

Package ‘GBASS’

May 7, 2026

Title Generalized Bayesian Adaptive Smoothing Splines

Version 2.0.1

Description Bayesian nonlinear regression under a range of likelihood models using generalized Bayesian adaptive smoothing splines. Robust regression with Student's t likelihoods, quantile regression, and related latent-scale models are included as special cases.

License BSD_3_clause + file LICENSE

Encoding UTF-8

RoxygenNote 7.3.2

Depends R (>= 3.5.0)

Imports Matrix, GIGrv, BASS

Suggests knitr, rmarkdown, lhs, testthat (>= 2.1.0)

NeedsCompilation no

Author Kellin Rumsey [aut, cre]

Maintainer Kellin Rumsey <knrumsey@lanl.gov>

Repository CRAN

Date/Publication 2026-04-24 20:10:16 UTC

Contents

GBASS-package	2
build_prior	2
gbass	3
gbass2bass	6
gsobol	6
hbass	7
nwbass	8
nw_est_mom	10
nw_gamma_prior	12
nw_triangle	13
plot.gbass	14

predict.gbass	14
print.gbass	15
qbass	16
rgig2	17
rmhn	18
tbass	18

Index	20
--------------	-----------

GBASS-package	<i>GBASS: Generalized Bayesian Adaptive Smoothing Splines</i>
---------------	---

Description

Bayesian nonlinear regression under a range of likelihood models using generalized Bayesian adaptive smoothing splines. Robust regression with Student's t likelihoods, quantile regression, and related latent-scale models are included as special cases.

Author(s)

Maintainer: Kellin Rumsey <knrumsey@lanl.gov>

build_prior	<i>Construct a prior specification for GBASS</i>
-------------	--

Description

Construct a prior specification for GBASS

Usage

```
build_prior(
  type = c("GIG", "GBP"),
  p,
  a,
  b,
  lower_bound = NULL,
  prop_sigma = NULL,
  adapt = NULL,
  adapt_delay = NULL,
  adapt_thin = NULL
)

build_GIG(
  p,
  a,
```

```

    b,
    lower_bound = NULL,
    prop_sigma = NULL,
    adapt = NULL,
    adapt_delay = NULL,
    adapt_thin = NULL
  )

  build_GBP(
    p,
    a,
    b,
    lower_bound = NULL,
    prop_sigma = NULL,
    adapt = NULL,
    adapt_delay = NULL,
    adapt_thin = NULL
  )

```

Arguments

type	"GIG" or "GBP"
p, a, b	prior hyperparameters
lower_bound	optional lower bound
prop_sigma	optional proposal sd (log scale)
adapt	logical; use adaptive MH?
adapt_delay	delay before adapting
adapt_thin	thinning for adaptation updates

Value

A named list containing the prior specification.

 gbass

Generalized Bayesian MARS

Description

Fits a generalized Bayesian MARS (GBASS) model for nonlinear regression under flexible latent-scale likelihoods.

Usage

```

gbass(
  X,
  y,
  w_prior = list(type = "GIG", p = 0, a = 0, b = 0),
  v_prior = list(type = "GIG", p = -15, a = 0, b = 30),
  maxInt = 3,
  maxBasis = 1000,
  npart = NULL,
  nmcmc = 10000,
  nburn = 9000,
  thin = 1,
  moveProbs = rep(1/3, 3),
  a_tau = 1/2,
  b_tau = NULL,
  a_lambda = 1,
  b_lambda = 1,
  m_beta = 0,
  s_beta = 0,
  scale = 1,
  Iw0 = rep(1, maxInt),
  Zw0 = rep(1, ncol(X)),
  verbose = TRUE
)

```

Arguments

<code>X</code>	An $N \times p$ numeric matrix of predictors. A numeric vector is treated as a single-column matrix.
<code>y</code>	A numeric response vector of length N .
<code>w_prior</code>	A named list specifying the prior for the global variance component. See Details.
<code>v_prior</code>	A named list specifying the shared prior for the local variance components. See Details.
<code>maxInt</code>	Integer giving the maximum interaction degree in proposed basis functions.
<code>maxBasis</code>	Maximum number of basis functions.
<code>npart</code>	Minimum number of nonzero points required for a proposed basis function. Defaults to $\min(20, 0.1 * N)$.
<code>nmcmc</code>	Total number of MCMC iterations.
<code>nburn</code>	Number of initial iterations discarded as burn-in.
<code>thin</code>	Thinning interval for retained draws.
<code>moveProbs</code>	A length-3 vector giving probabilities for birth, death, and mutation moves.
<code>a_tau</code>	Prior hyperparameter for tau.
<code>b_tau</code>	Prior hyperparameter for tau. Defaults to $N / 2$.
<code>a_lambda</code>	Prior hyperparameter for lambda.

