Package: evinf (via r-universe)

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Type Package

Title Inference with Extreme Value Inflated Count Data

Version 0.8.10

Description Allows users to model and draw inferences from extreme value inflated count data, and to evaluate these models and compare to non extreme-value inflated counterparts. The package is built to be compatible with standard presentation tools such as 'broom', 'tidy', and 'modelsummary'.

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Encoding UTF-8

LazyData true

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LinkingTo Rcpp, RcppArmadillo

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RdMacros Rdpack

Depends generics, R (>= 2.10)

URL https://github.com/Doktorandahl/evinf

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Author David Randahl [cre, aut], Johan Vegelius [aut]

Maintainer David Randahl <david.randahl@pcr.uu.se>

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`coefficient_extractor` *Bootstrap coefficient extractor*

Description

Bootstrap coefficient extractor

Usage

```
coefficient_extractor(object, ...)
```

Arguments

object	a fitted model with bootstraps of class evzinb, evinb, nbboot, or zinbboot
...	Component to be extracted (not for nbboot). Alternatives are 'nb','zi','evinf','pareto', and 'all'

Value

A tibble with coefficient values, one row per bootstrap and component

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
coefficient_extractor(model, component = 'all')
```

---

coefficient_extractor.evinb

Bootstrap coefficient extractor

Description

Bootstrap coefficient extractor

Usage

```
## S3 method for class 'evinb'
coefficient_extractor(
  object,
  component = c("nb", "evinf", "pareto", "all"),
  ...
)
```

Arguments

object	A fitted evinb model with bootstraps
component	Which component should be extracted
...	Not in use

Value

A tibble with coefficient values, one row per bootstrap and component

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
coefficient_extractor(model, component = 'all')
```

coefficient_extractor.evzinb

Bootstrap coefficient extractor

Description

Bootstrap coefficient extractor

Usage

```
## S3 method for class 'evzinb'
coefficient_extractor(
  object,
  component = c("nb", "zi", "evinf", "pareto", "all"),
  ...
)
```

Arguments

object	A fitted evzinb model with bootstraps
component	Which component should be extracted
...	Not in use

Value

A tibble with coefficient values, one row per bootstrap and component

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
coefficient_extractor(model, component = 'all')
```

coefficient_extractor.nbboot

Bootstrap coefficient extractor

Description

Bootstrap coefficient extractor

Usage

```
## S3 method for class 'nbboot'
coefficient_extractor(object, ...)
```

Arguments

object	A fitted nbboot model with bootstraps
...	Not in use

Value

A tibble with coefficient value, one row per bootstrap

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
zinb_comp <- compare_models(model)
coefficient_extractor(zinb_comp$nb)
```

---

coefficient_extractor.zinbboot

Bootstrap coefficient extractor

Description

Bootstrap coefficient extractor

Usage

```
## S3 method for class 'zinbboot'
coefficient_extractor(object, component = c("nb", "zi", "all"), ...)
```

Arguments

object	A fitted evinb model with bootstraps
component	Which component should be extracted
...	Not in use

Value

A tibble with coefficient values, one row per bootstrap and component

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps=10)
zinb_comp <- compare_models(model)
coefficient_extractor(zinb_comp$zinb)
```

compare_models *Function to compare evzinb or evinb models with zinb and nb models*

Description

Function to compare evzinb or evinb models with zinb and nb models

Usage

```
compare_models(
  object,
  nb_comparison = TRUE,
  zinb_comparison = TRUE,
  winsorize = FALSE,
  razorize = FALSE,
  cutoff_value = 10,
  init_theta = NULL,
  multicore = FALSE,
  ncores = NULL
)
```

Arguments

object	A fitted evzinb or evinb model object
nb_comparison	Should comparison be made with a negative binomial model?
zinb_comparison	Should comparions be made with the zinb model?
winsorize	Should winsorizing be done in the comparisons?
razorize	Should razorizing (trimming) be done in the comparisons?
cutoff_value	Integer: Which observation should be used as a basis for winsorizing/razorising. E.g. 10 means that everything larger than the 10th observation will be winsorized/razorised
init_theta	Optional initial value for theta in the NB specification
multicore	Logical: should multiple cores be used
ncores	Number of cores if multicore is used

Value

A list with the original model as the first object and compared models as the following objects

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
compare_models(model)
```

evinb *Running an extreme value inflated negative binomial model with bootstrapping*

