

Package ‘influence.ME’

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Title Tools for detecting influential data in mixed effects models

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Description influence.ME provides a collection of tools for calculating measures of influential data for mixed effects models. It analyses models that were estimated using lme4. The basic rationale behind identifying influential data is that when iteratively single units are omitted from the data, models based on these data should not produce substantially different estimates. To standardize the assessment of how influential a (single group of) observation(s) is, several measures of influence are common practice. First, DFBETAS is a standardized measure of the absolute difference between the estimate with a particular case included and the estimate without that particular case. Second, Cook’s distance provides an overall measurement of the change in all parameter estimates, or a selection thereof.

License GPL-3

URL <http://www.rensenieuwenhuis.nl/r-project/influenceme/>

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influence.ME-package

Influence.ME: Tools for detecting influential data in mixed effects models

Description

influence.ME calculates measures of influence for mixed effects models estimated with lme4. The basic rationale behind measuring influential cases is that when iteratively single units are omitted from the data, models based on these data should not produce substantially different estimates. To standardize the assessment of how influential a (single group of) observation(s) is, several measures of influence are common practice. First, DFBETAS is a standardized measure of the absolute difference between the estimate with a particular case included and the estimate without that particular case. Second, Cook's distance provides an overall measurement of the change in all parameter estimates, or a selection thereof.

Details

| | |
|-----------|--------------|
| Package: | influence.ME |
| Type: | Package |
| Version: | 0.7 |
| Date: | 2009-07-06 |
| License: | GPL-3 |
| LazyLoad: | yes |

Calculating measures of influential data on a mixed effects regression model entails the re-estimation of this model for each set of potentially influential data separately. The estex() function does this, and returns the altered estimates resulting from each re-estimation. These altered estimates can subsequently be entered to the ME.cook() and ME.dfbetas() functions, to calculate Cook's Distance and the DFBETAS (standardized difference of the beta) measures.

Author(s)

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References

Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.

Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.

Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the American Sociological Review.

See Also

[estex](#), [exclude.influence](#), [ME.cook](#), [ME.dfbetas](#), [ME.pchange](#)

Examples

```
data(school23)
model.a <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)
alt.est.a <- estex(model.a, "school.ID")
ME.cook(alt.est.a, plot=TRUE, tol=.17)

model.b <- exclude.influence(model.a, "school.ID", "7472")
alt.est.b <- estex(model.b, "school.ID")
ME.cook(alt.est.b, plot=TRUE, tol=.18)

model.c <- exclude.influence(model.b, "school.ID", "54344")
alt.est.c <- estex(model.c, "school.ID")
ME.cook(alt.est.c, plot=TRUE, tol=.19)
```

dp.ME.cook

Dotplot visualization of Cook's Distance

Description

This is a wrapper function to the `dotplot()` function in the `lattice`-package. It transforms the output from the `ME.cook()` function and calls `dotplot()` to provide the user with a visualization of the Cook's Distances.

Usage

```
dp.ME.cook(estex, parameters = 0, groups = 0, cutoff = 0, sort=FALSE, ...)
```

Arguments

| | |
|-------------------------|---|
| <code>estex</code> | An object as returned by the <code>estex()</code> function, containing the altered estimates of a mixed effects regression model. |
| <code>parameters</code> | Used to define a selection of parameters. If <code>parameters = 0</code> (default), Cook's Distance is calculated based on all parameters in the model. |

| | |
|--------|---|
| groups | Used to define a selection of nesting groups that should be visualized. If groups = 0 (default), the values of Cook's Distance for all nesting groups are shown. |
| cutoff | Values of Cook's Distance exceeding the specified (cutoff) value are plotted visually different from values not exceeding the cutoff. If cutoff=0 (default), no such differentiation is made in the way values are plotted. |
| sort | If sort=TRUE the values of Cook's Distance are ordered based on magnitude before visualization. If sort=FALSE (default) no sorting takes place. |
| ... | Further arguments passed on to the dotplot() function. |

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

See Also

[estex](#), [ME.cook](#), [dp.ME.dfbetas](#), [dp.ME.pchange](#)

Examples

```
data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
dp.ME.cook(alt.est)
dp.ME.cook(alt.est, cutoff=.17)
```

dp.ME.dfbetas

Dotplot visualization of DFBETAS

Description

This is a wrapper function to the dotplot() function in the lattice-package. It transforms the output from the ME.dfbetas() function and calls dotplot() to provide the user with a visualization of the DFBETAS.

