution plot. fh_par_dis = [];
- matlabtypesubstitute **fh_par_dis_2D** = "[ ]"
- matlabtypesubstitute **plot_type** = {"'parameter','posterior'"}
- matlabtypesubstitute n_max = 1e4

6.1.1 Detailed Description

PestoPlottingOptions is 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.

6.1.2 Constructor & Destructor Documentation

6.1.2.1 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

Definition at line 472 of file PestoPlottingOptions.m.

References draw_bounds, group_CI_by, hold_on, interval, mark_constraint, MCMC, n_max, and title.

6.1.3 Member Data Documentation

6.1.3.1 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.

6.1.3.2 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 x m matrix of m additional points
- .col: color used for additional points (default = [0,0,0]). This can also be a m x 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.

6.1.3.3 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 ar close to the boundaries in one or more dimensions.
- .eps: minimal distance from boundary for which points are consider to e close do 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.

6.1.3.4 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.

6.1.3.5 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
  - = `simultanous` ... point-wise confidence interval
  - = {`point-wise`,`simultanous`} ... 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.

6.1.3.6 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 PestoPlottingOptions().

6.1.3.7 PestoPlottingOptions::fh_logPost_trace = "[ ]"

figure handle for log-posterior trace plot

Default: "[ ]"

Definition at line 430 of file PestoPlottingOptions.m.

6.1.3.8 PestoPlottingOptions::fh_par_dis_1D = "[ ]"

figure handle for the parameter distribution plot. fh_par_dis = [];

Default: "[ ]"

Definition at line 448 of file PestoPlottingOptions.m.

6.1.3.9 PestoPlottingOptions::fh_par_trace = "[ ]"

figure handle for parameter trace plots.

Default: "[ ]"

Definition at line 439 of file PestoPlottingOptions.m.

6.1.3.10 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.

6.1.3.11 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 PestoPlottingOptions().

6.1.3.12 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 PestoPlottingOptions().

6.1.3.13 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 PestoPlottingOptions().

6.1.3.14 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.

6.1.3.15 PestoPlottingOptions::legend

Initial value:

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

Legend options.

- .color: background color (default = `none`).
- .box: legend outine (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.

6.1.3.16 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 PestoPlottingOptions().

6.1.3.17 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 PestoPlottingOptions().

6.1.3.18 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.

6.1.3.19 PestoPlottingOptions::n_max = 1e4

Note

This property has custom functionality when its value is changed.

Definition at line 462 of file PestoPlottingOptions.m.

Referenced by PestoPlottingOptions().

6.1.3.20 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.

6.1.3.21 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.

6.1.3.22 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.

6.1.3.23 PestoPlottingOptions::subplot_indexing_1D = "[ ]"

TODO.

Default: "[ ]"

Definition at line 96 of file PestoPlottingOptions.m.

6.1.3.24 PestoPlottingOptions::subplot_size_1D = "[ ]"

TODO from plotparameteruncertainty.

Default: "[ ]"

Definition at line 87 of file PestoPlottingOptions.m.

6.1.3.25 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 PestoPlottingOptions().

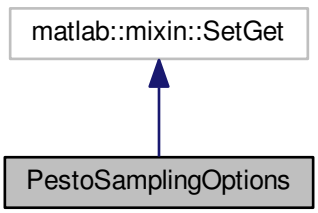
The documentation for this class was generated from the following file:

- @PestoPlottingOptions/PestoPlottingOptions.m

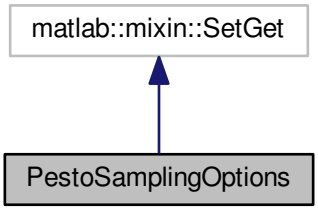
6.2 PestoSamplingOptions Class Reference

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

- mlhsInnerSubst< matlabtypesubstitute, obj > PestoOptions ()

  PestoSamplingOptions Construct a new PestoSamplingOptions object.
- mlhsInnerSubst< matlabtypesubstitute, new > **copy** ()

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 morereasonable 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 recommendet to set theta0 by taking into account the results from a preceeding 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 recommendet to set sigma0 by taking into account the results from a preceeding 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 ^opt.exponentT. (default 4) .alpha: Parameter which controlls 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 controlls 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.

