

# Package ‘flipscores’

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**Title** Robust Score Testing in GLMs, by Sign-Flip Contributions

**Description** Provides robust tests for testing in GLMs, by sign-flipping score contributions. The tests are robust against overdispersion, heteroscedasticity and, in some cases, ignored nuisance variables. See Hemerik, Goeman and Finos (2020) <[doi:10.1111/rssb.12369](https://doi.org/10.1111/rssb.12369)>.

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anova.flipscores	<i>anova.flipscores</i>
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### Description

This is the anova method for flipscores object. Remark: it performs type III deviance decomposition as in `car::Anova`.

### Usage

```
## S3 method for class 'flipscores'
anova(object, model1 = NULL, score_type = NULL, n_flips = 5000, id = NULL, ...)
```

### Arguments

object	(the object) <code>glm</code> (or <code>flipscores</code> ) object with the model under the null hypothesis (i.e. the covariates, the nuisance parameters).
model1	a <code>glm</code> (or <code>flipscores</code> ) or a matrix (or vector). If it is a <code>glm</code> object, it has the model under the alternative hypothesis. The variables in <code>model1</code> are the same variables in <code>object</code> plus one or more variables to be tested. Alternatively, if <code>model1</code> is a matrix, it contains the tested variables column-wise.
score_type	The type of score that is computed. It can be "orthogonalized", "effective" or "basic". Default is "orthogonalized". "effective" and "orthogonalized" take into account the nuisance estimation. The default is <code>NULL</code> , in this case the value is taken from <code>object</code> .
n_flips	The number of random flips of the score contributions. When <code>n_flips</code> is equal or larger than the maximum number of possible flips (i.e. $n^2$ ), all possible flips are performed. Default is 5000.
id	a vector identifying the clustered observations. If <code>NULL</code> (default) observations are assumed to be independent. NOTE: if <code>object</code> is a <code>flipscores</code> and <code>model1\$flip_param_call\$id</code> is not <code>NULL</code> , this is considered in the inference.
...	other parameters allowed in <code>stats::anova</code> .

### Examples

```
set.seed(1)
dt=data.frame(X=scale(rnorm(50)),
              Z=factor(rep(LETTERS[1:3],length.out=50)))
dt$Y=rpois(n=nrow(dt),lambda=exp(dt$X*(dt$Z=="C")))
mod0=flipscores(Y~Z+X,data=dt,family="poisson")
summary(mod0)
anova(mod0)

mod1=flipscores(Y~Z*X,data=dt,family="poisson")
summary(mod1)
anova(mod0,model1 = mod1)
```

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compute_scores	<i>compute_scores</i>
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### Description

Same usage as `anova.glm`. The parameter `id` is used too, if present in `model0` (with priority) or in `model1`.

### Usage

```
compute_scores(model0, model1, score_type = "standardized", ...)
```

### Arguments

<code>model0</code>	a <code>glm</code> object with the model under the null hypothesis (i.e. the covariates, the nuisance parameters).
<code>model1</code>	a <code>glm</code> or a matrix (or vector). If it is a <code>glm</code> object, it has the model under the alternative hypothesis. The variables in <code>model1</code> are the same variables in <code>model0</code> plus one or more variables to be tested. Alternatively, if <code>model1</code> is a matrix, it contains the tested variables column-wise.
<code>score_type</code>	The type of score that is computed. It is "orthogonalized", "effective" or "basic". "effective" and "orthogonalized" take into account the nuisance estimation.
<code>...</code>	other arguments.

### Author(s)

Jesse Hemerik, Riccardo De Santis, Vittorio Giatti, Jelle Goeman and Livio Finos

### Examples

```
set.seed(1)
Z=rnorm(20)
X=Z+rnorm(20)
Y=rpois(n=20,lambda=exp(Z+X))
mod0=glm(Y~Z,family="poisson")
X=data.frame(X=X)
scr0=compute_scores(model0 = mod0, model1 = X)
head(scr0)
```

flipscores

*Robust testing in GLMs, by sign-flipping score contributions***Description**

Provides robust tests for testing in GLMs, by sign-flipping score contributions. The tests are often robust against overdispersion, heteroscedasticity and, in some cases, ignored nuisance variables.

