Package ‘medExtractR’

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Title Extraction of Medication Information from Clinical Text
Description Function and support for medication and dosing information extraction from free-
text clinical notes. Medication entities for the basic medExtractR implementation that can be ex-
tracted include drug name, strength, dose amount, dose, frequency, in-
take time, dose change, and time of last dose. The extended medExtractR_tapering implementa-
tion is intended to extract dosing information for more tapering sched-
ules, which are far more complex. The tapering extension allows for the extraction of addi-
tional entities including dispense amount, refills, dose schedule, time keyword, transition,
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Medication Extraction With R

Description

Provides a function medExtractR for extracting dose attributes for medications within a given electronic health record (EHR) note.

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Examples

```r
note1 <- "Progrf Oral Capsule 1 mg 3 capsules by mouth twice a day - last dose at 10pm"
medExtractR(note1, c("prograf", "tacrolimus"), 60, "mg", 2, lastdose=TRUE)
medExtractR(note2, c("lamotrigine", "ltg"), 130, "mg", 1, strength_sep = "-")
```
Description

A dictionary with additional expressions that can be used to supplement the drug_list argument of medExtractR and medExtractR_tapering.

Usage

```r
addl_expr
```

Format

A data frame with the following variables:

- **expr** A character vector, additional optional expressions for the drug_list argument.
- **type** A character vector, what category the expression belongs to (e.g., symptom, lab name, medication abbreviation, or drug class).

Examples

```r
data(addl_expr)
```

---

**dosechange_vals** Keywords Specifying Dose Change

Description

A dictionary of words indicating a dose change, meaning that the associated drug regimen may not be current. This includes phrases such as increase, reduce, or switch. In the following example of clinical text, the word ‘increase’ represents a dose change keyword: “Increase prograf to 5mg bid.”

Usage

```r
dosechange_vals
```

Format

A data frame with dose change expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as dose change.

Examples

```r
data(dosechange_vals)
```
### doseschedule_vals

**Description**

A dictionary with words for indicating a tapering dosing schedule. These can explicitly refer to such a schedule with phrases like "tapering" or "wean". It also includes words indicating an alternating dose schedule (e.g., "alternate", "alt.", "even days", or "odd days") as well as stopping keywords indicating the patient is going completely off the medication (e.g., "done", "gone", "stop", "discontinue").

**Usage**

doseschedule_vals

**Format**

A data frame with dose schedule expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as dose schedule.

**Examples**

data(doseschedule_vals)

### duration_vals

**Description**

A dictionary with phrases indicating how long the patient should take a particular dose of the drug. Examples of duration expressions include "2 weeks", "14 days", "another 3 days", "through mid-April", or a specific date. The form of each duration is given as a regular expression.

**Usage**

duration_vals

**Format**

A data frame with duration expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as duration.

**Examples**

data(duration_vals)
extract_entities

Extract Medication Entities From Phrase

Description

This function searches a phrase for medication dosing entities of interest. It is called within medExtractR and generally not intended for use outside that function. The phrase argument containing text to search corresponds to an individual mention of the drug of interest.

Usage

extract_entities(
    phrase,
    p_start,
    p_stop,
    unit,
    frequency_fun = NULL,
    intaketime_fun = NULL,
    duration_fun = NULL,
    route_fun = NULL,
    strength_sep = NULL,
    ...
)

Arguments

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>phrase</td>
<td>Text to search.</td>
</tr>
<tr>
<td>p_start</td>
<td>Start position of phrase within original text.</td>
</tr>
<tr>
<td>p_stop</td>
<td>End position of phrase within original text.</td>
</tr>
<tr>
<td>unit</td>
<td>Unit of measurement for medication strength, e.g. ‘mg’.</td>
</tr>
<tr>
<td>frequency_fun</td>
<td>Function used to extract frequency.</td>
</tr>
<tr>
<td>intaketime_fun</td>
<td>Function used to extract intake time.</td>
</tr>
<tr>
<td>duration_fun</td>
<td>Function used to extract duration.</td>
</tr>
<tr>
<td>route_fun</td>
<td>Function used to extract route.</td>
</tr>
<tr>
<td>strength_sep</td>
<td>Delimiter for contiguous medication strengths.</td>
</tr>
</tbody>
</table>
| ...               | Parameter settings used in extracting frequency and intake time, including additional arguments to the <entity>_fun arguments. Use frequency_dict, intaketime_dict, duration_dict, and route_dict to identify custom frequency, intake time, duration, and route dictionaries, respectively.
Details