Usage

```
dp.ME.dfbetas(estex, parameters = 0, groups = 0, sort=FALSE, to.sort=NA, abs=FALSE,
```

Arguments

| | |
|------------|---|
| estex | An object as returned by the estex() function, containing the altered estimates of a mixed effects regression model. |
| parameters | Used to define a selection of parameters. If parameters = 0 (default), values for DFBETAS are visualized for parameters in the model. |

| | |
|---------|---|
| groups | Used to define a selection of nesting groups that should be visualized. If groups = 0 (default), the values of DFBETAS for all nesting groups are shown. |
| sort | If sort=TRUE the values of DFBETAS are ordered based on magnitude before visualization. If sort=FALSE (default) no sorting takes place. |
| to.sort | Specify on which variable the DFBETAS must be sorted. If only one variable present (either in the model, or due to the selection specified in parameters), this parameter can be omitted. If multiple variables are visualized, and sort=TRUE, specification of to.sort is required, or an error is returned. |
| abs | If abs=TRUE, the absolute values of DFBETAS are visualized, while if abs=FALSE (default), both positive and negative values are possible. If both abs=TRUE and sort=TRUE, the abs parameters precedes the sort parameter, and thus the absolute values of DFBETAS are sorted. |
| ... | Further arguments passed on to the dotplot() function. |

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

See Also

[estex](#), [ME.dfbetas](#), [dp.ME.cook](#)

Examples

```
data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
dp.ME.dfbetas(alt.est)
dp.ME.dfbetas(alt.est, layout=c(1,3))
```

dp.ME.pchange *Dotplot visualization of percentile change as measure of influential data*

Description

This is a wrapper function to the dotplot() function in the lattice-package. It transforms the output from the ME.pchange() function and calls dotplot() to provide the user with a visualization of the percentile changes in mixed models due to the neutralization of the influence of grouping factors.

Usage

```
dp.ME.pchange(estex, parameters = 0, groups = 0, sort=FALSE, to.sort=NA, abs=FALSE,
```

Arguments

| | |
|-------------------------|---|
| <code>estex</code> | An object as returned by the <code>estex()</code> function, containing the altered estimates of a mixed effects regression model. |
| <code>parameters</code> | Used to define a selection of parameters. If <code>parameters = 0</code> (default), values for percentile change are visualized for parameters in the model. |
| <code>groups</code> | Used to define a selection of nesting groups that should be visualized. If <code>groups = 0</code> (default), the percentile changes for all nesting groups are shown. |
| <code>sort</code> | If <code>sort=TRUE</code> the values of percentile change are ordered based on magnitude before visualization. If <code>sort=FALSE</code> (default) no sorting takes place. |
| <code>to.sort</code> | Specify on which variable the percentile changes must be sorted. If only one variable present (either in the model, or due to the selection specified in <code>parameters</code>), this parameter can be omitted. If multiple variables are visualized, and <code>sort=TRUE</code> , specification of <code>to.sort</code> is required, or an error is returned. |
| <code>abs</code> | If <code>abs=TRUE</code> , the absolute values of percentile change are visualized, while if <code>abs=FALSE</code> (default), both positive and negative percentages are possible. If both <code>abs=TRUE</code> and <code>sort=TRUE</code> , the <code>abs</code> parameter precedes the <code>sort</code> parameter, and thus the absolute values of percentile change are sorted. |
| <code>...</code> | Further arguments passed on to the <code>dotplot()</code> function. |

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

See Also

[estex](#), [ME.pchange](#)

Examples

```
data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
dp.ME.pchange(alt.est)
dp.ME.pchange(alt.est, layout=c(1,3))
```

`estex`

Estex returns mixed model estimates, iteratively excluding the influence of data nested within single grouping factors.