- matlabtypesubstitute PHS = "[ ]"

  % Parallel Hierarchical Sampling options PHS, struct containing the filds .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 supress strong ealy 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

- matlabtypesubstitute MALA = "[ ]"

  % 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 = "[ ]"

  % 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`

6.2.1 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.

6.2.2 Member Function Documentation

6.2.2.1 mlhsInnerSubst< matlabtypesubstitute, obj > PestoSamplingOptions::PestoOptions ( )

PestoSamplingOptions Construct a new PestoSamplingOptions object.

OPTS = PestoSamplingOptions() creates a set of options with each option set to itsdefault 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 259 of file PestoSamplingOptions.m.

6.2.3 Member Data Documentation

6.2.3.1 PestoSamplingOptions::DRAM = "[ ]"

% 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: "[ ]"

Definition at line 232 of file PestoSamplingOptions.m.

6.2.3.2 PestoSamplingOptions::MALA = "[ ]"

% 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: "[ ]"

Definition at line 213 of file PestoSamplingOptions.m.

6.2.3.3 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"

Definition at line 109 of file PestoSamplingOptions.m.

6.2.3.4 PestoSamplingOptions::nIterations = 1e5

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

Default: 1e5

Definition at line 67 of file PestoSamplingOptions.m.

6.2.3.5 PestoSamplingOptions::obj_type = "log-posterior"

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

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

Default: "log-posterior"

Definition at line 32 of file PestoSamplingOptions.m.

6.2.3.6 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

Definition at line 120 of file PestoSamplingOptions.m.

6.2.3.7 PestoSamplingOptions::PHS = "[ ]"

% Parallel Hierarchical Sampling options PHS, struct containing the filds .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 supress strong ealy 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: "[ ]"

Definition at line 185 of file PestoSamplingOptions.m.

6.2.3.8 PestoSamplingOptions::PT

Initial value:

```
= struct("'nTemps', 10,    \
            'exponentT', 4,    \
            'alpha', 0.51,    \
            'temperatureAlpha', 0.51,    \
            'memoryLength', 1,    \
            'regFactor', 1e-4,    \
            'temperatureAdaptionScheme', 'Lacki15'")
```

% 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 ^opt.exponentT. (default 4) .alpha: Parameter which controlls 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 controlls 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: struct("'nTemps', 10, \ 'exponentT', 4, \ 'alpha', 0.51, \ 'temperatureAlpha', 0.51, \ 'memoryLength', 1, \ 'regFactor', 1e-4, \ 'temperatureAdaptionScheme', 'Lacki15'")

Definition at line 137 of file PestoSamplingOptions.m.

6.2.3.9 PestoSamplingOptions::rndSeed = "shuffle"

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

Default: "shuffle"

Definition at line 47 of file PestoSamplingOptions.m.

6.2.3.10 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"

Definition at line 55 of file PestoSamplingOptions.m.

6.2.3.11 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 recommendet to set sigma0 by taking into account the results from a preceeding optimization.

Default: "[ ]"

Definition at line 94 of file PestoSamplingOptions.m.

6.2.3.12 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 recommendet 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

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

foldername	Name of folder from which results are collected.

Return values

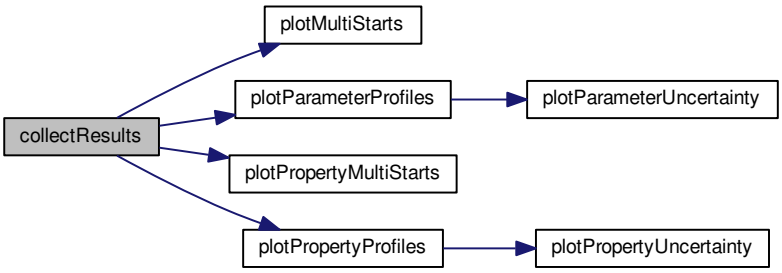
parameters	parameter struct.

