

**bayes: meglm** — Bayesian multilevel generalized linear model

|                                      |                                |                                      |                          |
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## 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   |
|--|---|
| <code>Model</code>                                       |   |
| <code><u>noconstant</u></code>                           | suppress constant term from the <a href="#">fixed-effects</a> equation        |
| <code><u>exposure</u>(<i>varname<sub>e</sub></i>)</code> | include $\ln(\text{varname}_e)$ in model with coefficient constrained to 1    |
| <code><u>offset</u>(<i>varname<sub>o</sub></i>)</code>   | include <i>varname<sub>o</sub></i> in model with coefficient constrained to 1 |
| <code>asis</code>  | retain perfect predictor variables  |

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

| <i>options</i>                  | Description   |
|---------------------------------|---|
| <b>Model</b>                    |   |
| <u>f</u> amily( <i>family</i> ) | distribution of <i>depvar</i> ; default is family(gaussian) |
| <u>l</u> ink( <i>link</i> )     | link function; default varies per family                    |
| <b>Reporting</b>                |   |
| eform                           | report exponentiated coefficients                           |
| irr                             | report incidence-rate ratios                                |
| or                              | report odds ratios  |
| <u>n</u> otable                 | suppress coefficient table                                  |
| <u>n</u> oheader                | suppress output header                                      |
| <u>n</u> ogroup                 | suppress table summarizing groups                           |
| <i>display_options</i>          | control spacing, line width, and base and empty cells       |
| <u>l</u> evel(#)                | set credible level; default is level(95)                    |

*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>                     |  |
| * <u>n</u> ormalprior(#)          | specify standard deviation of default normal priors for regression coefficients; default is normalprior(100)                         |
| * <u>i</u> gammaprior(# #)        | specify shape and scale of default inverse-gamma prior for variance components; default is igammaprior(0.01 0.01)                    |
| * <u>i</u> wishartprior(# [...])  | specify degrees of freedom and, optionally, scale matrix of default inverse-Wishart prior for unstructured random-effects covariance |
| <u>p</u> rior( <i>priorspec</i> ) | prior for model parameters; this option may be repeated  |
| <u>d</u> ryrun                    | show model summary without estimation  |

|   |   |
|---|---|
| <b>Simulation</b>                             |   |
| <u>n</u> chains(#)                            | number of chains; default is to simulate one chain                  |
| <u>m</u> cmcsizе(#)                           | MCMC sample size; default is mcmcsizе(10000)                        |
| <u>b</u> urnin(#)                             | burn-in period; default is burnin(2500)                             |
| <u>t</u> hinning(#)                           | thinning interval; default is thinning(1)                           |
| <u>r</u> seed(#)                              | random-number seed  |
| <u>e</u> xclude( <i>paramref</i> )            | specify model parameters to be excluded from the simulation results |
| <u>r</u> estubs( <i>restub1 restub2 ...</i> ) | specify stubs for random-effects parameters for all levels          |

|  |  |
|--|--|
| <b>Blocking</b>  |  |
| * <u>b</u> locksize(#)                                 | maximum block size; default is blocksize(50)                     |
| <u>b</u> lock( <i>paramref</i> [ , <i>blockopts</i> ]) | specify a block of model parameters; this option may be repeated |
| <u>b</u> locksummary                                   | display block summary  |
| * <u>n</u> oblocking                                   | do not block parameters by default                               |

### Initialization

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

### Adaptation

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

### Reporting

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

### Advanced

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

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\*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

[stata.com](http://stata.com)

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     | <code>{e.depvar:sigma2}</code>   | $\text{InvGamma}(0.01, 0.01)$  |
| Bernoulli/Binomial | None               | None   | None                           |
| Ordinal            | Cutpoints          | <code>{cut1}</code> , <code>{cut2}</code> , ...                                | Flat                           |
| Poisson            | None               | None   | None                           |
| Negative binomial  | Log-overdispersion | <code>{lnalpha}</code> (mean disp.)<br><code>{lndelta}</code> (constant disp.) | $N(0, 10000)$<br>$N(0, 10000)$ |
| Gamma              | Log-scale          | <code>{lnscale}</code>   | $N(0, 10000)$                  |

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 for bayesmh and the bayes prefix

[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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