

## 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 `x1` and `x2`, 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 devar [ indepvars ] [ if ] [ in ] [ weight ] [ , options ]
```

| <i>options</i>  | Description   |
|---|---|
| <b>Model</b>  |   |
| <code>noconstant</code>   | suppress constant term  |
| <code>dispersion(<u>m</u>ean)</code>  | parameterization of dispersion; the default   |
| <code>dispersion(<u>c</u>onstant)</code>  | constant dispersion for all observations  |
| <code>exposure(<i>varname</i><sub>e</sub>)</code>   | include $\ln(\text{varname}_e)$ 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  |
| <b>Reporting</b>  |   |
| <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 level(95)  |
| <p><i>indepvars</i> may contain factor variables; see [U] 11.4.3 <b>Factor variables</b>.</p> <p><i>devar</i>, <i>indepvars</i>, <i>varname</i><sub>e</sub>, and <i>varname</i><sub>o</sub> may contain time-series operators; see [U] 11.4.4 <b>Time-series varlists</b>.</p> <p>fweights are allowed; see [U] 11.1.6 <b>weight</b>.</p> <p>bayes: nbreg, level() is equivalent to bayes, clevel(): nbreg.</p> <p>For a detailed description of options, see <i>Options for nbreg</i> in [R] <b>nbreg</b>.</p> |   |
| <i>bayesopts</i>  | Description   |
| <b>Priors</b>   |   |
| * <code>normalprior(#)</code>   | specify standard deviation of default normal priors for regression coefficients and log-overdispersion parameter; default is normalprior(100) |
| <code>prior(<i>priorspec</i>)</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 mcmcsize(10000)  |
| <code>burnin(#)</code>  | burn-in period; default is burnin(2500)   |
| <code>thinning(#)</code>  | thinning interval; default is thinning(1)   |
| <code>rseed(#)</code>   | random-number seed  |
| <code>exclude(<i>paramref</i>)</code>   | specify model parameters to be excluded from the simulation results   |
| <b>Blocking</b>   |   |
| * <code>blocksize(#)</code>   | maximum block size; default is blocksize(50)  |
| <code>block(<i>paramref</i> [ , <i>blockopts</i> ])</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>initial(<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(cov)</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;<br>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>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#[ , 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>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 `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}` 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] `bayan`.

## 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 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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