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 arethen 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 arethen 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, varargout > **mtoc_subst_getMultiStarts_m_tsbus_cotm_obj** (matlabtypesubstitute varargin)
- mlhsInnerSubst< matlabtypesubstitute, varargout > **mtoc_subst_getMultiStarts_m_tsbus_cotm_obj_↩w_error_count** (matlabtypesubstitute varargin)
- mlhsInnerSubst< matlabtypesubstitute, stringTimePrediction > **mtoc_subst_getMultiStarts_m_tsbus_↩cotm_updateWaitBar** (matlabtypesubstitute timePredicted)
- noret::substitute **mtoc_subst_getMultiStarts_m_tsbus_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 arethen 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 arethen 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

parameters	parameter struct
objective_function	objective function to be optimized. This function should accept one input, the parameter vector.
varargin	`1 getMultiStarts ( ..., options )` *Required Parameters for varargin:* • options A PestoOptions object holding various options for the algorithm.

Return values

parameters	updated parameter object
fh	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 cummulative 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 cummulative 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 cummulative 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 cummulative 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

parameters	parameter struct
alpha	vector with desired confidence levels for the intervals

Return values

parameters	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

parameters	parameter struct
objective_function	objective function to be optimized. This function should accept one input, the parameter vector.
varargin	`1 getParameterProfiles ( ..., options )` *Required Parameters for varargin:* • options A PestoOptions object holding various options for the algorithm.

Return values

parameters	updated parameter struct
fh	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 of log-posterior values corresponding to the parameters listed in .par.
  - hessian: Hessian matrix (or approximation) at the optimal point

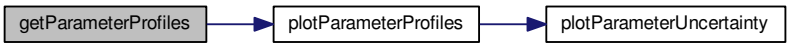
Generated fields of parameters:

- `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

— General — opt.rndSeed: Either a number or `shuffle` opt.nIterations: Number of iterations, e.g. 1e6 opt.↩ samplingAlgorithm: Specifies the code body which will be used.

Further options (details below) depend on the choice made here

`DRAM` for Delayed Rejection Adaptive Metropolis `MALA` for Metropolis Adaptive Langevin Algorithm `PT` (default) for Metropolis-Hastings, Adaptive Metropolis, Parallel Tempering `PHS`for Parallel Hierarchical Sampling opt.theta0: Initial points for all chains. If the algorithm uses multiple chains (as `PT`), one can specify multiple theta0 as in example: opt.theta0 = repmat([0.1,1.05,-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 recommendet to set theta0 by taking into account the results from a preceeding optimization. opt.sigma0: Initial covariance matrix 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 recommendet to set sigma0 by taking into account the results from a preceeding optimization.

— 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

— Metropolis Adaptive Langevin Algorithm — Note: This algorithm uses gradients & hessian either calculated by sensitivites or finite differences. opt.MALA.regFactor : This factor is used for regularization in cases where the proposal covariance matrices are ill conditioned. Larger values equal stronger regularization.

— Parallel Tempering — opt.PT.nTemps: Initial number of temperatures (default 10) opt.PT.exponentT: The initial temperatures are set by a power law to ^opt.exponentT. (default 4) opt.PT.alpha: Parameter which controlls the adaption degeneration velocity of the single-chain proposals. Value between 0 and 1. Default 0.51. No adaption for value = 0. opt.PT.temperatureAlpha: Parameter which controlls the adaption degeneration velocity of the temperature adaption. Value between 0 and 1. Default 0.51. No effect for value = 0. opt.PT.memoryLength: The higher the value the more it lowers the impact of early adaption steps. Default 1. opt.PT.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 opt.regFactor. opt.PT.temperatureAdaptionScheme: Follows the temperature adaption scheme from `Vousden16` or `Lacki15`. Can be set to `none` for no temperature adaption.

