

Package ‘r2dii.match’

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Title Tools to Match Corporate Lending Portfolios with Climate Data

Version 0.1.1

Description These tools implement in R a fundamental part of the software 'PACTA' (Paris Agreement Capital Transition Assessment), which is a free tool that calculates the alignment between financial portfolios and climate scenarios (<<https://2degrees-investing.org/>>). Financial institutions use 'PACTA' to study how their capital allocation decisions align with climate change mitigation goals. This package matches data from corporate lending portfolios to asset level data from market-intelligence databases (e.g. power plant capacities, emission factors, etc.). This is the first step to assess if a financial portfolio aligns with climate goals.

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crucial_lbk	<i>Crucial loanbook columns for match_name()</i>
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Description

This is a helper to select the minimum loanbook columns you need to run `match_name()`. Using more columns may use too much time and memory.

Usage

```
crucial_lbk()
```

Value

A character vector.

See Also

Other helpers: [prioritize_level\(\)](#)

Examples

```
crucial_lbk()
```

match_name	<i>Match a loanbook to asset-based company data (abcd) by the name_* columns</i>
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Description

match_name() scores the match between names in a loanbook dataset (columns can be name_direct_loantaker, name_intermediate_parent* and name_ultimate_parent) with names in an asset-based company data (column name_company). The raw names are first internally transformed, and aliases are assigned. The similarity between aliases in each of the loanbook and abcd is scored using [stringdist::stringsim\(\)](#).

Usage

```
match_name(
  loanbook,
  abcd,
  by_sector = TRUE,
  min_score = 0.8,
  method = "jw",
  p = 0.1,
  overwrite = NULL,
  ald = deprecated(),
  ...
)
```

Arguments

loanbook, abcd	data frames structured like r2dii.data::loanbook_demo and r2dii.data::abcd_demo .
by_sector	Should names only be compared if companies belong to the same sector?
min_score	A number between 0-1, to set the minimum score threshold. A score of 1 is a perfect match.
method	Method for distance calculation. One of <code>c("osa", "lv", "dl", "hamming", "lcs", "qgram", "cosine", "jaccard", "jw", "soundex")</code> . See stringdist::stringdist-metrics .
p	Prefix factor for Jaro-Winkler distance. The valid range for p is $0 \leq p \leq 0.25$. If $p=0$ (default), the Jaro-distance is returned. Applies only to <code>method='jw'</code> .
overwrite	A data frame used to overwrite the sector and/or name columns of a particular direct loantaker or ultimate parent. To overwrite only sector, the value in the name column should be NA and vice-versa. This file can be used to manually match loanbook companies to abcd.
ald	[Superseded] ald has been superseded by abcd.
...	Arguments passed on to stringdist::stringsim() .

Value

A data frame with the same groups (if any) and columns as loanbook, and the additional columns:

- `id_2dii` - an id used internally by `match_name()` to distinguish companies
- `level` - the level of granularity that the loan was matched at (e.g `direct_loantaker` or `ultimate_parent`)
- `sector` - the sector of the loanbook company
- `sector_abcd` - the sector of the abcd company
- `name` - the name of the loanbook company
- `name_abcd` - the name of the abcd company
- `score` - the score of the match (manually set this to 1 prior to calling `prioritize()` to validate the match)
- `source` - determines the source of the match. (equal to loanbook unless the match is from `overwrite`)

The returned rows depend on the argument `min_value` and the result of the column score for each loan: * If any row has score equal to 1, `match_name()` returns all rows where score equals 1, dropping all other rows. * If no row has score equal to 1, `match_name()` returns all rows where score is equal to or greater than `min_score`. * If there is no match the output is a 0-row tibble with the expected column names – for type stability.

Package options

`r2dii.match.sector_classifications`: Allows you to use your own `sector_classifications` instead of the default. This feature is experimental and may be dropped and/or become a new argument to `match_name()`.

Assigning aliases

The transformation process used to compare names between loanbook and abcd datasets applies best practices commonly used in name matching algorithms:

- Remove special characters.
- Replace language specific characters.
- Abbreviate certain names to reduce their importance in the matching.
- Spell out numbers to increase their importance.

Handling grouped data

This function ignores but preserves existing groups.

