

## Description

`bayes: meglm` fits a Bayesian multilevel generalized linear model to outcomes of different types such as continuous, binary, count, and so on; see [\[BAYES\] bayes](#) and [\[ME\] meglm](#) for details.

## Quick start

Bayesian two-level generalized linear model of `y` on `x1` and `x2` with random intercepts by `id`, using the Gaussian family and log link, and using default normal priors for regression coefficients and default inverse-gamma prior for the variance of random intercepts

```
bayes: meglm y x1 x2 || id:, family(gaussian) link(log)
```

Use a standard deviation of 10 instead of 100 for the default normal priors

```
bayes, normalprior(10): meglm y x1 x2 || id:, family(gaussian) link(log)
```

Use uniform priors for the slopes and a normal prior for the intercept

```
bayes, prior({y: x1 x2}, uniform(-10,10)) ///
prior({y:_cons}, normal(0,10)): ///
meglm y x1 x2 || id:, family(gaussian) link(log)
```

Save simulation results to `simdata.dta`, and use a random-number seed for reproducibility

```
bayes, saving(simdata) rseed(123): ///
meglm y x1 x2 || id:, family(gaussian) link(log)
```

Specify 20,000 Markov chain Monte Carlo (MCMC) samples, set length of the burn-in period to 5,000, and request that a dot be displayed every 500 simulations

```
bayes, mcmcsize(20000) burnin(5000) dots(500): ///
meglm y x1 x2 || id:, family(gaussian) link(log)
```

In the above, request that the 90% highest posterior density (HPD) credible interval be displayed instead of the default 95% equal-tailed credible interval

```
bayes, clevel(90) hpd
```

Fit a logit model and display results as odds ratios

```
bayes: meglm z x1 x2 || id:, family(binomial) eform
```

Display odds ratios on replay

```
bayes, eform
```

Also see [Quick start](#) in [\[BAYES\] bayes](#) and [Quick start](#) in [\[ME\] meglm](#).

## Menu

Statistics > Multilevel mixed-effects models > Bayesian regression > Generalized linear model (GLM)

## Syntax

```
bayes [ , bayesopts ] : meglm depvar fe_equation
[ || re_equation ] [ || re_equation ... ] [ , options ]
```

where the syntax of *fe\_equation* is

```
[ indepvars ] [ if ] [ in ] [ weight ] [ , fe_options ]
```

and the syntax of *re\_equation* is one of the following:

for random coefficients and intercepts

```
levelvar: [ varlist ] [ , re_options ]
```

for random effects among the values of a factor variable

```
levelvar: R. varname
```

*levelvar* either is a variable identifying the group structure for the random effects at that level or is `_all`, representing one group comprising all observations.

<i>fe_options</i>	Description
Model	
<code>noconstant</code>	suppress constant term from the <b>fixed-effects</b> equation
<code>exposure(<i>varname</i><sub>e</sub>)</code>	include $\ln(\textit{varname}_e)$ in model with coefficient constrained to 1
<code>offset(<i>varname</i><sub>o</sub>)</code>	include <i>varname</i> <sub>o</sub> in model with coefficient constrained to 1
<code>asis</code>	retain perfect predictor variables

<i>re_options</i>	Description
Model	
<code>covariance(<i>vartype</i>)</code>	variance–covariance structure of the <b>random effects</b> ; only structures independent, exchangeable, identity, and unstructured are supported
<code>noconstant</code>	suppress constant term from the random-effects equation

<i>options</i>	Description
<b>Model</b>	
<code>family(family)</code>	distribution of <i>depvar</i> ; default is <code>family(gaussian)</code>
<code>link(link)</code>	link function; default varies per family
<b>Reporting</b>	
<code>eform</code>	report exponentiated coefficients
<code>irr</code>	report incidence-rate ratios
<code>or</code>	report odds ratios
<code>notable</code>	suppress coefficient table
<code>noheader</code>	suppress output header
<code>nogroup</code>	suppress table summarizing groups
<code>display_options</code>	control spacing, line width, and base and empty cells
<code>level(#)</code>	set credible level; default is <code>level(95)</code>

*indepvars* may contain factor variables; see [U] 11.4.3 Factor variables.

