Package: drugprepr (via r-universe)

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Title Prepare Electronic Prescription Record Data to Estimate Drug Exposure

Version 0.0.5.9000

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BugReports https://github.com/belayb/drugprepr/issues

Description Prepare prescription data (such as from the Clinical Practice Research Datalink) into an analysis-ready format, with start and stop dates for each patient's prescriptions. Based on Pye et al (2018) <doi:10.1002/pds.4440>.

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Imports dplyr, doseminer, rlang, tidyr, sqldf, stringr, purrr, DescTools

Depends R (>= 2.10)

Suggests knitr, rmarkdown, testthat, kableExtra

VignetteBuilder knitr

Repository https://belayb.r-universe.dev

RemoteUrl https://github.com/belayb/drugprepr

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Contents

`clean_duration` *Clean implausibly-long prescription durations*

Description

Given a prescription length limit, truncate any prescriptions that appear to be longer than this, or mark them as missing.

Usage

```
clean_duration(data, max_months = Inf, method = c("truncate", "remove"))
```

Arguments

`data`	A data frame containing a column called `duration`
`max_months`	The maximum plausible prescription length in months
`method`	Either 'truncate' or 'remove'. See details

Details

The method 'truncate' causes any duration longer than `max_months` to be replaced with the value of `max_months` (albeit converted to days). The method 'remove' causes such durations to be replaced with `NA`. There is no explicit 'ignore' method, but if you want to 'do nothing', simply set `max_months` to an arbitrarily high number. By default, the maximum is infinite, so nothing should happen. (Of course, you could also just *not* run the function...)

Value

A data frame of the same structure as the input, possibly with some elements of the `duration` column changed

Note

Currently the variable name is hard-coded as 'duration', but in principle this could be parametrised for datasets where the column has a different name.

Examples

```
long_presc <- data.frame(duration = c(100, 300, 400, 800))
clean_duration(long_presc, 6)
clean_duration(long_presc, 12, 'remove')
```

`close_small_gaps` *Close small gaps between successive prescriptions*

Description

Given a series of prescriptions in `data`, if one prescription (for the same patient and drug) starts $\leq$ `min_gap` days after the previous one finishes, we extend the length of the previous prescription to cover the gap.

Usage

```
close_small_gaps(data, min_gap = 0L)
```

Arguments

`data`	A data frame containing columns `prodcode`, `patid`, `start_date` and `stop_date`
`min_gap`	Size of largest gaps to close. Default is zero, i.e. do nothing

Value

The input data frame `data`, possibly with some of the `stop_dates` changed.

Examples

```
gappy_data <- data.frame(
  patid = 1,
  prodcode = 'a',
  start_date = Sys.Date() + (0:6) * 7,
  stop_date = Sys.Date() + (0:6) * 7 + 4
)
close_small_gaps(gappy_data)
close_small_gaps(gappy_data, 7)
```

`compute_ndd` *Compute numerical daily dose from free-text prescribing instructions*

Description

The function calls the R package **doseminer** to extract dose information from free-text prescribing instructions, then computes the average numerical daily dose according to a given decision rule.

Usage

```
compute_ndd(data, dose_fn = mean, freq_fn = mean, interval_fn = mean)
```

Arguments

`data`	a data frame containing free-text prescribing instructions in a column called `text`.
`dose_fn`	function to summarise range of numbers by a single value
`freq_fn`	function to summarise range of frequencies by a single value
`interval_fn`	function to summarise range of intervals by a single value

Details

The general formula for computing numerical daily dose (ndd) is given by

$$\text{ndd} = \text{DF} \times \text{DN}/\text{DI},$$

where

DF is dose frequency, the number of dose 'events' per day

DN is dose number, or number of units of drug taken during each dose 'event'

DI is dose interval, or the number of days between 'dose days', where an interval of 1 means every day

Prescriptions can have a variable dose frequency or dose number, such as '2-4 tablets up to 3 times per day'. In this case, the user can choose to reduce these ranges to single values by taking the minimum, maximum or average of these endpoints.

Value

A data frame mapping the raw `text` to structured dosage information.

