SIPPI

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SIPPI

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About

SIPPI is a Matlab toolbox (compatible with GNU Octave) that allow sampling the solution of non-linear inverse problems with realistic a priori information.

In order to make use of SIPPI one has to

- Install and setup SIPPI
- Define the prior model, in form of the prior data structure
- Define the forward model, in form of the forward data structure, and the sippi_forward.m m-file
- Define the data and noise model, in form of the prior data structure
- Choose a method for sampling the a posteriori probability density.

Details about the implentation and the methods implemented in SIPPI can be found in [HCM12], [CHM12], [HCLM13a], [HCLM13b] and, [HCM14].

Chapter 1

Installation

1.1 SIPPI

Download the latest version of SIPPI from http://sippi.sourceforge.net.

Unpack ZIPPI.zip somewhere, for example to 'c:\Users\tmh\SIPPI'. Then setup the Matlab path to point to the appropriate SIPPI directories:

```
addpath c:\Users\tmh\SIPPI
sippi_set_path
```

1.1.1 SGeMS (optional)

To make use of the SISIM and SNESIM type priori models SGeMS needs to be available. Currently only SGeMS version 2.1 (download) for Windows is supported.

Chapter 2

Setting up SIPPI

This section contains information about how to use and control SIPPI, which requires one to

- Define the prior model, in form of the prior data structure
- Define the forward model, in form of the forward data structure, and the sippi_forward.m m-file
- Define the data and noise model, in form of the prior data structure

[For examples of how to apply SIPPI for different problems, see the section with examples].

2.1 prior: The a priori model

A priori information is defined by the prior Matlab structure. Any mumber of types of a priori models can be defined. For example a 1D uniform prior can be defined in prior{1}, and 2D Gaussian prior can be defined in prior{2}.

Once a prior data stricture has been defined, a realization from the prior model can be generated using

```
m=sippi_prior(prior);
```

The realization from the prior can be visualized using

```
sippi_plot_prior(prior);
sippi_plot_prior(prior,m);
```

A sample from the prior can be visualized using

```
m=sippi_plot_prior_sample(prior);
```

Each prior type is defined by setting a number field in the prior Matlab structure. For example, an decsriptive name (which is can be optionallyt set) decsribing the prior can be set in the name field, e.g.

```
prior{1}.name='My Prior';
```

2.1.1 Types of a priori models

5 types of a priori models are available, and can be selected by setting the type in the prior structure using e.q. prior{1}.type='gaussian'.

The GAUSSIAN type prior specifes a 1D generalized Gaussian model.

The FFTMA specifes 1D-3D Gaussian Gaussian modelm using efficient unconditional sampling,

The VISIM type prior model specifes a 1D-3D Gaussian Gaussian model, utilizing both sequential Gaussian simulation and direct sequential simulation, and conditioning the data of point support and linear average data.

The SNESIM type prior model specifes a 1D-3D multiple point statistical model, relying on traning images to infer a model multiple point statistics. This type of prior requires SGEMS to be installed.

The following section documents the properties of each type of prior model.

Examples of different types of (combinations of) a priori model can be found in the examples section.

2.1.1.1 1D Generalized Gaussian

A 1D generalized Gaussian prior model can be specified using the 'gaussian' type prior model

```
prior{1}.type='gaussian';
```

A simple 1D Gaussian distribution with mean 10, and standard deviation 2, can be specified using

```
ip=1;
prior{ip}.type='gaussian';
prior{ip}.m0=10;
prior{ip}.std=2;
```

The norm of a generalized Gaussian can be set using the 'norm' field. A generalized 1D Gaussian with mean 10, standard devation of 2, and a norm of 70, can be specified using (The norm is equivelent of the beta factor referenced in Wikipedia:Generalized_normal_distribution)

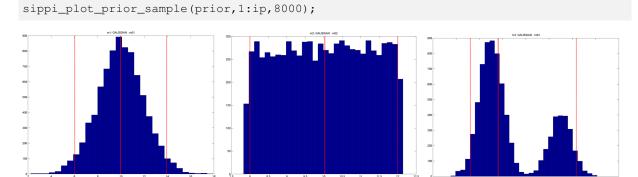
```
ip=2;
prior{ip}.type='gaussian';
prior{ip}.m0=10;
prior{ip}.std=2;
prior{ip}.norm=70;
```

A 1D distribution with an arbitrary distrution shape, can be defined by setting <code>d_target</code>, which must contain a sample of the distribtion that onw would like to replicate. For example, to generate a sample from a non-symmetric bimodal distrbution, one can use e.g.

```
% Create target distribution
N=10000;
prob_chan=0.3;
d1=randn(1,ceil(N*(1-prob_chan)))*.5+8.5;
d2=randn(1,ceil(N*(prob_chan)))*.5+11.5;
d_target=[d1(:);d2(:)];

% set the target distribution
ip=3;
prior{ip}.type='gaussian';
prior{ip}.d_target=d_target;
```

The following figure shows the 1D histrogram of a sample, consisting of 8000 realizations, generated using



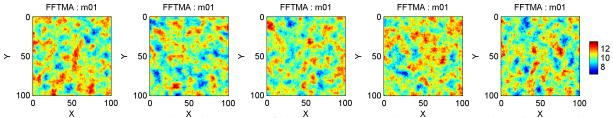
2.1.1.2 FFTMA - 3D Gaussian model

The FFT moving average method provides an efficient approach for computing unconditional realizations of a Gaussian random field.

The mean and the covariance model must be specified in the m0 and Cm fields. The format for describing the covariance model follows 'gstat'-type notation, and is described in more details in the mGstat manual.

A 2D covariance model with mean 10, and a Spherical type covariance model can be defined in a 101x101 size grid (1m between cells) using

```
im=1;
prior{im}.type='FFTMA';
prior{im}.x=[0:1:100];
prior{im}.y=[0:1:100];
prior{im}.m0=10;
prior{im}.Cm='1 Sph(10)';
```



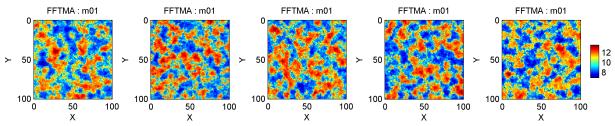
Optionally one can translate the output of the Gaussian simulation into an arbitrarily shaped 'target' distribution, using normal score transformation. Note that this transformation will ensure a certin distribution, but will alter the assumed covariance model, such the covariance model properties are no longer esnured. To ensure the covariance model properties are honored, make use of the VISIM type prior model.

```
im=1;
prior{im}.type='FFTMA';
prior{im}.x=[0:1:100];
prior{im}.y=[0:1:100];
prior{im}.m0=10;
prior{im}.Cm='1 Sph(10)';

% Create target distribution
N=10000;
prob_chan=0.5;
d1=randn(1,ceil(N*(1-prob_chan)))*.5+8.5;
d2=randn(1,ceil(N*(prob_chan)))*.5+11.5;
d_target=[d1(:);d2(:)];
prior{im}.d_target=d_target;
```

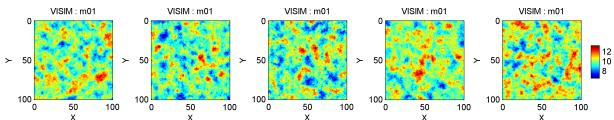
Alternatively, the normal score transformation can be defined manually to control tail behaviour using

```
N=10000;
prob_chan=0.5;
d1=randn(1,ceil(N*(1-prob_chan)))*.5+8.5;
d2=randn(1,ceil(N*(prob_chan)))*.5+11.5;
d_target=[d1(:);d2(:)];
[d_nscore,o_nscore]=nscore(d_target,1,1,min(d_target),max(d_target),0);
prior{im}.o_nscore=o_nscore;
```



2.1.1.3 VISIM

```
im=im+1;
prior{im}.type='VISIM';
prior{im}.x=[0:1:100];
prior{im}.y=[0:1:100];
prior{im}.m0=10;
prior{im}.Cm='1 Sph(10)';
```



As with the FFTMA type prior the VISIM type prior can make use of a target distribution. However, if a target distribution is set, the use of the VISIM type prior will utilize direct sequential simulation, which will ensure both histogram and covariance reproduction.

