

Package: bqmm (via r-universe)

July 11, 2026

Title Bayesian Multilevel Quantile Regression

Version 0.1.0

Description Fits Bayesian mixed-effects (multilevel) quantile regression models using the asymmetric Laplace working likelihood and Stan. Supports an 'lme4'-style formula interface with nested and crossed random effects, fitting one or several quantiles, post-hoc non-crossing rearrangement of fitted quantiles, and the Yang, Wang and He (2016) posterior-variance correction for valid frequentist inference from the (misspecified) asymmetric Laplace posterior.

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Encoding UTF-8

Roxygen list(markdown = TRUE, load = ``installed")

RoxygenNote 7.3.3

Biarch true

Depends R (>= 4.0.0)

Imports methods, stats, graphics, grDevices, lme4, Matrix, Rcpp (>= 0.12.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.0), rstantools (>= 2.4.0), posterior

LinkingTo BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.0), StanHeaders (>= 2.26.0)

Suggests bayesplot, ggplot2, lqmm, MASS, nlme, quantreg, testthat (>= 3.0.0), knitr, rmarkdown

SystemRequirements GNU make

Config/testthat/edition 3

Config/Needs/website pkgdown

VignetteBuilder knitr

URL <https://github.com/kvenkita/bqmm>, <https://kvenkita.github.io/bqmm/>

BugReports <https://github.com/kvenkita/bqmm/issues>

Config/pak/sysreqs cmake make
Repository https://kvenkita.r-universe.dev
Date/Publication 2026-06-06 22:12:33 UTC
RemoteUrl https://github.com/kvenkita/bqmm
RemoteRef HEAD
RemoteSha 5e923b38ebc3e9444ef76297b931a453a1d96327

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ald	<i>The asymmetric Laplace family for quantile regression</i>
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Description

A lightweight family object describing the asymmetric Laplace distribution (ALD) working likelihood used by `bqmm()`. It mirrors the role of a `stats::family` object but is intentionally minimal in this release.

Usage

```
ald(link = "identity")
```

Arguments

link Name of the link for the location (quantile) parameter. Only "identity" is supported in v0.1.

Value

An object of class "bqmm_family".

Examples

```
ald()
```

as.matrix.bqmm	<i>Coerce a bqmm fit to a matrix of posterior draws</i>
----------------	---

Description

Coerce a bqmm fit to a matrix of posterior draws

Usage

```
## S3 method for class 'bqmm'
as.matrix(x, ...)
```

Arguments

x A bqmm fit.
... Unused.

Value

A matrix of posterior draws (draws in rows, parameters in columns), using the raw Stan parameter names.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
dim(as.matrix(fit))
```

<code>as_draws.bqmm</code>	<i>Convert a bqmm fit to a posterior draws object</i>
----------------------------	---

Description

Convert a bqmm fit to a posterior draws object

Usage

```
## S3 method for class 'bqmm'
as_draws(x, ...)
```

Arguments

<code>x</code>	A bqmm fit.
<code>...</code>	Unused.

Value

A `draws_array` (from the `posterior` package) with tidy variable names: `b_<name>` for fixed effects, `sd_<component>` for random-effect SDs, and `sigma`.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
as_draws(fit)
```

<code>bqmm</code>	<i>Bayesian multilevel quantile regression</i>
-------------------	--

Description

Fits a Bayesian mixed-effects quantile regression model using the asymmetric Laplace working likelihood and Stan. The interface follows `lme4`: random effects are written inline in the formula, e.g. `y ~ x + (1 + x | group)`, and nested or crossed grouping factors are both supported.

Usage

```

bqmm(
  formula,
  data,
  tau = 0.5,
  family = ald(),
  prior = NULL,
  cov = c("diagonal", "unstructured"),
  adjust = TRUE,
  prior_only = FALSE,
  chains = 4,
  iter = 2000,
  warmup = floor(iter/2),
  cores = getOption("mc.cores", 1L),
  seed = NULL,
  control = list(adapt_delta = 0.95),
  ...
)