— Parallel Hierarchical Sampling — opt.PHS.nChains : Number of chains (1 `mother`-chain and opt.PHS.n↩ Chains-1 auxillary chains) opt.PHS.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. opt.PHS.memoryLength : Control parameter for adaption. Higher values supress strong ealy adaption. opt.PHS.regFactor : This factor is used for regularization in cases where the single-chain proposal covariance matrices are ill conditioned. nChainsarger values equal stronger regularization. opt.PHS.trainingTime : The iterations before the first chain swap is invoked

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

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 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 cummulative 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 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 cummulative 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 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 cummulative 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 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 cummulative 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

properties	property struct
alpha	vector with desired confidence levels for the intervals

Return values

properties	updated properties struct

Required fields of properties:

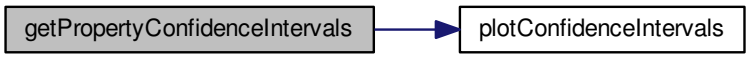
Generated fields of properties:

- `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 get↩ ParameterProfiles)
  - 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

properties	property struct containing at least:
parameters	parameter struct containing at least:
varargin	`1 getPropertyMultiStarts ( ..., MS, number, min, max, options )` *Required Parameters for varargin:* • MS information about multi-start optimization • number Number of properties • min lower bound for property values • 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. • options A PestoOptions object holding the options for the algorithm.

Return values

properties	updated parameter object containing:
fh	figure handle
MS	properties for multi-start optimization results • par(:,i): ith MAP • logPost(i): log-posterior for ith MAP • exitflag(i): exit flag of ith MAP • prop(j,i): values of jth property for ith MAP • prop_Sigma(:,:,i): covariance of properties for ith MAP

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_tsbus_cotm_obj** (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute type)
- mlhsInnerSubst< matlabtypesubstitute, varargout > **mtoc_subst_getPropertyProfiles_m_tsbus_cotm_obj_con** (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute fun_min, matlabtypesubstitute type)
- mlhsInnerSubst< matlabtypesubstitute, varargout > **mtoc_subst_getPropertyProfiles_m_tsbus_cotm_prop_fun** (matlabtypesubstitute theta, matlabtypesubstitute fun, matlabtypesubstitute prop_min, matlabtypesubstitute prop_max, matlabtypesubstitute s)
- mlhsInnerSubst< matlabtypesubstitute, varargout > **mtoc_subst_getPropertyProfiles_m_tsbus_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

properties	property struct
parameters	parameter struct
objective_function	objective function to be optimized. This function should accept one input, the parameter vector.
varargin	`1 getPropertyProfiles ( ..., options )`

Return values

properties	updated property struct
fh	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 of log-posterior values corresponding to the parameters listed in .par.
  - hessian: Hessian matrix (or approximation) at the optimal point

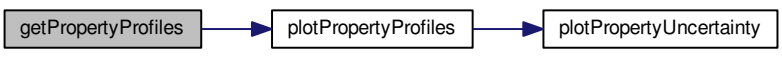
Generated fields of properties:

- `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

properties	property struct
parameters	parameter struct
varargin	`1 getPropertySamples ( ..., options )` *Required Parameters for varargin:* • options A PestoOptions object holding various options for the algorithm.

Return values

properties	updated parameter object
fh	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): ith samples parameter vector
  - logPost(i): log-posterior for ith samples parameter vector
  - prop(j,i): values of jth property for ith samples parameter vector
  - prop_Sigma(∗,∗,i): covariance of properties for ith 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 fun, 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

theta	parameter vector
fun	objective function handle
varargin	

Definition at line 17 of file meigoDummy.m.

7.11 plotConfidenceIntervals.m File Reference

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

Functions

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

  plotConfidenceIntervals.m visualizes confidence itervals stored in either the parameters or properties struct .CI
- mlhsInnerSubst< matlabtypesubstitute, methodsOut > **mtoc_subst_plotConfidenceIntervals_m_tsbus_cotm_checkMeth** (matlabtypesubstitute methodsIn, matlabtypesubstitute pStruct, matlabtypesubstitute boolWarning)

7.11.1 Detailed Description

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

USAGE

fh = plotParameterUncertainty(pStruct) fh = plotParameterUncertainty(pStruct, methods) fh = plotParameter↩ Uncertainty(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

pStruct	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 plotMultiStarts.m, getProfiles.m or plotSamples.m.
varargin	`1 plotConfidenceIntervals ( ..., method, options )` *Required Parameters for varargin:* • method integer array, from which method confidence intervals should be plotted: • options options of plotting as instance of PestoPlottingOptions

