

Package ‘ccoptimalmatch’

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Type Package

Title Implementation of Case-Control Optimal Matching

Version 0.1.0

Description

Cases are matched to controls in an efficient, optimal and computationally flexible way. It uses the idea of sub-sampling in the level of the case, by creating pseudo-observations of controls. The user can select between replacement and without replacement, the number of controls, and several covariates to match upon. See Mamouris (2021) <doi:10.1186/s12874-021-01256-3> for an overview.

Depends R (>= 2.10)

License GPL-2

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Imports dplyr, rlang

Suggests knitr, rmarkdown

VignetteBuilder knitr

NeedsCompilation no

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being_processed *Data for matching cases with controls*

Description

A dataset containing cases and controls using the Intego registry data. The variables are as follows:

Usage

```
data(being_processed)
```

Format

A data frame with 77110 rows and 11 variables

Details

- cluster_case: each case forms a cluster with all possible controls to be matched
- Patient_Id: Unique identifier for each patient
- case_control: binary, if case==Colorectal Cancer, else control
- case_ind: binary, if 1==case, else control
- JCG: Year of Contact
- entry_year: the year that the patient first entered the database
- CI: Comorbidity Index. Count of chronic diseases before index data
- age_diff: difference of age between cases and controls
- fup_diff: difference of follow-up between cases and controls
- total_control_per_case: total controls that are available to be pooled per case
- freq_of_controls: how many times the control is available to be matched for different cases

ccoptimalmatch *ccoptimalmatch: Optimal Case Control matching*

Description

Fast and optimal matching for cases and controls

Author(s)

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not_processed	<i>Not-processed data for matching cases with controls</i>
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Description

A dataset containing cases and controls using the Intego registry data. But not the final dataset. The variables are as follows:

Usage

```
data(not_processed)
```

Format

A data frame with 656506 rows and 9 variables

Details

- Patient_Id: Unique identifier for each patient
- JCG: Year of Contact
- Birth_Year: Patient's year of birth
- Gender: Patient's Gender
- Practice_Id: Patient's general practice
- case_control: binary, if case==Colorectal Cancer, else control
- entry_year: the year that the patient first entered the database
- fup_diff: difference of follow-up between cases and controls
- CI: Comorbidity Index. Count of chronic diseases before index data

optimal_matching	<i>optimal_matching</i>
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Description

optimal_matching is performing the optimal match between cases and controls in an iterative way and computational efficient way

Usage

```
optimal_matching(  
  total_database,  
  n_con,  
  cluster_var,  
  Id_Patient,  
  total_cont_per_case,  
  case_control,  
  with_replacement = FALSE  
)
```

Arguments

`total_database` a data frame that contains the cases and controls

`n_con` number of controls to be matched

`cluster_var` a variable that contains one case with all available controls to be pooled

`Id_Patient` Id of the patient

`total_cont_per_case`
total number of controls that are available for each case

`case_control` a variable containing "case" and "control"

`with_replacement`
Use replacement or not

Details

Here is where I should put all my details. This is where I should give more examples if necessary

Value

a data frame containing the cases and the corresponding number of controls

Examples

```
optimal_matching(being_processed, n_con=2, cluster_var=cluster_case,  
Id_Patient=Patient_Id, total_cont_per_case=total_control_per_case, case_control = case_control)
```

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