Description

Running an extreme value inflated negative binomial model with bootstrapping

Usage

```
evinb(
  formula_nb,
  formula_evi = NULL,
  formula_pareto = NULL,
  data,
  bootstrap = TRUE,
  n_bootstraps = 100,
  multicore = FALSE,
  ncores = NULL,
  block = NULL,
  boot_seed = NULL,
  max.diff.par = 0.01,
  max.no.em.steps = 500,
  max.no.em.steps.warmup = 5,
  c.lim = c(50, 1000),
  max.upd.par.pl.multinomial = 0.5,
  max.upd.par.nb = 0.5,
  max.upd.par.pl = 0.5,
  no.m.bfgs.steps.multinomial = 3,
  no.m.bfgs.steps.nb = 3,
  no.m.bfgs.steps.pl = 3,
  pdf.pl.type = "approx",
  eta.int = c(-1, 1),
  init.Beta.multinom.PL = NULL,
  init.Beta.NB = NULL,
  init.Beta.PL = NULL,
  init.Alpha.NB = 0.01,
  init.C = 200,
  verbose = FALSE
)
```

Arguments

formula_nb	Formula for the negative binomial (count) component of the model
formula_evi	Formula for the extreme-value inflation component of the model. If NULL taken as the same formula as nb

`formula_pareto` Formula for the pareto (extreme value) component of the model. If NULL taken as the same formula as nb

`data` Data to run the model on

`bootstrap` Should bootstrapping be performed. Needed to obtain standard errors and p-values

`n_bootstraps` Number of bootstraps to run. For use of bootstrapped p-values, at least 1,000 bootstraps are recommended. For approximate p-values, a lower number can be sufficient

`multicore` Should multiple cores be used?

`ncores` Number of cores if multicore is used. Default (NULL) is one less than the available number of cores

`block` Optional string indicating a case-identifier variable when using block bootstrapping

`boot_seed` Optional bootstrap seed to ensure reproducible results.

`max.diff.par` Tolerance for EM algorithm. Will be considered to have converged if the maximum absolute difference in the parameter estimates are lower than this value

`max.no.em.steps`
Maximum number of EM steps to run. Will be considered to not have converged if this number is reached and convergence is not reached

`max.no.em.steps.warmup`
Number of EM steps in the warmup rounds

`c.lim` Integer range defining the possible values of C

`max.upd.par.pl.multinomial`
Maximum parameter change step size in the extreme value inflation component

`max.upd.par.nb` Maximum parameter change step size in the count component

`max.upd.par.pl` Maximum parameter change step size in the pareto component

`no.m.bfgs.steps.multinomial`
Number of BFGS steps for the multinomial model

`no.m.bfgs.steps.nb`
Number of BFGS steps for the negative binomial model

`no.m.bfgs.steps.pl`
Number of BFGS steps for the pareto model

`pdf.pl.type` Probability density function type for the pareto component. Either 'approx' or 'exact'. 'approx' is adviced in most cases

`eta.int` Initial values for eta

`init.Beta.multinom.PL`
Initial values for beta parameters in the extreme value inflation component. Vector of same length as number of parameters in the extreme value inflation component or NULL (which gives starting values of 0)

`init.Beta.NB` Initial values for beta parameters in the count component. Vector of same length as number of parameters in the count component or NULL (which gives starting values of 0)

init.Beta.PL	Initial values for beta parameters in the pareto component. Vector of same length as number of parameters in the pareto component or NULL (which gives starting values of 0)
init.Alpha.NB	Initial value of Alpha NB, integer or NULL (giving a starting value of 0)
init.C	Initial value of C. Integer which should be within the C_lim range.
verbose	Should progress be printed for the first run of evinb

Value

An object of class 'evinb'

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
```

---

evzinb *Running an extreme value and zero inflated negative binomial model with bootstrapping*

Description

Running an extreme value and zero inflated negative binomial model with bootstrapping

Usage