**Usage**

```
flipscores(formula, family, data, score_type = "standardized",
n_flips = 5000, alternative = "two.sided", id = NULL,
seed = NULL, to_be_tested = NULL, flips = NULL,
precompute_flips = TRUE, output_flips = FALSE, ...)
```

**Arguments**

formula	see glm function. It can also be a model (usually generated by a call to glm); in this case, any other glm-related parameter (e.g. family, data, etc.) are discarded, the function will make use of the ones used to generate the model. (i.e. formula, family, data, etc) are not considered. It is NULL by default (i.e. not used).
family	see glm function.
data	see glm function.
score_type	The type of score that is computed. It can be "standardized" "orthogonalized", "effective" or "basic". Both "orthogonalized" and "effective" take into account the nuisance estimation and they provide the same test statistic. In case of small samples "effective score" might have a slight anti-conservative behaviour. "standardized effective score" gives a solution for this issue. "orthogonalized" has a similar intent, note however that in case of a big model matrix, it may be slow.
n_flips	The number of random flips of the score contributions. Overwritten with the nrow(flips) when flips is not NULL (see parameter flips for more details). When n_flips is equal or larger than the maximum number of possible flips (i.e. $n^2$ ), all possible flips are performed.
alternative	It can be "greater", "less" or "two.sided" (default)
id	a vector identifying the clustered observations. If NULL (default) observations are assumed to be independent. If id is not NULL, only score_type=="effective" is allowed, yet.
seed	NULL by default.
to_be_tested	vector of indices or names of coefficients of the glm model to be tested (it is faster than computing every scores and p-values of course).
flips	matrix fo +1 or -1, the matrix has n_flips rows and n (number of observations) columns

`precompute_flips` TRUE by default. Overwritten if `flips` is not NULL. If FALSE the matrix of flips is not computed and the flips are made 'on-the-fly' before computing the test statistics; it may be usefull when `flips` is very large (see parameter `flips` for more details).

`output_flips` FALSE by default. If TRUE the flips matrix is returned. Useful when the same flips are needed for more glms, for example in the case of multivariate glms where the joint distribution of test statistis if used for multivariate inference.

... see `glm` function.

### Details

`flipscores` borrows the same parameters from function `glm` (and `glm.nb`). See these helps for more details about parameters such as `formula`, `data`, `family`. Note: in order to use Negative Binomial family, `family` reference must have quotes (i.e. `family="negbinom"`). Furthermore, `flipscores` object contains two extra elements: `scores` – i.e. a matrix of `n` score contributions, one column for each tested coefficient – and `Tspace` – i.e. a matrix of size `n_flips` times `ncol(scores)`. The fist row of `Tspace` contains column-wise the test statistics generated by randomly flipping the score contributions, each column refers to the same column of scores, the vector of observed test statistics (i.e. no flips) is in the first row of `Tspace`.

### Value

an object of class `flipscores`. See also its methods (`summary.flipscores`, `anova.flipscores`, `print.flipscores`).

### Author(s)

Livio Finos, Riccardo De Santis, Jesse Hemerik and Jelle Goeman

### References

"Robust testing in generalized linear models by sign-flipping score contributions" by J.Hemerik, J.Goeman and L.Finos.

### See Also

[anova.flipscores](#), [summary.flipscores](#), [flip](#)

### Examples

```
set.seed(1)
dt=data.frame(X=rnorm(20),
  Z=factor(rep(LETTERS[1:3],length.out=20)))
dt$Y=rpois(n=20,lambda=exp(dt$Z=="C"))
mod=flipscores(Y~Z+X,data=dt,family="poisson",n_flips=1000)
summary(mod)

# Equivalent to:
model=glm(Y~Z+X,data=dt,family="poisson")
```

```
mod2=flipscores(model)
summary(mod2)
```

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flipscores-method      *Methods for flipscores objects*

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### Description

Methods for flipscores objects. The following are methods to extract and manipulate relevant information from a flipscores object.

### Usage

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

## S3 method for class 'flipscores'
summary(object, ...)
```

### Arguments

x	a flipscores object
...	additional arguments to be passed
object	a flipscores object

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make_flips	<i>It creates a n_flipsxn_obs matrix of random +1 and -1. The first row is made by ones (i.e. the observed test statistic is computed)</i>
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### Description

It creates a n\_flipsxn\_obs matrix of random +1 and -1. The first row is made by ones (i.e. the observed test statistic is computed)

### Usage

```
make_flips(n_obs, n_flips)
```

### Arguments

n_obs	number of observations
n_flips	number of flips

*make\_flips*

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### **Examples**

```
# example code  
make_flips(n_obs=10, n_flips=20)
```

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