Examples

```
compute_ndd(cprd, min, min, mean)
```

`cprd` *Example data from the Clinical Practice Research Datalink (CPRD).*

Description

A dataset containing prescription information for two individuals. The dataset is a hypothetical dataset resembling the real CPRD data.

Usage

```
cprd
```

Format

A data frame with 18 rows and 9 variables:

patid unique identifier given to a patient in CPRD GOLD

pracid unique identifier given to a practice in CPRD GOLD

start_date Beginning of the prescription period

prodcode CPRD unique code for the treatment selected by the GP

dosageid Identifier that allows dosage information on the event to be retrieved from a Common Dosages lookup table

text Prescription instruction for the prescribed product, as entered by the GP

qty Total quantity entered by the GP for the prescribed product

numdays Number of treatment days prescribed for a specific therapy event

dose_duration an estimated prescription duration, as entered by CPRD ...

Source

https://cprdcw.cprd.com/_docs/CPRD_GOLD_Full_Data_Specification_v2.0.pdf

See Also

`min_max_dat`

`decision_1` *Decision 1: impute implausible total quantities*

Description

A light wrapper around `impute_qty`.

Usage

```
decision_1(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings: **"a"** do nothing; leave implausible values as-is **"b"** set implausible values to missing **"c1"** set to mean for individual's prescriptions for that drug **"c2"** set to mean for practice's prescriptions for that drug **"c3"** set to mean for populations's prescriptions for that drug **"d1"** set to median for individual's prescriptions for that drug **"d2"** set to median for practice's prescriptions for that drug **"d3"** set to median for population's prescriptions for that drug **"e1"** set to mode for individual's prescriptions for that drug **"e2"** set to mode for practice's prescriptions for that drug **"e3"** set to mode for population's prescriptions for that drug **"f1"** use value of individual's next prescription **"f2"** use value of practice's next prescription **"f3"** use value of population's next prescription **"g1"** use value of individual's previous prescription **"g2"** use value of practice's previous prescription **"g3"** use value of population's previous prescription

Note

Decisions `f` and `g` are not yet implemented.

See Also

Other decision functions: `decision_10()`, `decision_2()`, `decision_3()`, `decision_4()`, `decision_5()`, `decision_6()`, `decision_7()`, `decision_8()`, `decision_9()`, `drug_prep()`

decision_10 *Decision 10: close small gaps between successive prescriptions*

Description

Where one prescription (for the same drug and patient) starts only a short time after the previous finishes, this function can close the gap, as if the prescription was continuous over the entire period.

Usage

```
decision_10(data, decision = "a")
```

Arguments

data	a data frame
decision	one of the following strings: **"a"** do nothing **"b_15"** change stop date of first prescription to start date of next if gap is ≤ 15 days **"b_30"** change stop date of first prescription to start date of next if gap is ≤ 30 days **"b_60"** change stop date of first prescription to start date of next if gap is ≤ 60 days

Details

The underlying function is called `close_small_gaps`

See Also

Other decision functions: `decision_1()`, `decision_2()`, `decision_3()`, `decision_4()`, `decision_5()`, `decision_6()`, `decision_7()`, `decision_8()`, `decision_9()`, `drug_prep()`

decision_2 *Decision 2: impute missing total quantities*

Description

A light wrapper around `impute_qty`.

Usage

```
decision_2(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings:

"a" Leave as missing (implicitly drop this prescription)
"b1" set to mean for individual's prescriptions for that drug
"b2" set to mean for practice's prescriptions for that drug
"b3" set to mean for populations's prescriptions for that drug
"c1" set to median for individual's prescriptions for that drug
"c2" set to median for practice's prescriptions for that drug
"c3" set to median for population's prescriptions for that drug
"d1" set to mode for individual's prescriptions for that drug
"d2" set to mode for practice's prescriptions for that drug
"d3" set to mode for population's prescriptions for that drug
"e1" use value of individual's next prescription
"e2" use value of practice's next prescription
"e3" use value of population's next prescription
"f1" use value of individual's previous prescription
"f2" use value of practice's previous prescription
"f3" use value of population's previous prescription

Note

Decisions `e` and `f` are not yet implemented.