Various medication dosing entities are extracted within this function including the following:

- **strength**: The amount of drug in a given dosage form (i.e., tablet, capsule).
- **dose amount**: The number of tablets, capsules, etc. taken at a given intake time.
- **dose strength**: The total amount of drug given intake. This quantity would be equivalent to strength \times dose amount, and appears similar to strength when dose amount is absent.
- **frequency**: The number of times per day a dose is taken, e.g., ‘once daily’ or ‘2x/day’.
- **intake time**: The time period of the day during which a dose is taken, e.g., ‘morning’, ‘lunch’, ‘in the pm’.
- **duration**: How long a patient is on a drug regimen, e.g., ‘2 weeks’, ‘mid-April’, ‘another 3 days’.
- **route**: The administration route of the drug, e.g., ‘by mouth’, ‘IV’, ‘topical’.

Note that extraction of the entities drug name, dose change, and time of last dose are not handled by the extract_entities function. Those entities are extracted separately and appended to the extract_entities output within the main medExtractR function. Strength, dose amount, and dose strength are primarily numeric quantities, and are identified using a combination of regular expressions and rule-based approaches. Frequency, intake time, route, and duration, on the other hand, use dictionaries for identification.

By default and when an argument `<entity>_fun` is `NULL`, the extract_generic function will be used to extract that entity. This function can also inherit user-defined entity dictionaries, supplied as arguments `<entity>_dict` to medExtractR or medExtractR_tapering (see documentation files for main function(s) for details).

The strength_sep argument is `NULL` by default, but can be used to identify shorthand for morning and evening doses. For example, consider the phrase “Lamotrigine 300-200” (meaning 300 mg in the morning and 200 mg in the evening). The argument `strength_sep = ‘-‘` identifies the full expression 300-200 as dose strength in this phrase.

Value

data.frame with entities information. At least one row per entity is returned, using NA when no expression was found for a given entity.

The “entity” column of the output contains the formatted label for that entity, according to the following mapping.

- strength: “Strength”
- dose amount: “DoseAmt”
- dose strength: “DoseStrength”
- frequency: “Frequency”
- intake time: “IntakeTime”
- duration: “Duration”
- route: “Route”

Sample output for the phrase “Lamotrigine 200mg bid” would look like:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
</tr>
</thead>
<tbody>
<tr>
<td>IntakeTime</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>Strength</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>DoseAmt</td>
<td>&lt;NA&gt;</td>
</tr>
</tbody>
</table>
**extract_entities_tapering**

<table>
<thead>
<tr>
<th>Route</th>
<th>&lt;NA&gt;</th>
</tr>
</thead>
<tbody>
<tr>
<td>Duration</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>Frequency</td>
<td>bid;19:22</td>
</tr>
<tr>
<td>DoseStrength</td>
<td>200mg;13:18</td>
</tr>
</tbody>
</table>

**Examples**

```r
note <- "Lamotrigine 25 mg tablet - 3 tablets oral twice daily"
extract_entities(note, 1, nchar(note), "mg")
# A user-defined dictionary can be used instead of the default
my_dictionary <- data.frame(c("daily", "twice daily"))
extract_entities(note, 1, 53, "mg", frequency_dict = my_dictionary)
```

**Description**

This function searches a phrase for medication dosing entities of interest. It is called within `medExtractR_tapering` and generally not intended for use outside that function.

