Classes for record linkage of big data sets

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As of version 0.3, the package RecordLinkage includes extensions to overcome the problem of high memory consumption that arises when processing a large number of records (i.e. building record pairs out of ≥ 1000 records without blocking). In versions 0.3_x, this was achieved by blockwise on-demand creation of comparison patterns in an embedded SQLite database (through package RSQLite). Package version 0.4 replaces this mechanism by using file-based data structures from package ff. This approach restricts the amount of data pairs to the available disk space but speeds up execution and facilitates the implementation of methods that need to process the whole set of record pairs (e.g. calculation of optimal classification thresholds).

The interface to the big data methods has is compatible to code written for version 0.3_x, so users familiar with these can stick to their existing workflow (unless access to internal structures like object slots is involved). Therefore, the following text sticks to the vignette already included in versions before 0.4 and only technical details are changed to reflect the different implementation.

In order to facilitate a tidier design, S4 classes and methods were used to implement the extensions. In favor of backward compatibility and development time, plans of a complete transition to S4 were dismissed. Nevertheless, the existing functions were joined with their new counterparts, resulting in methods which dispatch on the new S4 as well as on the existing S3 classes. This approach combines two advantages: First, existing code using the package still works, second, the new classes and methods offer (nearly) the same interface, i.e. the necessary function calls for a linkage task differ only slightly. An exception is getPairs, whose arguments differ from the existing version (see man page).

1 Defining data and comparison parameters

The existing S3 class "RecLinkData" is supplemented by the S4 classes "RLBigDataLinkage" and "RLBigDataDedup" for linking two datasets and deduplication of one dataset respectively. Both share the common abstract superclass "RLBigData".

```
library(RecordLinkage)
showClass("RLBigData")

## Virtual Class "RLBigData" [package "RecordLinkage"]
##
## Slots:
##
```

```
## Name: frequencies blockFld excludeFld strcmpFld
## Class: numeric list numeric numeric
##
## Name: strcmpFun phoneticFld phoneticFun
                                           pairs
## Class: character numeric character
                                            ffdf
##
## Name: Wdata
                   WdataInd M
## Class: ff_vector ff_vector ff_vector
##
## Known Subclasses: "RLBigDataDedup", "RLBigDataLinkage"
showClass("RLBigDataDedup")
## Class "RLBigDataDedup" [package "RecordLinkage"]
##
## Slots:
##
                   identity frequencies
## Name:
             data
                                         blockFld
## Class: data.frame
                     factor numeric
                                          list
## Name: excludeFld strcmpFld strcmpFun phoneticFld
## Class: numeric numeric character numeric
##
## Name: phoneticFun pairs Wdata WdataInd
## Class: character ffdf ff_vector ff_vector
##
## Name:
                     U
## Class: ff_vector ff_vector
## Extends: "RLBigData"
showClass("RLBigDataLinkage")
## Class "RLBigDataLinkage" [package "RecordLinkage"]
##
## Slots:
##
## Name: data1 data2 identity1 identity2
## Class: data.frame data.frame
                                factor
                                          factor
## Name: frequencies blockFld excludeFld strcmpFld
                    list numeric numeric
## Class: numeric
##
## Name: strcmpFun phoneticFld phoneticFun
                                           pairs
## Class: character numeric character
                   WdataInd M
          Wdata
## Name:
## Class: ff_vector ff_vector ff_vector
##
## Extends: "RLBigData"
```

For the two non-virtual classes, the constructor-like function RLBigDataDedup and RLBigDataLinkage exist, which correspond to compare.dedup and compare.linkage for the S3 classes and share most of their arguments.

The following example shows the basic usage of the constructors, for details consult their documentation.

2 Supervised classification

The existing function classifySupv was transformed to a S4 method which handles the old S3 object ("RecLinkData") as well as the new classes. However, at the moment a classificator can only be trained with an object of class "RecLinkData".

