

Package ‘msprog’

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Type Package

Title Compute MS Progression from Longitudinal Data

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compute_delta	<i>Definition of progression deltas for different tests.</i>
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Description

compute_delta() returns the minimum delta to be considered as a valid change from baseline of an outcome measure (EDSS, NHPT, T25FW, or SDMT).

Usage

```
compute_delta(baseline, outcome = "edss")
```

Arguments

baseline	Outcome value at baseline.
outcome	One of: <ul style="list-style-type: none"> 'edss' (Extended Disability Status Scale, default); 'nhpt' (Nine-Hole Peg Test); 't25fw' (Timed 25-Foot Walk); 'sdmt' (Symbol Digit Modalities Test).

Value

Minimum delta corresponding to valid change from the provided baseline value. Specifically:

- EDSS: 1.5 if baseline==0, 1 if 0<baseline<=5, 0.5 if baseline>=5.5;
- NHPT and T25FW: 20% of baseline;
- SDMT: either 3 points or 10% of baseline.

Examples

```
compute_delta(4.5) # default outcome is 'edss'
compute_delta(55, outcome='sdmt')
```

criteria_text.MSprogOutput	<i>Textual description of criteria used to compute disability progression.</i>
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Description

criteria_text method for class 'MSprogOutput'.

Usage

```
## S3 method for class 'MSprogOutput'
criteria_text(object)
```

Arguments

object An object of class 'MSprogOutput' (result of a call to [MSprog\(\)](#)).

Details

The method prints out a short paragraph describing the set of criteria used to obtain the output.

Examples

```
# EDSS progression
output <- MSprog(toydata_visits, 'id', 'EDSS', 'date', 'edss',
  relapse=toydata_relapses, conf_weeks=12, conf_tol_days=30,
  event='multiple', baseline='roving', verbose=2)
criteria_text(output) # textual description of parameters used to obtain output
```

```
event_count.MSprogOutput
```

Event count for disability progression results.

Description

event_count method for class 'MSprogOutput'.

Usage

```
## S3 method for class 'MSprogOutput'
event_count(object)
```

Arguments

object An object of class 'MSprogOutput' (result of a call to [MSprog\(\)](#)).

Value

A data.frame object containing the sequence of events for each subject, as well as the event count separated by event type (improvement, progression, RAW, PIRA, undefined progression).

Examples

```
# EDSS progression
output <- MSprog(toydata_visits, 'id', 'EDSS', 'date', 'edss',
  relapse=toydata_relapses, conf_weeks=12, conf_tol_days=30,
  event='multiple', baseline='roving', verbose=2)
print(event_count(output)) # event sequence and count for each subject
```

is_event	<i>Compare value to reference.</i>
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Description

is_event() checks if an outcome value determines a valid progression, or improvement, or change, from a given reference value.

Usage

```
is_event(
  x,
  baseline,
  type,
  outcome = "edss",
  worsening = NULL,
  delta_fun = NULL,
  sub_threshold = FALSE
)
```

Arguments

x	Outcome value to test.
baseline	Outcome value at baseline.
type	One of: <ul style="list-style-type: none"> • 'prog' (progression); • 'impr' (improvement); • 'change' (any valid change).
outcome	One of: <ul style="list-style-type: none"> • 'edss' (Extended Disability Status Scale, default); • 'nhpt' (Nine-Hole Peg Test); • 't25fw' (Timed 25-Foot Walk); • 'sdmt' (Symbol Digit Modalities Test); • NULL (only accepted when specifying the direction of worsening).
worsening	The direction of worsening ('increase' if higher values correspond to worse disease course, 'decrease' otherwise). This argument is only used when outcome is set to NULL. If outcome is specified, worsening is automatically set to 'increase' for EDSS, NHPT, T25FW, and to 'decrease' for SDMT.
delta_fun	Custom function specifying the minimum shift corresponding to a valid change from the provided baseline value. If none is specified (default), compute_delta() for the specified outcome is used.
sub_threshold	If TRUE, any confirmed progression, or improvement, or change in outcome measure is valid, regardless of delta_fun.