```

Arguments

<code>formula</code>	An lme4-style model formula.
<code>data</code>	A data frame containing the variables in formula.
<code>tau</code>	Quantile level(s) in (0, 1). Scalar or vector.
<code>family</code>	A <code>bqmm_family</code> object; currently only <code>ald()</code> .
<code>prior</code>	A <code>bqmm_prior()</code> object, or NULL for data-scaled defaults.
<code>cov</code>	Random-effect covariance structure. "diagonal" (default) models independent random effects and supports any number of nested or crossed terms. "unstructured" adds an LKJ-correlated covariance but currently requires exactly one random-effects term (e.g. $y \sim x + (1 + x \mid g)$).
<code>adjust</code>	Logical; compute the Yang-Wang-He (2016) variance correction so that <code>vcov(fit, adjusted = TRUE)</code> returns valid fixed-effect uncertainty. Default TRUE.
<code>prior_only</code>	Logical; sample from the prior predictive distribution.
<code>chains, iter, warmup, cores, seed</code>	Passed to <code>rstan::sampling()</code> .
<code>control</code>	A list of sampler control parameters (e.g. <code>adapt_delta</code>). Defaults raise <code>adapt_delta</code> to 0.95 because ALD posteriors are sharp.
<code>...</code>	Additional arguments forwarded to <code>rstan::sampling()</code> .

Details

One or several quantiles may be requested through `tau`. A scalar returns a single `bqmm` fit; a vector fits each quantile independently and returns a `bqmm_multi` container.

Value

A `bqmm` object (single `tau`) or a `bqmm_multi` object (vector `tau`).

Examples

```
# A minimal fit; raise chains/iter for real analyses.
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
summary(fit)
```

bqmm_prior

Priors for a Bayesian quantile mixed model

Description

Builds the list of prior hyperparameters passed to Stan. Defaults are weakly informative and scaled to the data (see `bqmm_default_priors()`); any element supplied here overrides the default.

Usage

```
bqmm_prior(
  beta_mean = 0,
  beta_sd = NULL,
  sigma_scale = NULL,
  re_scale = NULL,
  lkj = 2
)
```

Arguments

beta_mean	Numeric scalar or vector: prior mean(s) for the fixed-effect coefficients. Recycled to the number of columns of the design matrix. Default 0.
beta_sd	Positive scalar or vector: prior SD(s) for the fixed-effect coefficients. NULL (default) uses a data-scaled value.
sigma_scale	Positive scalar: half-normal scale for the ALD scale sigma. NULL (default) uses a data-scaled value.
re_scale	Positive scalar: half-normal scale for the random-effect standard deviations. NULL (default) uses a data-scaled value.
lkj	Positive scalar: LKJ shape parameter for the random-effect correlation matrix (used only when <code>cov = "unstructured"</code>). 2 favours weak correlations; 1 is uniform over correlation matrices.

Value

An object of class "bqmm_prior".

Examples

```
bqmm_prior(beta_sd = 5)
```

coef.bqmm	<i>Extract model coefficients</i>
-----------	-----------------------------------

Description

Alias for `fixef()`; returns the posterior-median fixed effects.

Usage

```
## S3 method for class 'bqmm'
coef(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

A named numeric vector of posterior-median fixed-effect coefficients.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
coef(fit)
```

coef.bqmm_multi	<i>Coefficient-versus-tau matrix for a bqmm_multi fit</i>
-----------------	---

Description

Coefficient-versus-tau matrix for a bqmm_multi fit

Usage

```
## S3 method for class 'bqmm_multi'
coef(object, ...)
```

Arguments

object	A bqmm_multi fit.
...	Unused.

Value

A tau-by-coefficient matrix of posterior-median fixed effects, with one row per quantile.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = c(0.25, 0.75), chains = 1, iter = 250,
            refresh = 0, seed = 1)
coef(fit)
```

confint.bqmm

Confidence (credible) intervals for the fixed effects

Description

Wald-type intervals built from the posterior-median estimates and the (optionally misspecification-corrected) fixed-effect covariance.

Usage