*depvar*, *indepvars*, and *varlist* may contain time-series operators; see [U] 11.4.4 Time-series varlists.

*fweights* are allowed; see [U] 11.1.6 weight.

`bayes: meglm`, `level()` is equivalent to `bayes, clevel(): meglm`.

For a detailed description of *options*, see *Options* in [ME] `meglm`.

<i>bayesopts</i>	Description
<b>Priors</b>	
* <code>normalprior(#)</code>	specify standard deviation of default normal priors for regression coefficients; default is <code>normalprior(100)</code>
* <code>igammaprior(##)</code>	specify shape and scale of default inverse-gamma prior for variance components; default is <code>igammaprior(0.01 0.01)</code>
* <code>iwishartprior(#[...])</code>	specify degrees of freedom and, optionally, scale matrix of default inverse-Wishart prior for unstructured random-effects covariance
<code>prior(priorspec)</code>	prior for model parameters; this option may be repeated
<code>dryrun</code>	show model summary without estimation
<b>Simulation</b>	
<code>nchains(#)</code>	number of chains; default is to simulate one chain
<code>mcmcsize(#)</code>	MCMC sample size; default is <code>mcmcsize(10000)</code>
<code>burnin(#)</code>	burn-in period; default is <code>burnin(2500)</code>
<code>thinning(#)</code>	thinning interval; default is <code>thinning(1)</code>
<code>rseed(#)</code>	random-number seed
<code>exclude(paramref)</code>	specify model parameters to be excluded from the simulation results
<code>restubs(restub1 restub2 ...)</code>	specify stubs for random-effects parameters for all levels
<b>Blocking</b>	
* <code>blocksize(#)</code>	maximum block size; default is <code>blocksize(50)</code>
<code>block(paramref[, blockopts])</code>	specify a block of model parameters; this option may be repeated
<code>blocksummary</code>	display block summary
* <code>noblocking</code>	do not block parameters by default

#### Initialization

<u>initial</u> ( <i>initspec</i> )	specify initial values for model parameters with a single chain
<u>init#</u> ( <i>initspec</i> )	specify initial values for #th chain; requires <code>nchains()</code>
<u>initall</u> ( <i>initspec</i> )	specify initial values for all chains; requires <code>nchains()</code>
<u>nomleinitial</u>	suppress the use of maximum likelihood estimates as starting values
<u>initrandom</u>	specify random initial values
<u>initsummary</u>	display initial values used for simulation
* <u>noisily</u>	display output from the estimation command during initialization

#### Adaptation

<u>adaptation</u> ( <i>adaptopts</i> )	control the adaptive MCMC procedure
<u>scale</u> (#)	initial multiplier for scale factor; default is <code>scale(2.38)</code>
<u>covariance</u> ( <i>cov</i> )	initial proposal covariance; default is the identity matrix

#### Reporting

<u>clevel</u> (#)	set credible interval level; default is <code>clevel(95)</code>
<u>hpd</u>	display HPD credible intervals instead of the default equal-tailed credible intervals
* <u>irr</u>	report incidence-rate ratios
* <u>or</u>	report odds ratios
<u>eform</u> [ ( <i>string</i> ) ]	report exponentiated coefficients and, optionally, label as <i>string</i>
<u>remargl</u>	compute log marginal-likelihood
<u>batch</u> (#)	specify length of block for batch-means calculations; default is <code>batch(0)</code>
<u>saving</u> ( <i>filename</i> [ , <i>replace</i> ])	save simulation results to <i>filename.dta</i>
<u>nomodelsummary</u>	suppress model summary
<u>nomesummary</u>	suppress multilevel-structure summary
<u>chainsdetail</u>	display detailed simulation summary for each chain
[ <u>no</u> ] <u>dots</u>	suppress dots or display dots every 100 iterations and iteration numbers every 1,000 iterations; default is <code>dots</code>
<u>dots</u> (#[ , <u>every</u> (#) ])	display dots as simulation is performed
[ <u>no</u> ] <u>show</u> ( <i>paramref</i> )	specify model parameters to be excluded from or included in the output
<u>showeffects</u> [ ( <i>reref</i> ) ]	specify that all or a subset of random-effects parameters be included in the output
<u>melabel</u>	display estimation table using the same row labels as <code>meglm</code>
<u>nogroup</u>	suppress table summarizing groups
<u>notable</u>	suppress estimation table
<u>noheader</u>	suppress output header
<u>title</u> ( <i>string</i> )	display <i>string</i> as title above the table of parameter estimates
<u>display_options</u>	control spacing, line width, and base and empty cells