See Also

Other decision functions: `decision_10()`, `decision_1()`, `decision_3()`, `decision_4()`, `decision_5()`, `decision_6()`, `decision_7()`, `decision_8()`, `decision_9()`, `drug_prep()`

`decision_3` *Decision 3: impute implausible daily doses*

Description

A light wrapper around `impute_ndd`.

Usage

```
decision_3(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings:

"a" do nothing; leave implausible values as-is
"b" set implausible values to missing
"c1" set to mean for individual's prescriptions for that drug
"c2" set to mean for practice's prescriptions for that drug
"c3" set to mean for populations's prescriptions for that drug
"d1" set to median for individual's prescriptions for that drug
"d2" set to median for practice's prescriptions for that drug
"d3" set to median for population's prescriptions for that drug
"e1" set to mode for individual's prescriptions for that drug
"e2" set to mode for practice's prescriptions for that drug
"e3" set to mode for population's prescriptions for that drug
"f1" use value of individual's next prescription
"f2" use value of practice's next prescription
"f3" use value of population's next prescription
"g1" use value of individual's previous prescription
"g2" use value of practice's previous prescription
"g3" use value of population's previous prescription

Note

Decisions `f` and `g` are not yet implemented.

See Also

Other decision functions: `decision_10()`, `decision_1()`, `decision_2()`, `decision_4()`, `decision_5()`, `decision_6()`, `decision_7()`, `decision_8()`, `decision_9()`, `drug_prep()`

`decision_4` *Decision 4: impute missing daily doses*

Description

A light wrapper around `impute_ndd`.

Usage

```
decision_4(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings: **"a"** Leave as missing (implicitly drop this prescription) **"b1"** set to mean for individual's prescriptions for that drug **"b2"** set to mean for practice's prescriptions for that drug **"b3"** set to mean for populations's prescriptions for that drug **"c1"** set to median for individual's prescriptions for that drug **"c2"** set to median for practice's prescriptions for that drug **"c3"** set to median for population's prescriptions for that drug **"d1"** set to mode for individual's prescriptions for that drug **"d2"** set to mode for practice's prescriptions for that drug **"d3"** set to mode for population's prescriptions for that drug **"e1"** use value of individual's next prescription **"e2"** use value of practice's next prescription **"e3"** use value of population's next prescription **"f1"** use value of individual's previous prescription **"f2"** use value of practice's previous prescription **"f3"** use value of population's previous prescription

Note

Decisions `e` and `f` are not yet implemented.

See Also

Other decision functions: `decision_10()`, `decision_1()`, `decision_2()`, `decision_3()`, `decision_5()`, `decision_6()`, `decision_7()`, `decision_8()`, `decision_9()`, `drug_prep()`

`decision_5` *Decision 5: impute implausible prescription durations*

Description

A light wrapper around `clean_duration`.

Usage

```
decision_5(data, decision = "a")
```

Arguments

data	a data frame
decision	one of the following strings: **"a"** leave duration as-is **"b_6"** set to missing if > 6 months **"b_12"** set to missing if > 12 months **"b_24"** set to missing if > 24 months **"c_6"** set to 6 months if > 6 months **"c_12"** set to 12 months if > 12 months **"c_24"** set to 24 months if > 24 months

See Also

Other decision functions: decision_10(), decision_1(), decision_2(), decision_3(), decision_4(), decision_6(), decision_7(), decision_8(), decision_9(), drug_prep()

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decision_6 *Decision 6: choose method of calculating prescription duration*

Description

This is just shorthand for defining a column equal to one of the specified formulae. If the column(s) corresponding to decision are missing, an error will be thrown. If you have already calculated or obtained the column duration from elsewhere, this step is not necessary.

Usage

```
decision_6(data, decision = "c")
```

Arguments

data	a data frame
decision	one of the following strings: **"a"** numdays **"b"** dose_duration **"c"** qty / ndd

Note

This step actually takes place *before* decision_5.

See Also

Other decision functions: decision_10(), decision_1(), decision_2(), decision_3(), decision_4(), decision_5(), decision_7(), decision_8(), decision_9(), drug_prep()

decision_7 *Decision 7: impute missing prescription durations*

Description

A light wrapper around `impute_duration`.