Description

`estex()` is the workhorse function of the `influence.ME` package. Based on a priorly estimated mixed effects regression model (estimated using `lme4`), the `estex()` function iteratively modifies the mixed effects model to neutralize the effect a grouped set of data has on the parameters, and which returns returns the fixed parameters of these iteratively modified models. These are used to compute measures of influential data.

Usage

```
estex(model, group, select = 0, gf="single", count = FALSE, delete=FALSE, ...)
```

Arguments

| | |
|--------|---|
| model | Mixed effects model of class 'mer' |
| group | Grouping factor in model of which iteratively levels are neutralized |
| select | Defines the selection of grouping factors that should be omitted. Defaults to 0, resulting in each level of the grouping factor being omitted iteratively. When a selection is defined, model parameters for the full model, and the altered model are returned. The selection can be a vector of multiple levels of the grouping factor. |
| gf | Indicates from which of the model's grouping factors the influence of the specified grouping factor is to be neutralized. If <code>gf="single"</code> (default), the levels of the specified grouping factor are only neutralized regarding the grouping factor specified in <code>group</code> . In its present form, <code>gf="single"</code> only works on mixed models with a maximum of 2 grouping factors. If <code>gf="all"</code> , the influence from the levels of <code>group</code> is neutralized regarding all grouping factors in the model. This option only applies to models with more than a single grouping factor. |
| count | If <code>count=TRUE</code> , the remaining number of grouping factors that still need to be omitted are printed. |
| delete | If <code>delete=FALSE</code> (default), the influence of higher level groups is excluded from the model by setting the intercept-vector for the observations nested within these groups to 0, and by adding a dummy-variable indicating these observations (Langford and Lewis, 1998). If <code>delete=TRUE</code> , the influence is excluded by simply deleting the observations nested within the higher level group. |
| ... | Optional arguments that are passed on to the <code>lmer/glmer</code> function |

Details

The basic rationale behind measuring influential cases is that when iteratively single units are omitted from the data, models based on these data should not produce substantially different estimates. To apply this logic to mixed effects models one has to measure the influence of a particular higher level unit on the estimates of a higher level predictor. This means that the mixed effects model has to be adjusted to neutralize the unit's influence on that estimate, while at the same time allowing the unit's lower-level cases to help estimate the effects of the lower-level predictors in the model. This procedure is based on a modification of the intercept and the addition of a dummy variable for the cases that might be influential.

`estex()` is the workhorse function of this likewise called package. Based on a priorly estimated mixed effects regression model (of the 'mer' class), the `estex()` function iteratively modifies the mixed effects model by neutralizing the effect a grouped set of data has on the parameters, and which returns returns the fixed parameters of these iteratively modified models.

The returned object (see 'value') contains information which is required for functions computing various measures of influential data.

Value

The object returned by `estex()` of class "alt.est" contains the 'altered estimates' required by several other functions to calculate measures of influential data. A list containing six elements is returned:

| | |
|------------------------|--|
| <code>or.fixed</code> | Fixed estimates of the original model (based on the full data) |
| <code>or.se</code> | Standard Error of the estimates of the original model |
| <code>or.vcov</code> | Variance / Covariance matrix of the original model |
| <code>alt.fixed</code> | Matrix of the fixed parameters estimate, after iteratively subsets of data are removed. Altered estimates associated with the deletion of data nested within each grouping factor are provided. |
| <code>alt.se</code> | Matrix of the standard errors of the fixed parameter estimates, after iteratively subsets of data are removed. Altered estimates associated with the deletion of data nested within each grouping factor are provided. |
| <code>alt.vcov</code> | Variance / Covariance matrix of the altered models, after iteratively subsets of data are removed. Altered estimates associated with the deletion of data nested within each grouping factor are provided. |

Note

Please note that in its present form, the `estex` function only works on mixed effects regression models of class `mer` that have been estimated using the functions in the `lme4` package.

Also, it is required that the `mer` model was estimated using a factor variable to indicate group levels. When using something similar to `+ (1 | as.factor(variable))`, the function is not able of identifying the correct grouping factors, and returns an error.