b_lambda	Prior hyperparameter for lambda.
m_beta	Prior mean for beta.
s_beta	Prior standard deviation for beta.
scale	Fixed variance-scale parameter. Defaults to 1.
Iw0	Vector of positive nominal weights for interaction order in proposed basis functions. Must have length maxInt.
Zw0	Vector of positive nominal weights for variable selection in proposed basis functions. Must have length ncol(X).
verbose	Logical; should progress be printed?

Details

The priors for w and v_i must belong to the generalized inverse Gaussian ("GIG") or generalized beta prime ("GBP") families.

Each prior list may contain:

1. type: either "GIG" or "GBP".
2. p, a, b: prior hyperparameters.
3. lower_bound: optional lower bound for the parameter support. For backward compatibility, lb is also accepted.
4. prop_sigma: proposal standard deviation on the log scale for Metropolis updates, when applicable.
5. adapt, adapt_delay, adapt_thin: optional controls for adaptive Metropolis updates when applicable.

For w , prop_sigma and the adaptive Metropolis fields are mainly relevant when $w_prior\$type = "GBP"$. In the "GIG" case with nonzero beta, w is sampled using the modified half-normal formulation rather than a random-walk Metropolis step.

Retained draws are taken at iterations $nburn + 1$, $nburn + 1 + thin$, \dots . Thus $nburn$ is interpreted as the number of initial iterations discarded as burn-in.

Value

An object of class "gbass" containing posterior draws and fitted model information.

Note

Current implementation notes:

1. Basis function parameters are stored as lists.
2. Knot locations use continuous uniform proposals.
3. Basis coefficients use a ridge-type prior.

Examples

```
ff1 <- function(x) 10.391*((x[1]-0.4)*(x[2]-0.6) + 0.36)

n <- 100
p <- 4
X <- matrix(runif(n * p), nrow = n)
y <- apply(X, 1, ff1)
mod <- gbass(X, y, nmcmc = 1000, nburn = 900, thin = 2)
```

gbass2bass
Convert GBASS object to BASS-like object

Description

Converts an object of class `gbass` to an object with class `bass`, so that downstream BASS utilities such as Sobol decomposition can be used.

Usage

```
gbass2bass(gm)

gm2bm(gm)
```

Arguments

`gm` An object of class `gbass`.

Value

An object with class `c("bass", "gbass")`.

gsobol
Sobol decomposition for GBASS models

Description

Computes Sobol sensitivity indices for a fitted `gbass` model by first converting it to a compatible BASS object with `gbass2bass()`, and then calling `BASS::sobol()`.

Usage

```
gsobol(object, ...)
```

Arguments

`object` A fitted object of class `"gbass"`.
`...` Additional arguments passed to `BASS::sobol()`.

Details

This is a thin wrapper around `BASS::sobol()` for GBASS models. Users who want direct access to the converted BASS object can call `gbass2bass()` explicitly and then use `BASS::sobol()` themselves.

Value

An object of class "bassSob" returned by `BASS::sobol()`.

See Also

[gbass2bass](#), [sobol](#)

Examples

```
ff1 <- function(x) 10.391*((x[1]-0.4)*(x[2]-0.6) + 0.36)

n <- 100
p <- 4
X <- matrix(runif(n * p), nrow = n)
y <- apply(X, 1, ff1)

mod <- gbass(X, y, nmcmc = 1000, nburn = 900)

# Direct wrapper
sob <- gsobol(mod)

# Equivalent manual conversion
bm <- gbass2bass(mod)
sob2 <- BASS::sobol(bm)
```

hbass

HBASS - Bayesian MARS with Horseshoe or Strawderman-Berger likelihood

Description

Wrapper around `gbass()` using generalized beta prime latent-scale priors.

