

Package ‘baorista’

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Title Bayesian Aoristic Analyses

Version 0.1.3

Description Provides an alternative approach to aoristic analyses for archaeological datasets by fitting Bayesian parametric growth models and non-parametric random-walk Intrinsic Conditional Autoregressive (ICAR) models on time frequency data (Crema (2024)<[doi:10.1111/arcm.12984](https://doi.org/10.1111/arcm.12984)>). It handles event typochronology based timespans defined by start/end date as well as more complex user-provided vector of probabilities.

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Imports stats,coda,graphics

Suggests knitr, rmarkdown

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createProbMat	<i>Creates a probMat class object from user data</i>
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Description

Converts either a data.frame with the start and end date of each event or matrix of probabilities values into a probMat class object.

Usage

```
createProbMat(x = NULL, pmat = NULL, timeRange = NULL, resolution = NULL)
```

Arguments

x	A data.frame containing the start and end date of the timespan of each event. Dates should be in BP, with the first column defining the start and the second column defining the end of the timespan.
pmat	A matrix of aoristic weights (probabilities), with row representing events and column representing timeblocks.
timeRange	A vector of two numerical values representing the start and end of the window of analysis in BP.
resolution	Resolution of the timeblock. Ignored if pmat is provided.

Value

An object of class probMat.

expfit	<i>Estimate Exponential Growth rate from Aoristic data</i>
--------	--

Description

Fits an exponential growth model to ProbMat class objects.

Usage

```

expfit(
  x,
  niter = 1e+05,
  nburnin = 50000,
  thin = 10,
  nchains = 4,
  rPrior = "dnorm(mean=0,sd=0.05)",
  rSampler = NULL,
  parallel = FALSE,
  seeds = 1:4
)

```

Arguments

x	A ProbMat class object
niter	Number of MCMC iterations. Default is 500,000.
nburnin	Number of iterations discarded for burn-in. Default is 250,000.
thin	Thinning interval
nchains	Number of MCMC chains
rPrior	A string defining prior for the growth parameter r. Default is 'dnorm(mean=0,sd=0.05)'.
rSampler	A list containing settings for the MCMC sampler. Default is null and employs nimble's Default sampler (RW sampler).
parallel	Logical specifying whether the chains should be run in parallel or not.
seeds	Random seed for each chain. Default is 1:4.

Details

The function fits a discrete bounded exponential growth model on the observed data using MCMC as implemented by the nimble package. The Bayesian model consists of a single growth rate parameter (r), and users can define suitable priors using character strings for the argument `rPrior` (for details on how this should be specified please consult the nimble manual). Please note that the function returns posterior of the growth rate normalised by the resolution defined in the ProbMat class object. MCMC settings such as the choice the sampler, number of iterations, chains, etc can also be specified.

Value

A `fittedExp` class object containing the original ProbMat class object, posterior of the growth rate, along with its Gelman Rubin statistic and effective sample sizes.

 icarfit

Fits a random walk ICAR model to Aoristic data

Description

Estimates parameters of a multinomial probability distribution that describes the shape of the of the time-frequency distribution of an observed sets of events with chronological uncertainty. The function is wrapper for fitting a 1D random walk ICAR model via nimble.

Usage

```
icarfit(
  x,
  niter = 1e+05,
  nburnin = 50000,
  thin = 10,
  nchains = 4,
  sigmaPrior = "dexp(1)",
  sigmaSampler = NULL,
  parallel = FALSE,
  seeds = 1:4
)
```

Arguments

x	A ProbMat class object
niter	Number of MCMC iterations. Default is 500,000.
nburnin	Number of iterations discarded for burn-in. Default is 250,000.
thin	Thinning interval
nchains	Number of MCMC chains
sigmaPrior	A string defining prior for the sigma parameter. Default is 'dexp(1)'.
sigmaSampler	A list containing settings for the MCMC sampler. Default is null and employs nimble's Default sampler (RW sampler).
parallel	Logical specifying whether the chains should be run in parallel or not.
seeds	Random seed for each chain. Default is 1:4.

Details

The function estimates a vector temporally autocorrelated probability values on the observed data using MCMC as implemented by the nimble package. The model is effectively non-parametric, and at its core it is an implementation of a 1D random ICAR model. Users can specify the prior for the variance parameter through the argument `sigmaPrior`. Different settings for this parameter can greatly influence the estimates of the probability vectors. For example using `sigmaPrior=dexp(100)` instead of the default `sigmaPrior=dexp(1)` would lead to 'flatter' time-series with higher temporal autocorrelation. Please consult the nimble package manual for the syntax required in specifying the

prior. MCMC settings such as the choice the sampler, number of iterations, chains, etc can also be specified. Please note that the function is computationally intensive and might require a larger number of iterations to reach satisfactory MCMC convergence.