```
## S3 method for class 'bqmm'
confint(
  object,
  parm,
  level = 0.95,
  adjusted = TRUE,
  method = c("ywh", "ij"),
  cluster = TRUE,
  ...
)
```

Arguments

object	A bqmm fit.
parm	Optional subset of coefficients (names or indices) to return.
level	Interval coverage (default 0.95).
adjusted	Logical; if TRUE (default) use the corrected covariance from vcov.bqmm() , otherwise the naive posterior covariance.
method	Correction to use when adjusted = TRUE; see vcov.bqmm() .
cluster	Logical; use the cluster-robust form (default TRUE).
...	Unused.

Value

A matrix with one row per coefficient and lower/upper interval columns.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
confint(fit)
```

fitted.bqmm

*Linear predictor (conditional tau-quantile) at the posterior median***Description**

Linear predictor (conditional tau-quantile) at the posterior median

Usage

```
## S3 method for class 'bqmm'
fitted(object, ...)
```

Arguments

```
object      A bqmm fit.
...         Unused.
```

Value

Numeric vector of fitted conditional quantiles.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
head(fitted(fit))
```

fixef.bqmm

*Posterior-median fixed effects***Description**

Posterior-median fixed effects

Usage

```
## S3 method for class 'bqmm'
fixef(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

A named numeric vector of posterior-median fixed-effect coefficients.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
fixef(fit)
```

log_lik.bqmm	<i>Pointwise log-likelihood draws</i>
--------------	---------------------------------------

Description

Pointwise log-likelihood draws

Usage

```
## S3 method for class 'bqmm'
log_lik(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

An S x N matrix of pointwise log-likelihood values, suitable for use with the loo package.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
dim(log_lik(fit))
```

nobs.bqmm	<i>Number of observations used in the fit</i>
-----------	---

Description

Number of observations used in the fit

Usage

```
## S3 method for class 'bqmm'
nobs(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

An integer, the number of observations.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
nobs(fit)
```

plot.bqmm	<i>Plot a bqmm fit</i>
-----------	------------------------

Description

Default plot shows fixed-effect posterior intervals. With bayesplot installed, richer MCMC plots are available via [as_draws\(\)](#).

Usage

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

Arguments

x	A bqmm fit.
...	Unused.

Value

Invisibly, x.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
plot(fit)
```

plot.bqmm_multi	<i>Plot coefficient-versus-tau paths for a bqmm_multi fit</i>
-----------------	---

Description

Plot coefficient-versus-tau paths for a bqmm_multi fit

Usage

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

Arguments

x	A bqmm_multi fit.
...	Unused.

Value

Invisibly, x.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = c(0.25, 0.75), chains = 1, iter = 250,
            refresh = 0, seed = 1)
plot(fit)
```

posterior_epred.bqmm *Draws of the expected response (conditional tau-quantile)*

Description

Draws of the expected response (conditional tau-quantile)

Usage

```
## S3 method for class 'bqmm'
posterior_epred(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

An S x N matrix of posterior draws of the linear predictor, where S is the number of posterior draws and N the number of observations.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
dim(posterior_epred(fit))
```

posterior_predict.bqmm *Draws from the posterior predictive distribution*

Description

Draws from the posterior predictive distribution

Usage

```
## S3 method for class 'bqmm'
posterior_predict(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

An S x N matrix of posterior predictive draws of the response, where S is the number of posterior draws and N the number of observations.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
dim(posterior_predict(fit))
```

predict.bqmm	<i>Predictions from a bqmm fit</i>
--------------	------------------------------------

Description

Predictions from a bqmm fit

Usage

```
## S3 method for class 'bqmm'
predict(
  object,
  newdata = NULL,
  re.form = NULL,
  noncrossing = c("none", "rearrange"),
  ...
)
```

Arguments

object	A bqmm fit.
newdata	Optional data frame; if omitted, training data are used.
re.form	NULL includes random effects (training data only); NA gives population-level predictions.
noncrossing	One of "none" or "rearrange". Rearrangement only has an effect for bqmm_multi objects (multiple quantiles).
...	Unused.

Value

Numeric vector of predicted conditional quantiles.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
head(predict(fit, re.form = NA))
```

ranef.bqmm	<i>Posterior-median random effects</i>
------------	--

Description

Posterior-median random effects

Usage

```
## S3 method for class 'bqmm'
ranef(object, ...)
```

Arguments

object	A bqmm fit.
...	Unused.