Except for the type the use of a target distribtion is similar to that of the FFTMA type prior

```
clear all; close all;
im=1;
prior(im).type='VISIM';
prior{im}.x=[0:1:40];
prior{im}.y=[0:1:40];
prior(im).m0=10;
prior{im}.Cm='1 Sph(10)';
% Create target distribution
N=10000;
prob_chan=0.5;
d1=randn(1,ceil(N*(1-prob_chan)))*.5+8.5;
d2=randn(1,ceil(N*(prob_chan)))*.5+11.5;
d_target=[d1(:);d2(:)];
prior(im).d_target=d_target;
       VISIM: m01
                          VISIM: m01
                                             VISIM: m01
                                                                 VISIM: m01
                                                                                    VISIM: m01
  10
                     10
                                        10
 20
                     20
                                        20
                                                                                                  10
  30
                     30
                                        30
                                                            30
                                                                               30
                     40
                                        40
                                                                               40
                                                            40
                 40
                       0
                             20
                                    40
                                          0
                                                20
                                                       40
                                                             0
                                                                    20
                                                                           40
```

2.1.1.4 SNESIM

2.1.2 Sampling the prior

Once the prior data structure has been defined a sample from the prior distribution can be generated using

```
m=sippi_prior(prior);
```

'm' is a Matlab data structure of the same size as the 'prior' data structure. Thus, if two prior distributions have been defined in 'prior{1}' and 'prior{2}', then 'm{1}' will hold a realization of 'prior{1}', and 'm{2}' will hold a realization of 'prior{2}'.

Each time 'm=sippi_prior(prior)' is called, a new independant realization of the prior will be generated.

2.1.3 Sequential Gibbs sampling / Conditional Resampling

All the available a priori types available allow perturbing one realization of a prior into a new realization of prior, in the vicinity of the first one. To do this we make use of sequential Gibbs sampling [HCM12]. Sequential Gibbs in essence is a type of conditional resampling. From a current realization of a prior, a number of model parameters are discarded and treated as unknown, and the simulated conditional to the fixed values of the model parameters.

In order to generate a new realization 'm2' in the vicinity of the realization 'm1' use

```
ml=sippi_prior(prior);
[m2,prior]=sippi_prior(prior,m1);
```

If this process is iterated, then a random walk in the space of a priori acceptable models will be perform. And, the collection of realization obatined, will represent a sample from prior distribution.

2.1.3.1 Controlling sequential Gibbs sampling / Conditional Resampling

All properties related to sequential Gibbs sampling can be set in the 'seq_gibbs' data struture, for each prior type. The follwing two parameters determined how the a current model is perturbed

```
prior{m}.seq_gibbs.step=1;
prior{m}.seq_gibbs.type=2;
```

2.2 data: The data and the noise

data is Matlab structure that defines any number of data and a corresponding noise model.

 $data\{1\}$ defines the first data set (which must always be defined), and ny number of additional data sets can be defined in $data\{2\}$, $data\{3\}$, ...

This allow to consider for example seismic data in data {1}, and electromagnetic data in data {2}.

For each set of data, a Gaussian noise model (both correlated and uncorrelated) can be specified). The noise model for different data types (e.g. data{1} and data{2} are independent).

Once the noise model has been defined the log-likelihood related to any model, m, with the corresponding forward response, d, can be computed using

```
logL=sippi_likelihood(data,d)
```

where d is the output of sippi_forward.

The specification of the noise model can be divided into a description of the measurement noise (mandatory) and the modeling error (optional).

2.2.1 Gaussian measurement noise

2.2.1.1 Uncorrelated Gaussian measurement noise

To define a set of observed data, [0,1,2], with an associated uncertainty defined by a Gaussian model with mean 0 and standard deviation 2, use

```
data{1}.d_obs=[0 1 2]';
data{1}.d_std=[2 2 2]';
```

which is equivalent to (as the noise model for each data the same, and independent)

```
data{1}.d_obs=[0 1 2]';
data{1}.d_std=2;
```

One can also choose to define the uncertainty using a variance as opposed to the standard deviation

```
data{1}.d_obs=[0 1 2]';
data{1}.d_var=4;
```

2.2.1.2 correlated Gaussian measurement noise

Correlated Gaussian measurement uncertainty can be specified using the Cd field, as for example

```
data{1}.Cd=[4 1 0 ; 1 4 1 ; 0 1 4];
```

Note that $data\{1\}$.Cd must be of size [NDxND], where ND is the the number of data in $data\{1\}$. d obs.

2.2.2 Gaussian modeling error

The modeling error refer to errors caused by using for example an imperfect forward model, see [HCM14]. A Gaussian model of the modeling error can is specified by the mean, dt, and the covariance, Ct. For example

```
data{1}.dt=[0 0 0];
data{1}.Ct=[4 4 4; 4 4 4; 4 4 4];
```

is equivalent to

```
data{1}.Ct=4
```

which implies a zero mean modeling error with a coavraince model where all model paremeters has a covariace of 4.

See the tomography example, for an example of accounting for correlated modeling errors.

2.3 forward: The forward model

The specification of the prior and data is intended to be generic, applicable to any inverse problem considered. The forward problem, on the other hand, is typically specific for each different inverse problem.

In order to make use of SIPPI to sample the posterior distribtion, the solution to the forward problem, must be embedded in a Matlab function with the following input and output arguments:

```
[d, forward, prior, data] = sippi_forward(m, forward, prior, data, id)
```

m is a realization of the prior model, and prior and data are the Matlab structures defining the prior and the noise model (see Prior and Data)

id is optional, and can be used to compute the forward response of a subset of the diffrent types of data available (i.e. data{1}, data{2},...)

The forward variable is a Matlab stucture that can contain any information needed to solve the forward problem. Thus, the parameters for the The forward structure is problem dependant. One option, forward_function is though generic, and point to the m-file that implements the forward problem.

The output variable d is a Matlab stucture of the same size of data. Thus, if 4 types of data have been specified, then d must also be a structures of size 4.

```
length(data) == length(d);
```

Further, d{i} must refer to an array of the same size as data{i}.d_obs.

An example of an implementation of the forward problem related to a simple line fitting problem can be:

This implementation requires that the 'x'-locations, for which the y-values of the straight line is to be computed, is specified through forward.x. Say some some y-data has been observed at locations x=[1,5,8], with the values [2,4,9], and a standard devation of 1 specifying the uncertainty, the forward stucture must be set as

```
forward.forward_function='sippi_forward_linefit';
forward.x=[1,5,8];
```

while the data structure will be

```
data{1}.d_obs=[2 4 9]
data{1}.d_std=1;
```

This implementation also requires that the prior model consists of two 1D prior types, such that

```
m=sippi_prior(prior)
```

returns the intercept in $m\{1\}$ and the gradient in $m\{2\}$.