**Usage**

```r
extract_entities_tapering(
  phrase,
  p_start,
  d_stop,
  unit,
  frequency_fun = NULL,
  intaketime_fun = NULL,
  duration_fun = NULL,
  route_fun = NULL,
  doseschedule_fun = NULL,
  preposition_fun = NULL,
  timekeyword_fun = NULL,
  transition_fun = NULL,
  dosechange_fun = NULL,
  strength_sep = NULL,
  ...
)
```

**Arguments**

- **phrase**: Text to search.
- **p_start**: Start position of phrase within original text.
Details

Various medication dosing entities are extracted within this function including the following:

- **strength**: The amount of drug in a given dosage form (i.e., tablet, capsule).
- **dose amount**: The number of tablets, capsules, etc. taken at a given intake time.
- **dose strength**: The total amount of drug given intake. This quantity would be equivalent to strength \( \times \) dose amount, and appears similar to strength when dose amount is absent.
- **frequency**: The number of times per day a dose is taken, e.g., “once daily” or ‘2x/day’.
- **intaketime**: The time period of the day during which a dose is taken, e.g., ‘morning’, ‘lunch’, ‘in the pm’.
- **duration**: How long a patient is on a drug regimen, e.g., ‘2 weeks’, ‘mid-April’, ‘another 3 days’.
- **route**: The administration route of the drug, e.g., ‘by mouth’, ‘IV’, ‘topical’.
- **dose change**: Whether the dosage of the drug was changed, e.g., ‘increase’, ‘adjust’, ‘reduce’.
- **dose schedule**: Keywords which represent special dosing regimens, such as tapering schedules, alternating doses, or stopping keywords, e.g., ‘weaning’, ‘even days’ or ‘odd_days’, ‘discontinue’.
- **time keyword**: Whether the dosing regimen is a past dose, current dose, or future dose, e.g., ‘currently’, ‘remain’, ‘yesterday’.
- **transition**: Words or symbols that link consecutive doses of a tapering regimen, e.g., ‘then’, ‘followed by’, or a comma ‘,’.
- **preposition**: Prepositions that occur immediately next to another identified entity, e.g., ‘to’, ‘until’, ‘for’.
- **dispense amount**: The number of pills prescribed to the patient.
- **refill**: The number of refills allowed for the patient’s prescription.
extract_entities_tapering

Similar to the basic implementation, drug name and and time of last dose are not handled by the extract_entities_tapering function. Those entities are extracted separately and appended to the extract_entities_tapering output within the main medExtractR_tapering function. In the tapering extension, however, dose change is treated the same as other dictionary-based entities and extracted within extract_entities_tapering. Strength, dose amount, dose strength, dispense amount, and refill are primarily numeric quantities, and are identified using a combination of regular expressions and rule-based approaches. All other entities use dictionaries for identification. For more information about the default dictionary for a specific entity, view the documentation file for the object <entity>_vals.

By default and when an argument <entity>_fun is NULL, the extract_generic function will be used to extract that entity. This function can also inherit user-defined entity dictionaries for each entity, supplied as arguments <entity>_dict to medExtractR or medExtractR_tapering (see documentation files for main function(s) for details).

Note that extract_entities_tapering has the argument d_stop. This differs from extract_entities, which uses the end position of the full search window. This is a consequence of medExtractR using a fixed search window length and medExtractR_tapering dynamically constructing a search window.

**Value**

data.frame with entities information. At least one row per entity is returned, using NA when no expression was found for a given entity.

