

Package ‘lddmm’

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Type Package

Title Longitudinal Drift-Diffusion Mixed Models (LDDMM)

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Description Implementation of the drift-diffusion mixed model for category learning as described in Paulon et al. (2021).

Depends R (>= 3.5.0)

Language en-US

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Author Giorgio Paulon [aut, cre],
Abhra Sarkar [aut, ctb]

Maintainer Giorgio Paulon <giorgio.paulon@utexas.edu>

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| B_basis | <i>Spline Basis Functions</i> |
|---------|-------------------------------|

Description

Construct the J basis functions for the splines evaluated on a grid.

Usage

B_basis(xgrid, knots)

Arguments

| | |
|-------|--|
| xgrid | grid where we want to evaluate the spline functions (vector of length n) |
| knots | vector of knots for the splines (vector of length K) |

Value

n x (K+1) - matrix representing the value of each basis function evaluated on xgrid

| | |
|------|------------------------|
| data | <i>Example dataset</i> |
|------|------------------------|

Description

A toy dataset in the correct format for the LDDMM function call. This dataset has two possible response categories.

Usage

data

Format

A data frame with 24,254 rows and 6 columns

Details

- subject: vector of size n containing the participant labels
- block: vector of size n containing the training blocks (longitudinal units)
- s: vector of size n containing the stimuli
- d: vector of size n containing the decisions
- r_time: vector of size n containing the response times
- cens: vector of size n containing the censoring indicators (1 censored, 0 non censored)

extract_post_draws *Parameter posterior draws*

Description

Function to extract the posterior draws of the parameters of interest from a lddmm fit object.

Usage

```
extract_post_draws(data, fit, par = c("drift", "boundary"))
```

Arguments

| | |
|------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
| fit | fit from the lddmm function |
| par | parameter to output ('drift', or 'boundary') |

Value

Matrix with the following columns:

- subject: participant labels
- block: training blocks
- draw: iteration of the MCMC estimates
- par_s_d, ...: posterior draws for the requested parameters

| | |
|-------------------|----------------------------------|
| extract_post_mean | <i>Parameter posterior means</i> |
|-------------------|----------------------------------|

Description

Function to extract the posterior means of the parameters of interest from a lddmm fit object.

Usage

```
extract_post_mean(data, fit, par = c("drift", "boundary"))
```

Arguments

| | |
|------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
| fit | fit from the lddmm function |
| par | parameter to output ('drift', or 'boundary') |

Value

Matrix with the following columns:

- subject: participant labels
- block: training blocks
- par_s_d, ...: posterior means for the requested parameters

| | |
|--------|---------------------|
| H_ball | <i>Hamming Ball</i> |
|--------|---------------------|

Description

Computes the Hamming Ball centered at x with radius r.

Usage

```
H_ball(x, S, r)
```

Arguments

| | |
|---|----------------------------|
| x | center of the Hamming Ball |
| S | number of states |
| r | radius of the Hamming Ball |

Value

Hamming Ball

LDDMM

Drift Diffusion Model Fit

Description

Main function for the Gibbs sampler for the drift-diffusion model.

Usage

```
LDDMM(
  data,
  hypers,
  fix_boundary = FALSE,
  Niter = 5000,
  burnin = 2000,
  thin = 5
)
```

Arguments

| | |
|--------------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
| hypers | hyperparameters of the MCMC: list containing "s_sigma_mu" and "s_sigma_b", which are the smoothness parameters for drifts and boundaries, respectively) |
| fix_boundary | whether to fix the boundary parameters to a single scalar or not |
| Niter | total number of iterations |
| burnin | burnin of the chain |
| thin | thinning factor |

Value

List with the following MCMC posterior samples:

- post_mean_delta: posterior samples for the population offset parameters
- post_mean_mu: posterior samples for the population drift parameters
- post_mean_b: posterior samples for the population boundary parameters
- post_ind_delta: posterior samples for the individual offset parameters
- post_ind_mu: posterior samples for the individual drift parameters
- post_ind_b: posterior samples for the individual boundary parameters
- sigma2_mu_us: posterior samples for the random effects drift smoothness parameters
- sigma2_mu_ua: posterior samples for the random effects drift variance parameters
- sigma2_b_us: posterior samples for the random effects boundary smoothness parameters
- sigma2_b_ua: posterior samples for the random effects boundary variance parameters
- sigma2_1_mu: posterior samples for the drift smoothness parameters
- sigma2_1_b: posterior samples for the boundary smoothness parameters
- pred_ans: predicted population-level categories
- pred_time: predicted population-level response times
- pred_ans_ind: predicted individual-level categories
- pred_time_ind: predicted individual-level response times

| | |
|----------------|-----------------------------------|
| log_likelihood | <i>Log-likelihood computation</i> |
|----------------|-----------------------------------|

Description

Compute the log-likelihood for the drift-diffusion model, including the censored data contribution.

Usage

```
log_likelihood(tau, mu, b, delta, cens, D, log)
```

Arguments

| | |
|-------|--|
| tau | vector of size n containing the response times |
| mu | matrix of size (n x d1) containing the drift parameters corresponding to the n response times for each possible d1 decision |
| b | matrix of size (n x d1) containing the boundary parameters corresponding to the n response times for each possible d1 decision |
| delta | vector of size n containing the offset parameters corresponding to the n response times |

| | |
|------|---|
| cens | vector of size n containing censoring indicators (1 censored, 0 not censored) corresponding to the n response times |
| D | (n x 2) matrix whose first column has the n input stimuli, and whose second column has the n decision categories |
| log | should the results be returned on the log scale? |

plot_accuracy *Descriptive plots*

Description

Plot the accuracy of the raw data.

Usage

```
plot_accuracy(data)
```

Arguments

| | |
|------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
|------|---|

Value

Individual and population level raw accuracies

plot_post_pars *Plot posterior estimates*

Description

Function to plot the posterior mean and credible intervals of the parameters of interest from a lddmm fit object.

Usage

```
plot_post_pars(data, fit, par = c("drift", "boundary"))
```

Arguments

| | |
|------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
| fit | fit from the lddmm function |
| par | parameter to output ('drift', or 'boundary') |

Value

Posterior mean and 95% CI

| | |
|---------|--------------------------|
| plot_RT | <i>Descriptive plots</i> |
|---------|--------------------------|

Description

Plot the mean response times of the raw data.

Usage

```
plot_RT(data)
```

Arguments

| | |
|------|---|
| data | dataframe with the following columns: <ul style="list-style-type: none"> • subject: vector of size n containing the participant labels • block: vector of size n containing the training blocks (longitudinal units) • s: vector of size n containing the stimuli • d: vector of size n containing the decisions • r_time: vector of size n containing the response times • cens: vector of size n containing the censoring indicators (1 censored, 0 non censored) |
|------|---|

Value

Population level raw response times

P_smooth1

Spline Penalty Matrix

Description

Construct the covariance matrix P of the smoothness inducing prior for the spline coefficients

Usage

P_smooth1(K)

Arguments

K Number of spline knots

Value

Covariance of the smoothness inducing prior (penalizing first differences in the spline coefficients)

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