icd9: working with ICD-9 codes and comorbidities in R

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Welcome to the icd9 package for interpretation of ICD-9 codes and assignment of comorbidities. Suggestions for improvements are welcome. Please cite this package if you find it useful for your published work. citation(package='icd9')

1 Introduction

This package is designed to be used with a variety of input data, including multiple possible formats of ICD-9 codes, but some assumptions are made. There are many ways of misinterpreting ICD-9 codes, especially when dealing with ranges. The code in this package carefully considers a wide range of possibilities. ICD-9 codes are not numeric. Using numeric values for either decimal or non-decimal form will cause serious problems, hence the predominantly string-based processing here, and a robust set of unit tests.

When calcuating which patients have which comorbidities, the input data is typically structured as follows:

```
patientData
##
     visitId icd9 poa
## 1
        1000 27801
## 2
        1000 7208
                      N
## 3
        1000 25001
                      Y
## 4
        1001 34400
                      N
## 5
        1001
               4011
                      Υ
## 6
        1002
               4011
```

Only the visitId column is propogated to the results. If 'present-on-arrival' is needed, it must be separated out first. The implicit default, therefore, is to ignore it, and give ICD-9 code regardless of POA status.

The comorbidities can be determined as follows:

```
icd9Comorbidities(icd9df = patientData)[, 1:5]

## visitId CHF VALVE PULMCIRC PERIVASC
## 1 1000 FALSE FALSE FALSE FALSE
## 2 1001 FALSE FALSE FALSE FALSE
## 3 1002 FALSE FALSE FALSE FALSE
```

The following shows the same code with default options written out:

2 Converting ICD-9 codes between types

These functions were designed with the problem of incorrectly or bizarrely formatted ICD-9 codes in mind.

leadingZeroes

allows the result (when syntactically valid) to be prefixed by zeroes. For short codes, this is only true for top-level codes (integers 1 to 999), even then, "010" could be ambiguous (is it "1.0" or "10"?). These functions make the assumption that short codes of three or fewer characters must be describing only the 'major' part.

keepLoneDecimal

allows retention of the decimal plan

```
icd9DecimalToShort(c("10.20", "100", "123.45"))
## [1] "01020" "100" "12345"

icd9ShortToDecimal(icd9DecimalToShort(c("10.20", "100", "123.45")))
## [1] "10.20" "100" "123.45"

# note that only a subset of short codes can suffer dropping of leading zeroes.
icd9DecimalToShort(c("1", "22", "22.44", "100"), leadingZeroes = TRUE)

## [1] "001" "022" "02244" "100"
icd9DecimalToShort(c("1", "22", "22.44", "100"), leadingZeroes = FALSE)
```

```
## [1] "1" "22" "02244" "100"
icd9ShortToDecimal(icd9DecimalToShort(c("1", "1.2", "123.45")), leadingZeroes = TRUE)

## [1] "001" "001.2" "123.45"
icd9ShortToDecimal(icd9DecimalToShort(c("1", "1.2", "123.45")), leadingZeroes = FALSE)

## [1] "1" "1.2" "123.45"
icd9ShortToDecimal(icd9DecimalToShort(c("1", "2.2", "100")), keepLoneDecimal = FALSE)

## [1] "1" "2.2" "100"
icd9ShortToDecimal(icd9DecimalToShort(c("1", "2.2", "100")), keepLoneDecimal = TRUE)

## [1] "1." "2.2" "100."
```

3 Validation of ICD-9 codes

```
icd9ValidDecimal("V10.2")
## [1] TRUE
icd9ValidShort(c("099.17", "-1"))
## [1] FALSE FALSE
```

Validation forces the package user to provide character format ICD-9 codes. If great care is taken, passing some integers could be valid, but given the high chance of mistakes, and the simplicity of dealing entirely with character input, character is enforced:

```
icd9ValidShort(100) # throws an error
```

4 Ranges of ICD-9 codes

These functions generate syntactically valid ICD-9 codes, without including parent codes when the range limit would subset the parent. E.g. "100.99" %i9d% "101.01" does not include "100" or "100.0", both of which imply large subsets than requested by the range command.

```
"10099" %i9s% "10101"

## [1] "10099" "10100" "10101"

"V10" %i9d% "V10.02"

## [1] "V10" "V10.0" "V10.00" "V10.01" "V10.02"

# "E987" %i9d% "E988.9"
```