An example of computing the forward response using an intercept of 0, and a gradients of 2 is then

```
m{1}=0;
m{2}=2;
d=sippi_forward(m, forward)
```

and the correspnding log-likelihood of m, can be computed using

```
logL=sippi_likelihood(data,d);
```

[see more deatils and example related to polynomial line fitting at polynomial line fitting]. The Examples section contains more example of implementation of different forward problems.

2.4 Validating prior, data, and forward

A simple way to test the validity of prior, data, and forward is test if the following sequence can be eavlauted without errors:

```
% Generate a realization, m, of the prior model
m=sippi_prior(prior);
% Compute the forward response
d=sippi_forward(m, forward, prior, data);
% Evaluate the log-likelihood of m
logL=sippi_likelihood(data,d);
```

Chapter 3

The a posteriori distribution

3.1 Sampling the a posteriori probability density

Once the prior, data, and forward data structures have been defined, the associated a posteriori probability can be sampled using the rejection sampler and the extended Metropolis sampler.

3.1.1 The rejection sampler

The rejection sampler provides a simples, and also in many cases inefficient, approach to sample the posterior distribution.

At each iteration of the rejection sample an independent realization, m_pro , of the prior is generated, and the model is accepted as a realization of the posterior with probability $Pacc = L(m_pro)/L_max$. It can be initiated using

```
options.mcmc.nite=400000; % Number of iteration, defaults to 1000 options.mcmc.i_plot=500; % Number of iteration between visual updates, defaults ← to 500 options=sippi_rejection(data,prior,forward,options);
```

By default the rejection sampler is run assuming a maximum likelihood of 1 (i.e. L_max = 1). If L_max is known, then it can be set using in the options.Lmax or options.logLmax fields

```
options.mcmc.Lmax=1e-9;
options=sippi_rejection(data,prior,forward,options);
```

or

```
options.mcmc.logLmax=log(1e-9);
options=sippi_rejection(data,prior,forward,options);
```

Alternatively, L_max can be automatically adjusted to reflect the maximum likelihood found while running the rejection sampler using

```
options.mcmc.adaptive_rejection=1
options=sippi_rejection(data,prior,forward,options);
```

An alternative to rejection sampling, also utilizing independant realizations of the prior, that does not require one to set L_max is the independant extended metropolis sampler, which may be computatinoally superior to the rejection sampler,

3.1.2 The extended Metropolis sampler

The extended Metropolis algorithm is in general a mcuh more efficient algroirthm for sampling the a posteriori probability

The extended Metropolis sampler can be run using

```
options.mcmc.nite=40000; % number of iterations, default nite=30000 options.mcmc.i_sample=50; % save the current model for every 50 iterations, \leftarrow default, i_sample=500
```

```
options.mcmc.i_plot=1000; % plot progress of the Metropolis sampler for every ←
100 iterations
% default i_plot=50;
options.txt='case_line_fit'; % descriptive name appended to output foldername, ←
default txt='';

[options, data, prior, forward, m_current]=sippi_metropolis(data, prior, forward, ←
options)
```

One can choose to accept all steps in the Metropolis sampler, which will result in an algorithm sampling the prior model, using

```
options.mcmc.accept_all=1; % default [0]
```

One can choose to accept models that lead to an improvement in the likelihood, which results in an optimization like algorithm using

```
options.mcmc.accept_only_improvements=1; % default [0]
```

See sippi_metropolis for more details.

3.1.2.1 Controling the step length

One optionally, as part of running the extended Metropolis sampler, automatically update the 'step'-length of the sequential Gibbs sampler in order to ensure a specific approximate acceptance ratio of the Metropolios sampler. See [CHM12] for details.

The default parameters for adjusting the step length, as given below, are set in the 'prior.seq_gibbs' structure. These parameters will be set the first time 'sippi_prior' is called with the 'prior' structure as output. The default parameters.

```
prior{m}.seq_gibbs.step_min=0;
prior{m}.seq_gibbs.step_min=1;
prior{m}.seq_gibbs.i_update_step=50
prior{m}.seq_gibbs.i_update_step_max=1000
prior{m}.seq_gibbs.n_update_history=50
prior{m}.seq_gibbs.P_target=0.3000
```

By default, adjustment of the step length, in order to achieve an acceptance ratio of 0.3 ('prior{m}.seq_gibbs.P_target'), will be performed for every 50 ('prior{m}.seq_gibbs.i_update_step') iterations, using the acceptance ratio observed in the last 50 ('prior{m}.seq_gibbs.i_update_history') iterations.

Adjustment of the step length will be performed only in the first 1000 ('prior{m}.seq_gibbs.i_update_step_max') iterations.

In order to disable automatiuc adjustment of the step length simply set

```
prior{m}.seq_gibbs.i_update_step_max=0; % disable automatic step length
```

3.1.2.2 The independent extended Metropolis sampler

The 'independent' extended Metropolis sampler, in which each proposed model is independant of the previsouly visited model, can be chosen by forcing the 'step'-length to be 1 (i.e. leading to independant samples from the prior), using e.g.

3.1.2.3 Annealing schedule

Simulated annealing like behaviour can be controlled in the options.mcmc.anneal structure. By default annealing is disabled.

Annealing consist of multiplying the the noise level using an exponentially decerasing noise factor from options.mcmc.anneal.fac_begin to options.mcmc.anneal.fac_end, from iteration number options.mcmc.anneal.i_begin to options.mcmc.anneal.i_end.

The annealing schedule can be used start a Metropolis sampler that allow to explore more of the model space in the beginning. Recall though that the posterior is not sampled until (at least) the annealing has been ended at iteration, options.mcmc.anneal.i_end, if the options.mcmc.anneal.fac_end=1. This can potentially help not to get trapped in a local minima.

To use this type of annealing, where the annealing stops after 10000 iterations, after which the algorothm performs like a regular Metropolis sampler, use for example

```
options.mcmc.anneal.i_begin=1; % default, iteration number when annealing begins options.mcmc.anneal.i_end=10000; % iteration number when annealing stops
```

which is equivalent to

```
options.mcmc.anneal.i_begin=1; % default, iteration number when annealing begins options.mcmc.anneal.i_end=10000; % iteration number when annealing stops options.mcmc.anneal.fac_begin=20; % default, noise is scaled by fac_begin at ← iteration i_begin options.mcmc.anneal.fac_end=1; % default, noise is scaled by fac_end at iteration ← i_end
```

3.2 Simulated Annealing

Simulated annealing type optimization can be setup using an annealing schedule that is enable to the entire run og the Metropolis sampler, and that ends by a noise scaling factor less than 1. This can be obtained using e.g.

```
options.mcmc.anneal.i_begin=1; % default, iteration number when annealing begins options.mcmc.anneal.i_end=options.mcmc.nite; % iteration number when annealing \leftrightarrow stops options.mcmc.anneal.fac_begin=20; % default, noise is scaled by fac_begin at \leftrightarrow iteration i_begin options.mcmc.anneal.fac_end=0.01; % 1/100 of the noise level
```

Chapter 4

Examples

SIPPI can be used as a convenient approach for unconditional an conditional simulation.

In order to use SIPPI to solve inverse problems, one must provide the solution to the forward problem. Essentially this amounts to implementing a Matlab function that solves the forward problem in using a specific input/output format. If a solution to the forward problem already exist, this can be quite easily done simply using a Matlab wrapper function.