```
evzinb(
  formula_nb,
  formula_zi = NULL,
  formula_evi = NULL,
  formula_pareto = NULL,
  data,
  bootstrap = TRUE,
  n_bootstraps = 100,
  multicore = FALSE,
  ncores = NULL,
  block = NULL,
  boot_seed = NULL,
  max.diff.par = 0.01,
  max.no.em.steps = 500,
  max.no.em.steps.warmup = 5,
  c.lim = c(50, 1000),
  max.upd.par.zc.multinomial = 0.5,
  max.upd.par.pl.multinomial = 0.5,
  max.upd.par.nb = 0.5,
  max.upd.par.pl = 0.5,
```

```
  no.m.bfgs.steps.multinomial = 3,
  no.m.bfgs.steps.nb = 3,
  no.m.bfgs.steps.pl = 3,
  pdf.pl.type = "approx",
  eta.int = c(-1, 1),
  init.Beta.multinom.ZC = NULL,
  init.Beta.multinom.PL = NULL,
  init.Beta.NB = NULL,
  init.Beta.PL = NULL,
  init.Alpha.NB = 0.01,
  init.C = 200,
  verbose = FALSE
)
```

Arguments

formula_nb	Formula for the negative binomial (count) component of the model
formula_zi	Formula for the zero-inflation component of the model. If NULL taken as the same formula as nb
formula_evi	Formula for the extreme-value inflation component of the model. If NULL taken as the same formula as nb
formula_pareto	Formula for the pareto (extreme value) component of the model. If NULL taken as the same formula as nb
data	data to run the model on
bootstrap	Should bootstrapping be performed. Needed to obtain standard errors and p-values
n_bootstraps	Number of bootstraps to run. For use of bootstrapped p-values, at least 1,000 bootstraps are recommended. For approximate p-values, a lower number can be sufficient
multicore	Should multiple cores be used?
ncores	Number of cores if multicore is used. Default (NULL) is one less than the available number of cores
block	Optional string indicating a case-identifier variable when using block bootstrapping
boot_seed	Optional bootstrap seed to ensure reproducible results.
max.diff.par	Tolerance for EM algorithm. Will be considered to have converged if the maximum absolute difference in the parameter estimates are lower than this value
max.no.em.steps	Maximum number of EM steps to run. Will be considered to not have converged if this number is reached and convergence is not reached
max.no.em.steps.warmup	Number of EM steps in the warmup rounds
c.lim	Integer range defining the possible values of C
max.upd.par.zc.multinomial	Maximum parameter change step size in the zero inflation component

`max.upd.par.pl.multinomial`
: Maximum parameter change step size in the extreme value inflation component

`max.upd.par.nb`
: Maximum parameter change step size in the count component

`max.upd.par.pl`
: Maximum parameter change step size in the pareto component

`no.m.bfgs.steps.multinomial`
: Number of BFGS steps for the multinomial model

`no.m.bfgs.steps.nb`
: Number of BFGS steps for the negative binomial model

`no.m.bfgs.steps.pl`
: Number of BFGS steps for the pareto model

`pdf.pl.type`
: Probability density function type for the pareto component. Either 'approx' or 'exact'. 'approx' is adviced in most cases

`eta.int`
: Initial values for eta

`init.Beta.multinom.ZC`
: Initial values for beta parameters in the zero value inflation component. Vector of same length as number of parameters in the zero value inflation component or NULL (which gives starting values of 0)

`init.Beta.multinom.PL`
: Initial values for beta parameters in the extreme value inflation component. Vector of same length as number of parameters in the extreme value inflation component or NULL (which gives starting values of 0)

`init.Beta.NB`
: Initial values for beta parameters in the count component. Vector of same length as number of parameters in the count component or NULL (which gives starting values of 0)

`init.Beta.PL`
: Initial values for beta parameters in the pareto component. Vector of same length as number of parameters in the pareto component or NULL (which gives starting values of 0)

`init.Alpha.NB`
: Initial value of Alpha NB, integer or NULL (giving a starting value of 0)

`init.C`
: Initial value of C. Integer which should be within the C_lim range.

`verbose`
: Logical: should progress of the full run of the model be tracked?

Value

An object of class 'evzinb'

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
```

genevzinb *Simulated data from the EVZBINB distribution*

Description

A simulated dataset of 1,000 observations with one dependent and three dependent variables generated using the EVZINB distribution