#### Advanced

<u>search</u> ( <i>search_options</i> )	control the search for feasible initial values
<u>corrlag</u> (#)	specify maximum autocorrelation lag; default varies
<u>corrtol</u> (#)	specify autocorrelation tolerance; default is <code>corrtol(0.01)</code>

\* Starred options are specific to the bayes prefix; other options are common between bayes and bayesmh.

Options prior() and block() may be repeated.

priorspec and paramref are defined in [BAYES] bayesmh.

paramref may contain factor variables; see [U] 11.4.3 Factor variables.

collect is allowed; see [U] 11.1.10 Prefix commands.

See [U] 20 Estimation and postestimation commands for more capabilities of estimation commands.

Model parameters are regression coefficients {*depvar: indepvars*}, parameters as described in *Additional model parameters*, random effects {*rename*}, and either variance components {*rename: sigma2*} or, if option covariance(unstructured) is specified, matrix parameter {*restub: Sigma, matrix*}; see *Likelihood model* in [BAYES] bayes for how *renames* and *restub* are defined. Use the dryrun option to see the definitions of model parameters prior to estimation.

For a detailed description of *bayesopts*, see *Options* in [BAYES] bayes.

## Remarks and examples

For a general introduction to Bayesian analysis, see [BAYES] Intro. For a general introduction to Bayesian estimation using an adaptive Metropolis–Hastings algorithm, see [BAYES] bayesmh. For remarks and examples specific to the bayes prefix, see [BAYES] bayes. For details about the estimation command, see [ME] meglm.

For a simple example of the bayes prefix, see *Introductory example* in [BAYES] bayes. For multilevel examples, see *Multilevel models* in [BAYES] bayes. Also see *Crossed-effects model* in [BAYES] bayes.

## Additional model parameters

In addition to regression coefficients {*depvar: indepvars*}, bayes: meglm defines extra parameters that depend on the chosen family; see table 1 below.

Table 1. Additional model parameters defined by bayes: meglm

Family	Parameter	Model parameter	Default prior
Gaussian	Error variance	{ <i>e.depvar: sigma2</i> }	InvGamma(0.01, 0.01)
Bernoulli/Binomial	None	None	None
Ordinal	Cutpoints	{ <i>cut1</i> }, { <i>cut2</i> }, ...	Flat
Poisson	None	None	None
Negative binomial	Log-overdispersion	{ <i>lnalpha</i> } (mean disp.) { <i>lndelta</i> } (constant disp.)	$N(0, 10000)$ $N(0, 10000)$
Gamma	Log-scale	{ <i>lnscale</i> }	$N(0, 10000)$

Use the dryrun option with the bayes prefix to see the definitions of model parameters prior to estimation.

## Stored results

See *Stored results* in [BAYES] bayes.

## Methods and formulas

See *Methods and formulas* in [BAYES] [bayesmh](#).

## Also see

[BAYES] [bayes](#) — Bayesian regression models using the bayes prefix

[ME] [meglm](#) — Multilevel mixed-effects generalized linear models

[BAYES] [Bayesian postestimation](#) — Postestimation tools after Bayesian estimation

[BAYES] [Bayesian estimation](#) — Bayesian estimation commands

[BAYES] [Bayesian commands](#) — Introduction to commands for Bayesian analysis

[BAYES] [Intro](#) — Introduction to Bayesian analysis

[BAYES] [Glossary](#)

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