The “entity” column of the output contains the formatted label for that entity, according to the following mapping:

- strength: “Strength”
- dose amount: “DoseAmt”
- dose strength: “DoseStrength”
- frequency: “Frequency”
- intake time: “IntakeTime”
- duration: “Duration”
- route: “Route”
- dose change: “DoseChange”
- dose schedule: “DoseScheule”
- time keyword: “TimeKeyword”
- transition: “Transition”
- preposition: “Preposition”
- dispense amount: “DispenseAmt”
- refill: “Refill”

Sample output for the phrase “Lamotrigine 200mg bid for 14 days” would look like:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
</tr>
</thead>
<tbody>
<tr>
<td>IntakeTime</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>Strength</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>DoseAmt</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>DoseChange</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>DoseSchedule</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>TimeKeyword</td>
<td>&lt;NA&gt;</td>
</tr>
<tr>
<td>Transition</td>
<td>&lt;NA&gt;</td>
</tr>
</tbody>
</table>
Preposition       <NA>
DispenseAmt       <NA>
Refill            <NA>
Frequency         bid;19:22
DoseStrength      200mg;13:18
Preposition       for;23:26
Duration          14 days;27:34

Examples

```r
note <- "prednisone 20mg daily tapering to 5mg daily over 2 weeks"
extract_entities_tapering(note, 1, 11, "mg")
# A user-defined dictionary can be used instead of the default
my_dictionary <- data.frame(c("daily", "twice daily"))
extract_entities(note, 1, 11, "mg", frequency_dict = my_dictionary)
```

**extract_generic**  
*Extract Generic Entities From Phrase*

**Description**

This function searches a phrase for the position and length of expressions specified in a dictionary. This is called within other main functions of the package and generally not intended for use on its own.

**Usage**

```r
extract_generic(phrase, dict)
```

**Arguments**

- `phrase`: Text to search.
- `dict`: data.frame, the first column should contain expressions to find. These can be regular expressions or exact phrases.

**Details**

`extract_generic` is used to extract entities that are identified with an associated dictionary of phrases or regular expressions, such as dose change, frequency, intake time, route, or duration in `medExtractR` and `medExtractR_tapering`, as well as dose schedule, time keyword, transition, and preposition in `medExtractR_tapering`. This function is called within `extract_entities`.

**Value**

A numeric matrix with position and expression length.
extract_lastdose

Examples

data(frequency_vals)
extract_generic("take two every day", dict = frequency_vals)
extract_generic("take two every morning",
    dict = data.frame(c("morning", "every morning")))

extract_lastdose Extract Last Dose Time From Phrase

Description

This function searches a phrase for the expression and position of the time at which the last dose of a drug was taken. It is called within medExtractR and generally not intended for use outside that function.

Usage

extract_lastdose(phrase, p_start, d_start, d_stop, time_exp = "default")

Arguments

phrase Text to search.
p_start Start position of phrase in the overall text (e.g., the full clinical note).
d_start Start position of drug name in larger text.
d_stop End position of drug name in larger text.
time_exp Vector of regular expressions to identify time expressions.

Details

This function identifies the time at which the last dose of a drug of interest was taken. The arguments p_start, d_start, and d_stop represent global start or stop positions for the phrase or drug. These arguments are used to determine the position of any found last dose time expressions relative to the overall clinical note, not just within phrase.

The time_exp argument contains regular expressions for numeric or text representations of last dose time. See time_regex for more information about the default regular expressions used in medExtractR.

Value

data.frame with last dose time entity information. This output format is consistent with the output of extract_entities, and the formatted label for the time of last dose entity is "LastDose.”
Sample output for the phrase “Last prograf at 5pm” would look like:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
</tr>
</thead>
<tbody>
<tr>
<td>LastDose</td>
<td>5pm;17:20</td>
</tr>
</tbody>
</table>
Examples

# Suppose this phrase begins at character 120 in the overall clinical note
extract_lastdose("took aspirin last night at 8pm", p_start = 120,
            d_start = 125, d_stop = 131)

---

**frequency_vals**

**Keywords Specifying Frequency**

**Description**

A dictionary mapping frequency expressions to numeric values representing the corresponding number of doses per day. Example expressions include "q12 hours", "bid", "daily", and "three times a day". The form of each frequency is given as a regular expression.

**Usage**

frequency_vals

**Format**

A data frame with frequency expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as frequency.
- **value** A numeric vector, numeric value of frequency represented as number of doses taken per day. For example, “bid” and “twice a day” would both have a numeric value of 2.