Usage

```
genevzinb
```

Format

'genevzinb' A tibble with 1,000 rows and 4 columns:

y Dependent variable following EVZINB distribution

x1, x2, x3 Continuous independent variables following the random normal distribution

genevzinb2 *Simulated data from the EVZBINB distribution*

Description

A simulated dataset of 100 observations with one dependent and three dependent variables generated using the EVZINB distribution

Usage

```
genevzinb2
```

Format

'genevzinb2' A tibble with 100 rows and 4 columns:

y Dependent variable following EVZINB distribution

x1, x2, x3 Continuous independent variables following the random normal distribution

glance.evinb *EVZINB and EVINB glance functions*

Description

EVZINB and EVINB glance functions

Usage

```
## S3 method for class 'evinb'
glance(x, ...)
```

Arguments

x	An EVZINB or EVINB object
...	Further arguments to be passed to glance()

Value

An EVZINB glance function

See Also

glance

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
glance(model)
```

glance.evzinb *EVZINB and EVINB glance functions*

Description

EVZINB and EVINB glance functions

Usage

```
## S3 method for class 'evzinb'
glance(x, ...)
```

Arguments

x	An EVZINB or EVINB object
...	Further arguments to be passed to glance()

Value

An EVZINB glance function

See Also

glance

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
glance(model)
```

---

glance.nbboot *zinbboot and nboot glance functions*

Description

zinbboot and nboot glance functions

Usage

```
## S3 method for class 'nbboot'
glance(x, ...)
```

Arguments

x	An nbboot or zinbboot object
...	Further arguments to be passed to glance()

Value

An nbboot glance function

See Also

glance

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
zinb_comp <- compare_models(model)
glance(zinb_comp$nb)
```

`glance.zinbboot` *zinbboot and nboot glance functions*

Description

zinbboot and nboot glance functions

Usage

```
## S3 method for class 'zinbboot'
glance(x, ...)
```

Arguments

`x`	An nbboot or zinbboot object
`...`	Further arguments to be passed to glance()

Value

An nbboot glance function

See Also

`glance`

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
zinb_comp <- compare_models(model)
glance(zinb_comp$zinb)
```

gm_evzinb — *A goodness-of-fit gof tibble for GOF metrics when using modelsummary*

Description

A goodness-of-fit gof tibble for GOF metrics when using modelsummary. The GM tibble can be used to obtain correct table output when making regression tables with modelsummary

Usage

```
gm_evzinb
```

Format

'gm_evzinb' A tibble with 7 rows and 3 columns:

raw The modelsummary/broom internal name for the statistic

clean The table output for the statistic

fmt The number of decimals reported for each statistic by default (can be adapted)

hks — *Replication data for Hultman, Kathman, and Shannon (2013) United Nations Peacekeeping and Civilian Protection in Civil War*

Description

A reduced replication data set from Hultman et al. (2013) United Nations Peacekeeping and Civilian Protection in Civil War. Used to reproduce the the results from Randahl and Vegelius (2023). Note, to reproduce any other results from Hultman et al. (2013) please download the original replication dataset using the link under source.

Usage

```
hks
```

Format

A tibble with 3746 rows and 12 columns:

conflict_id The Uppsala Conflict Data Programme conflict ID for the conflict

osvAll The number of observed fatalities from one-sided violence against civilian in the specified conflict-month

troopLag The number of UN military troops in thousands of troops (lagged)

policeLag The number of UN police in thousands of troops (lagged)

militaryobserversLag The number of UN military troops in thousands of troops (lagged)

epduration The number of months the current conflict-episode has been ongoing

lntpop The natural logarithm of the population of the country in which the conflict takes place

lnbrv_AllLag The natural logarithm of the total number of battle related deaths in the conflict in the previous month

osvAllLagDum A dummy variable taking the value 1 if any one-sided violence against civilians took place in the previous conflict month

incomp A dummy variable taking the value 1 if the conflict is about government and 0 otherwise

lntroopLag The log1p logarithm of troopLag

lnepdur The log1p logarithm of the episode duration

Source

https://dataverse.harvard.edu/dataset.xhtml?persistentId=doi:10.7910/DVN/6EBCGA

References

Hultman L, Kathman J, Shannon M (2013). "United Nations peacekeeping and civilian protection in civil war." *American Journal of Political Science*, **57**(4), 875–891.