Another way of specifying ranges are to use function calls. These are exactly equivalent to the %i9s% and %i9d% range operators. This example shows the result when the user specifies a range which would include parents but not all their children:

```
icd9ExpandRangeShort("V100", "V1002")
## [1] "V100" "V1000" "V1001" "V1002"
```

Although V100 would include ten children, the range only returns 4 values. In all other cases, parents are omitted to avoid the range returning overly broad classifications than intended. A planned feature is to optionally enable returning these parent codes, which would then follow a more numerical pattern (although still distinguishing trailing zeroes).

5 Human-readable ICD-9

```
icd9ExplainDecimal("1.0")
                                                       Description
##
     TCD9
                               Diagnosis
## 1 0010 Cholera due to vibrio cholerae Cholera d/t vib cholerae
icd9Explain("1.0", short = FALSE)
                               Diagnosis
                                                       Description
## 1 0010 Cholera due to vibrio cholerae Cholera d/t vib cholerae
icd9ExplainDecimal("001.1")
##
     ICD9
                                       Diagnosis
                                                            Description
## 1 0011 Cholera due to vibrio cholerae el tor Cholera d/t vib el tor
icd9ExplainDecimal(icd9ShortToDecimal("0019"))
     ICD9
                     Diagnosis Description
## 1 0019 Cholera, unspecified Cholera NOS
```

```
icd9ExplainShort("0019")
##
    ICD9
                     Diagnosis Description
## 1 0019 Cholera, unspecified Cholera NOS
# named list(s) of codes
icd9ExplainDecimal(list(cholera = c("001", "001.0", "001.1", "001.9")))
## $cholera
## ICD9
                                       Diagnosis
                                                              Description
                 Cholera due to vibrio cholerae Cholera d/t vib cholerae
## 1 0010
## 2 0011 Cholera due to vibrio cholerae el tor Cholera d/t vib el tor
## 3 0019
                           Cholera, unspecified
                                                              Cholera NOS
# same using decimal codes without a list
icd9ExplainDecimal(c("001", "001.0", "001.1", "001.9"))
##
    ICD9
                                       Diagnosis
                                                              Description
## 1 0010
                 Cholera due to vibrio cholerae Cholera d/t vib cholerae
## 2 0011 Cholera due to vibrio cholerae el tor
                                                   Cholera d/t vib el tor
## 3 0019
                           Cholera, unspecified
                                                              Cholera NOS
# 001/cholera doesn't itself have an explanation: TODO walk down children to get next level
icd9ExplainDecimal(list(cholera = "001", rheumatic_heart = "390"))
## $cholera
##
    ICD9
                                       Diagnosis
                                                              Description
## 1 0010
                 Cholera due to vibrio cholerae Cholera d/t vib cholerae
## 2 0011 Cholera due to vibrio cholerae el tor
                                                   Cholera d/t vib el tor
## 3 0019
                           Cholera, unspecified
                                                              Cholera NOS
##
## $rheumatic_heart
    ICD9
                                                       Diagnosis
## 1 390 Rheumatic fever without mention of heart involvement
                   Description
## 1 Rheum fev w/o hrt involv
  Now try to explain on a non-existent (but 'valid') ICD-9 code:
```

```
icd9ExplainDecimal("001.5")

## [1] ICD9    Diagnosis   Description
## <0 rows> (or 0-length row.names)
```

icd9ExplainDecimal(list(cholera=c("001.0", "001.1", "001.9"))) @

6 AHRQ comorbidity classification

The AHRQ keeps an updated version of the Elixhauser classification of ICD-9-CM codes into comorbidities, useful for research. They provide the data in the form of SAS code. This package provides just enough code to parse the SAS source code provided by the AHRQ (but probably not much other SAS code), and generate a list of ICD-9 codes for each comorbidity.

```
ahrqComorbid <- parseAhrqSas(save = F)</pre>
```

Here are a couple of the shorter ICD-9 groups listed in ahrqComorbid:

