

Package ‘BIKER’

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Title Bayesian Inference/Inversion of the K600 Evasion Rate

Version 0.0.0.9000

Description Uses Bayesian inference + SWOT observations to estimate the normalized gas transfer velocity from rivers, no in situ info needed.

License file LICENSE

Encoding UTF-8

LazyData true

Biarch true

Depends R (>= 3.4.0)

Imports methods, Rcpp (>= 0.12.0), rstan (>= 2.18.1), rstantools (>= 2.0.0), dplyr, reshape2, rlang, settings

LinkingTo BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), rstan (>= 2.18.1), StanHeaders (>= 2.18.0)

SystemRequirements GNU make

RoxygenNote 7.1.2

URL <https://github.com/craigbrinkerhoff/BIKER>

BugReports <https://github.com/craigbrinkerhoff/BIKER/issues>

NeedsCompilation yes

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Archs i386, x64

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BIKER-package	<i>The 'BIKER' package.</i>
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Description

Uses Bayesian Inference + SWOT observations to estimate normalized gas transfer velocity from large rivers

References

Stan Development Team (2020). RStan: the R interface to Stan. R package version 2.19.3. <https://mc-stan.org>

biker_check_args	<i>Performs the following checks: - types: - everything else matrix - dimensions: - all matrices have same dims</i>
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Description

Performs the following checks: - types: - everything else matrix - dimensions: - all matrices have same dims

Usage

biker_check_args(datalist)

Arguments

datalist A list of biker data inputs

biker_check_nas	<i>Add missing-data inputs to data list</i>
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Description

Binary matrices indicating where data are/aren't missing are added to the data list. This is required in order to run ragged-array data structures in the stanfile.

Usage

```
biker_check_nas(datalist)
```

Arguments

datalist	a list of BIKER inputs
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Details

Previously this function omitted any times with missing data, but now that ragged arrays are accommodated in the stanfile the operations are entirely different.

biker_data	<i>Preprocess data for BIKER estimation</i>
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Description

Produces a bikerdata object that can be passed to biker_estimate function

Usage

```
biker_data(w, s, dA, priorQ, max_xs = 30L, seed = NULL)
```

Arguments

w	Matrix (or data frame) of widths: time as columns, space as rows
s	Matrix of slopes: time as columns, space as rows
dA	Matrix of area above base area: time as columns, space as rows
priorQ	Mean annual flow prior for Q
max_xs	Maximum number of cross-sections to allow in data. Used to reduce sampling time. Defaults to 30.
seed	RNG seed to use for sampling cross-sections, if nx > max_xs.

biker_estimate	<i>Estimate BIKER</i>
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Description

Fits a BIKER model of one of several variants using Hamiltonian Monte Carlo.

Usage

```
biker_estimate(
  bikerdata,
  bikerpriors,
  cores = getOption("mc.cores", default = parallel::detectCores()),
  meas_error = FALSE,
  chains = 3L,
  iter = 1000L,
  CI = 0.95,
  chainExtract = "all",
  pars = NULL,
  include = FALSE,
  suppressOutput = 1,
  ...
)
```

Arguments

bikerdata	A bikerdata object, as produced by <code>biker_data()</code>
bikerpriors	A bikerpriors object. If none is supplied, defaults are used from calling <code>biker_priors(bikerdata)</code> (with no other arguments).
cores	Number of processing cores for running chains in parallel. See <code>?rstan::sampling</code> . Defaults to <code>parallel::detectCores()</code> .
meas_error	Should we run with latent variables accounting for uncertainty in SWOT measurements. LEAVE THIS OFF, IT IS IN ACTIVE DEVELOPMENT
chains	A positive integer specifying the number of Markov chains. The default is 3.
iter	Number of iterations per chain (including warmup). Defaults to 1000.
CI	A positive integer between 0 and 1 indicating the confidence interval to return with the estimate. Default is 0.95.
chainExtract	Either an integer specifying which chain(s) to extract statistics from, or "all" (the default), in which case they are extracted from all chains.
pars	(passed to <code>rstan::sampling()</code>) A vector of character strings specifying parameters of interest to be returned in the stanfit object. If not specified, a default parameter set is returned.
include	(passed to <code>rstan::sampling()</code>) Defaults to FALSE, which excludes parameters specified in <code>pars</code> from the returned model.
suppressOutput	(passed to <code>rstan::sampling()</code>) Determines whether stan sampling updates are printed to the console. Defaults to 1. Any value ≤ 0 will suppress output.
...	Other arguments passed to <code>rstan::sampling()</code> for customizing the Monte Carlo sampler

biker_priors	<i>Establish prior hyperparameters for BIKER estimation</i>
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Description