Value

A fittedICAR class object containing the original ProbMat class object, posteriors of the probabilities for each time-block and the variance parameter along with their MCMC diagnostics (Gelman Rubin statistic and effective sample size).

logisticfit	<i>Fits a Logistic growth model on Aoristic data</i>
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Description

Fits an exponential growth model to ProbMat class objects.

Usage

```
logisticfit(
  x,
  niter = 1e+05,
  nburnin = 50000,
  thin = 10,
  nchains = 4,
  rPrior = "dexp(1/0.001)",
  mPrior = "dunif(1,z)",
  rSampler = NULL,
  mSampler = NULL,
  parallel = FALSE,
  seeds = 1:4
)
```

Arguments

x	A ProbMat class object
niter	Number of MCMC iterations. Default is 100,000.
nburnin	Number of iterations discarded for burn-in. Default is 50,000.
thin	Thinning interval
nchains	Number of MCMC chains
rPrior	A string defining prior for the growth parameter r. Default is 'dexp(1/0.01)'.
mPrior	A string defining prior for the point of maximum growth rate m. Default is 'dunif(1,z)', where 'z' is the number of time-blocks.
rSampler	A list containing settings for the MCMC sampler for the parameter 'r'. Default is null and employs nimble's Default sampler (RW sampler).

mSampler	A list containing settings for the MCMC sampler for the parameter 'm'. Default is null and employs nimble's Default sampler (RW sampler).
parallel	Logical specifying whether the chains should be run in parallel or not.
seeds	Random seed for each chain. Default is 1:4.

Details

The function fits a discrete bounded logistic growth model on the observed data using MCMC as implemented by the nimble package. The Bayesian model consists of two parameters, a growth rate (r) and a midpoint (m) defining the inflection point of the growth curve. Priors of the two parameters can be defined by the arguments `rPrior` and `mPrior`. In the latter case the object `z` is a placeholder for the number of blocks (e.g. the default `'dunif(1,z)'` is a uniform across all blocks). Priors are defined by character strings following the syntax used by nimble. Please note that the function returns posterior of the growth rate normalised by the resolution defined in the `ProbMat` class object. MCMC settings such as the choice the sampler, number of iterations, chains, etc can also be specified.

Value

A `fittedLogistic` class object containing the original `ProbMat` class object, posteriors of the growth rate and midpoint and their MCMC diagnostics (i.e. Gelman Rubin statistic and effective sample sizes).

plot.fittedExp	<i>Plot exponential model fitted to aoristic data</i>
----------------	---

Description

Plot posterior estimates of `fittedExp` class objects.

Usage

```
## S3 method for class 'fittedExp'
plot(
  x,
  hpd = c(0.5, 0.9),
  minortick = NULL,
  ylim = NULL,
  xlab = NULL,
  ylab = "Probability Mass",
  calendar = "BP",
  col = "black",
  lwd = 1,
  lty = 2,
  col1 = "steelblue",
  col2 = "lightblue",
  pch = 20,
```

```

    plot.legend = TRUE,
    legend.arg = NULL,
    ...
)

```

Arguments

x	An <code>fittedExp</code> class object
hpd	A vector with two values defining the highest posterior density interval to display. Default is 0.5 and 0.9.
minortick	Interval for minor ticks in the x-axis label. Default is estimated based on timescale.
ylim	Limits of the y-axis. Default estimated from posterior ranges.
xlab	Label for the x-axis. Default based on calendar.
ylab	Label for the y-axis. Default is "Probability Mass".
calendar	Either 'BP' or 'BCAD'. Indicate whether the x-axis should be displayed in BP or BC/AD. Default is 'BP'.
col	Color of posterior mean. Default is black.
lwd	Line width posterior mean. Default is 1.
lty	Line type posterior mean. Default is 2.
col1	Fill color for the first (inner) HPD interval. Default is 'steelblue'.
col2	Fill color for the second (outer) HPD interval. Default is 'lightblue'.
pch	Point symbol used to display mean posteriors. Default is 20.
plot.legend	Logical indicating whether to display a legend or not (default is TRUE).
legend.arg	List containing arguments to be directed to the <code>legend()</code> function.
...	Additional arguments affecting the plot.

Value

No return value (plot function)

plot.fittedICAR	<i>Plot 1D ICAR model fitted to aoristic data</i>
-----------------	---

Description

Plot posterior estimates of `fittedICAR` class objects.