A few implementations of solutions to forward problems are included as examples as part of SIPPI. These will be demonstrated in the following

4.1 Examples of A priori models

A prior model consisting of three independent 1D distributions (a Gaussian, Laplace, and Uniform distribution) can be defined using

```
ip=1;
prior{ip}.type='GAUSSIAN';
prior{ip}.name='Gaussian';
prior{ip}.m0=10;
prior{ip}.std=2;
ip=2;
prior{ip}.type='GAUSSIAN';
prior{ip}.name='Laplace';
prior{ip}.m0=10;
prior(ip).std=2;
prior(ip).norm=1;
prior{ip}.type='GAUSSIAN';
prior{ip}.name='Uniform';
prior(ip).m0=10;
prior{ip}.std=2;
prior{ip}.norm=60;
m=sippi_prior(prior);
    [14.3082]
                  [9.4436]
                              [10.8294]
```

1D histograms of a sample (consisting of 1000 realizations) of the prior models can be visualized using ...

```
sippi_plot_prior_sample(prior);
```

4.2 Polynomial line fitting

Here follows simple polynomial (of order 0, 1 or 2) line-fitting is considered. Example m-files can be found in the SIPPI/examples/case_linefit folder.

First, the forward problem is defined. Then examples of stochastic inversion using SIPPI is demonstrated using a a synthetic data set.

4.2.1 The forward problem

The forward problem consists of computing the y-value as a function of the x-position of the data, and the polynomial coefficients determining the line. sippi_forward_linefit.m:

```
% sippi_forward_linefit Line fit forward solver for SIPPI
%
    [d, forward, prior, data] = sippi_forward_linefit (m, forward, prior, data);
%
function [d, forward, prior, data] = sippi_forward_linefit (m, forward, prior, data);

if length (m) ==1;
    d{1} = forward.x*m{1};
elseif length (m) == 2;
    d{1} = forward.x*m{1} + m{2};
else
    d{1} = forward.x.^2*m{1} + forward.x*m{2} + m{3};
end
```

the forward.x must be an array of the x-locations, for which the y-values of the corresponding line will be evaluated.

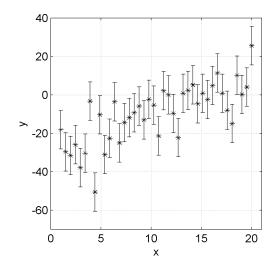
Note that the prior must be defined such that prior{1} refer to the intercept, prior{2} to the gradient, and prior{3} to the 2nd order polynomial coefficient.

If only one prior type is defined then the forward response will just be a constant, and if two prior types are defined, then the forward response will be a straight line.

4.2.2 Reference data, data, forward

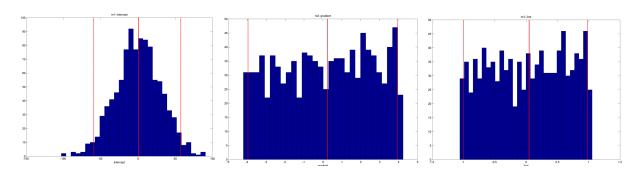
A reference data set can be computed using

```
clear all; close all;
rand('seed',1);randn('seed',1);
%% Select reference model
m_ref{1}=-30;
m_ref{2}=2;
m_ref{3}=0;
%% Setup the forward model in the 'forward' structure
forward.x=linspace(1,20,nd);
forward.forward_function='sippi_forward_linefit';
%% Compute a reference set of observed data
d=sippi_forward(m_ref, forward);
d_obs=d{1};
d_std=10;
d_obs=d_obs+randn(size(d_obs)).*d_std;
data{1}.d_obs=d_obs;
data{1}.d_std=d_std;
```



4.2.3 The prior model

```
%% Setting up the prior model
% the intercept
im=1;
prior{im}.type='gaussian';
prior(im).name='intercept';
prior(im).m0=0;
prior(im).std=30;
prior(im).m_true=m_ref(1);
% 1st order, the gradient
im=2;
prior{im}.type='gaussian';
prior(im).name='gradient';
prior(im).m0=0;
prior(im).std=4;
prior(im).norm=80;
prior{im}.m_true=m_ref{2};
% 2nd order
im=3;
prior(im).type='gaussian';
prior(im).name='2nd';
prior(im).m0=0;
prior(im).std=1;
prior(im).norm=80;
prior(im).m_true=m_ref(3);
sippi_plot_prior_sample(prior);
```



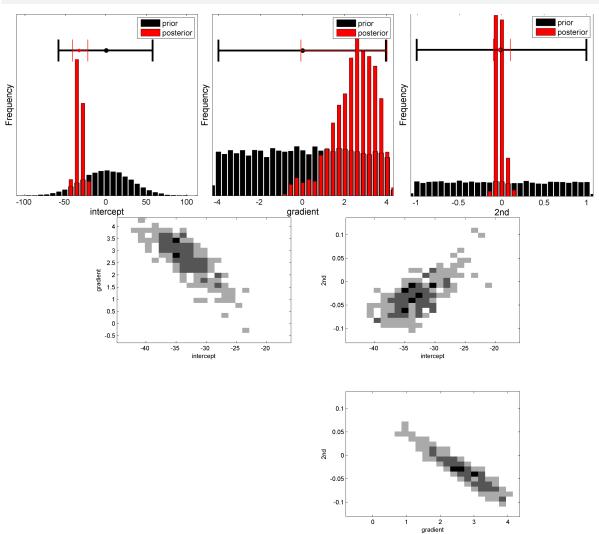
4.2.4 Setup and run the Metropolis sampler

Now, information about the model parameters can be inferred by running the extended Metropolis sampler using

```
options.mcmc.nite=40000; % Run for 40000 iterations
options.mcmc.i_sample=50; % Save every 50th visited model to disc
options.mcmc.i_plot=2500; % Plot the progress information for every 2500 ↔
    iterations
options.txt='case_line_fit_2nd_order'; % descriptive name for the output folder

[options]=sippi_metropolis(data,prior,forward,options);

% plot posterior statistics, such as 1D and 2D marginals from the prior and ↔
    posterior distributions
sippi_plot_prior_sample(options.txt);
sippi_plot_posterior(options.txt);
20140521_1644_sippi_metropolis_case_line_fit_2nd_order_m1_3_posterior_sample.png
```



4.2.5 Setup and run the rejection sampler

In a similar manner the rejection sampler can be setup and run using

```
options.mcmc.adaptive_rejection=1; % automatically adjust the normalizing ←
    likelihood
options.mcmc.nite=100000;
options=sippi_rejection(data,prior,forward,options);
```

4.3 Cross hole tomography

For now, please see [HCLM13b] for example of using SIPPI to sample the posterior for cross hole tomographic inverse problems.

Chapter 5

Bibliography

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Accepted for publication in Geophysics, xx, 2014.

Chapter 6

Reference

6.1 SIPPI

6.1.1 getinunits

```
Get object properties in specified units
V = GETINUNITS(H, PROP, UNITS) returns the object property
in the specified UNITS. It will leave the 'Units' and 'FontUnits'
property unchanged afterwards.
H is the handle of the object. If it is an M-element array of handles,
the function will return an M-by-1 cell array. PROP can be a string or
a cell array of strings. If it is a 1-by-N or N-by-1 cell array, the
function will return an M-by-N cell array of values. UNITS can be a
string or a cell array. If it is a cell array, then PROP must also be a
cell array with the same size as UNITS, and each cell element of UNITS
corresponds to a cell element of PROP.
V = GETINUNITS(H, PROP) is the same as GET(H, PROP)
Examples:
  V = GETINUNITS(H, 'Position', 'Pixels')
  V = GETINUNITS(H, {'FontSize', 'Position'}, 'Normalized')
V = GETINUNITS(H, {'FontSize', 'Position'}, {'Points', 'Pixels'})
See also GET, SET
```