Usage

```
decision_7(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings: **"a"** Leave missing durations as-is (implicitly drop the prescription) **"b"** Use mean prescription duration for that drug, for that individual **"c"** Use mean prescription duration for that drug, for the population **"d"** Use individual mean duration; if not available use population mean

See Also

Other decision functions: `decision_10()`, `decision_1()`, `decision_2()`, `decision_3()`, `decision_4()`, `decision_5()`, `decision_6()`, `decision_8()`, `decision_9()`, `drug_prep()`

decision_8 *Decision 8: disambiguate prescriptions with the same start date*

Description

A light wrapper around `impute_duration`, followed by removing duplicate rows with the same combination of `prodcode`, `patid` and `start_date`.

Usage

```
decision_8(data, decision = "a")
```

Arguments

`data`	a data frame
`decision`	one of the following strings **"a"** do nothing **"b"** replace with a prescription of duration equal to the mean **"c"** choose the shortest prescription **"d"** choose longest prescription **"e"** replace with a prescription of duration equal to the sum

See Also

Other decision functions: decision_10(), decision_1(), decision_2(), decision_3(), decision_4(), decision_5(), decision_6(), decision_7(), decision_9(), drug_prep()

---

decision_9 *Decision 9: handle overlapping prescription periods*

Description

In situations where one prescription starts before another (for the same patient and drug) finishes, this function will either implicitly sum the doses (i.e. do nothing) or it will divide the intervals into non-overlapping subsets, shifting these sub-intervals forward in time until there is no overlap.

Usage

```
decision_9(data, decision = "a")
```

Arguments

data	a data frame
decision	one of the following strings:
	"a" allow overlapping prescriptions (implicitly sum doses)
	"b" move later prescription to next available time that this product is not prescribed

Details

The underlying algorithm for shifting overlapping intervals is implemented by the internal function shift_interval.

See Also

Other decision functions: decision_10(), decision_1(), decision_2(), decision_3(), decision_4(), decision_5(), decision_6(), decision_7(), decision_8(), drug_prep()

drug_prep *Run drug preparation algorithm*

Description

Run drug preparation algorithm

Usage

```
drug_prep(data, plausible_values, decisions = rep("a", 10))
```

Arguments

data	data frame containing prescription data
plausible_values	data frame containing variables prodcode, min_qty, max_qty, min_ndd, max_ndd describing plausible ranges for values for each drug
decisions	character vector of length 10

Value

A data frame including estimated stop_date for each prescription

See Also

Other decision functions: decision_10(), decision_1(), decision_2(), decision_3(), decision_4(), decision_5(), decision_6(), decision_7(), decision_8(), decision_9()

Examples

```
plausible_values <- data.frame(
  prodcode = c('a', 'b', 'c'),
  min_qty = 0,
  max_qty = c(50, 100, 200),
  min_ndd = 0,
  max_ndd = c(10, 20, 30)
)
drug_prep(example_therapy,
          plausible_values,
          decisions = c('a', 'a', 'a', 'a', 'a',
                        'c', 'a', 'a', 'a', 'a'))
```

example_therapy *Example electronic prescription dataset*

Description

Based on a hypothetical 'therapy' file from the Clinical Practical Research Datalink (CPRD), a UK database of primary care records.

Usage

```
example_therapy
```

Format

An object of class `data.frame` with 30 rows and 6 columns.

Note

This dataset is now generated deterministically, so it will not vary between sessions.

get_mode *Get the mode (most common value) of a vector*

Description

Get the mode (most common value) of a vector

Usage

```
get_mode(v, na.rm = TRUE)
```

Arguments

v	a vector
na.rm	Logical. If TRUE (the default), find mode of non-NA values

`impute` *Impute missing or implausible values*

Description

This is a workhorse function used by `impute_ndd`, `impute_qty` and others.

Usage