Since `estex()` entails the re-estimation of the provided mixed effects model for each level of the specified grouping factor (after alteration of the data), executing this procedure can be computationally highly demanding.

To facilitate the use of `estex()` with more complex models (i.e. models based on large amounts of data and / or with a large number of groups).

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

References

- Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.
- Langford, I. H. and Lewis, T. (1998). Outliers in multilevel data. *Journal of the Royal Statistical Society: Series A (Statistics in Society)*, 161:121-160.
- Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.
- Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the *American Sociological Review*.

See Also

[ME.dfbetas](#), [ME.cook](#)

Examples

```
data(school23)
model.a <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)
alt.est.a <- estex(model.a, "school.ID")
alt.est.b <- estex(model.a, "school.ID", "7472")
alt.est.c <- estex(model.a, "school.ID", c("7472", "62821"))

data(Penicillin)
model.b <- lmer(diameter ~ (1|plate) + (1|sample), Penicillin)
alt.est.d <- estex(model.b, "plate")
alt.est.e <- estex(model.b, "sample")
alt.est.f <- estex(model.b, "sample", gf="all")
```

`exclude.influence` *Exclude the influence of a grouped set of observations in mixed effects models.*

Description

Using mixed effects regression models, `exclude.influence` excludes the influence of a group of cases grouped within a single grouping factor, or a set of grouping factors. The function returns a model in which the influence a grouped set of observations has on both the variance and point-estimate of the (random) intercept.

Usage

```
exclude.influence(model, grouping, level, gf="single", delete=FALSE)
```

Arguments

| | |
|-----------------------|--|
| <code>model</code> | A mixed effects regression model |
| <code>grouping</code> | The grouping factor of which one or more groupings levels are to be 'neutralized' |
| <code>level</code> | Vector of character strings, indicating either a single level or a set of grouping levels the influence of which is to be neutralized |
| <code>gf</code> | Indicates from which of the model's grouping factors the influence of the specified grouping factor is to be neutralized. If <code>gf="single"</code> (default), the levels of the specified grouping factor are only neutralized from the grouping factor specified in <code>group</code> . In its present form, <code>gf="single"</code> only works on mixed models with a maximum of 2 grouping factors. If <code>gf="all"</code> , the influence from the levels of <code>group</code> is neutralized regarding all grouping factors in the model. This option only applies to models with more than a single grouping factor. |

`delete` If `delete=FALSE` (default), the influence of higher level groups is excluded from the model by setting the intercept-vector for the observations nested within these groups to 0, and by adding a dummy-variable indicating these observations (Langford and Lewis, 1998). If `delete=TRUE`, the influence is excluded by simply deleting the observations nested within the higher level group.

Details

To apply the basic logic of influential cases to mixed effects models one has to measure the influence of a particular higher level unit on the estimates of a higher level predictor. This means that the mixed effects model has to be adjusted to neutralize the unit's influence on that estimate, while at the same time allowing the unit's lower-level cases to help estimate the effects of the lower-level predictors in the model. This procedure is based on a modification of the intercept and the addition of a dummy variable for the cases that might be influential.

The model that is returned by `exclude.influence` thus contains a modified intercept, and one or more additional dummy variables. To help identify this model as modified (which is required when in a later stage the influence of additional grouping levels is excluded), the intercept is renamed to 'intercept.alt'. The additional dummy variables, indicating the observations associated with the grouping factor levels of which the influence was neutralized, are labeled starting with 'estex.', combined with the label of the neutralized grouping level.

Value

Mixed effects regression model of class 'mer', with a modified random intercept and dummy variables indicating the estimates of the neutralized influence of selected grouping levels.

Note

Please note that in its present form, the `exclude.influence` function only works on mixed effects regression models of class `mer` that have been estimated using the functions in the `lme4` package.

Also, it is required that the `mer` model was estimated using a factor variable to indicate group levels. When using something similar to `+ (1 | as.factor(variable))`, the function is not able of identifying the correct grouping factors, and returns an error.