**Examples**

data(frequency_vals)

---

**intaketime_vals**

**Keywords Specifying Intake Time**

**Description**

A dictionary with intake time expressions representing the approximate time of day when a dose should be taken. Example expressions include "in the morning", "with lunch", "at bedtime", and "qpm". The form of each intake time is given as a regular expression.

**Usage**

intaketime_vals
**medExtractR**

**Format**

A data frame with intake time expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as intake time.

**Examples**

```r
data(intaketime_vals)
```

---

**Description**

This function identifies medication entities of interest and returns found expressions with start and stop positions.

**Usage**

```r
medExtractR(
  note,  
  drug_names, 
  window_length, 
  unit, 
  max_dist = 0, 
  drug_list = "rxnorm", 
  lastdose = FALSE, 
  lastdose_window_ext = 1.5, 
  strength_sep = NULL, 
  flag_window = 30, 
  dosechange_dict = "default", 
  ...
)
```

**Arguments**

- **note** Text to search.
- **drug_names** Vector of drug names of interest to locate.
- **window_length** Length (in number of characters) of window after drug in which to look.
- **unit** Strength unit to look for (e.g., 'mg').
- **max_dist** Numeric - edit distance to use when searching for drug_names.
- **drug_list** Vector of known drugs that may end search window. By default calls `rxnorm_druglist`.
- **lastdose** Logical - whether or not last dose time entity should be extracted.
- **lastdose_window_ext** Numeric - multiplicative factor by which `window_length` should be extended when identifying last dose time.
**strength_sep**  Delimiter for contiguous medication strengths (e.g., ‘-’ for “LTG 200-300”).

**flag_window**  How far around drug (in number of characters) to look for dose change keyword - default fixed to 30. See ‘Details’ section below for further explanation.

**dosechange_dict**  List of keywords used to determine if a dose change entity is present.

... Parameter settings used in extracting frequency, intake time, route, and duration. Potentially useful parameters include freq_dict, intaketime_dict, route_dict, and duration_dict (see ... argument in extract_entities) to specify frequency or intake time dictionaries, as well as ‘freq_fun’, ‘intake-time_fun’, ‘route_fun’, and ‘duration_fun’ for user-specified extraction functions. If no additional arguments are provided, medExtractR_tapering will use extract_generic and the default dictionary for each entity. See extract_entities documentation for details.

**Details**

This function uses a combination of regular expressions, rule-based approaches, and dictionaries to identify various drug entities of interest. Specific medications to be found are specified with drug_names, which is not case-sensitive or space-sensitive (e.g., ‘lamotrigine XR’ is treated the same as ‘lamotrigineXR’). Entities to be extracted include drug name, strength, dose amount, dose, frequency, intake time, route, duration, and time of last dose. See extract_entities and extract_lastdose for more details.

When searching for medication names of interest, fuzzy matching may be used. The max_dist argument determines the maximum edit distance allowed for such matches. If using fuzzy matching, any drug name with less than 5 characters will only allow an edit distance of 1, regardless of the value of max_dist.

The purpose of the drug_list argument is to reduce false positives by removing information that is likely to be related to a competing drug, not our drug of interest. By default, this is “rxnorm” which calls data(rxnorm_druglist). A custom drug list in the form of a character string can be supplied instead, or can be appended to rxnorm_druglist by specifying drug_list = c("rxnorm",custom_drug_list). medExtractR then uses this list to truncate the search window at the first appearance of an unrelated drug name. This uses publicly available data courtesy of the U.S. National Library of Medicine (NLM), National Institutes of Health, Department of Health and Human Services; NLM is not responsible for the product and does not endorse or recommend this or any other product. See rxnorm_druglist documentation for details.