Usage

```
hbass(
  X,
  y,
  w_prior = list(type = "GBP", p = 1, a = 1/2, b = 1/2),
  likelihood = "h",
  ...
)
```

Arguments

<code>X</code>	An $N \times p$ numeric matrix of predictor variables.
<code>y</code>	A numeric response vector of length N .
<code>w_prior</code>	A named list specifying the prior for the global variance component. See Details.
<code>likelihood</code>	Character string specifying the likelihood family. Use "h" for Horseshoe or "sb" for Strawderman-Berger.
<code>...</code>	Additional arguments passed to <code>gbass()</code> .

Details

The Horseshoe and Strawderman-Berger likelihoods can be expressed using generalized beta prime latent-scale distributions. This function provides a convenient wrapper for these likelihoods. For additional flexibility, use `gbass()` directly.

Value

An object of class `c("nwbass", "gbass")` containing posterior draws and fitted model information.

nwbass

Generalized Bayesian MARS with a Normal-Wald likelihood

Description

Fits a generalized BMARS model under a Normal-Wald likelihood. This provides flexible nonlinear regression with a unimodal, potentially skewed error model.

Usage

```
nwbass(
  X,
  y,
  w_prior = list(type = "GIG", p = -0.1, a = 0, b = 0.1),
  maxInt = 3,
  maxBasis = 1000,
  npart = NULL,
  nmcmc = 10000,
  nburn = 9000,
  thin = 1,
  moveProbs = rep(1/3, 3),
  a_tau = 1/2,
  b_tau = NULL,
  a_lambda = 1,
  b_lambda = 1,
  m_beta = 0,
  s_beta = 0,
  lag_beta = 1001,
```

```

    m_gamma = 1,
    s_gamma = 0,
    scale = 1,
    Iw0 = rep(1, maxInt),
    Zw0 = rep(1, ncol(X)),
    verbose = TRUE
)

nwbass2(...)

```

Arguments

<code>X</code>	An $N \times p$ numeric matrix of predictor variables. A numeric vector is treated as a single-column matrix.
<code>y</code>	A numeric response vector of length N .
<code>w_prior</code>	A named list specifying the prior for the global variance component. See Details.
<code>maxInt</code>	Integer giving the maximum degree of interaction in spline basis functions. Defaults to 3.
<code>maxBasis</code>	Maximum number of basis functions.
<code>npart</code>	Minimum number of nonzero points required for a proposed basis function. Defaults to $\min(20, 0.1 * N)$.
<code>nmcmc</code>	Total number of MCMC iterations.
<code>nburn</code>	Number of initial MCMC iterations discarded as burn-in.
<code>thin</code>	Thinning interval for retained draws.
<code>moveProbs</code>	A length-3 vector giving probabilities for birth, death, and mutation moves.
<code>a_tau</code>	Prior hyperparameter for tau.
<code>b_tau</code>	Prior hyperparameter for tau. Defaults to $N / 2$.
<code>a_lambda</code>	Prior hyperparameter for lambda.
<code>b_lambda</code>	Prior hyperparameter for lambda.
<code>m_beta</code>	Prior mean for beta.
<code>s_beta</code>	Prior standard deviation for beta.
<code>lag_beta</code>	Number of initial iterations for which beta is fixed at <code>m_beta</code> . This is often used to stabilize early sampling.
<code>m_gamma</code>	Prior mean for gamma.
<code>s_gamma</code>	Prior standard deviation for gamma.
<code>scale</code>	Fixed variance-scale parameter. Defaults to 1.
<code>Iw0</code>	Vector of positive nominal weights for interaction order in proposed basis functions. Must have length <code>maxInt</code> .
<code>Zw0</code>	Vector of positive nominal weights for variable selection in proposed basis functions. Must have length <code>ncol(X)</code> .
<code>verbose</code>	Logical; should progress be printed?
<code>...</code>	(for backwards compatability)

Details

The latent local-scale prior is fixed internally to the Normal-Wald form $v_i \sim \text{GIG}(-1/2, \gamma^2, 1)$. Unlike `gbass()`, `nwbass()` does not expose a user-specified `v_prior`.

The `w_prior` list should contain:

1. `type`: either "GIG" or "GBP".
2. `p`, `a`, `b`: hyperparameters for the prior.
3. `lower_bound`: optional lower bound for `w`. For backward compatibility, `lb` is also accepted.
4. `prop_sigma`: proposal standard deviation on the log scale for Metropolis updates of `w`. This is only used when `w` is updated by Metropolis-Hastings, such as the "GBP" case.
5. `adapt`, `adapt_delay`, `adapt_thin`: optional controls for adaptive Metropolis updates of `w` when applicable.

