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## Description

`bayes: meologit` fits a Bayesian multilevel ordered logistic regression to an ordinal outcome; see [\[BAYES\] bayes](#) and [\[ME\] meologit](#) for details.

## Quick start

Bayesian two-level ordered logistic regression of `y` on `x1` and `x2` with random intercepts by `id`, using default normal priors for regression coefficients, flat priors for cutpoints, and default inverse-gamma prior for the variance of random intercepts

```
bayes: meologit y x1 x2 || id:
```

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

```
bayes, normalprior(10): meologit y x1 x2 || id:
```

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)): meologit y x1 x2 || id:
```

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

```
bayes, saving(simdata) rseed(123): meologit y x1 x2 || id:
```

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, mcmcsz(20000) burnin(5000) dots(500): meologit y x1 x2 || id:
```

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
```

Display odds ratios instead of coefficients

```
bayes: meologit y x1 x2 || id: , or
```

Display odds ratios on replay

```
bayes, or
```

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

## Menu

Statistics > Multilevel mixed-effects models > Bayesian regression > Ordered logistic regression

## Syntax

```
bayes [ , bayesopts ] : meologit 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
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Model	
<code><u>offset</u>(<i>varname</i>)</code>	include <i>varname</i> in model with coefficient constrained to 1

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

<i>options</i>	Description
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Reporting	
<code><u>or</u></code>	report odds ratios
<code><u>notable</u></code>	suppress coefficient table
<code><u>noheader</u></code>	suppress output header
<code><u>nogroup</u></code>	suppress table summarizing groups
<code><i>display_options</i></code>	control spacing, line width, and base and empty cells
<code><u>level</u>(#)</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: meologit, `level()` is equivalent to bayes, `clevel()`: meologit.  
For a detailed description of *options*, see *Options* in [ME] **meologit**.

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

<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>* or</code>	report coefficients as odds ratios
<code>eform(string)</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(filename[, replace])</code>	save simulation results to <i>filename.dta</i>
<code>nomodelsummary</code>	suppress model summary
<code>nomesummary</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(paramref)</code>	specify model parameters to be excluded from or included in the output
<code>showeffects(reref)</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>meologit</code>
<code>nogroup</code>	suppress table summarizing groups
<code>notable</code>	suppress estimation table
<code>noheader</code>	suppress output header
<code>title(string)</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(search_options)</code>	control the search for feasible initial values
<code>corrlag(#)</code>	specify maximum autocorrelation lag; default varies
<code>corrto1(#)</code>	specify autocorrelation tolerance; default is <code>corrto1(0.01)</code>

\* Starred options are specific to the `bayan` prefix; other options are common between `bayan` 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}`, cutpoints `{cut1}`, `{cut2}`, and so on, 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.

Flat priors, `flat`, are used by default for cutpoints.

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] [meologit](#).

For a simple example of the bayes prefix, see *Introductory example* in [BAYES] [bayes](#). For multilevel examples, see *Multilevel models* in [BAYES] [bayes](#).

## 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] [meologit](#) — Multilevel mixed-effects ordered logistic regression

[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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