Most medication entities are searched for in a window after the drug. The dose change entity, or presence of a keyword to indicate a non-current drug regimen, may occur before the drug name. The flag_window argument adjusts the width of the pre-drug window. Both flag_window and dosechange_dict are not default arguments to the extended function medExtractR_tapering since that extension uses a more flexible search window and extraction procedure. In the tapering extension, entity extraction is more flexible, and any entity can be extracted either before or after the drug mention. Thus functionality for dose change identification is identical to all other dictionary-based entities.

The strength_sep argument is NULL by default, but can be used to identify shorthand for morning and evening doses. For example, consider the phrase ‘Lamotrigine 300-200’ (meaning 300 mg in the morning and 200 mg in the evening). The argument strength_sep = ‘-’ identifies the full expression 300-200 as dose strength in this phrase.
Value

data.frame with entity information. Only extractions from found entities are returned. If no dosing information for the drug of interest is found, the following output will be returned:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
<th>pos</th>
</tr>
</thead>
<tbody>
<tr>
<td>NA</td>
<td>NA</td>
<td>NA</td>
</tr>
</tbody>
</table>

The “entity” column of the output contains the formatted label for that entity, according to the following mapping:
drug name: “DrugName”
strength: “Strength”
dose amount: “DoseAmt”
dose strength: “DoseStrength”
frequency: “Frequency”
intake time: “IntakeTime”
duration: “Duration”
route: “Route”
dose change: “DoseChange”
time of last dose: “LastDose”

Sample output:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
<th>pos</th>
</tr>
</thead>
<tbody>
<tr>
<td>DoseChange</td>
<td>decrease</td>
<td>66:74</td>
</tr>
<tr>
<td>DrugName</td>
<td>Prograf</td>
<td>78:85</td>
</tr>
<tr>
<td>Strength</td>
<td>2 mg</td>
<td>86:90</td>
</tr>
<tr>
<td>DoseAmt</td>
<td>1</td>
<td>91:92</td>
</tr>
<tr>
<td>Route</td>
<td>by mouth</td>
<td>100:108</td>
</tr>
<tr>
<td>Frequency</td>
<td>bid</td>
<td>109:112</td>
</tr>
<tr>
<td>LastDose</td>
<td>2100</td>
<td>129:133</td>
</tr>
</tbody>
</table>

References


Examples

```r
note1 <- "Progrf Oral Capsule 1 mg 3 capsules by mouth twice a day - last dose at 10pm"
medExtractR(note1, c("prograf", "tacrolimus"), 60, "mg", 2, lastdose=TRUE)
note2 <- "Currently on lamotrigine 150-200, but will increase to lamotrigine 200mg bid"
medExtractR(note2, c("lamotrigine", "ltg"), 130, "mg", 1, strength_sep = "-")
```
medExtractR\_tapering Extract Medication Entities From Clinical Note - Extension of medExtractR for Tapering applications

Description
This function identifies medication entities of interest and returns found expressions with start and stop positions.

Usage
medExtractR\_tapering(
  note,  
  drug\_names,  
  unit,  
  max\_dist = 0,  
  drug\_list = "rxnorm",  
  lastdose = FALSE,  
  strength\_sep = NULL,  
  ...  
)

Arguments

  note           Text to search.  
  drug\_names    Vector of drug names of interest to locate.  
  unit           Strength unit to look for (e.g., 'mg').  
  max\_dist      Numeric - edit distance to use when searching for drug\_names.  
  drug\_list     Vector of known drugs that may end search window. By default calls rxnorm\_druglist. Can be supplemented with expressions in addl\_expr.  
  lastdose       Logical - whether or not last dose time entity should be extracted. See ‘Details’ section below for more information.  
  strength\_sep   Delimiter for contiguous medication strengths (e.g., '-' for “LTG 200-300”).  
  ...            Parameter settings used in dictionary-based entities. For each dictionary-based entity, the user can supply the optional arguments <entity>_fun and <entity>_dict to provide custom extraction functions and dictionaries, respectively. If no additional arguments are provided, medExtractR\_tapering will use extract\_generic and the default dictionary for each entity. See extract\_entities\_tapering documentation for details.

