

**bayes: nbreg** — Bayesian negative binomial regression

[Description](#)[Remarks and examples](#)[Quick start](#)[Stored results](#)[Menu](#)[Methods and formulas](#)[Syntax](#)[Also see](#)

## Description

`bayes: nbreg` fits a Bayesian negative binomial regression to a nonnegative count outcome; see [\[BAYES\] bayes](#) and [\[R\] nbreg](#) for details.

## Quick start

Bayesian negative binomial regression of  $y$  on  $x_1$  and  $x_2$ , using default normal priors for regression coefficients and log-overdispersion parameter

```
bayes: nbreg y x1 x2
```

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

```
bayes, normalprior(10): nbreg y x1 x2
```

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)): nbreg y x1 x2
```

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

```
bayes, saving(simdata) rseed(123): nbreg y x1 x2
```

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): nbreg y x1 x2
```

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 incidence-rate ratios instead of coefficients

```
bayes: nbreg y x1 x2, irr
```

Display incidence-rate ratios on replay

```
bayes, irr
```

Also see [Quick start](#) in [\[BAYES\] bayes](#) and [Quick start](#) in [\[R\] nbreg](#).

## Menu

Statistics > Count outcomes > Bayesian regression > Negative binomial regression

## Syntax

```
bayes [ , bayesopts ] : nbreg deivar [indepvars] [if] [in] [weight] [ , options ]
```

| <i>options</i> | Description |
|----------------|-------------|
|----------------|-------------|

### Model

|  |  |
|--|--|
| <code>noconstant</code>                    | suppress constant term   |
| <code>dispersion(mean)</code>              | parameterization of dispersion; the default                                |
| <code>dispersion(constant)</code>          | constant dispersion for all observations                                   |
| <code>exposure(varname<sub>e</sub>)</code> | include $\ln(\text{varname}_e)$ in model with coefficient constrained to 1 |
| <code>offset(varname<sub>o</sub>)</code>   | include $\text{varname}_o$ in model with coefficient constrained to 1      |

### Reporting

|                              |   |
|------------------------------|---|
| <code>irr</code>             | report incidence-rate ratios                          |
| <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**.

*deivar*, *indepvars*, *varname<sub>e</sub>*, and *varname<sub>o</sub>* may contain time-series operators; see [U] 11.4.4 **Time-series varlists**.

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

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

For a detailed description of options, see *Options for nbreg* in [R] **nbreg**.

| <i>bayesopts</i> | Description |
|------------------|-------------|
|------------------|-------------|

### Priors

|                               |  |
|-------------------------------|--|
| * <code>normalprior(#)</code> | specify standard deviation of default normal priors for regression coefficients and log-overdispersion parameter; default is <code>normalprior(100)</code> |
| <code>prior(priorspec)</code> | prior for model parameters; this option may be repeated  |
| <code>dryrun</code>           | show model summary without estimation  |

### Simulation

|                                |   |
|--------------------------------|---|
| <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 |

### Blocking

|  |  |
|--|--|
| * <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

|  |  |
|--|--|
| <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>initransom</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>eform[ (<i>string</i>) ]</code>                  | report exponentiated coefficients and, optionally, label as <i>string</i>   |
| <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>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>nodots</code> |
| <code>dots(#[, <i>every</i>(#)])</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>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> |

\*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 `{devar:indepvars}` and log-overdispersion parameter `{lnalpha}` with mean dispersion or `{lndelta}` with constant dispersion. 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 [\[R\] nbreg](#).

For a simple example of the `bayes` prefix, see *Introductory example* 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

[\[R\] nbreg](#) — Negative binomial regression

[\[BAYES\] 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](#)