6.1.2 logdet

```
LOGDET Computation of logarithm of determinant of a matrix

v = logdet(A);
    computes the logarithm of determinant of A.

Here, A should be a square matrix of double or single class.
    If A is singular, it will returns -inf.

Theoretically, this function should be functionally equivalent to log(det(A)). However, it avoids the overflow/underflow problems that are likely to happen when applying det to large matrices.

The key idea is based on the mathematical fact that the determinant of a triangular matrix equals the product of its diagonal elements. Hence, the matrix's log-determinant is equal to the sum of their logarithm
```

```
values. By keeping all computations in log-scale, the
    problem of underflow/overflow caused by product of
    many numbers can be effectively circumvented.
    The implementation is based on LU factorization.
v = logdet(A, 'chol');
    If A is positive definite, you can tell the function
    to use Cholesky factorization to accomplish the task
    using this syntax, which is substantially more efficient
    for positive definite matrix.
Remarks
   logarithm of determinant of a matrix widely occurs in the
   context of multivariate statistics. The log-pdf, entropy,
   and divergence of Gaussian distribution typically comprises
    a term in form of log-determinant. This function might be
   useful there, especially in a high-dimensional space.
   Theoretially, LU, QR can both do the job. However, LU
    factorization is substantially faster. So, for generic
   matrix, LU factorization is adopted.
    For positive definite matrices, such as covariance matrices,
    Cholesky factorization is typically more efficient. And it
    is STRONGLY RECOMMENDED that you use the chol (2nd syntax above)
    when you are sure that you are dealing with a positive definite
   matrix.
Examples
    % compute the log-determinant of a generic matrix
   A = rand(1000);
   v = logdet(A);
    % compute the log-determinant of a positive-definite matrix
    A = rand(1000);
   C = A * A';
                  % this makes C positive definite
    v = logdet(C, 'chol');
```

6.1.3 pathdef

```
PATHDEF Search path defaults.

PATHDEF returns a string that can be used as input to MATLABPATH

in order to set the path.
```

6.1.4 plotboxpos

```
PLOTBOXPOS Returns the position of the plotted axis region

pos = plotboxpos(h)

This function returns the position of the plotted region of an axis, which may differ from the actual axis position, depending on the axis limits, data aspect ratio, and plot box aspect ratio. The position is returned in the same units as the those used to define the axis itself. This function can only be used for a 2D plot.

Input variables:
```

```
h: axis handle of a 2D axis (if ommitted, current axis is used).

Output variables:

pos: four-element position vector, in same units as h
```

6.1.5 sippi_adjust_step_size

```
sippi_adjust_step_size Adjust step length length for Metropolis sampler in 
SIPPI

Call:
    step=sippi_adjust_step_size(step,P_average,P_target);

step: current step
P_current: Current acceptance ratio
P_target: preferred acceptance ratio (def=0.3);

See also sippi_compute_acceptance_rate, sippi_prior_set_steplength
```

6.1.6 sippi_anneal_adjust_noise

```
sippi_anneal_adjust_noise : Adjust noise level in annealing schedul

Call:
    [data_adjust,mcmc]=sippi_anneal_adjust_noise(data,i,mcmc,prior);

See also: sippi_metropolis, sippi_anneal_factor
```

6.1.7 sippi_anneal_factor

```
sippi_anneal_factor : compute simple noise multiplication factor for
annealing type sampling
See also sippi_metropolis, sippi_anneal_adjust_noise
```

6.1.8 sippi_colormap

```
sippi_colormap Default colormap for sippi

Call :
    sippi_colormap; % the same as sippi_colormap(3);

or :
    sippi_colormap(1) - Red Green Black
    sippi_colormap(2) - Red Green Blue Black
    sippi_colormap(3) - Jet
```

6.1.9 sippi_compute_acceptance_rate

```
sippi_compute_acceptance_rate Computes acceptance rate for the Metropolis 
    sampler in SIPPI

Call:
    P_acc=sippi_compute_acceptance_rate(acc,n_update_history);
```

6.1.10 sippi_forward

```
sippi_forward Simple forward wrapper for SIPPI

Assumes that the actual forward solver has been defined by forward.forward_function

Call:
   [d,forward,prior,data]=sippi_forward(m,forward,prior,data,id,im)
```

6.1.11 sippi_forward_traveltime

```
call :
    [d,forward_prior,data]=sippi_forward_traveltime(m,forward,prior,data,id,im)

forward.type determines the method used to compute travel times
forward.type='ray';
forward.type='fat';
forward.type='eikonal';
forward.type='eikonal';
forward.type='born';
```

6.1.12 sippi_get_sample

```
sippi_get_sample Get a posterior sample

Call :
  [reals,etype_mean,etype_var]=sippi_get_sample(data,prior,id,im,n_reals,options ↔
  );
```

6.1.13 sippi_least_squares

6.1.14 sippi_likelihood

```
sippi_likelihood Compute likelihood given an observed dataset
Call
  [logL, L, data] = sippi_likelihood(d, data);
data{1}.d_obs [N_data,1] N_data data observations
data{1}.d_std [N_data,1] N_data uncorrelated Gaussian STD
data{1}.d_var [N_data,1] N_data uncorrelated Gaussian variances
Gaussian modelization error, N(dt,Ct), is specified as
data{1}.dt [N_data,1] : Bias/mean of modelization error
data{1}.Ct [N_data, N_data] : Covariance of modelization error
data{1}.Ct [1,1] : Constant Covariance of modelization error
                    imples data{1}.Ct=ones(N_data.N_data)*data{1}.Ct;
data{id}.recomputeCD [default=0], if '1' then data{1}.iCD is recomputed
each time sippi_likelihood is called. This should be used if the noise model
changes between each call to sippi_likelihood.
data{id}.full_likelihood [default=]0; if '1' the the full likelihood
(including the determinant) is computed. This not needed if the data
civariance is constant, but if it changes, then use
data{id}.full_likelihood=1;
```

6.1.15 sippi_mcmc_init

```
sippi_mcmc_init Initialize McMC options for Metropolis and rejection sampling 
in SIPPI

Call:
   options=sippi_mcmc_init(options, prior);
```

6.1.16 sippi_metropolis

```
sippi_metropolis Extended Metropolis sampling in SIPPI

Metropolis sampling.
   See e.g. Hansen, T. M., Cordua, K. S., and Mosegaard, K., 2012.
        Inverse problems with non-trivial priors - Efficient solution through ← Sequential Gibbs Sampling.
        Computational Geosciences. doi:10.1007/s10596-011-9271-1.

Call:
        [options, data, prior, forward, m_current] = sippi_metropolis(data, prior, forward, ← options)

Input:
        data: sippi data structure
        prior: sippi prior structure
        forward: sippi forward structure
```

```
options :
  options.txt [string] : string to be used as part of all output files
  options.mcmc.nite [1] : Number if iterations
  options.mcmc.i_plot [1]: Number of iterations between updating plots
  options.mcmc.i_sample=: Number of iterations between saving model to disk
  options.mcmc.m_init : Manually chosen starting model
  options.mcmc.m\_ref : Reference known target model
  options_mcmc.accept_only_improvements [0] : Optimization
 %% PERTUBATION STRATEGY
 options.mcmc.pert_strategy.perturb_all=1; % Perturb all priors in each
                                      % iteration. def = [0]
  %% SIMULATED ANNEALING
  options.mcmc.anneal.i_end=100000; % iteration number when annealing stops
  options.mcmc.anneal.fac_begin=20; % default, noise is scaled by fac_begin at \hookleftarrow
      iteration i_begin
  iteration i_end
See also sippi_rejection
```

6.1.17 sippi_plot_current_model

```
sippi_plot_current_model Plots the current model during Metropolis sampling
Call :
    sippi_plot_current_model(mcmc,data,d,m_current,prior);
```