SAS source code has a strong whiff of the 1970s about it. A fragment of a recent AHRQ SAS comorbidity mapping SAS FORMAT is as follows. Note the mix of character and numeric-only ranges, isolated values, all in 'short' ICD-9 code form.

```
PROC FORMAT LIB=library fmtlib;
   VALUE $RCOMFMT
      "2780 ",
      "27800",
      "27801",
      "27803",
      "64910"-"64914".
      "V8530"-"V8539",
      "V8541"-"V8545",
      "V8554",
      "79391"
                       = "OBESE"
                                      /* Obesity
                                                        */
      "3004 ",
      "30112",
      "3090 ",
      "3091 ".
      "311 "
                       = "DEPRESS"
```

parseAhrqSas()

resulting in a named list. here is an extract.

```
ahrqComorbid[c("OBESE", "DEPRESS")]
## $OBESE
## [1] "2780" "27800" "27801" "27802" "27803" "27804"
    [7] "27805" "27806" "27807" "27808" "27809" "V8554"
## [13] "79391" "64910" "64911" "64912" "64913" "64914"
## [19] "V8530" "V8531" "V8532" "V8533" "V8534" "V8535"
## [25] "V8536" "V8537" "V8538" "V8539" "V8541" "V8542"
  [31] "V8543" "V8544" "V8545"
##
##
## $DEPRESS
    [1] "3004" "30040" "30041" "30042" "30043" "30044"
##
    [7] "30045" "30046" "30047" "30048" "30049" "30112"
##
   [13] "3090" "30900" "30901" "30902" "30903" "30904"
    [19] "30905" "30906" "30907" "30908" "30909" "3091"
##
    [25] "30910" "30911" "30912" "30913" "30914" "30915"
    [31] "30916" "30917" "30918" "30919" "311"
##
    [37] "3111" "3112" "3113" "3114" "3115" "3116"
    [43] "3117" "3118" "3119" "31100" "31110" "31120"
##
    [49] "31130" "31140" "31150" "31160" "31170" "31180"
    [55] "31190" "31101" "31111" "31121" "31131" "31141"
##
##
    [61] "31151" "31161" "31171" "31181" "31191" "31102"
    [67] "31112" "31122" "31132" "31142" "31152" "31162"
##
    [73] "31172" "31182" "31192" "31103" "31113" "31123"
##
    [79] "31133" "31143" "31153" "31163" "31173" "31183"
    [85] "31193" "31104" "31114" "31124" "31134" "31144"
    [91] "31154" "31164" "31174" "31184" "31194" "31105"
    [97] "31115" "31125" "31135" "31145" "31155" "31165"
## [103] "31175" "31185" "31195" "31106" "31116" "31126"
## [109] "31136" "31146" "31156" "31166" "31176" "31186"
## [115] "31196" "31107" "31117" "31127" "31137" "31147"
## 「121<sup>1</sup> "31157" "31167" "31177" "31187" "31197" "31108"
## [127] "31118" "31128" "31138" "31148" "31158" "31168"
## [133] "31178" "31188" "31198" "31109" "31119" "31129"
## [139] "31139" "31149" "31159" "31169" "31179" "31189"
## [145] "31199"
lapply(ahrqComorbid[c("OBESE", "DEPRESS")], icd9ChildrenShort)
## $OBESE
## [1] "2780" "27800" "27801" "27802" "27803" "27804"
    [7] "27805" "27806" "27807" "27808" "27809" "V8554"
```

```
[13] "79391" "64910" "64911" "64912" "64913" "64914"
   [19] "V8530" "V8531" "V8532" "V8533" "V8534" "V8535"
   [25] "V8536" "V8537" "V8538" "V8539" "V8541" "V8542"
  [31] "V8543" "V8544" "V8545"
##
## $DEPRESS
     [1] "3004" "30040" "30041" "30042" "30043" "30044"
##
     [7] "30045" "30046" "30047" "30048" "30049" "30112"
##
    [13] "3090" "30900" "30901" "30902" "30903" "30904"
##
##
    [19] "30905" "30906" "30907" "30908" "30909" "3091"
    [25] "30910" "30911" "30912" "30913" "30914" "30915"
    [31] "30916" "30917" "30918" "30919" "311"
##
    [37] "3111" "3112" "3113" "3114" "3115" "3116"
##
    [43] "3117" "3118" "3119" "31100" "31110" "31120"
##
    [49] "31130" "31140" "31150" "31160" "31170" "31180"
##
    [55] "31190" "31101" "31111" "31121" "31131" "31141"
##
    [61] "31151" "31161" "31171" "31181" "31191" "31102"
##
    [67] "31112" "31122" "31132" "31142" "31152" "31162"
##
    [73] "31172" "31182" "31192" "31103" "31113" "31123"
    [79] "31133" "31143" "31153" "31163" "31173" "31183"
##
    [85] "31193" "31104" "31114" "31124" "31134" "31144"
##
##
    [91] "31154" "31164" "31174" "31184" "31194" "31105"
    [97] "31115" "31125" "31135" "31145" "31155" "31165"
   [103] "31175" "31185" "31195" "31106" "31116" "31126"
##
   [109] "31136" "31146" "31156" "31166" "31176" "31186"
   [115] "31196" "31107" "31117" "31127" "31137" "31147"
  [121] "31157" "31167" "31177" "31187" "31197" "31108"
  [127] "31118" "31128" "31138" "31148" "31158" "31168"
   [133] "31178" "31188" "31198" "31109" "31119" "31129"
  [139] "31139" "31149" "31159" "31169" "31179" "31189"
  [145] "31199"
```