Return values

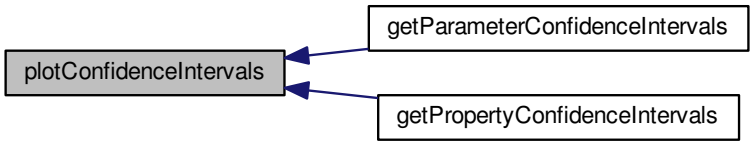
fh	figure handle

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 = plotMCM↩ Cdiagnosis(parameters,type,fh) fh = plotMCMCdiagnosis(parameters,type,fh,I) fh = plotMCMCdiagnosis(parameters,type,fh,↩ I,options)

History

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

Parameters

parameters	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.S). This structures is the output of plotMultiStarts.m, getProfiles.m or plotSamples.m.
varargin	

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, matlabtypesubstitute 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

parameters	parameter struct containing information about parameters and log-posterior.
varargin	`1 plotMultiStarts ( ..., fh, options )` *Required Parameters for varargin:* • fh handle of figure in which profile likelihood is plotted. If no figure handle is provided, a new figure is opened. • options options of plotting as instance of PestoPlottingOptions

Return values

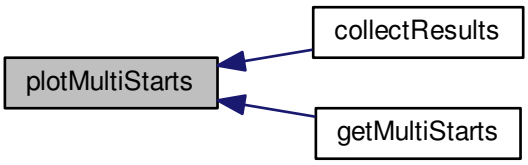
fh	figure handle

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 = plotParameter↩ Profiles(parameters,type,fh) fh = plotParameterProfiles(parameters,type,fh,I) fh = plotParameterProfiles(parameters,type,fh,↩ I,options)

History

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

Parameters

parameters	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of plotMultiStarts.m, getProfiles.m or plotSamples.m.
varargin	`1 plotParameterProfiles ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` or `2D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of parameters which are updated. If no index is provided all parameters are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

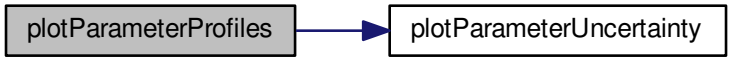
fh	figure handle

Definition at line 17 of file plotParameterProfiles.m.

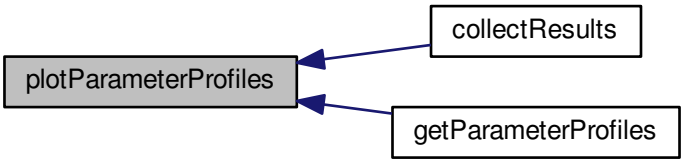
References plotParameterUncertainty().

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 = plotParameter↩ Samples(parameters,type,fh) fh = plotParameterSamples(parameters,type,fh,I) fh = plotParameter↩ Samples(parameters,type,fh,I,options)

History

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

Parameters

parameters	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of plotMultiStarts.m, getProfiles.m or plotSamples.m.
varargin	`1 plotParameterSamples ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` or `2D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of parameters which are updated. If no index is provided all parameters are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

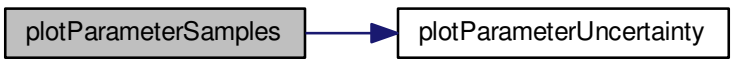
fh	figure handle

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 mlhsInnerSubst< 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 = plotParameterUncertainty(parameters,type,fh) fh = plotParameterUncertainty(parameters,type,fh,I) fh = plotParameterUncertainty(parameters,type,fh,I,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

parameters	parameter struct containing information about parameters and results of optimization (.MS) and uncertainty analysis (.P and .S). This structures is the output of plotMultiStarts.m, getProfiles.m or plotSamples.m.
varargin	`1 plotParameterUncertainty ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` or `2D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of parameters which are updated. If no index is provided all parameters are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

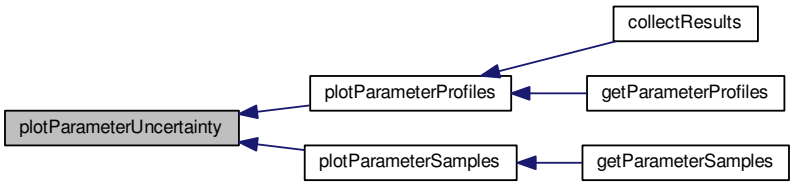
fh	figure handle

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

properties	property struct containing information about properties and log-posterior.
varargin	`1 plotPropertyMultiStarts ( ..., fh, options )` *Required Parameters for varargin:* • fh handle of figure in which profile likelihood is plotted. If no figure handle is provided, a new figure is opened. • options options of plotting as instance of PestoPlottingOptions