6.1.18 sippi_plot_data

```
sippi_plot_data plot data in SIPPI

Call.
    sippi_plot_data(d,data);
```

6.1.19 sippi_plot_loglikelihood

```
sippi_plot_loglikelihood Plot loglikelihood time series

Call :
    acc=sippi_plot_loglikelihood(logL,i_acc,N,itext)
```

6.1.20 sippi_plot_model

```
Call :
    sippi_plot_model (prior, m, im_array);

prior : Matlab structure for SIPPI prior model
    m : Matlab structure for SIPPI realization
    im_array : integer array of type of models to plot (typically 1)

Example
    m=sippi_prior(prior);
    sippi_plot_model(prior, m);

m=sippi_prior(prior);
    sippi_plot_model(prior, m, 2);

See also sippi_plot_prior
```

6.1.21 sippi_plot_movie

```
sippi_plot_movie plot movie of prior and posterior realizations

Call:
    sippi_plot_movie(fname);
    sippi_plot_movie(fname,im_array,n_frames,skip_burnin);
        fname : name of folder with results (e.g. options.txt)
        im_array : array of indexes of model parameters to make into movies
        n_frames [200] : number of frames in movie
        skip_burnin [200] : start movie after burn_in;

Ex:
    sippi_plot_movie('20130812_Metropolis');
    sippi_plot_movie(options.txt);

%% 1000 realization including burn-in, for prior number 1
    sippi_plot_movie('20130812_Metropolis',1,1000,0);
```

6.1.22 sippi_plot_posterior

```
sippi_plot_posterior Plot statistics from posterior sample

Call :
    sippi_plot_posterior(fname,im_arr,prior,options,n_reals);

See also sippi_plot_prior
```

6.1.23 sippi_plot_prior

```
sippi_plot_prior Plot a sample of the prior in SIPPI

Call :
    sippi_plot_prior(prior,ip,n_reals,cax,supt);

See also sippi_plot_posterior, sippi_plot_model
```

6.1.24 sippi_prior

```
sippi_prior A priori models for SIPPI
 To generate a realization of the prior model defined by the prior structure use \leftrightarrow
   [m_propose, prior] = sippi_prior(prior);
To generate a realization of the prior model defined by the prior structure,
 in the vicinity of a current model (using sequential Gibbs sampling) use:
   [m_propose, prior] = sippi_prior(prior, m_current);
The following types of a priori models can be used
   SNESIM [1D-3D] : based on a multiple point statistical model inferref from a \leftrightarrow
       training images. Relies in the SNESIM algorithm
   SISIM [1D-3D] : based on Sequential indicator SIMULATION
          [1D-3D] : based on Sequential Gaussian and Direct Sequential \leftrightarrow
   VISIM
      simulation
   FFTMA [1D-3D]: based on the FFT-MA method (Multivariate Gaussian)
            [1D] : 1D generalized gaussian model
%%% SIMPLE EXAMPLE %%%
% A simple 2D multivariate Gaissian based prior model based on the
% FFT-MA method, can be defined using
  im=1;
  prior(im).type='FFTMA';
  prior(im).name='A SIMPLE PRIOR';
  prior{im}.x=[0:1:100];
  prior{im}.y=[0:1:100];
  prior(im).m0=10;
  prior{im}.Va='1 Sph(10)';
  prior=sippi_prior_init(prior);
% A realization from this prior model can be generated using
  m=sippi_prior(prior);
% This realization can now be plotted using
   sippi_plot_prior(m, prior);
% or
   imagesc(prior{1}.x,prior{1}.y,m{1})
%%% A PRIOR MODEL WITH SEVERAL 'TYPES OF A PRIORI MODEL'
   im=1;
   prior(im).type='GAUSSIAN';
  prior(im).m0=100;
  prior(im).std=50;
  prior(im).norm=100;
  im=2:
  prior{im}.type='FFTMA';
  prior{im}.x=[0:1:100];
  prior{im}.y=[0:1:100];
  prior(im).m0=10;
  prior{im}.Cm='1 Sph(10)';
  im=3;
  prior{im}.type='SISIM';
   prior{im}.x=[0:1:100];
  prior{im}.y=[0:1:100];
  prior(im).m0=10;
  prior{im}.Cm='1 Sph(10)';
  im=4;
  prior(im).type='SNESIM';
   prior{im}.x=[0:1:100];
 prior{im}.y=[0:1:100];
```

```
sippi_plot_model(prior);
%% Sequential Gibbs sampling
  All a priori model types can be perturbed, such that a new realization
  is generated in the vicinity of a current model.
  To do this Sequential Gibbs Sampling is used.
  For more information, see <a href="matlab:web('http://dx.doi.org/10.1007/ \leftrightarrow
      s10596-011-9271-1')">Hansen, T. M., Cordua, K. S., and Mosegaard, K.,
      2012. Inverse problems with non-trivial priors - Efficient solution \,\,\leftrightarrow\,\,
      through Sequential Gibbs Sampling. Computational Geosciences</a>.
  The type of sequential Gibbs sampling can be controlled in the
  'seq_gibbs' structures, e.g. prior{1}.seq_gibbs
  prior(im).type='SNESIM';
  prior{im}.x=[0:1:100];
  prior{im}.y=[0:1:100];
  [m,prior] = sippi_prior(prior);
  prior{1}.seq_gibbs.step=1; % Large step--> independant realizations
  prior{1}.seq_gibbs.step=.1; % Smaller step--> Dependant realizations
  for i=1:30:
      [m,prior]=sippi_prior(prior,m); % One iteration of Sequential Gibbs
      sippi_plot_model(prior,m);
   end
See also: sippi_prior_init, sippi_plot_prior, sippi_prior_set_steplength.m
TMH/2012
```

6.1.25 sippi_prior_fftma

```
sippi_prior A priori models for SIPPI
 To generate a realization of the prior model defined by the prior structure use \leftrightarrow
   [m_propose, prior] = sippi_prior(prior);
 To generate a realization of the prior model defined by the prior structure,
 in the vicinity of a current model (using sequential Gibbs sampling) use:
   [m_propose, prior] = sippi_prior(prior, m_current);
 The following types of a priori models can be used
   SNESIM [1D-3D] : based on a multiple point statistical model inferref from a \leftrightarrow
       training images. Relies in the SNESIM algorithm
  SISIM [1D-3D] : based on Sequential indicator SIMULATION
  VISIM [1D-3D]: based on Sequential Gaussian and Direct Sequential \leftrightarrow
      simulation
  FFTMA [1D-3D]: based on the FFT-MA method (Multivariate Gaussian)
  GAUSSIAN [1D] : 1D generalized gaussian model
%%% SIMPLE EXAMPLE %%%
% A simple 2D multivariate Gaissian based prior model based on the
% FFT-MA method, can be defined using
   prior{id}.type='FFTMA';
  prior{id}.name='A SIMPLE PRIOR';
```

```
prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior(id).m0=10;
  prior{id}.Va='1 Sph(10)';
  prior=sippi_prior_init(prior);
% A realization from this prior model can be generated using
  m=sippi_prior(prior);
% This realization can now be plotted using
  sippi_plot_prior(m,prior);
  imagesc(prior{1}.x,prior{1}.y,m{1})
%%% A PRIOR MODEL WITH SEVERAL 'TYPES OF A PRIORI MODEL'
  id=1;
  prior{id}.type='FFTMA';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
  prior{id}.Cm='1 Sph(10)';
  id=2;
  prior{id}.type='SISIM';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
  prior{id}.Cm='1 Sph(10)';
  id=3;
  prior{id}.type='GAUSSIAN';
  prior{id}.m0=100;
  prior{id}.std=50;
  prior{id}.norm=100;
  prior=sippi_prior_init(prior);
  sippi_plot_model(prior);
%% Sequential Gibbs sampling
% For more information, see <a href="matlab:web('http://dx.doi.org/10.1007/ \leftrightarrow
   s10596-011-9271-1')">Hansen, T. M., Cordua, K. S., and Mosegaard, K., 2012. \leftrightarrow
   Sequential Gibbs Sampling. Computational Geosciences</a>.
See also: sippi_prior_init, sippi_plot_prior, sippi_prior_set_steplength.m
TMH/2012
```