```
impute(
  data,
  variable,
  method = c("ignore", "mean", "median", "mode", "replace", "min", "max", "sum"),
  where = is.na,
  group,
  ...,
  replace_with = NA_real_
)
```

Arguments

`data`	A data frame containing columns `prodcode`, `pracid`, `patid`
`variable`	Unquoted name of the column in `dataset` to be imputed
`method`	Method for imputing the values. See details.
`where`	Logical vector, or function applied to `variable` returning such a vector, indicating which elements to impute. Defaults to `is.na`
`group`	Level of structure for imputation. Defaults to whole study population.
`...`	Extra arguments, currently ignored
`replace_with`	if the method 'replace' is selected, which value should be inserted?

- `ignore`. Do nothing, leaving input unchanged.
- `mean`. Replace values with the mean by `group`
- `median`. Replace values with the median by `group`
- `mode`. Replace values with the most common value by `group`
- `replace`. Replace values with `replace_with`, which defaults to `NA` (i.e. mark as missing)
- `min`. Replace with minimum value.
- `max`. Replace with maximum value.
- `sum`. Replace with sum of values.

Details

The argument `where` indicates which values are to be imputed. It can be specified as either a vector or as a function. Thus you can specify, for example, `is.na` to impute all missing values, or you can pass in a vector, if it depends on something else rather than just the current values of the variable to imputed. This design may change in future. In particular, if we want to impute implausible values and impute missing values separately, it's important that these steps are independent.

Value

A data frame of the same structure as `data`, with values imputed

---

`impute_duration` *Replace missing or implausible prescription durations*

Description

Instead of replacing missing stop dates, we impute the durations and then infer the stop dates from there.

Usage

```
impute_duration(
  data,
  method,
  where = is.na,
  group = c("patid", "start_date"),
  ...
)
```

Arguments

`data`	A data frame containing columns `prodcode`, `pracid`, `patid`
`method`	Method for imputing the values. See details.
`where`	Logical vector, or function applied to `variable` returning such a vector, indicating which elements to impute. Defaults to `is.na`
`group`	Level of structure for imputation. Defaults to whole study population.
`...`	Extra arguments, currently ignored

Details

We can fix clashing start dates by setting `group` to `start_date` and `patid`, i.e. average over groups with more than one member; any metric should return the original values if the group size is one.

Value

A data frame of the same structure as `data`, with values imputed

Examples

```
example_duration <- transform(example_therapy, duration = qty / ndd)
impute_duration(example_duration, method = 'mean', group = 'patid')
```

impute_ndd *Replace implausible or missing numerical daily doses (NDD)*

Description

Replace implausible or missing numerical daily doses (NDD)

Usage

```
impute_ndd(data, method, where = is.na, group = "population", ...)
```

Arguments

data	A data frame containing columns prodcode, pracid, patid
method	Method for imputing the values. See details.
where	Logical vector, or function applied to variable returning such a vector, indicating which elements to impute. Defaults to is.na
group	Level of structure for imputation. Defaults to whole study population.
...	Extra arguments, currently ignored

Value

A data frame of the same structure as data, with values imputed

Examples

```
impute_ndd(example_therapy, 'mean')
```

impute_qty *Find implausible entries Replace implausible or missing prescription quantities*

Description

Find implausible entries Replace implausible or missing prescription quantities

Usage

```
impute_qty(data, method, where = is.na, group = "population", ...)
```

Arguments

data	A data frame containing columns prodcode, pracid, patid
method	Method for imputing the values. See details.
where	Logical vector, or function applied to variable returning such a vector, indicating which elements to impute. Defaults to is.na
group	Level of structure for imputation. Defaults to whole study population.
...	Extra arguments, currently ignored

Value

A data frame of the same structure as data, with values imputed

Examples

```
impute_qty(example_therapy, 'mean')
```

---

isolate_overlaps *Separating overlapping prescription periods*

Description

Run this function and then you can either simply discard overlapping intervals or shift them around using an appropriate algorithm.

Usage

```
isolate_overlaps(data)
```

Arguments

data	A data frame including variables patid, start_date, stop_date and prodcode

Details

The older implementation used isolateoverlaps from the intervalaverage package and Overlap from the DescTools package. Here we refactor it using functions from tidyverse instead.

Value

A data frame of patid, prodcode, start_date and stop_date, where intervals are either exactly overlapping or mutually non-overlapping (but not partially overlapping), such that the union of such intervals is equivalent to those originally provided in data

Note

This function currently doesn't use any keys except `patid` and `prodcode`. It may be desirable to add a row ID, for matching each partial interval back to the original interval from which it was derived. This may be relevant to models using weighted dosages.

See Also

`intervalaverage::isolateoverlaps`, `foverlaps`

Examples