Value

A boolean value specifying if a valid event was found.

Examples

```
is_event(x=4.5, baseline=4, type='prog', outcome='edss')
is_event(x=50, baseline=57, type='prog', outcome='sdtmt')
```

MSprog

*Compute multiple sclerosis progression from longitudinal data.***Description**

MSprog() detects and characterises the progression (or improvement) events of an outcome measure (EDSS, NHPT, T25FW, or SDMT) for one or more subjects, based on repeated assessments through time and on the dates of acute episodes. Several qualitative and quantitative options are given as arguments that can be set by the user and reported as a complement to the results to ensure reproducibility.

Usage

```
MSprog(
  data,
  subj_col,
  value_col,
  date_col,
  outcome,
  subjects = NULL,
  date_format = NULL,
  relapse = NULL,
  rsubj_col = NULL,
  rdate_col = NULL,
  delta_fun = NULL,
  worsening = NULL,
  conf_weeks = 12,
  conf_tol_days = 30,
  conf_unbounded_right = FALSE,
  require_sust_weeks = 0,
  relapse_to_bl = 30,
  relapse_to_event = 0,
  relapse_to_conf = 30,
  relapse_assoc = 90,
  event = "firstprog",
  baseline = "fixed",
  relapse_indep = NULL,
  sub_threshold = FALSE,
  relapse_rebl = FALSE,
  min_value = NULL,
  prog_last_visit = FALSE,
  include_dates = FALSE,
  include_value = FALSE,
  include_stable = TRUE,
  verbose = 1,
  devtest_conf = FALSE
)
```

Arguments

<code>data</code>	<code>data.frame</code> containing longitudinal data, including: subject ID, outcome value, date of visit.
<code>subj_col</code>	Name of data column with subject ID.
<code>value_col</code>	Name of data column with outcome value.
<code>date_col</code>	Name of data column with date of visit.
<code>outcome</code>	Specifies the outcome type. Must be one of the following: <ul style="list-style-type: none"> 'edss' (Expanded Disability Status Scale); 'nhpt' (Nine-Hole Peg Test); 't25fw' (Timed 25-Foot Walk); 'sdmt' (Symbol Digit Modalities Test); NULL (only accepted when specifying a custom <code>delta_fun</code>)
<code>subjects</code>	Subset of subjects (list of IDs). If none is specified, all subjects listed in <code>data</code> are included.
<code>date_format</code>	Format of dates in the input data. If not specified, it will be inferred by function as.Date .
<code>relapse</code>	<code>data.frame</code> containing longitudinal data, including: subject ID and relapse date.
<code>rsubj_col</code>	Name of subject ID column for relapse data, if different from outcome data.
<code>rdate_col</code>	Name of date column for relapse data, if different from outcome data.
<code>delta_fun</code>	Custom function specifying the minimum shift corresponding to a valid change from the provided reference value. It must take a numeric value (reference) as input, and return a numeric value corresponding to the minimum shift from baseline. If none is specified (default), function compute_delta() for the specified outcome is used.
<code>worsening</code>	The direction of worsening ('increase' if higher values correspond to worse disease course, 'decrease' otherwise). This argument is only used when outcome is set to NULL. Otherwise, worsening is automatically set to 'increase' if outcome is set to 'edss', 'nhpt', 't25fw', and to 'decrease' if outcome is set to 'sdmt'.
<code>conf_weeks</code>	Period before confirmation (weeks).
<code>conf_tol_days</code>	Tolerance window for confirmation visit (days); can be an integer (same tolerance on left and right) or list-like of length 2 (different tolerance on left and right). In all cases, the right end of the interval is ignored if <code>conf_unbounded_right</code> is set to TRUE.
<code>conf_unbounded_right</code>	If TRUE, confirmation window is unbounded on the right.
<code>require_sust_weeks</code>	Minimum number of weeks over which a confirmed change must be sustained to be retained as an event. Events sustained for the entire follow-up are retained regardless of follow-up duration. Setting <code>require_sust_weeks=Inf</code> , events are only retained if sustained for the entire follow-up duration.
<code>relapse_to_bl</code>	Minimum distance from last relapse (days) for a visit to be used as baseline (otherwise the next available visit is used as baseline).
<code>relapse_to_event</code>	Minimum distance from last relapse (days) for an event to be considered as such.