6.1.26 sippi_prior_init

```
sippi_prior_init Initialize PRIOR structure for SIPPI

Call
    prior=sippi_prior_init(prior);

See also sippi_prior
```

6.1.27 sippi_prior_new

```
sippi_prior A priori models for SIPPI
```

```
To generate a realization of the prior model defined by the prior structure use \hookleftarrow
   [m_propose,prior] = sippi_prior(prior);
To generate a realization of the prior model defined by the prior structure,
in the vicinity of a current model (using sequential Gibbs sampling) use:
   [m_propose, prior] = sippi_prior(prior, m_current);
The following types of a priori models can be used
  SNESIM [1D-3D] : based on a multiple point statistical model inferref from a \leftrightarrow
       training images. Relies in the SNESIM algorithm
         [1D-3D] : based on Sequential indicator SIMULATION
         [1D-3D] : based on Sequential Gaussian and Direct Sequential \leftrightarrow
  VISIM
      simulation
  FFTMA
         [1D-3D] : based on the FFT-MA method (Multivariate Gaussian)
            [1D] : 1D generalized gaussian model
%%% SIMPLE EXAMPLE %%%
% A simple 2D multivariate Gaissian based prior model based on the
% FFT-MA method, can be defined using
  id=1;
  prior{id}.type='FFTMA';
  prior{id}.name='A SIMPLE PRIOR';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
  prior{id}.Va='1 Sph(10)';
  prior=sippi_prior_init(prior);
% A realization from this prior model can be generated using
  m=sippi_prior(prior);
% This realization can now be plotted using
  sippi_plot_prior(m,prior);
% or
  imagesc(prior{1}.x,prior{1}.y,m{1})
%%% A PRIOR MODEL WITH SEVERAL 'TYPES OF A PRIORI MODEL'
  id=1:
  prior{id}.type='FFTMA';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
  prior{id}.Cm='1 Sph(10)';
  id=2;
  prior{id}.type='SISIM';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
  prior{id}.Cm='1 Sph(10)';
  id=3;
  prior{id}.type='GAUSSIAN';
  prior{id}.m0=100;
  prior{id}.std=50;
  prior{id}.norm=100;
  prior=sippi_prior_init(prior);
  sippi_plot_model(prior);
%% Sequential Gibbs sampling
% For more information, see <a href="matlab:web('http://dx.doi.org/10.1007/ \leftrightarrow
   s10596-011-9271-1')">Hansen, T. M., Cordua, K. S., and Mosegaard, K., 2012. \leftrightarrow
```

```
Sequential Gibbs Sampling. Computational Geosciences</a>.

See also: sippi_prior_init, sippi_plot_prior, sippi_prior_set_steplength.m

TMH/2012
```

6.1.28 sippi_prior_old

```
sippi_prior A priori models for SIPPI
 To generate a realization of the prior model defined by the prior structure use \leftrightarrow
   [m_propose, prior] = sippi_prior(prior);
 To generate a realization of the prior model defined by the prior structure,
 in the vicinity of a current model (using sequential Gibbs sampling) use:
   [m_propose, prior] = sippi_prior(prior, m_current);
The following types of a priori models can be used
   SNESIM [1D-3D] : based on a multiple point statistical model inferref from a \leftrightarrow
        training images. Relies in the SNESIM algorithm
         [1D-3D] : based on Sequential indicator SIMULATION
  VISIM [1D-3D] : based on Sequential Gaussian and Direct Sequential \leftrightarrow
      simulation
   FFTMA [1D-3D]: based on the FFT-MA method (Multivariate Gaussian)
             [1D] : 1D generalized gaussian model
%%% SIMPLE EXAMPLE %%%
% A simple 2D multivariate Gaissian based prior model based on the
% FFT-MA method, can be defined using
  id=1;
  prior{id}.type='FFTMA';
  prior{id}.name='A SIMPLE PRIOR';
  prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
   prior{id}.Va='1 Sph(10)';
   prior=sippi_prior_init(prior);
% A realization from this prior model can be generated using
  m=sippi_prior(prior);
% This realization can now be plotted using
   sippi_plot_prior(m, prior);
% or
   imagesc(prior{1}.x,prior{1}.y,m{1})
%%% A PRIOR MODEL WITH SEVERAL 'TYPES OF A PRIORI MODEL'
  id=1;
   prior{id}.type='FFTMA';
   prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
  prior{id}.m0=10;
   prior{id}.Cm='1 Sph(10)';
  id=2:
  prior{id}.type='SISIM';
   prior{id}.x=[0:1:100];
  prior{id}.y=[0:1:100];
 prior(id).m0=10;
```

```
prior{id}.Cm='1 Sph(10)';
id=3;
prior{id}.type='GAUSSIAN';
prior{id}.m0=100;
prior{id}.m0=100;
prior{id}.std=50;
prior{id}.norm=100;
prior=sippi_prior_init(prior);

sippi_plot_model(prior);

% Sequential Gibbs sampling
% For more information, see <a href="matlab:web('http://dx.doi.org/10.1007/ \cup s10596-011-9271-1')">Hansen, T. M., Cordua, K. S., and Mosegaard, K., 2012. \cup Inverse problems with non-trivial priors - Efficient solution through \cup Sequential Gibbs Sampling. Computational Geosciences</a>.
See also: sippi_prior_init, sippi_plot_prior, sippi_prior_set_steplength.m

TMH/2012
```

6.1.29 sippi_prior_set_steplength

```
sippi_prior_set_steplength Set step length for Metropolis sampler in SIPPI

Call
    prior=sippi_prior_set_steplength(prior,mcmc,im);
```

6.1.30 sippi_rejection

6.1.31 sippi_set_path

```
sippi_set_path Set paths for running sippi
```

6.2 SIPPI toolbox: Traveltime tomography

6.2.1 calc Cd

```
Calc_cd Setup a covariance model to account for borehole imperfections
Call: Cd=calc_Cd(ant_pos, var_uncor, var_cor1, var_cor2, L)
This function sets up a data covariance matrix that accounts for static
(i.e. correlated) data errors.
Inputs:
\star ant_pos: A N x 4 array that contains N combinations of transmitter/source
and receiver positions. The first two columns are the x- and y-coordinates
of the transmitter/source position. The last two columns are the x- and
y-coordiantes of the receiver position.
* var_uncor: The variance of the uncorrelated data errors.
* var_corl: The variance of the correlated data errors
related to the transmitter/source positions.
\star var_cor2: The variance of the correlated data errors
related to the receiver positions.
\star L: The correlation length for the correlation between the individual
transmitter/source or receiver positions using an exponential covariance
function. For typical static errors the correlation length is set to a
small number (e.g. 10^-6).
For more details and practical examples see:
Cordua et al., 2008 in Vadose zone journal.
Cordua et al., 2009 in Journal of applied geophysics.
Knud S. Cordua (2012)
```