See Also

Other main functions: [prioritize\(\)](#)

Examples

```

library(r2dii.data)
library(tibble)

# Small data for examples
loanbook <- head(loanbook_demo, 50)
abcd <- head(abcd_demo, 50)

match_name(loanbook, abcd)

match_name(loanbook, abcd, min_score = 0.9)

# Use your own `sector_classifications`
your_classifications <- tibble(
  sector = "power",
  borderline = FALSE,
  code = "3511",
  code_system = "XYZ"
)

restore <- options(r2dii.match.sector_classifications = your_classifications)

loanbook <- tibble(
  sector_classification_system = "XYZ",
  sector_classification_direct_loantaker = "3511",
  id_ultimate_parent = "UP15",
  name_ultimate_parent = "Alpine Knits India Pvt. Limited",
  id_direct_loantaker = "C294",
  name_direct_loantaker = "Yuamen Xinneng Thermal Power Co Ltd"
)

abcd <- tibble(
  name_company = "alpine knits india pvt. limited",
  sector = "power"
)

match_name(loanbook, abcd)

# Cleanup
options(restore)

```

prioritize

Pick rows where score is 1 and level per loan is of highest priority

Description

When multiple perfect matches are found per loan (e.g. a match at `direct_loantaker` level and `ultimate_parent` level), we must prioritize the desired match. By default, the highest priority is the most granular match (i.e. `direct_loantaker`).

Usage

```
prioritize(data, priority = NULL)
```

Arguments

<code>data</code>	A data frame like the validated output of <code>match_name()</code> . See <i>Details</i> on how to validate data.
<code>priority</code>	One of: <ul style="list-style-type: none"> • <code>NULL</code>: defaults to the default level priority as returned by <code>prioritize_level()</code>. • A character vector giving a custom priority. • A function to apply to the output of <code>prioritize_level()</code>, e.g. <code>rev</code>. • A quosure-style lambda function, e.g. <code>~ rev(.x)</code>.

Details**How to validate data**

Write the output of `match_name()` into a `.csv` file with:

```
# Writing to current working directory
matched %>%
  readr::write_csv("matched.csv")
```

Compare, edit, and save the data manually:

- Open *matched.csv* with any spreadsheet editor (Excel, Google Sheets, etc.).
- Compare the columns `name` and `name_abcd` manually to determine if the match is valid. Other information can be used in conjunction with just the names to ensure the two entities match (sector, internal information on the company structure, etc.)
- Edit the data:
 - If you are happy with the match, set the `score` value to 1.
 - Otherwise set or leave the `score` value to anything other than 1.
- Save the edited file as, say, *valid_matches.csv*.

Re-read the edited file (validated) with:

```
# Reading from current working directory
valid_matches <- readr::read_csv("valid_matches.csv")
```

Value

A data frame with a single row per loan, where `score` is 1 and priority level is highest.

Handling grouped data

This function ignores but preserves existing groups.

See Also

[match_name\(\)](#), [prioritize_level\(\)](#).

Other main functions: [match_name\(\)](#)

Examples

```
library(dplyr)

# styler: off
matched <- tribble(
  ~sector, ~sector_abcd, ~score, ~id_loan, ~level,
  "coal", "coal", 1, "aa", "ultimate_parent",
  "coal", "coal", 1, "aa", "direct_loantaker",
  "coal", "coal", 1, "bb", "intermediate_parent",
  "coal", "coal", 1, "bb", "ultimate_parent",
)
# styler: on

prioritize_level(matched)

# Using default priority
prioritize(matched)

# Using the reverse of the default priority
prioritize(matched, priority = rev)

# Same
prioritize(matched, priority = ~ rev(.x))

# Using a custom priority
bad_idea <- c("intermediate_parent", "ultimate_parent", "direct_loantaker")

prioritize(matched, priority = bad_idea)
```

prioritize_level	<i>Arrange unique level values in default order of priority</i>
------------------	---

Description

Arrange unique level values in default order of priority

Usage

```
prioritize_level(data)
```

Arguments

data A data frame, commonly the output of [match_name\(\)](#).

Value

A character vector of the default level priority per loan.

See Also

Other helpers: [crucial_lbk\(\)](#)

Examples

```
matched <- tibble::tibble(  
  level = c(  
    "intermediate_parent_1",  
    "direct_loantaker",  
    "direct_loantaker",  
    "direct_loantaker",  
    "ultimate_parent",  
    "intermediate_parent_2"  
  )  
)  
prioritize_level(matched)
```


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