

# Example session for Weight-based deduplication

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This document shows an example session using the package *RecordLinkage*. A single data set is deduplicated using an EM algorithm for weight calculation. Conducting linkage of two data sets differs only in the step of generating record pairs.

## 1 Generating record pairs

The data to be deduplicated is expected to reside in a data frame or matrix, each row containing one record. Example data sets of 500 and 10000 records are included in the package as `RLdata500` and `RLdata10000`.

```
data(RLdata500)
RLdata500[1:5,]

##   fname_c1 fname_c2 lname_c1 lname_c2   by bm bd
## 1  CARSTEN   <NA>    MEIER    <NA> 1949 7 22
## 2    GERD    <NA>    BAUER    <NA> 1968 7 27
## 3  ROBERT    <NA>  HARTMANN  <NA> 1930 4 30
## 4  STEFAN    <NA>    WOLFF    <NA> 1957 9  2
## 5    RALF    <NA>  KRUEGER    <NA> 1966 1 13
```

For deduplication, `compare.dedup` is to be used. In this example, blocking is set to return only record pairs which agree in at least two components of the subdivided date of birth, resulting in 810 pairs. The argument `identity` preserves the true matching status for later evaluation.

```
pairs=compare.dedup(RLdata500,identity=identity.RLdata500,
  blockfld=list(c(5,6),c(6,7),c(5,7)))
summary(pairs)

##
## Deduplication Data Set
##
## 500 records
## 571 record pairs
##
## 49 matches
## 522 non-matches
## 0 pairs with unknown status
```

## 2 Weight calculation

Weights are calculated by means of an EM algorithm. This step is computationally intensive and might take a while. The histogram shows the resulting weight distribution.

```
pairs=emWeights(pairs)
```

```
hist(pairs$Wdata, plot=FALSE)
```

```
## $breaks
## [1] -15 -10 -5 0 5 10 15 20 25 30 35 40 45
##
## $counts
## [1] 352 13 0 0 5 26 42 123 9 0 0 1
##
## $density
## [1] 0.1232924694 0.0045534151 0.0000000000 0.0000000000
## [5] 0.0017513135 0.0091068301 0.0147110333 0.0430823117
## [9] 0.0031523643 0.0000000000 0.0000000000 0.0003502627
##
## $mids
## [1] -12.5 -7.5 -2.5 2.5 7.5 12.5 17.5 22.5 27.5
## [10] 32.5 37.5 42.5
##
## $xname
## [1] "pairs$Wdata"
##
## $equidist
## [1] TRUE
##
## attr("class")
## [1] "histogram"
```

## 3 Classification

For determining thresholds, record pairs within a given range of weights can be printed using `getPairs`<sup>1</sup>. In this case, 24 is set as upper and -7 as lower threshold, dividing links, possible links and non-links. The summary shows the resulting contingency table and error measures.

```
getPairs(pairs,30,20)
```

```
##      id fname_c1 fname_c2 lname_c1 lname_c2   by bm bd
## 23 457   URSULA   BIRGIT  MUELLER    <NA> 1940 6 15
```

---

<sup>1</sup>The output of `getPairs` is shortened in this document.

```

## 24
## 25 467    ULRIKE    NICOLE    BECKRR    <NA> 1982  8  4
## 26 472    ULRIKE    NICOLE    BECKER    <NA> 1982  8  4
## 27
## 28 183    ULRICH      <NA>    MUELLER    <NA> 1962  6 19
## 29 444      SILKE      <NA>    MUELLER    <NA> 1962  6 14
## 30
## 31  25 MATTHIAS      <NA>      HAAS      <NA> 1955  7  8
## 32 107 MATTHIAS      <NA>      HAAS      <NA> 1955  8  8
## 33
## 34 106      ANDRE      <NA>    MUELLER    <NA> 1976  2 25
## 35 175      ANDRE      <NA>    MUELLER    <NA> 1976  1 25
## 36
##      Weight
## 23 25.14137
## 24
## 25
## 26 25.14137
## 27
## 28
## 29 24.20333
## 30
## 31
## 32 24.11923
## 33
## 34
## 35 24.11923
## 36

```

```

pairs=emClassify(pairs, threshold.upper=24, threshold.lower=-7)
summary(pairs)

```

```

##
## Deduplication Data Set
##
## 500 records
## 571 record pairs
##
## 49 matches
## 522 non-matches
## 0 pairs with unknown status
##
##
## Weight distribution:
##
## [-15,-10]  (-10,-5]    (-5,0]    (0,5]    (5,10]  (10,15]
##          352         13         0         0         5         26
##  (15,20]   (20,25]   (25,30]   (30,35]   (35,40]  (40,45]
##          42         123         9         0         0         1

```

```
##
## 15 links detected
## 198 possible links detected
## 358 non-links detected
##
## alpha error: 0.000000
## beta error: 0.002786
## accuracy: 0.997319
##
##
## Classification table:
##
##           classification
## true status  N   P   L
##      FALSE 358 163   1
##      TRUE   0  35  14
```

Review of the record pairs denoted as possible links is facilitated by `getPairs`, which can be forced to show only possible links via argument `show`. A list with the ids of linked pairs can be extracted from the output of `getPairs` with argument `single.rows` set to `TRUE`.

```
possibles <- getPairs(pairs, show="possible")
possibles[1:6,]

##      id  fname_c1 fname_c2 lname_c1 lname_c2   by bm bd
## 1  17 ALEXANDER      <NA>  MUELLER      <NA> 1974  9  9
## 2 193 CHRISTIAN      <NA>  MUELLER      <NA> 1974  8  9
## 3
## 4  61      ANDRE      <NA>  FISCHER      <NA> 1943  6 25
## 5 254  STEFANIE      <NA>  FISCHER      <NA> 1943 11 25
## 6
##           Weight
## 1
## 2 21.691086
## 3
## 4
## 5 21.691086
## 6

links=getPairs(pairs,show="links", single.rows=TRUE)
link_ids <- links[, c("id1", "id2")]
link_ids

##      id1 id2
## 290 290 466
##  50  50 234
##  87  87 117
## 145 145 240
## 286 286 383
```

```
## 289 289 399
## 297 297 388
## 357 357 414
## 313 313 457
## 467 467 472
## 183 183 444
## 25 25 107
## 106 106 175
## 370 370 478
## 127 127 142
```