6.2.2 eikonal

```
eikonal Traveltime computation by solving the eikonal equation
tmap=eikonal(x,y,z,V,Sources,type);
 x,y,z: arrays defining the x, y, and z axis
 V: velocity field, with size (length(y),length(x),length(z));
 Sources [ndata,ndim] : Source positions
 type (optional): type of eikonal solver: [1]:Fast Marching(default), [2]:FD
 tmap [size(V)]: travel times computed everywhere in the velocity grid
%Example (2D):
  x=[1:1:100];
  y=1:1:100;
  z=1;
  V=ones(100,100); V(:,1:50)=2;
  Sources = [10 50;75 50];
  t=eikonal(x,y,z,V,Sources);
  subplot(1,2,1); imagesc(x,y,t(:,:,1,1)); axis image; colorbar
  subplot(1,2,2); imagesc(x,y,t(:,:,1,2)); axis image; colorbar
See also eikonal_traveltime
```

6.2.3 eikonal_raylength

```
eikonal_raylength : Computes the raylength from S to R using the eikonal ←
        equaiton

Call:
    raylength=eikonal_raylength(x,y,v,S,R,tS,doPlot)
```

6.2.4 eikonal_traveltime

```
eikonal_traveltime Computes traveltime between sources and receivers by solving \leftrightarrow
     the eikonal equation
t=eikonal_traveltime(x,y,z,V,Sources,Receivers,iuse,type);
 x,y,z: arrays defining the x, y, and z axis
 V: velocity field, with size (length(y),length(x),length(z));
 Sources [ndata,ndim] : Source positions
 Receivers [ndata,ndim] : Receiver positions
 iuse (optional): optionally only use subset of data. eg.g i_use=[1 2 4];
 type (optional): type of eikonal solver: [1]:Fast Marching(default), [2]:FD
 tmap\ [size(V)]: travel times computed everywhere in the velocity grid
%Example (2%
Example 2d traveltime compuation
Example (2D):
  x=[1:1:100];
  y=1:1:100;
  z=1;
  V=ones(100,100); V(:,1:50)=2;
  S=[50 50 1;50 50 1];
  R=[90 90 1; 90 80 1];
  t=eikonal_traveltime(x,y,z,V,S,R)
Example (3D):
  nx=50; ny=50; nz=50;
  x=1:1:nx;
  y=1:1:ny;
  z=1:1:nz;
  V=ones(ny,nx,nz);V(:,1:50,:)=2;
  S=[10 10 1;10 10 1;10 9 1];
  R=[40 40 40; 40 39 40; 40 40 40];
  t=eikonal_traveltime(x,y,z,V,S,R)
See also eikonal
```

6.2.5 kernel buursink 2d

```
CALL :
    % specify a source trace (dt, wf_trace):
    [kernel,L,L1_all,L2_all]=kernel_buursink_2d(model,x,z,S,R,dt,wf_trace);
    % Use a ricker wavelet with center frequency 'f0'
    [kernel,L,L1_all,L2_all]=kernel_buursink_2d(model,x,z,S,R,f0));
Knud Cordua, 2009,
Thomas Mejer Hansen (small edits, 2009)
```

6.2.6 kernel finite 2d

```
kernel_finite_2d 2D sensitivity kernels

Call:
   [Knorm, K, dt, options] = kernel_finite_2d(v_ref, x, y, S, R, freq, options);
```

6.2.7 kernel fresnel 2d

```
kernel_fresnel_2d Sensitivity kernel for amplitude and first arrival

Call:
    [kernel_t,kernel_a,P_omega,omega]=kernel_fresnel_2d(v,x,y,S,R,omega,P_omega);

Based on Liu, Dong, Wang, Zhu and Ma, 2009, Sensitivity kernels for seismic Fresenl volume Tomography, Geophysics, 75(5), U35-U46

See also kernel_fresnel_monochrome_2d

Run with no argument for an example.
```

6.2.8 kernel_fresnel_monochrome_2d

```
kernel_fresnel_monochrome_2d 2D monchrome kernel for amplitude and first 
arrival

Call:
    [kernel_t,kernel_a]=kernel_fresnel_monochrome_2d(v,x,y,S,R,omega);
or
    [kernel_t,kernel_a]=kernel_fresnel_monochrome_2d(v,x,y,S,R,omega,L,L1,L2);

Based on Liu, Dong, Wang, Zhu and Ma, 2009, Sensitivity kernels for seismic Fresenl volume Tomography, Geophysics, 75(5), U35-U46

See also, kernel_fresnel_2d
```

6.2.9 kernel_multiple

kernel_multiple Computes the sensitivity kernel for a wave traveling from S to R.

```
CALL :
  Knorm);
IN :
  Vel [ny,nx] : Velocity field
  x [1:nx] :
  y [1:ny]
  z [1:nz] :
  S [1,3] : Location of Source
  R [1,3] : Location of Receiver
  T : Donminant period
  alpha: controls exponential decay away ray path
  Knorm [1] : normaliztion of K [0]:none, K:[1]:vertical
  K : Sensitivity kernel
  R : Ray sensitivity kernel (High Frequency approx)
  timeS : travel computed form Source
  timeR : travel computed form Receiver
  raypath [nraydata,ndim] : the center of the raypath
The sensitivity is the length travelled in each cell.
See also : fast_fd_2d
TMH/2006
```

6.2.10 kernel_slowness_to_velocity

```
kernel_slowness_to_velocity Converts from slowness to velocity 
    parameterizations

G : kernel [1,nkernels]
V : Velocity field (

CALL:
    G_vel=kernel_slowness_to_velocity(G,V);
or
    [G_vel,v_obs]=kernel_slowness_to_velocity(G,V,t);
or
    [G_vel,v_obs,Cd_v]=kernel_slowness_to_velocity(G,V,t,Cd);
```

6.2.11 mspectrum

```
mspectrum : Amplitude and Power spectrum
Call :
    function [A,P,smoothP,kx]=mspectrum(x,dx)

1D (A)mplitude and (P)owerspectrum of x-series with spacing dx
```

6.2.12 munk_fresnel_2d

```
2D frechet kernel, First Fresnel Zone

See Jensen, Jacobsen, Christensen-Dalsgaard (2000) Solar Physics 192.

Call:
S=munk_fresnel_2d(T,dt,alpha,As,Ar,K);

T: dominant period
dt:
alpha: degree of cancellation
As: Amplitude fo the wavefield propagating from the source
Ar: Amplitude fo the wavefield propagating from the receiver
K: normalization factor
```

6.2.13 munk_fresnel_3d

```
3D frechet kernel, First Fresnel Zone

See Jensen, Jacobsen, Christensen-Dalsgaard (2000) Solar Physics 192.

Call:
```

6.2.14 tomography_kernel

```
tomography_kernel Computes the sensitivity kernel for a wave traveling from S \,\,\leftrightarrow\,
   to R.
CALL :
   [K,RAY,Gk,Gray,timeS,timeR,raypath]=tomography_kernel(Vel,x,y,z,S,R,T,alpha, \leftrightarrow
  Vel [ny,nx] : Velocity field
   x [1:nx] :
   y [1:ny] :
   z [1:nz] :
  S [1,3] : Location of Source
  R [1,3] : Location of Receiver
  T : Donminant period
  alpha: controls exponential decay away ray path
  Knorm [1] : normaliztion of K [0]:none, K:[1]:vertical
OUT :
  K : Sensitivity kernel
  R : Ray sensitivity kernel (High Frequency approx)
  timeS : travel computed form Source
   timeR : travel computed form Receiver
  raypath [nraydata,ndim] : the center of the raypath
The sensitivity is the length travelled in each cell.
```