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

References

- Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.
- Langford, I. H. and Lewis, T. (1998). Outliers in multilevel data. *Journal of the Royal Statistical Society: Series A (Statistics in Society)*, 161:121-160.
- Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.

Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the American Sociological Review.

See Also

[estex](#)

Examples

```
data(school23)
model.a <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)
summary(model.a)
model.b <- exclude.influence(model.a, "school.ID", "7472")
summary(model.b)
model.c <- exclude.influence(model.a, "school.ID", c("7472", "62821"))
summary(model.c)

data(Penicillin)
model.d <- lmer(diameter ~ (1|plate) + (1|sample), Penicillin)
summary(model.d)
model.e <- exclude.influence(model.d, "sample", "A", gf="all")
summary(model.e)
```

| | |
|-----------------|--|
| grouping.levels | <i>Returns the levels of a grouping factor in a mixed effects regression model</i> |
|-----------------|--|

Description

Helper function returning all the levels of a grouping factor in a mixed effects regression model. This is used in combination with the influence.ME function, the divide long runs into multiple shorter ones.

Usage

```
grouping.levels(model, group)
```

Arguments

| | |
|-------|---|
| model | Mixed effects model of class 'mer' |
| group | Grouping factor of 'model' of which the levels are returned |

Details

Please note that at times different results may be obtained by using nesting.levels(), compared with deriving the levels of the grouping factor directly from the (original) data. This is because nesting.levels() only extracts the nesting levels that were de facto used in the model. Due to missing values, this may diverge from those present in the actual data.

Value

Returns a character vector containing all the names / labels of levels of the grouping factor.

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

Examples

```
# Penicillin data originates from the lme4 package.
model <- lmer(diameter ~ (1|plate) + (1|sample), Penicillin)

grouping.levels(model, "plate")
grouping.levels(model, "sample")
```

ME.cook

Compute the Cook's distance measure of influential data on mixed effects models

Description

Cook's Distance is a measure indicating to what extent model parameters are influenced by (a set of) influential data on which the model is based. This function computes the Cook's distance based on the information returned by the `estex()` function.

Usage

```
ME.cook(estex, parameters = 0, plot=FALSE, sort=FALSE, ...)
```

Arguments

| | |
|-------------------------|--|
| <code>estex</code> | An object as returned by the <code>estex()</code> function, containing the altered estimates of a mixed effects regression model |
| <code>parameters</code> | Used to define a selection of parameters. If <code>parameters=0</code> (default), Cook's Distance is calculated based on all parameters in the model |
| <code>plot</code> | If <code>plot=TRUE</code> , the results from the <code>ME.cook()</code> function are forwarded to the <code>dp.ME.cook()</code> function, which creates a visual representation of the Cook's Distances. |
| <code>sort</code> | If <code>sort=TRUE</code> the values of Cook's Distance are ordered based on magnitude. If <code>sort=FALSE</code> (default) no sorting takes place. |
| <code>...</code> | Further arguments passed on to the <code>dp.ME.cook()</code> function. |

Value

A one-column matrix is returned containing values for the Cook's Distance based on the selected (fixed) parameters of the model. Each row shows the Cook's Distance associated with each evaluated set of influential data (data nested within each evaluated level of the grouping factor).

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

References

Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.

Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.

Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the American Sociological Review.

See Also

[estex](#), [ME.dfbetas](#), [dp.ME.cook](#)

Examples

```
data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
ME.cook(alt.est)
ME.cook(alt.est, plot=TRUE, cutoff=.17)
```

ME.dfbetas

Compute the DFBETAS measure of influential data

Description

DFBETAS (standardized difference of the beta) is a measure that standardizes the absolute difference in parameter estimates between a (mixed effects) regression model based on a full set of data, and a model from which a (potentially influential) subset of data is removed. A value for DFBETAS is calculated for each parameter in the model separately. This function computes the DFBETAS based on the information returned by the `estex()` function.