Randahl D, Vegelius J (2023). "Inference with Extremes: Accounting for Extreme Values in Count Regression Models." *International Studies Quarterly*, **x**(x), x.

`lr_test` *Likelihood ratio test for individual variables of evzinb*

Description

Likelihood ratio test for individual variables of evzinb

Usage

```
lr_test(
  object,
  vars,
  single = TRUE,
  bootstrap = FALSE,
  multicore = FALSE,
  ncores = NULL,
  verbose = FALSE
)
```

Arguments

object	EVZINB or EVZINB object to perform likelihood ratio test on
vars	Either a list of character vectors with variable names which to be restricted in the LR test or a character vector of variable names. If a list, each character vector of the list will be run separately, allowing for multiple variables to be restricted as once. If a character vector, parameter 'single' can be used to determine whether all variables in the vector should be restricted at once (single = FALSE) or if the variables should be restricted one by one (single = TRUE)
single	Logical. Determining whether variables in 'vars' should be restricted individually (single = TRUE) or all at once (single = FALSE)
bootstrap	Should LR tests be conducted on each bootstrapped sample or only on the original sample.
multicore	Logical. Should the function be run in parallel?
ncores	Number of cores to use if multicore = TRUE
verbose	Logical. Should the function be verbose?

Value

A tibble with one row per performed LR test

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
 lr_test(model,'x1')
```

---

oob_evaluation *Out of bag predictive performance of EVZINB and EVINB models*

Description

Out of bag predictive performance of EVZINB and EVINB models

Usage

```
oob_evaluation(
  object,
  predict_type = c("harmonic", "explog"),
  metric = c("rmsle", "rmse", "mse", "mae")
)
```

Arguments

object	A fitted evzinb or evinb with bootstraps on which to conduct out-of-bag evaluation
predict_type	What type of prediction should be made? Harmonic mean, or exp(log(prediction))?
metric	What metric should be used for the out of bag evaluation? Default options include rmsle, rmse, mse, and mae. Can also take a user supplied function of the form function(y_pred,y_true) which returns a single value

Value

A vector of oob evaluation metrics of the length of the number of bootstraps in the evzinb/evinb object.

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
oob_evaluation(model)
```

---

predict.evinb *Predictions from evinb object*

Description

Predictions from evinb object

Usage

```
## S3 method for class 'evinb'
predict(
  object,
  newdata = NULL,
  type = c("harmonic", "explog", "counts", "pareto_alpha", "evinf", "count_state",
    "states", "all", "quantile"),
  pred = c("original", "bootstrap_median", "bootstrap_mean"),
  quantile = NULL,
  confint = FALSE,
  conf_level = 0.9,
  multicore = FALSE,
  ncores = NULL,
  ...
)
```

Arguments

object	An evinb object for which to produce predicted values
newdata	Optional new data (tibble) to produce predicted values from
type	Character string, 'harmonic' for the harmonic mean and 'explog' for exponentiated expected log, 'counts' for predicted count of the negative binomial component, 'pareto_alpha' for the predicted pareto alpha value, 'states' for the predicted component states (prior), 'count_state' for predicted probability of the count state, 'evinf' for predicted probability of the pareto state, 'all' for all predicted values, and 'quantile' for quantile prediction.
pred	Type of prediction to be used, defaults to the original prediction from the fitted model, with alternatives being the bootstrapped median or mean. Note that bootstrap mean may yield infinite values, especially when doing quantile prediction
quantile	Quantile for which to produce quantile prediction
confint	Should confidence intervals be made for the predictions? Note: only available for vector type predictions and not 'states' and 'all'.
conf_level	What confidence level should be used for confidence intervals
multicore	Should multicore be used when calculating quantile prediction? Often it is enough to run quantile prediction on a single core, but in cases of large data or very skewed distributions it may be useful to run multicore
ncores	Number of cores to be used for multicore.
...	Other arguments passed to predict function