relapse_to_conf	Minimum distance from last relapse (days) for a visit to be a valid confirmation visit.
relapse_assoc	Maximum distance from last relapse (days) for a progression event to be considered as RAW.
event	Specifies which events to detect. Must be one of the following: <ul style="list-style-type: none"> • 'firstprog' (first progression, default); • 'first' (only the very first event - improvement or progression); • 'firstteach' (first improvement and first progression - in chronological order); • 'firstprogtype' (first progression of each kind - PIRA, RAW, and undefined, in chronological order); • 'firstPIRA' (first PIRA); • 'firstRAW' (first RAW); • 'multiple' (all events in chronological order).
baseline	Specifies the baseline scheme. Must be one of the following: <ul style="list-style-type: none"> • 'fixed' (first valid outcome value, default); • 'roving_impr' (updated every time the value is lower than the previous measure and confirmed at the following visit; suitable for a first-progression setting to discard fluctuations around baseline); • 'roving'``}{ (updated after each event to last valid confirmed outcome value; suitable for 'firstprogtype' - or when searching for a specific type of progression - i.e., when event is set to 'firstPIRA' or 'firstRAW').
relapse_indep	Specifies relapse-free intervals for PIRA definition. Must be given in the form produced by function <code>relapse_indep_from_bounds()</code> by calling <code>relapse_indep_from_bounds(b0, b1, e0, e1, c0, c1)</code> to specify the intervals around baseline (b0 and b1), event (e0 and e1), and confirmation (c0 and c1). For instance: <ul style="list-style-type: none"> • [Muller JAMA Neurol 2023] No relapses within event-90dd->event+30dd and within confirmation-90dd->confirmation+30dd: <code>relapse_indep <- relapse_indep_from_bounds(0, 0, 90, 30, 90, 30)</code> (default); • [Muller JAMA Neurol 2023](high-specificity definition) No relapses between baseline and confirmation: <code>\coderelapse_indep <- relapse_indep_from_bounds(0, NULL, NULL, NULL, NULL, 0);</code> • [Kappos JAMA Neurol 2020] No relapses within baseline->event+30dd and within confirmation+-30dd: <code>relapse_indep <- relapse_indep_from_bounds(0, NULL, NULL, 30, 30, 30)</code>
sub_threshold	If TRUE - and only if baseline is 'roving' or 'roving_impr' - move roving baseline at any sub-threshold confirmed event (i.e. any confirmed change in outcome measure, regardless of delta_fun).
relapse_rebl	If TRUE, re-baseline after every relapse to search for PIRA events.
min_value	Only consider progressions events where the outcome is >= value.
prog_last_visit	If TRUE, include progressions occurring at last visit (i.e. with no confirmation). If a numeric value N is passed, unconfirmed events are included only if occurring within N weeks of follow up (e.g., in case of early discontinuation).
include_dates	If TRUE, report dates of events.

include_value If TRUE, report value of outcome at event.

include_stable If TRUE, subjects with no events are included in extended output data.frame, with time2event = total follow up.

verbose One of:

- 0 (print no info);
- 1 (print concise info, default);
- 2 (print extended info).

devtest_conf (to be removed soon) Temporary test - for developer's use only.

Details

The events are detected sequentially by scanning the outcome values in chronological order. Progression events are classified as relapse-associated or relapse-independent based on their relative timing with respect to the relapses. Specifically, relapse-associated worsening (RAW) events are defined as confirmed progression events occurring within the influence of a relapse, while progression independent of relapse activity (PIRA) is established when the progression event occurs out of relapse influence, and with no relapses between baseline and confirmation.

Value

An object of class 'MSprogOutput'.