Details
This function uses a combination of regular expressions, rule-based approaches, and dictionaries to identify various drug entities of interest, with a particular focus on drugs administered with a tapering schedule. Specific medications to be found are specified with drug\_names, which is not case-sensitive or space-sensitive (e.g., ‘lamotrigine XR’ is treated the same as ‘lamotrigineXR’). Entities
to be extracted include drug name, strength, dose amount, dose strength, frequency, intake time, route, duration, dose schedule, time keyword, preposition, transition, dispense amount, refill, and time of last dose. While it is still an optional entity in medExtractR_tapering, if lastdose=TRUE then medExtractR_tapering will search for time of last dose in the same search window used for all other entities. As a result, there is no need for the lastdose_window_ext argument. See extract_entities_tapering and extract_lastdose for more details.

When searching for medication names of interest, fuzzy matching may be used. The max_dist argument determines the maximum edit distance allowed for such matches. If using fuzzy matching, any drug name with less than 7 characters will force an exact match, regardless of the value of max_dist. The default value of 7 was selected based on a set of training notes for the drug prednisone, and differs slightly from the default values of 5 for medExtractR. The tapering extension does not use the window_length argument to define the search window, since tapering schedules can be much longer than a static regimens. Instead, medExtractR_tapering dynamically generates the search window based on competing drug names or phrases, and the distance between consecutive entities. The strength_sep argument is NULL by default, and operates in the same manner as it does in medExtractR.

By default, the drug_list argument is “rxnorm” which calls data(rxnorm_druglist). A custom drug list in the form of a character string can be supplied instead, or can be appended to rxnorm_druglist by specifying drug_list = c("rxnorm",custom_drug_list). This uses publicly available data courtesy of the U.S. National Library of Medicine (NLM), National Institutes of Health, Department of Health and Human Services; NLM is not responsible for the product and does not endorse or recommend this or any other product. See rxnorm_druglist documentation for details.

Value
data.frame with entity information. If no dosing information for the drug of interest is found, the following output will be returned:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
<th>pos</th>
</tr>
</thead>
<tbody>
<tr>
<td>NA</td>
<td>NA</td>
<td>NA</td>
</tr>
</tbody>
</table>

The “entity” column of the output contains the formatted label for that entity, according to the following mapping.

- drug name: “DrugName”
- strength: “Strength”
- dose amount: “DoseAmt”
- dose strength: “DoseStrength”
- frequency: “Frequency”
- intake time: “IntakeTime”
- duration: “Duration”
- route: “Route”
- dose change: “DoseChange”
- dose schedule: “DoseSchedule”
- time keyword: “TimeKeyword”
- transition: “Transition”
- preposition: “Preposition”
dispense amount: “DispenseAmt”
refill: “Refill”
time of last dose: “LastDose”
Sample output:

<table>
<thead>
<tr>
<th>entity</th>
<th>expr</th>
<th>pos</th>
</tr>
</thead>
<tbody>
<tr>
<td>DoseChange</td>
<td>decrease</td>
<td>66:74</td>
</tr>
<tr>
<td>DrugName</td>
<td>Prograf</td>
<td>78:85</td>
</tr>
<tr>
<td>Strength</td>
<td>2 mg</td>
<td>86:90</td>
</tr>
<tr>
<td>DoseAmt</td>
<td>1</td>
<td>91:92</td>
</tr>
<tr>
<td>Frequency</td>
<td>bid</td>
<td>101:104</td>
</tr>
<tr>
<td>LastDose</td>
<td>2100</td>
<td>121:125</td>
</tr>
</tbody>
</table>

References


Description

A dictionary with preposition expressions. Such expressions often represent a relationship with an adjacent entity. Since most expressions in this dictionary are very short, we require word boundaries (any character other than a letter or number) to appear on either side of the expression. Example expressions include “for”, “to”, “until”, and “in”.