Retained draws are taken at iterations `nburn + 1`, `nburn + 1 + thin`, `...`. Thus `nburn` is interpreted as the number of initial iterations discarded as burn-in.

Value

An object of class `c("nwbass", "gbass")` containing posterior draws and fitted model information.

Examples

```
n <- 200
X <- lhs::maximinLHS(n, 2)
f <- 20 * apply(X, 1, function(x) sin(pi * x[1]) + x[2]^2)
eps <- rgamma(n, 3, 1.5) - 2
y <- f + eps

mod <- nwbass(X, y,
              m_beta=0, s_beta=10,
              m_gamma = nw_gamma_prior()[1], s_gamma = nw_gamma_prior()[2])
```

nw_est_mom

Method-of-moments estimation for the Normal-Wald distribution

Description

Estimates Normal-Wald parameters from sample moments or from a supplied vector of moments.

Usage

```
nw_est_mom(
  data = NULL,
  stats = NULL,
  mu = NA,
  delta = NA,
```

```

    beta = NA,
    alpha = NA,
    triangle = FALSE,
    ...
  )

```

Arguments

<code>data</code>	Optional data vector. If supplied, empirical moments are computed from data and stats is ignored.
<code>stats</code>	Optional numeric vector of length 4 containing mean, variance, skewness, and kurtosis. Entries may be NA when the corresponding parameter is fixed.
<code>mu</code>	Location parameter, if fixed.
<code>delta</code>	Scale parameter, if fixed.
<code>beta</code>	Skewness parameter, if fixed.
<code>alpha</code>	Tail parameter, if fixed.
<code>triangle</code>	Logical; if TRUE, return only the asymmetry and steepness summary parameters.
<code>...</code>	Additional arguments passed to <code>optim</code> .

Details

This function computes method-of-moments estimates for the Normal-Wald distribution. If data is supplied, sample mean, variance, skewness, and kurtosis are computed automatically. Otherwise, the user may provide these moments directly through `stats`.

If some parameters are fixed, only the remaining parameters are estimated. When `triangle = TRUE`, the returned values are the transformed summary quantities $\text{steepness} = (1 + |\gamma|)^{-1/2}$ and a corresponding asymmetry measure.

Value

If `triangle = FALSE`, a named numeric vector with entries `mu`, `delta`, `beta`, and `alpha`. If `triangle = TRUE`, a named numeric vector with entries `asymmetry` and `steepness`.

Examples

```

n <- 500
y <- rgamma(n, 3, 1.5) + rlnorm(n, 1, 0.5)
z <- (y - mean(y)) / sd(y)
skew <- mean(z^3)
kurt <- mean(z^4)

nw_est_mom(stats = c(NA, NA, skew, kurt), mu = 0, delta = 1, triangle = TRUE)
nw_est_mom(stats = c(NA, var(y), skew, kurt), mu = 0, triangle = TRUE)

```

nw_gamma_prior

Calibrate a prior for the Normal-Wald gamma parameter

Description

Selects prior hyperparameters for gamma in the Normal-Wald model using probability statements about the steepness parameter $\xi = (1 + \gamma)^{-1/2}$.

Usage

```
nw_gamma_prior(
  q1 = 0.1,
  q2 = 0.9,
  p1 = 0.5,
  p2 = 0.05,
  par0 = NULL,
  lambda = 0
)
```

Arguments

q1	Lower reference value for the steepness parameter. Default is 0.1.
q2	Upper reference value for the steepness parameter. Default is 0.9.
p1	Target probability associated with q1. Default is 0.5.
p2	Target probability associated with q2. Default is 0.05.
par0	Optional starting values for the numerical optimization.
lambda	Optional ridge penalty used in the optimization. Default is 0.

Details

This function chooses hyperparameters for a prior on gamma by solving a simple calibration problem based on the steepness parameter $(1 + \gamma)^{-1/2}$. The optimization is carried out with `optim` using the Nelder-Mead method.

Value

A named numeric vector with entries `m_gamma` and `s_gamma`.