Examples

```
# EDSS progression
output_edss <- MSprog(toydata_visits, 'id', 'EDSS', 'date', 'edss',
  relapse=toydata_relapses, conf_weeks=12, conf_tol_days=30,
  event='multiple', baseline='roving', verbose=1)
print(results(output_edss)) # extended info on each event for all subjects
print(event_count(output_edss)) # summary of event sequence for each subject
# SDMT progression
output_sdmtd <- MSprog(toydata_visits, 'id', 'SDMT', 'date', 'sdmt',
  relapse=toydata_relapses, conf_weeks=12, conf_tol_days=30,
  event='multiple', baseline='roving', verbose=1)
print(results(output_sdmtd)) # extended info on each event for all subjects
print(event_count(output_sdmtd)) # summary of event sequence for each subject
```

relapse_indep_from_bounds

Define relapse-free intervals for PIRA definition.

Description

relapse_indep_from_bounds() organises the given interval bounds around baseline, event, and confirmation into a named list to be given as argument relapse_indep to function [MSprog\(\)](#).

Usage

```
relapse_indep_from_bounds(b0, b1, e0, e1, c0, c1)
```


Arguments

b0	Days before baseline (≥ 0).
b1	Days after baseline (≥ 0), or NULL.
e0	Days before event (≥ 0), or NULL.
e1	Days after event (≥ 0), or NULL.
c0	Days before confirmation (≥ 0), or NULL.
c1	Days after confirmation (≥ 0).

Details

If the right end is NULL, the interval is assumed to extend up to the left end of the next interval. If the left end is NULL, the interval is assumed to extend up to the right end of the previous interval.

Value

A named list to be given as argument `relapse_indep` to function [MSprog\(\)](#)

Examples

```
# [Muller JAMA Neurol 2023](high-specificity definition)
# No relapses between baseline and confirmation:
relapse_indep <- relapse_indep_from_bounds(0,NULL,NULL,NULL,NULL,0)
# [Muller JAMA Neurol 2023]
# No relapses within event-90dd->event+30dd
# and within confirmation-90dd->confirmation+30dd:
relapse_indep <- relapse_indep_from_bounds(0,0,90,30,90,30)
# [Kappos JAMA Neurol 2020]
# No relapses within baseline->event+30dd and within confirmation+-30dd:
relapse_indep <- relapse_indep_from_bounds(0,NULL,NULL,30,30,30)
```

results.MSprogOutput *Extended disability progression results.*

Description

results method for class 'MSprogOutput'.

Usage

```
## S3 method for class 'MSprogOutput'
results(object)
```

Arguments

object An object of class 'MSprogOutput' (result of a call to [MSprog\(\)](#)).

Value

A data.frame object containing an extended report of all events detected by function [MSprog\(\)](#) for each subject.

Examples

```
# EDSS progression
output <- MSprog(toydata_visits, 'id', 'EDSS', 'date', 'edss',
  relapse=toydata_relapses, conf_weeks=12, conf_tol_days=30,
  event='multiple', baseline='roving', verbose=2)
print(results(output)) # extended event info for each subject
```

toydata_relapses	<i>Synthetic Relapse Data</i>
------------------	-------------------------------

Description

Artificially generated relapse dates for some example patients in [toydata_visits](#) to illustrate the use of the package.

Usage

```
data(toydata_relapses)
```

Format

An object of class `data.frame`

id Subject ID

date The relapse date

References

This data set was artificially created for the `msprog` package.

Examples

```
data(toydata_relapses)
head(toydata_relapses)
```

toydata_visits	<i>Synthetic Longitudinal EDSS and SDMT Data</i>
----------------	--

Description

Small, artificially generated toy data set providing Extended Disability Status Scale (EDSS) and Symbol Digit Modalities Test (SDMT) information on 4 example patients to illustrate the use of the package.

Usage

```
data(toydata_visits)
```

Format

An object of class `data.frame`

id Subject ID

date The visit date

EDSS A value between 0 and 10

SDMT A value between 0 and 110

References

This data set was artificially created for the `msprog` package.

Examples

```
head(toydata_visits)
```

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