Usage

preposition_vals

Format

A data frame with preposition expressions (exact and/or regular expressions).

expr A character vector, expressions to consider as preposition.

Examples

data(preposition_vals)
### route_vals

**Keywords Specifying Route**

**Description**
A dictionary mapping route expressions to standardized forms, specifying the way in which a medication is administered. Example expressions include "oral", "topical", "IV", and "intravenous".

**Usage**
```
route_vals
```

**Format**
A data frame with route expressions (exact and/or regular expressions).

- `expr` A character vector, expressions to consider as route.
- `value` A standardized version of the raw expression. For example, "orally" and "by mouth" both have the standardized form "orally".

**Examples**
```
data(route_vals)
```

### rxnorm_druglist

**List of Medications**

**Description**
A dictionary that contains a vector of medication names, primarily derived from RxNorm.

**Usage**
```
rxnorm_druglist
```

**Format**
A vector with character strings for competing drug names.

**Details**
RxNorm is provided by the U.S. National Library of Medicine. This dictionary uses the February 1, 2021 RxNorm files directly downloaded from [https://www.nlm.nih.gov/research/umls/rxnorm/docs/rxnormfiles.html](https://www.nlm.nih.gov/research/umls/rxnorm/docs/rxnormfiles.html).

This list contains ingredient and brand names, cleaned to remove expressions that likely are ambiguous (e.g., ‘today’ or ‘date’). This product uses publicly available data courtesy of the U.S. National Library of Medicine (NLM), National Institutes of Health, Department of Health and Human Services; NLM is not responsible for the product and does not endorse or recommend this or any other product.
References

Examples

data(rxnorm_druglist)

timekeyword_vals

<table>
<thead>
<tr>
<th>expr</th>
<th>Keywords Specifying Time Keyword</th>
</tr>
</thead>
</table>

Description
A dictionary with time keyword expressions representing whether the dosing regimen is past, current, or future. Example expressions include "currently", "remain","not taking", "yesterday", and "past".

Usage
timekeyword_vals

Format
A data frame with time keyword expressions (exact and/or regular expressions).

expr A character vector, expressions to consider as time keyword.

Examples
data(timekeyword_vals)

<table>
<thead>
<tr>
<th>expr</th>
<th>Keywords Specifying Time Expressions</th>
</tr>
</thead>
</table>

Description
A vector of regular expressions to identify different forms of time expressions for last dose time. These are the default values used in link(extract_lastdose).

Usage
time_regex
transition_vals

Format
A vector with 5 regular expressions for the following categories.

- **am/pm** Time is indicated by the presence of ‘am’ or ‘pm’ following a numeric expression.
- **military** Time is given in military time, for unambiguous times of 13:00-23:59.
- **qualifier_after** Am/pm indication is implicit through a qualifying term like ‘last night’ or ‘this morning’. The qualifier occurs after the time, e.g., ‘10 last night.’
- **qualifier_before** Am/pm indication is implicit through a qualifying term like ‘last night’ or ‘this morning’. The qualifier occurs before the time, e.g., ‘last night at 10.’
- **duration** Time (in hours) between the last dose and most recent lab value

Details
Certain expressions which might be considered ambiguous are excluded from the regular expressions presented here. For instance, expressions such as ‘600’ could refer to either 6am or 6pm.

Examples
```
data(time_regex)
```

<table>
<thead>
<tr>
<th>transition_vals</th>
<th>Keywords/Symbols Specifying Transition</th>
</tr>
</thead>
</table>

Description
A dictionary with transition symbols and expressions representing a break between consecutive doses within a tapering regimen. This dictionary includes the expressions "then" and "followed by", as well as the punctuation ",(?!\s?then)" or ";(?!\s?then)" (i.e., a comma or semicolon not followed by the word "then").

Usage
```
transition_vals
```

Format
A data frame with transition expressions (exact and/or regular expressions).

- **expr** A character vector, expressions to consider as transitions.

Examples
```
data(transition_vals)
```
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