Usage

```
ME.dfbetas(estex, parameters = 0, plot=FALSE, sort=FALSE, to.sort=NA, abs=FALSE, ..
```

Arguments

| | |
|-------------------------|---|
| <code>estex</code> | An object as returned by the <code>estex()</code> function, containing the altered estimates of a mixed effects regression model |
| <code>parameters</code> | Used to define a selection of parameters. If <code>parameters=0</code> (default), DFBETAS is calculated for all parameters in the model |
| <code>plot</code> | If <code>plot=TRUE</code> , the results from the <code>ME.dfbetas()</code> function are forwarded to the <code>dp.ME.dfbetas()</code> function, which creates a visual representation of the values for DFBETAS |
| <code>sort</code> | If <code>sort=TRUE</code> the values of DFBETAS are ordered based on magnitude. If <code>sort=FALSE</code> (default) no sorting takes place. |
| <code>to.sort</code> | Specify on which variable the DFBETAS must be sorted. If only one variable present (either in the model, or due to the selection specified in <code>parameters</code>), this parameter can be omitted. If DFBETAS is calculated for multiple variables, and <code>sort=TRUE</code> , specification of <code>to.sort</code> is required, or an error is returned. |
| <code>abs</code> | If <code>abs=TRUE</code> , the absolute values of DFBETAS are returned, while if <code>abs=FALSE</code> (default), both positive and negative values are possible. If both <code>abs=TRUE</code> and <code>sort=TRUE</code> , the <code>abs</code> parameter precedes the <code>sort</code> parameter, and thus the absolute values of DFBETAS are sorted. |
| <code>...</code> | Further arguments passed on to the <code>dp.ME.dfbetas()</code> function. |

Value

A matrix is returned, containing DFBETAS-values for each (selected) fixed parameter of the model, and separately for each evaluated set of influential data.

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

References

- Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.
- Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.
- Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the American Sociological Review.

See Also

[estex](#), [ME.cook](#)

Examples

```

data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
ME.dfbetas(alt.est)
ME.dfbetas(alt.est, plot=TRUE, layout=c(1,3))

```

ME.pchange

*Compute the percentage change, as measure of influential data***Description**

Compute the percentile change, as measure of influential data. This unstandardized measure can serve to help interpret the magnitude of the influence single or combined grouping levels exert on mixed effects models. The percentage change in parameter estimates between a (mixed effects) regression model based on a full set of data, and a model from which a (potentially influential) subset of data is removed. A value of percentage change is calculated for each parameter in the model separately, based on the information returned by the `estex()` function.

Usage

```
ME.pchange(estex, parameters = 0, plot=FALSE, sort=FALSE, to.sort=NA, abs=FALSE, ...)
```

Arguments

| | |
|-------------------------|--|
| <code>estex</code> | An object as returned by the <code>estex()</code> function, containing the altered estimates of a mixed effects regression model |
| <code>parameters</code> | Used to define a selection of parameters. If <code>parameters=0</code> (default), percentage change are calculated for all parameters in the model |
| <code>plot</code> | If <code>plot=TRUE</code> , the results from the <code>ME.pchange()</code> function are forwarded to the <code>dp.ME.pchange()</code> function, which creates a visual representation of the percentage changes. |
| <code>sort</code> | If <code>sort=TRUE</code> the values of percentage change are ordered based on magnitude. If <code>sort=FALSE</code> (default) no sorting takes place. |
| <code>to.sort</code> | Specify on which variable the percentage changes must be sorted. If only one variable present (either in the model, or due to the selection specified in <code>parameters</code>), this parameter can be omitted. If percentage changes are calculated for multiple variables, and <code>sort=TRUE</code> , specification of <code>to.sort</code> is required, or an error is returned. |
| <code>abs</code> | If <code>abs=TRUE</code> , the absolute values of percentage change are returned, while if <code>abs=FALSE</code> (default), both positive and negative values are possible. If both <code>abs=TRUE</code> and <code>sort=TRUE</code> , the <code>abs</code> parameter precedes the <code>sort</code> parameter, and thus the absolute values of percentage change are sorted. |
| <code>...</code> | Further arguments passed on to the <code>dp.ME.pchange()</code> function. |

Value

A matrix is returned, containing values of percentage change for each (selected) fixed parameter estimate of the model, and separately for each evaluated set of influential data.