Examples

```
nw_gamma_prior()
nw_gamma_prior(q1 = 0.2, q2 = 0.8, p1 = 0.4, p2 = 0.1)
```

nw_triangle	<i>Plot Normal-Wald shape parameters on the asymmetry-steepness triangle</i>
-------------	--

Description

Visualizes posterior draws of the Normal-Wald shape parameters in terms of asymmetry and steepness. The plotted coordinates are steepness = $(1 + \gamma)^{-1/2}$ and asymmetry = $\beta(\beta^2 + \gamma^2)^{-1/2} \times$ steepness.

Usage

```
nw_triangle(obj, add = FALSE, details = FALSE, ...)
```

Arguments

obj	A fitted object containing components beta and gamma, typically an object returned by <code>nwbass()</code> .
add	Logical; if FALSE, initialize a new triangle plot. If TRUE, add points to the current plot.
details	Logical; if TRUE and add = FALSE, add a few reference distributions to the plot.
...	Additional graphical arguments passed to points .

Details

This plot provides a simple geometric summary of the shape of the Normal-Wald likelihood. The upper boundary corresponds to symmetric models, while points away from zero on the horizontal axis indicate asymmetry.

Value

Invisibly returns a data frame with columns asymmetry and steepness.

Examples

```
# mod is a fitted nwbass model

# Simple example
n <- 200
X <- lhs::maximinLHS(n, 2)
f <- 20 * apply(X, 1, function(x) sin(pi * x[1]) + x[2]^2)
eps <- rgamma(n, 3, 1.5) - 2
y <- f + eps

mod <- nwbass(X, y,
             m_beta=0, s_beta=10,
             m_gamma = nw_gamma_prior()[1], s_gamma = nw_gamma_prior()[2])
nw_triangle(mod)
```

plot.gbass	<i>Plot method for gbass objects</i>
------------	--------------------------------------

Description

Plot method for gbass objects

Usage

```
## S3 method for class 'gbass'  
plot(x, ...)
```

Arguments

x	A fitted object of class "gbass".
...	Additional graphical arguments.

Value

No return value, called for its side effects.

predict.gbass	<i>Predict method for GBASS objects</i>
---------------	---

Description

Returns posterior draws of either: (i) the linear predictor / mean surface, or (ii) the full posterior predictive distribution.

Usage

```
## S3 method for class 'gbass'  
predict(  
  object,  
  newdata = NULL,  
  mcmc.use = NULL,  
  predictive = TRUE,  
  bias_correct = FALSE,  
  samples = 1,  
  ...  
)
```

Arguments

object	an object of class "gbass" (including subclasses like "tbass", "qbass", "nwbass")
newdata	a matrix of predictor variables. Defaults to training inputs.
mcmc.use	optional vector indicating which posterior draws to use.
predictive	logical. If TRUE, return posterior predictive draws. If FALSE, return draws of the linear predictor.
bias_correct	logical. Ignored unless predictive = FALSE. If TRUE, return the posterior mean response rather than just the linear predictor.
samples	Integer giving the number of predictive samples to generate per retained MCMC draw when predictive = TRUE. Ignored when predictive = FALSE. Default is 1.
...	Additional graphical arguments (not used)

Details

If predictive = FALSE and bias_correct = FALSE, this returns draws of $B(x)a$.

If predictive = FALSE and bias_correct = TRUE, this returns draws of $B(x)a + E(\text{error} \mid \text{posterior draw})$, i.e. the mean response under the fitted GBASS error model.

If predictive = TRUE, this returns posterior predictive draws by simulating a fresh latent local variance v_{new} and Gaussian draw for each posterior sample.

For qbass objects, bias_correct = TRUE is usually not what you want, because qbass is typically being used for quantile regression rather than mean regression.

Currently, posterior predictive draws are implemented for GIG-based models. If the fitted object uses a GBP prior for v , predictive = TRUE will stop.

Value

a matrix with rows corresponding to posterior draws and columns corresponding to rows of newdata.

print.gbass	<i>Print a gbass object</i>
-------------	-----------------------------

Description

Print a gbass object

Usage

```
## S3 method for class 'gbass'
print(x, ...)
```

Arguments

x	An object of class "gbass".
...	Unused.

Value

The input object, invisibly.

 qbass

QBASS - Bayesian MARS with an asymmetric Laplace likelihood

Description

Wrapper around `gbass()` for quantile regression.