Value

A vector of predicted values for type 'harmonic', 'explog', 'counts', 'pareto_alpha','evinf', 'count_state', and 'quantile' or a tibble of predicted values for type 'states' and 'all' or if confint=T

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
predict(model)
predict(model, type='all') # Getting all of the available predicted values
```

---

predict.evzinb *Predictions from evzinb object*

Description

Predictions from evzinb object

Usage

```
## S3 method for class 'evzinb'
predict(
  object,
  newdata = NULL,
 type = c("harmonic", "explog", "counts", "pareto_alpha", "zi", "evinf", "count_state",
    "states", "all", "quantile"),
  pred = c("original", "bootstrap_median", "bootstrap_mean"),
  quantile = NULL,
  confint = FALSE,
  conf_level = 0.9,
  multicore = FALSE,
  ncores = NULL,
  ...
)
```

Arguments

object	An evzinb object for which to produce predicted values
newdata	Optional new data (tibble) to produce predicted values from
type	Character string, 'harmonic' for the harmonic mean and 'explog' for exponentiated expected log, 'counts' for predicted count of the negative binomial component, 'pareto_alpha' for the predicted pareto alpha value, 'states' for the predicted component states (prior), 'count_state' for predicted probability of the count state, 'evinf' for predicted probability of the pareto state,'zi' for the predicted probability of the zero state, 'all' for all predicted values, and 'quantile' for quantile prediction.
pred	Type of prediction to be used, defaults to the original prediction from the fitted model, with alternatives being the bootstrapped median or mean. Note that bootstrap mean may yield infinite values, especially when doing quantile prediction
quantile	Quantile for which to produce quantile prediction
confint	Should confidence intervals be made for the predictions? Note: only available for vector type predictions and not 'states' and 'all'.
conf_level	What confidence level should be used for confidence intervals
multicore	Should multicore be used when calculating quantile prediction? Often it is enough to run quantile prediction on a single core, but in cases of large data or very skewed distributions it may be useful to run multicore
ncores	Number of cores to be used for multicore.
...	Other arguments passed to predict function

Value

A vector of predicted values for type 'harmonic', 'explog', 'counts', 'pareto_alpha','zi','evinf', 'count_state', and 'quantile' or a tibble of predicted values for type 'states' and 'all' or if confint=T

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
predict(model)
predict(model, type='all') # Getting all of the available predicted values
```

`predict.zinbboot` *Prediction for zinbboot*

Description

Prediction for zinbboot

Usage

```
## S3 method for class 'zinbboot'
predict(
  object,
  newdata = NULL,
  type = c("predicted", "counts", "zi", "evinf", "count_state", "states", "all",
    "quantile"),
  pred = c("original", "bootstrap_median", "bootstrap_mean"),
  quantile = NULL,
  confint = FALSE,
  conf_level = 0.9,
  ...
)
```

Arguments

`object`	a fitted zinbboot object
`newdata`	Data to make predictions on
`type`	What prediction should be computed?
`pred`	Prediction type, 'original', 'bootstra_median', or 'bootstrap_mean'
`quantile`	Quantile for quantile prediction
`confint`	Should confidence intervals be created?
`conf_level`	Confidence level when predicting with CIs
`...`	Not used

Value

Predictions from zinbboot

print.evinb *EVINB print function*

Description

EVINB print function

Usage

```
## S3 method for class 'evinb'
print(x, ...)
```

Arguments

x	A fitted evinb model
...	Not used

Value

An evinb print function

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
print(model)
```

print.evzinb *EVZINB print function*

Description

EVZINB print function

Usage

```
## S3 method for class 'evzinb'
print(x, ...)
```

Arguments

x	A fitted evzinb model
...	Not used

Value

An evzinb print function

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
print(model)
```

`print.evzinbcomp` *Compare_models print function*

Description

Compare_models print function

Usage

```
## S3 method for class 'evzinbcomp'
print(x, ...)
```

Arguments

`x`	A fitted evinb model
`...`	Not used

Value

An evinb print function

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10)
print(model)
```

revinb_fit *Random draws from a fitted evinb model*

Description

Random draws from a fitted evinb model

Usage