7 Elixhauser co-morbidities

Elixhauser originally devleoped this set of co-morbidities to predict long term mortality based on hospital ICD-9-CM coding records. The AHRQ comorbidities are an updated version of this, however the original Elixhauser have been used in many publications. The ICD-9-CM codes have changed slightly over the years.

```
names(elixhauserComorbid)
## [1] "chf" "arrhythmia" "valve"
## [4] "pulm.circ" "pvd" "htn"
```

```
[7] "htncx"
                                        "neuro.other"
##
                        "paralysis"
## [10] "chronic.pulm" "dm.uncomp"
                                        "dm.comp"
                        "renal"
## [13] "hypothyroid"
                                        "liver"
## [16] "pud"
                        "hiv"
                                        "lymphoma"
## [19] "mets"
                        "solid.tumor"
                                        "rheum"
##
  [22] "coag"
                        "obesity"
                                        "wt.loss"
## [25] "lytes"
                        "anemia.loss"
                                        "anemia.def"
## [28] "etoh"
                        "drugs"
                                        "psychoses"
## [31] "depression"
```

8 Quan

Quan's paper looked at indices using both ICD-10 and ICD-9-CM. Quan generated updated ICD-9-CM codes for all 30 of Elixhauser and all 17 of Charlson/Deyo's co-morbidities. Thus there are two 'Quan' comorbidity mappings.

```
names(quanDeyoComorbid)
    [1] "Myocardial Infarction"
##
##
    [2] "Congestive Heart Failure"
    [3] "Periphral Vascular Disease"
##
##
    [4] "Cerebrovascular Disease"
    [5] "Dementia"
##
##
    [6] "Chronic Pulmonary Disease"
##
    [7] "Connective Tissue Disease-Rheumatic Disease"
    [8] "Peptic Ulcer Disease"
##
##
   [9] "Mild Liver Disease"
## [10] "Diabetes without complications"
## [11] "Diabetes with complications"
  [12] "Paraplegia and Hemiplegia"
##
## [13] "Renal Disease"
## [14] "Cancer"
  [15] "Moderate or Severe Liver Disease"
  [16] "Metastatic Carcinoma"
## [17] "AIDS/HIV"
names(quanElixhauserComorbid)
    [1] "chf"
                        "arrhythmia"
                                        "valve"
##
##
    [4] "pulm.circ"
                        "pvd"
                                        "htn"
##
    [7] "htncx"
                        "paralysis"
                                       "neuro.other"
## [10] "chronic.pulm" "dm.uncomp"
                                       "dm.comp"
## [13] "hypothyroid"
                                        "liver"
                       "renal"
```

```
## [16] "pud"
                     "hiv"
                                   "lymphoma"
## [19] "mets"
                                   "rheum"
                     "solid.tumor"
## [22] "coag"
                     "obesity"
                                   "wt.loss"
                     "anemia.loss"
## [25] "lytes"
                                   "anemia.def"
                     "drugs"
## [28] "etoh"
                                   "psychoses"
## [31] "depression"
```