

# Package ‘ccoptimalmatch’

July 22, 2025

**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](https://doi.org/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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**Repository** CRAN

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being\_processed      *Data for matching cases with controls*

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**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

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ccoptimalmatch      *ccoptimalmatch: Optimal Case Control matching*

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**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

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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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