```
revinb_fit(object, newdata = NULL, n_draws = 1)
```

Arguments

object	A fitted EVINB object
newdata	Optional newdata
n_draws	Number of random draws to make

Value

A vector of randomly drawn values from the fitted evinb if n_draws == 1, or a list of length n_draws with random drawn values if n_draws > 1

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3, data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
revinb_fit(model)
```

revzinb_fit *Random draws from a fitted evzinb model*

Description

Random draws from a fitted evzinb model

Usage

```
revzinb_fit(object, newdata = NULL, n_draws = 1)
```

Arguments

object	A fitted EVZINB object
newdata	Optional newdata
n_draws	Number of random draws to make

Value

A vector of randomly drawn values from the fitted evzinb if n_draws == 1, or a list of length n_draws with random drawn values if n_draws > 1

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3, data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
revzinb_fit(model)
```

`summary.evinb` *EVINB summary function*

Description

EVINB summary function

Usage

```
## S3 method for class 'evinb'
summary(
  object,
  coef = c("original", "bootstrapped_mean", "bootstrapped_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "both", "none"),
  bootstrapped_props = c("none", "mean", "median"),
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  ...
)
```

Arguments

`object`	an EVINB object with bootstraps
`coef`	Type of coefficients. Original are the coefficient estimates from the non-bootstrapped version of the model. 'bootstrapped_mean' are the mean coefficients across bootstraps, and 'bootstrapped_median' are the median coefficients across bootstraps
`standard_error`	Should standard errors be computed?
`p_value`	What type of p_values should be computed? 'bootstrapped' are bootstrapped p_values through confidence interval inversion. 'approx' are p-values based on the t-value produced by dividing the coefficient with the standard error.
`bootstrapped_props`	Type of bootstrapped proportions of component proportions to be returned
`approx_t_value`	Should approximate t-values be returned

symmetric_bootstrap_p
: Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.

...
: Additional arguments passed to the summary function

Value

An EVINB summary object

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
summary(model)
```

---

summary.evzinb *EVZINB summary function*

Description

EVZINB summary function

Usage

```
## S3 method for class 'evzinb'
summary(
  object,
  coef = c("original", "bootstrapped_mean", "bootstrapped_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "both", "none"),
  bootstrapped_props = c("none", "mean", "median"),
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  ...
)
```

Arguments

object
: an EVZINB object with bootstraps

coef
: Type of coefficients. Original are the coefficient estimates from the non-bootstrapped version of the model. 'bootstrapped_mean' are the mean coefficients across bootstraps, and 'bootstrapped_median' are the median coefficients across bootstraps

standard_error
: Should standard errors be computed?

p_value	What type of p_values should be computed? 'bootstrapped' are bootstrapped p_values through confidence interval inversion. 'approx' are p-values based on the t-value produced by dividing the coefficient with the standard error.
bootstrapped_props	Type of bootstrapped proportions of component proportions to be returned
approx_t_value	Should approximate t-values be returned
symmetric_bootstrap_p	Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.
...	Additional arguments passed to the summary function

Value

An EVZINB summary object

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
summary(model)
```

---

tidy.evinb *EVINB tidy function*

Description

EVINB tidy function

Usage

```
## S3 method for class 'evinb'
tidy(
  x,
  component = c("evi", "count", "pareto", "all"),
  coef_type = c("original", "bootstrap_mean", "bootstrap_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "none"),
  confint = c("none", "bootstrapped", "approx"),
  conf_level = 0.95,
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  ...
)
```

Arguments

x	An evinb object
component	Which component should be shown?
coef_type	Type of coefficients. Original are the coefficient estimates from the non-bootstrapped version of the model. 'bootstrapped_mean' are the mean coefficients across bootstraps, and 'bootstrapped_median' are the median coefficients across bootstraps
standard_error	Should standard errors be computed?
p_value	What type of p_values should be computed? 'bootstrapped' are bootstrapped p_values through confidence interval inversion. 'approx' are p-values based on the t-value produced by dividing the coefficient with the standard error.
confint	What type of confidence should be computed. Same options as p_value
conf_level	What confidence level should be used for the confidence interval
approx_t_value	Should approximate t-values be returned
symmetric_bootstrap_p	Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.
...	Other arguments passsed to tidy function

Value

An EVINB tidy function

Examples