Usage

```
qbass(
  X,
  y,
  q = 0.5,
  w_prior = list(type = "GIG", p = -0.1, a = 0, b = 0.1),
  ...
)
```

Arguments

<code>X</code>	An $N \times p$ numeric matrix of predictor variables.
<code>y</code>	A numeric response vector of length N .
<code>q</code>	Quantile of interest. Default is 0.5 for median regression.
<code>w_prior</code>	Prior for the global variance factor.
<code>...</code>	Additional arguments passed to <code>gbass()</code> .

Details

Performs quantile regression for quantile q using the asymmetric Laplace representation. For many quantiles, fitting separate models in parallel may be convenient.

Value

An object of class `c("qbass", "gbass")` containing posterior draws and fitted model information.

`rgig2`*Generalized Inverse Gaussian Generator*

Description

This function generates samples from the GIG(p , a , b) distribution with density function $f(x) = cx^{(p-1)}\exp(-1/2*(a*x + b/x))$

Usage

```
rgig2(p, a, b)
```

Arguments

<code>p</code>	a real valued parameter
<code>a</code>	a non-negative parameter. If $a=0$, then p must be negative.
<code>b</code>	a non-negative parameter. If $b=0$, then p must be positive.

Details

A uniformly bounded rejection sample based on Hörmann et. al. (2014). Special cases include Gamma ($b=0$, $p>0$), Inverse Gamma ($a=0$, $p<0$) and Inverse Gaussian ($p=-1/2$).

Value

A numeric vector of random draws from the generalized inverse Gaussian distribution.

References

Hörmann, Wolfgang, and Josef Leydold. "Generating generalized inverse Gaussian random variates." *Statistics and Computing* 24.4 (2014): 547-557.

Examples

```
x <- rep(NA, 1000)
for(i in 1:1000) x[i] <- rgig2(2, 3, 0.5)
hist(x)
```

 rmhn

Random generation from the modified half-normal distribution

Description

Generates random samples from the modified half-normal distribution with density proportional to $x^{\alpha-1} \exp(-\beta x^2 + \gamma x)$ for $x > 0$.

Usage

```
rmhn(n = 1, alpha, beta, gamma)
```

Arguments

n	Number of random draws.
alpha	Positive shape parameter.
beta	Positive rate parameter multiplying x^2 .
gamma	Real-valued linear parameter multiplying x .

Details

This distribution arises in latent-scale updates for the generalized Bayesian adaptive smoothing spline model. The function is intended mainly for internal model fitting, but it may also be useful on its own.

Value

A numeric vector of length n containing random draws from the modified half-normal distribution.

Examples

```
x <- rmhn(1000, alpha = 3, beta = 1, gamma = 0.5)
hist(x, breaks = 30, freq = FALSE)
```

 tbass

TBASS - Bayesian MARS with a Student's t likelihood

Description

Wrapper around `gbass()` for a Student's t error model.

Usage

```
tbass(X, y, df = 5, ...)
```

Arguments

X	An $N \times p$ numeric matrix of predictor variables.
y	A numeric response vector of length N .
df	Degrees of freedom. Default is 5.
\dots	Additional arguments passed to <code>gbass()</code> .

Details

Uses an inverse-gamma latent-scale representation through a GIG prior on the local variance components. If $df > 2$, the scale is set to $(df - 2) / df$ so that the marginal variance matches w .

Value

An object of class `c("tbass", "gbass")` containing posterior draws and fitted model information.

Index

[build_GBP \(build_prior\)](#), [2](#)
[build_GIG \(build_prior\)](#), [2](#)
[build_prior](#), [2](#)

[GBASS \(GBASS-package\)](#), [2](#)
[gbass](#), [3](#)
[GBASS-package](#), [2](#)
[gbass2bass](#), [6](#), [7](#)
[gm2bm \(gbass2bass\)](#), [6](#)
[gsobol](#), [6](#)

[hbass](#), [7](#)

[nw_est_mom](#), [10](#)
[nw_gamma_prior](#), [12](#)
[nw_triangle](#), [13](#)
[nwbass](#), [8](#)
[nwbass2 \(nwbass\)](#), [8](#)

[optim](#), [11](#), [12](#)

[plot.gbass](#), [14](#)
[points](#), [13](#)
[predict.gbass](#), [14](#)
[print.gbass](#), [15](#)

[qbass](#), [16](#)

[rgig2](#), [17](#)
[rmhn](#), [18](#)

[sobol](#), [7](#)

[tbass](#), [18](#)