Return values

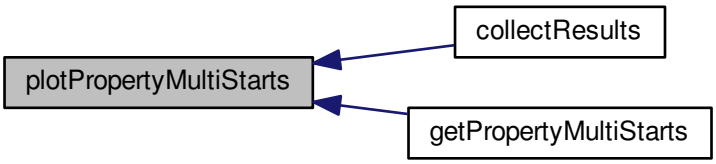
fh	figure handle

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 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.

USAGE

fh = plotPropertyProfiles(properties) fh = plotPropertyProfiles(properties,type) fh = plotProperty↩ Profiles(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

properties	property struct containing information about properties and results of optimization (.MS) and uncertainty analysis (.P and .S).
varargin	`1 plotPropertyProfiles ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` or `2D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of properties which are updated. If no index is provided all properties are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

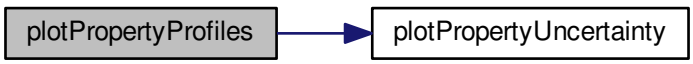
fh	figure handle

Definition at line 17 of file plotPropertyProfiles.m.

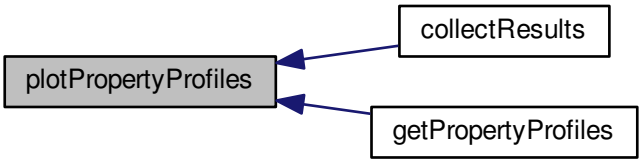
References plotPropertyUncertainty().

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

properties	property struct containing information about properties and results of optimization (.MS) and uncertainty analysis (.P and .S).
varargin	`1 plotPropertySamples ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` or `2D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of properties which are updated. If no index is provided all properties are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

fh	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,I) fh = plotPropertyUncertainty(properties,type,fh,I,options)

History

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

Parameters

properties	properties struct.
varargin	`1 plotPropertyUncertainty ( ..., type, fh, I, options )` *Required Parameters for varargin:* • type string indicating the type of visualization: `1D` • fh handle of figure. If no figure handle is provided, a new figure is opened. • I index of properties which are updated. If no index is provided all parameters are updated. • options options of plotting as instance of PestoPlottingOptions

Return values

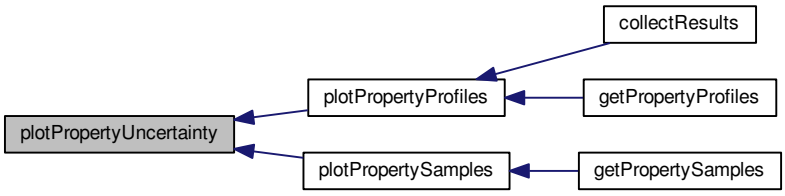
fh	figure handle

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_tsbus_cotm_error_plot** (matlabtypesubstitute g1, matlabtypesubstitute g2, matlabtypesubstitute ee)
- noret::substitute **mtoc_subst_testGradient_m_tsbus_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+eps∗e_i) - f(theta))/eps
forward differences: g_fd_b = (f(theta) - f(theta-eps∗e_i))/eps
central differences: g_fd_c = (f(theta+eps∗e_i) - f(theta-eps∗e_i))/(2∗eps)
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

varargin	`1 testGradient ( theta, fun, eps, il, ig )` *Required Parameters for varargin:* • theta parameter vector at which gradient is evaluated. • fun function of theta for which gradients are checked. • eps epsilon used for finite difference approximation of gradient (eps = 1e-4). • il argout index/fieldname at which function values are returned (default = 1). • ig argout index/fieldname at which gradient values are returned (default = 2).

Return values

g	gradient computed by f
g_fd_f	backward differences
g_fd_b	forward differences
g_fd_c	central differences

Definition at line 17 of file testGradient.m.

Index