Usage

```
## S3 method for class 'fittedICAR'
plot(
  x,
  hpd = c(0.5, 0.9),
  minortick = NULL,
  ylim = NULL,
  xlab = NULL,
  ylab = "Probability Mass",
  calendar = "BP",
  col1 = "steelblue",
  col2 = "lightblue",
  pch = 20,
  plot.legend = TRUE,
  legend.arg = NULL,
  ...
)
```

Arguments

<code>x</code>	An <code>fittedICAR</code> class object
<code>hpd</code>	A vector with two values defining the highest posterior density interval to display. Default is 0.5 and 0.9.
<code>minortick</code>	Interval for minor ticks in the x-axis label. Default is estimated based on timescale.
<code>ylim</code>	Limits of the y-axis. Default estimated from posterior ranges.
<code>xlab</code>	Label for the x-axis. Default based on calendar.
<code>ylab</code>	Label for the y-axis. Default is "Probability Mass".
<code>calendar</code>	Either 'BP' or 'BCAD'. Indicate whether the x-axis should be displayed in BP or BC/AD. Default is 'BP'.
<code>col1</code>	Fill color for the first (inner) HPD interval. Default is 'steelblue'.
<code>col2</code>	Fill color for the second (outer) HPD interval. Default is 'lightblue'.
<code>pch</code>	Point symbol used to display mean posteriors. Default is 20.
<code>plot.legend</code>	Logical indicating whether to display a legend or not (default is TRUE).
<code>legend.arg</code>	List containing arguments to be directed to the <code>legend()</code> function.
<code>...</code>	Additional arguments affecting the plot.

Value

No return value (plot function)

plot.fittedLogistic *Plot logistic model fitted to aoristic data*

Description

Plot posterior estimates of fittedLogistic class objects.

Usage

```
## S3 method for class 'fittedLogistic'
plot(
  x,
  hpd = c(0.5, 0.9),
  minortick = NULL,
  ylim = NULL,
  xlab = NULL,
  ylab = "Probability Mass",
  calendar = "BP",
  col = "black",
  lwd = 1,
  lty = 2,
  col1 = "steelblue",
  col2 = "lightblue",
  pch = 20,
  plot.legend = TRUE,
  legend.arg = NULL,
  ...
)
```

Arguments

x	An fittedExp class object
hpd	A vector with two values defining the highest posterior density interval to display. Default is 0.5 and 0.9.
minortick	Interval for minor ticks in the x-axis label. Default is estimated based on timescale.
ylim	Limits of the y-axis. Default estimated from posterior ranges.
xlab	Label for the x-axis. Default based on calendar.
ylab	Label for the y-axis. Default is "Probability Mass".
calendar	Either 'BP' or 'BCAD'. Indicate whether the x-axis should be displayed in BP or BC/AD. Default is 'BP'.
col	Color of posterior mean. Default is black.
lwd	Line width posterior mean. Default is 1.
lty	Line type posterior mean. Default is 2.
col1	Fill color for the first (inner) HPD interval. Default is 'steelblue'.

col2	Fill color for the second (outer) HPD interval. Default is 'lightblue'.
pch	Point symbol used to display mean posteriors. Default is 20.
plot.legend	Logical indicating whether to display a legend or not (default is TRUE).
legend.arg	List containing arguments to be directed to the legend() function.
...	Additional arguments affecting the plot.

Value

No return value (plot function)

plot.probMat	<i>Plot Aoristic Sums</i>
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Description

Plot summed probabilities of aoristic weights.

Usage

```
## S3 method for class 'probMat'
plot(
  x,
  xlab = NULL,
  ylab = NULL,
  type = "asum",
  calendar = "BP",
  lwd = 1,
  col = 1,
  minortick = NULL,
  add = FALSE,
  ...
)
```

Arguments

x	probMat class object generated using the generateProbMat().
xlab	Label for the x-axis. Default based on calendar.
ylab	Label for the y-axis. Default is 'Summed Probability' (if type='asum') or 'Probability Mass' (when type='dens').
type	Either 'asum' for Aoristic Sum, 'dens' for probability mass. Default is 'asum'.
calendar	Either 'BP' or 'BCAD'. Indicate whether the x-axis should be displayed in BP or BC/AD. Default is 'BP'.
lwd	Line width. Default is 1.
col	Line col. Default is 'black'
minortick	Interval for minor ticks in the x-axis label. Default is estimated based on timescale
add	if set to TRUE adds the line and point graph on existing plot.
...	Additional arguments affecting the plot.

Value

No return value (plot function)

sampledf	<i>Sample aoristic data (data.frame)</i>
----------	--

Description

Sample datasets to illustrate data formats required for createProbMat().

Usage

```
sampledf
```

Format

A data.frame class object with two columns defining the start and the end of each even (sample.df)

Examples

```
data(sampledf)
x <- createProbMat(x=sampledf,timeRange = c(6500,4001),resolution= 100)
```

samplemat	<i>Sample aoristic data (matrix)</i>
-----------	--------------------------------------

Description

Sample datasets to illustrate data formats required for createProbMat().

Usage

```
samplemat
```

Format

A matrix class object storing the probability of each event (row) in each time-block (column)

Examples

```
data(samplemat)
x <- createProbMat(pmat=samplemat,timeRange = c(5000,3001),resolution=100)
plot(x)
```

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