```
set.seed(1)
overlapping_data <- data.frame(
  rowid = 1:20,
  patid = 1:2,
  prodcode = 'a',
  start_date = Sys.Date() + c(round(rexp(19, 1/7)), -20),
  qty = rpois(20, 64),
  ndd = sample(seq(.5, 12, by = .5), 20, replace = TRUE),
  stringsAsFactors = FALSE
)
overlapping_data <- transform(overlapping_data,
  stop_date = start_date + qty / ndd
)
isolate_overlaps(overlapping_data)
```

---

make_decisions *Human-friendly interface to the drug prep algorithm*

Description

A helper function that allows specifying decision rules using English words rather than alphanumeric codes. Translates the rules into the corresponding codes and then passes them to `drug_prep` functions.

Usage

```
make_decisions(
  implausible_qty,
  missing_qty,
  implausible_ndd,
  missing_ndd,
  implausible_duration,
  calculate_duration,
  missing_duration,
  clash_start,
  overlapping,
```

```
  small_gaps
)
```

Arguments

`implausible_qty`	implausible total drug quantities
`missing_qty`	missing total drug quantities
`implausible_ndd`	implausible daily dosage
`missing_ndd`	missing daily dosage
`implausible_duration`	overly-long prescription durations
`calculate_duration`	formula or variable to compute prescription duration
`missing_duration`	missing prescription duration
`clash_start`	how to disambiguate prescriptions that start on the same date
`overlapping`	how to handle prescription periods that overlap with one another
`small_gaps`	how to handle short gaps between successive prescriptions

The argument `decision_phrases` may contain the following terms (without brackets, separated with spaces). Additional or incorrectly-named elements will be ignored.

implausible_qty (ignore|missing|mean|median|mode|next|previous) (individual|practice|population)
implausible_ndd (ignore|missing|mean|median|mode|next|previous) (individual|practice|population)
implausible_duration (ignore|missing|truncate) (6|12|24)
missing_qty (ignore|mean|median|mode|next|previous) (individual|practice|population)
missing_ndd (ignore|mean|median|mode|next|previous) (individual|practice|population)
missing_duration (ignore|mean) (individual|population|both)
calculate_duration (numdays|dose_duration|qty/ndd)
clash_start (ignore|mean|shortest|longest|sum)
overlapping (allow|shift)
small_gaps (ignore|close) (15|30|60)

Value

A character vector suitable for passing to the `decisions` argument of the `drug_prep` function.

Examples

```
make_decisions('ignore',
               'mean population',
               'missing',
               'mean practice',
               'truncate 6',
               'qty / ndd',
```

```
'mean individual',
'mean',
'allow',
'close 15')
```

`min_max_dat` *Example plausible ranges for prescription quantity and dosage*

Description

Minimum and maximum plausible values for total prescription quantity and numerical daily dose of prescriptions given in the `cprd` dataset. Both datasets are hypothetical.

Usage

```
min_max_dat
```

Format

A data frame with 2 rows and 5 variables:

prodcode CPRD unique code for the treatment selected by the GP

max_qty maximum possible quantity to be prescribed for the product

min_qty minimum possible quantity to be prescribed for the product

max_ndd maximum possible number of daily dose to be prescribed for the product

min_ndd minimum possible number of daily dose to be prescribed for the product ...

See Also

`cprd`

`outside_range` *Do values fall outside a specified 'plausible' range?*

Description

A utility function for indicating if elements of a vector are implausible.

Usage

```
outside_range(x, lower, upper, open = TRUE)
```

Arguments

x	numeric vector
lower	minimum plausible value
upper	maximum plausible value
open	logical. If TRUE, values exactly equal to lower or upper are also considered implausible

Details

Though the function between already exists, it is not vectorised over the bounds.

---

shift_interval *Shift time intervals until they no longer overlap*

Description

This is a function used by decision_9.

Usage

```
shift_interval(x)
```

Arguments

x	a data frame containing variables start_date, stop_date and patid

Value

A data frame with time intervals moved such that they no longer overlap

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