The result is an object of class "RLResult" which contains the classification result along with the data object.

```
showClass("RLResult")

## Class "RLResult" [package "RecordLinkage"]

##

## Slots:

##

## Name: data prediction

## Class: RLBigData ff_vector
```

A contingency table can be viewed via getTable, various error measures are calculated by getErrorMeasures.

```
getTable(result)
            classification
## true status N P
                                L
                      0
     0 124696
                                4
##
                         0
                                48
            1
getErrorMeasures(result)
## $alpha
## [1] 0.04
##
## $beta
## [1] 3.207698e-05
##
## $accuracy
## [1] 0.9999519
##
## $precision
## [1] 0.9230769
## $sensitivity
## [1] 0.96
##
## $specificity
## [1] 0.9999679
##
## $ppv
## [1] 0.9230769
##
## $npv
## [1] 0.999984
```

3 Weight-based classification

As with "RecLinkData" objects, weight-based classification with "RLBigData*" classes includes weight calculation and classification based on one or two thresholds, dividing links, non-links and, if desired, possible links. The following example applies classification with Epilink (see documentation of epiWeights for details):

```
rpairs1 <- epiWeights(rpairs1)
result <- epiClassify(rpairs1, 0.5)
getTable(result)
## classification</pre>
```

```
## true status N P L ## 0 124699 0 1 ## 1 4 0 46
```

4 Evaluation and results

In addition to getTable and getErrorMeasures, getPairs, which was redesigned as a versatile S4 method, is an important tool to inspect data and linkage results. For example, the following code extracts all links with weights greater or equal than 0.7 from the result set obtained in the last example:

```
getPairs(result, min.weight=0.7, filter.link="link")
##
      id fname_c1 fname_c2 lname_c1 lname_c2
                                                 by bm bd
## 1 290
            HELGA ELFRIEDE
                              BERGER
                                          <NA> 1989
                                                     1 18
## 2 466
            HELGA ELFRIEDE
                              BERGER
                                          <NA> 1989
                                                      1 28
## 3
## 4 313
           URSULA
                                          <NA> 1940
                                                      6 15
                     BIRGIT
                             MUELLRR
## 5 457
           URSULA
                     BIRGIT
                             MUELLER
                                          <NA> 1940
                                                      6 15
## 6
## 7 467
           ULRIKE
                     NICOLE
                              BECKRR
                                          <NA> 1982
                                                      8
                                                         4
## 8 472
           ULRIKE
                     NICOLE
                              BECKER
                                          <NA> 1982
                                                      8
## 9
##
     is_match Class
                        Weight
## 1
## 2
         TRUE
                   L 0.7786012
## 3
## 4
## 5
         TRUE
                   L 0.7293529
## 6
## 7
## 8
         TRUE
                   L 0.7293529
## 9
```

A frequent use case is to inspect misclassified record pairs; for this purpose two shortcuts are included that call getPairs with appropriate arguments:

```
getFalsePos(result)
## ==============
     id fname_c1 fname_c2 lname_c1 lname_c2
                                             by bm bd
## 1 388
          ANDREA
                    <NA>
                            WEBER.
                                      <NA> 1945 5 20
## 2 408
          ANDREA
                     <NA>
                          SCHMIDT
                                      <NA> 1945 2 20
## 3
    is_match Class
##
                     Weight
## 1
## 2
       FALSE
                 L 0.5067013
## 3
```

```
getFalseNeg(result)
id fname_c1 fname_c2 lname_c1 lname_c2 by bm bd
## 1 353 INGE <NA> SEIDEL <NA> 1949 9 4
## 2 355
         INGEU
                               <NA> 1949 8 4
                <NA> SEIDEL
## 3
## 4 285
        ERIKA <NA>
ERIKA <NA>
                             <NA> 1995 2 1
<NA> 1992 2 29
                       WEBER
## 5 379
                       WEBER
## 6
         KARL <NA> KLEIN <NA> 2002 6 20
## 7 127
## 8 142
          KARL
                <NA> KLEIBN <NA> 2002 6 29
## 9
## 10 37 HARTMHUT <NA> HOFFMSNN <NA> 1929 12 29
                 <NA> HOFFMANN
## 11 72 HARTMUT
                               <NA> 1929 12 29
## 12
## is_match Class
                 Weight
## 1
## 2
       TRUE N 0.4948059
## 3
## 4
## 5
       TRUE
              N 0.4782410
## 6
## 7
## 8
       TRUE
              N 0.4692532
## 9
## 10
## 11
      TRUE N 0.4081096
## 12
```