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

References

Belsley, D.A., Kuh, E. & Welsch, R.E. (1980). *Regression Diagnostics. Identifying Influential Data and Source of Collinearity*. Wiley.

Snijders, T.A. & Bosker, R.J. (1999). *Multilevel Analysis, an introduction to basic and advanced multilevel modeling*. Sage.

Van Der Meer, T., Te Grotenhuis, M. & Pelzer, B. *Influential cases in multi-level modeling. A methodological comment on 'National context, religiosity, and volunteering' by Ruiter and De Graaf*. Current status: Accepted for publication in the American Sociological Review.

See Also

[estex](#), [ME.cook](#), [ME.dfbetas](#)

Examples

```
data(school23)
model <- lmer(math ~ structure + SES + (1 | school.ID), data=school23)

alt.est <- estex(model, "school.ID")
ME.pchange(alt.est)
ME.pchange(alt.est, plot=TRUE, layout=c(1,3))
```

school23

Math test performance in 23 schools

Description

The `school23` data contains information on students' performance on a math test, as well as several explanatory variables. These data are subset of the NELS-88 data (National Education Longitudinal Study of 1988). Both a selected number of variables and a selected number of observations are given here.

Usage

```
data(school23)
```

Format

A data frame with 519 observations on the following 15 variables.

school.ID a factor with 23 levels, representing the 23 schools within which students are nested.

SES a numeric vector, representing the socio-economic status

mean.SES a numeric vector, representing the mean socio-economic status per school

homework a factor representing the time spent on math homework each week, with levels None, Less than 1 hour, 1 hour, 2 hours, 3 hours, 4-6 hours, 7-9 hours, and 10 or more

parented a factor representing the parents' highest education level, with levels Did not finish H.S., H.S. grad or GED, GT H.S. and LT 4yr degree, College graduate, M.A. or equivalent, and Ph.D., M.D., other

ratio a numeric vector, representing the student-teacher ratio

perc.minor a factor representing the percent minority in school, with levels None, 1-5, 6-10, 11-20, 21-40, 41-60, 61-90, and 91-100

math a numeric vector, representing the number of correct answers on a mathematics test

sex a factor with levels Male and Female

race a factor with levels Asian, Hispanic, Black, White, and American Indian

school.type a factor representing the school type, with levels Public school, Catholic school, Private, other religious affiliation, and Private, no religious affiliation

structure a numeric vector representing the degree to which the classroom environment is structured. High values represent higher levels of (accurate) classroom environment structure

school.size a factor representing the total school enrollment, with levels 1-199 Students, 200-399, 400-599, 600-799, 800-999, 1000-1199, and 1200+

urban a factor with levels Urban, Suburban, and Rural

region a factor with levels Northeast, North Central, South, and West

Details

Labels for the factors were found in an appendix in Kreft & De Leeuw (1998). All labels were designated, although in some cases not all possible values are represented in the variable (i.e. *region*). This is probably due to the fact that this is only a subsample from the full NELS-88 data.

Also, some of the variable names were changed.

Source

These data are used in the examples given in Kreft & De Leeuw (1998). Both the examples and the data are publicly available from the internet: <http://www.ats.ucla.edu/stat/examples/imm/>. Data reproduced with permission from Jan de Leeuw.

References

Kreft, I. and De Leeuw, J. (1998). *Introducing Multilevel Modeling*. Sage Publications.

Examples

```
data(school23)
model <- lmer(math ~ structure + (1 | school.ID), data=school23)
```

se.fixef *Standard errors of fixed estimates*

Description

Returns the standard errors of the fixed estimates in a mixed effects model.

Usage

```
se.fixef(model)
```

Arguments

model Mixed effects regression model of class 'mer'

Value

A vector with the standard errors of the fixed parameters of the model.

Note

This is a small helper-function to the influence.ME package. For more elaborate functionality, refer to the se.fixef function in the 'car' package.

Author(s)

Rense Nieuwenhuis, Ben Pelzer, Manfred te Grotenhuis

Examples

```
data(school23)
model <- lmer(math ~ homework + structure + (1 | school.ID), data=school23)
summary(model)
se.fixef(model)
```

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