```
data(genevzinb2)
model <- evinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
tidy(model)
```

---

tidy.evzinb *EVZINB tidy function*

Description

EVZINB tidy function

Usage

```
## S3 method for class 'evzinb'
tidy(
  x,
  component = c("zi", "evi", "count", "pareto", "all"),
  coef_type = c("original", "bootstrap_mean", "bootstrap_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "none"),
  confint = c("none", "bootstrapped", "approx"),
  conf_level = 0.95,
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  ...
)
```

Arguments

x	An evzinb object
component	Which component should be shown?
coef_type	Type of coefficients. Original are the coefficient estimates from the non-bootstrapped version of the model. 'bootstrapped_mean' are the mean coefficients across bootstraps, and 'bootstrapped_median' are the median coefficients across bootstraps
standard_error	Should standard errors be computed?
p_value	What type of p_values should be computed? 'bootstrapped' are bootstrapped p_values through confidence interval inversion. 'approx' are p-values based on the t-value produced by dividing the coefficient with the standard error.
confint	What type of confidence should be computed. Same options as p_value
conf_level	What confidence level should be used for the confidence interval
approx_t_value	Should approximate t-values be returned
symmetric_bootstrap_p	Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.
...	Other arguments passsed to tidy function

Value

An EVZINB tidy function

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
tidy(model)
```

`tidy.nbboot` *Tidy function for nbboot*

Description

Tidy function for nbboot

Usage

```
## S3 method for class 'nbboot'
tidy(
  x,
  coef_type = c("original", "bootstrap_mean", "bootstrap_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "none"),
  confint = c("none", "bootstrapped", "approx"),
  conf_level = 0.95,
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  include_ylev = FALSE,
  ...
)
```

Arguments

`x`	A fitted bootstrapped zero-inflated model
`coef_type`	What type of coefficient should be reported, original, bootstrapped mean, or bootstrapped median
`standard_error`	Should bootstrapped standard errors be reported?
`p_value`	What type of p-value should be reported? Bootstrapped p_values, approximate p-values, or none?
`confint`	What type of confidence intervals should be reported? Bootstrapped p_values, approximate p-values, or none?
`conf_level`	Confidence level for confidence intervals
`approx_t_value`	Should approximate t_values be reported
`symmetric_bootstrap_p`	Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.
`include_ylev`	Logical. Should y.lev be included in the tidy output? Makes for nicer tables when using modelsummary
`...`	Other arguments to be passed to tidy

Value

A tidy function for a bootstrapped nb model

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
zinb_comp <- compare_models(model)
tidy(zinb_comp$nb)
```

---

tidy.zinbboot *Tidy function for zinbboot*

Description

Tidy function for zinbboot

Usage

```
## S3 method for class 'zinbboot'
tidy(
  x,
  component = c("zi", "count", "all"),
  coef_type = c("original", "bootstrap_mean", "bootstrap_median"),
  standard_error = TRUE,
  p_value = c("bootstrapped", "approx", "none"),
  confint = c("none", "bootstrapped", "approx"),
  conf_level = 0.95,
  approx_t_value = TRUE,
  symmetric_bootstrap_p = TRUE,
  ...
)
```

Arguments

x	A fitted bootstrapped zero-inflated model
component	Which component should be shown?
coef_type	What type of coefficient should be reported, original, bootstrapped mean, or bootstrapped median
standard_error	Should bootstrapped standard errors be reported?
p_value	What type of p-value should be reported? Bootstrapped p_values, approximate p-values, or none?
confint	What type of confidence intervals should be reported? Bootstrapped p_values, approximate p-values, or none?

conf_level	Confidence level for confidence intervals
approx_t_value	Should approximate t_values be reported
symmetric_bootstrap_p	Should bootstrap p-values be computed as symmetric (leaving alpha/2 percent in each tail)? FALSE gives non-symmetric, but narrower, intervals. TRUE corresponds most closely to conventional p-values.
...	Other arguments to be passed to tidy

Value

A tidy function for a bootstrapped zinb model

Examples

```
data(genevzinb2)
model <- evzinb(y~x1+x2+x3,data=genevzinb2, n_bootstraps = 10, multicore = TRUE, ncores = 2)
zinb_comp <- compare_models(model)
tidy(zinb_comp$zinb)
```

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