Produces a bikerpriors object that can be passed to biker_estimate function

Usage

```
biker_priors(bikerdata, ...)
```

Arguments

bikerdata	An object of class bikerdata, as returned by biker_data
...	Optional manually set parameters. Unquoted expressions are allowed, e.g. logk_sd = cv2sigma(0.8). Additionally, any variables present in bikerdata may be referenced, e.g. lowerbound_logk = log(m

classify_func	<i>Classify river for expert framework</i>
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Description

Classify river for expert framework

Usage

```
classify_func(Wobs)
```

Arguments

Wobs	observed widths matrix
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cv2sigma	<i>Convert coefficient of variation to sigma parameter of lognormal distribution</i>
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Description

Convert coefficient of variation to sigma parameter of lognormal diistribution

Usage

```
cv2sigma(cv)
```

Arguments

cv	Coefficient of variation
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estimate_A0SD	<i>Estimate base cross-sectional area SD using bam dat</i>
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Description

Estimate base cross-sectional area SD using bam dat

Usage

```
estimate_A0SD(Wobs)
```

Arguments

Wobs	Observed W,as a space-down, time-across matrix.
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estimate_logA0	<i>Estimate base cross-sectional area using bam data</i>
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Description

Estimate base cross-sectional area using bam data

Usage

```
estimate_logA0(Wobs)
```

Arguments

Wobs	Observed W,as a space-down, time-across matrix
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estimate_logk	<i>Estimate k_hat using biker data for k600 model</i>
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Description

Estimate k_hat using biker data for k600 model

Usage

```
estimate_logk(Sobs, priorQ)
```

Arguments

Sobs	Observed S,as a space-down, time-across matrix
priorQ	mean annual flow estimate, single number

estimate_logksd	<i>Estimate k sd prior using biker data for k600 model</i>
-----------------	------------------------------------------------------------

Description

Estimate k sd prior using biker data for k600 model

Usage

```
estimate_logksd(Sobs)
```

Arguments

Sobs	Observed S, as a space-down, time-across matrix
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estimate_logn	<i>Estimate manning's n using bam data</i>
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Description

Estimate manning's n using bam data

Usage

```
estimate_logn(Wobs, Sobs)
```

Arguments

Wobs	Observed W, as a space-down, time-across matrix
Sobs	Observed s, as a space-down, time-across matrix

estimate_lognSD	<i>Estimate manning's n SD using bam dat</i>
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Description

Estimate manning's n SD using bam dat

Usage

```
estimate_lognSD(Wobs)
```

Arguments

Wobs	Observed W, as a space-down, time-across matrix.
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`estimate_lowerboundA0` *Estimate base cross-sectional area lowerbound using bam data*

Description

Estimate base cross-sectional area lowerbound using bam data

Usage

`estimate_lowerboundA0(Wobs)`

Arguments

`Wobs` Observed W , as a space-down, time-across matrix.

`estimate_upperboundA0` *Estimate base cross-sectional area upperbound using bam data*

Description

Estimate base cross-sectional area upperbound using bam data

Usage

`estimate_upperboundA0(Wobs)`

Arguments

`Wobs` Observed W , as a space-down, time-across matrix.

`prior_settings` *Options manager for BIKER default prior settings for k600 model*

Description

Options manager for BIKER default prior settings for k600 model

Usage

`prior_settings(..., __defaults = FALSE, __reset = FALSE)`

Arguments

`...` (Optional) named settings to query or set.
`__defaults` See `?settings::option_manager`
`__reset` See `?settings::option_manager`

sample_xs	<i>Take a random sample of a bikerdata object's cross-sections.</i>
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Description

Take a random sample of a bikerdata object's cross-sections.

Usage

```
sample_xs(bikerdata, n, seed = NULL)
```

Arguments

bikerdata	a bikerdata object, as returned by <code>biker_data()</code>
n	Number of cross-sections to
seed	option RNG seed, for reproducibility.

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