Value

A numeric vector of posterior-median random effects aligned with the columns of the random-effects design matrix Z, or NULL if the model has no random effects.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
ranef(fit)
```

rearrange_quantiles	<i>Rearrange fitted quantiles to remove crossing</i>
---------------------	--

Description

Post-hoc monotonisation of estimated quantile curves (Chernozhukov, Fernandez-Val and Galichon, 2010): at any covariate point, sorting the fitted values across quantile levels into increasing order yields a valid, non-crossing set of quantiles and never increases estimation error. This is the v0.1 non-crossing strategy; joint constrained estimation is deferred.

Usage

```
rearrange_quantiles(preds)
```

Arguments

`preds` A numeric matrix of fitted quantiles with one column per quantile level, ordered by increasing τ (rows = observations).

Value

A matrix of the same shape with each row sorted increasingly.

Examples

```
m <- rbind(c(1, 0.5, 2), c(0, 1, 0.8)) # some crossings
rearrange_quantiles(m)
```

residuals.bqmm	<i>Residuals from a bqmm fit</i>
----------------	----------------------------------

Description

Residuals from a bqmm fit

Usage

```
## S3 method for class 'bqmm'
residuals(object, ...)
```

Arguments

`object` A bqmm fit.
`...` Unused.

Value

Numeric vector of response-minus-fitted residuals.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
head(residuals(fit))
```

summary.bqmm

*Summarize a bqmm fit***Description**

Produces a fixed-effect coefficient table (estimate, standard error and interval) together with random-effect standard deviations.

Usage

```
## S3 method for class 'bqmm'
summary(
  object,
  level = 0.95,
  adjusted = TRUE,
  method = c("ywh", "ij"),
  cluster = TRUE,
  ...
)
```

Arguments

object	A bqmm fit.
level	Interval coverage (default 0.95).
adjusted	Logical; if TRUE (default) use the corrected covariance from vcov.bqmm() for the standard errors and intervals.
method	Correction to use when adjusted = TRUE; see vcov.bqmm() .
cluster	Logical; use the cluster-robust form (default TRUE).
...	Unused.

Value

An object of class summary.bqmm.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
summary(fit)
```

summary.bqmm_multi	<i>Summarize a bqmm_multi fit</i>
--------------------	-----------------------------------

Description

Summarize a bqmm_multi fit

Usage

```
## S3 method for class 'bqmm_multi'
summary(object, ...)
```

Arguments

object	A bqmm_multi fit.
...	Passed to the per-quantile summary() method for each fit.

Value

A list of summary.bqmm objects, one per quantile.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = c(0.25, 0.75), chains = 1, iter = 250,
            refresh = 0, seed = 1)
summary(fit)
```

VarCorr.bqmm	<i>Random-effect standard deviations and correlations</i>
--------------	---

Description

Random-effect standard deviations and correlations

Usage

```
## S3 method for class 'bqmm'
VarCorr(x, sigma = 1, ...)
```

Arguments

x	A bqmm fit.
sigma	Ignored; present for compatibility with the generic.
...	Unused.

Value

A named numeric vector of posterior-median random-effect standard deviations (with a posterior-median correlation matrix attached as the "correlation" attribute for unstructured models), or NULL if the model has no random effects.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
VarCorr(fit)
```

vcov.bqmm	<i>Variance-covariance of the fixed effects</i>
-----------	---

Description

Variance-covariance of the fixed effects

Usage

```
## S3 method for class 'bqmm'
vcov(object, adjusted = TRUE, method = c("ywh", "ij"), cluster = TRUE, ...)
```

Arguments

object	A bqmm fit.
adjusted	Logical; if TRUE (default) return a misspecification- corrected covariance (chosen by method), otherwise the naive posterior covariance.
method	Correction to use when adjusted = TRUE: "ywh" (default) is the Yang-Wang-He posterior-covariance sandwich (compute_ywh_multiplicative()); "ij" is the Infinitesimal Jackknife (ij_vcov()). Both are cluster-robust by default for a mixed model.
cluster	Logical; use the cluster-robust form (default TRUE).
...	Unused.

Value

A K x K covariance matrix for the fixed effects.

Examples

```
fit <- bqmm(distance ~ age + (1 | Subject), data = nlme::Orthodont,
            tau = 0.5, chains = 1, iter = 300, refresh = 0, seed = 1)
vcov(fit)
```

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