K clustering on diabetes dataset



Install and load libraries: mlr

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install.packages("mlr", dependencies = TRUE)

	binary <chr></chr>	source <chr></chr>	needs_compilation <lgl></lgl>
future	1.19.1	1.20.1	FALSE
backports	1.1.10	1.2.0	TRUE
2 rows			

Install and load libraries: tidyverse and mclust

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install.packages("tidyverse")

	binary <chr></chr>	source <chr></chr>	needs_compilation <lgl></lgl>
dbplyr	1.4.4	2.0.0	FALSE
1 row			

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install.packages("mclust")

Load the diabetes data:

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data(diabetes, package = "mclust")

Use the installed libs:

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library(mlr)

Hide

library(tidyverse)

```
Registered S3 methods overwritten by 'dbplyr':
 method
                 from
  print.tbl_lazy
  print.tbl_sql
-- Attaching packages ----- tidyverse 1.3.0 --
v ggplot2 3.3.2
                 v purrr 0.3.4
v tibble 3.0.4 v dplyr 1.0.2
v tidyr 1.1.2 v stringr 1.4.0
                  v forcats 0.5.0
v readr
        1.4.0
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
                  masks stats::lag()
```

Format and display data:

Hide

```
diabetesTib <- as_tibble(diabetes)
diabetesTib</pre>
```

class <fctr></fctr>	glucose <dbl></dbl>	insulin <dbl></dbl>	sspg <dbl></dbl>
Normal	80	356	124
Normal	97	289	117
Normal	105	319	143
Normal	90	356	199
Normal	90	323	240
Normal	86	381	157
Normal	100	350	221
Normal	85	301	186
Normal	97	379	142
Normal	97	296	131
1-10 of 145 rows	Previous 1	2 3 4 5 6	6 15 Next

Get some stats of the data:

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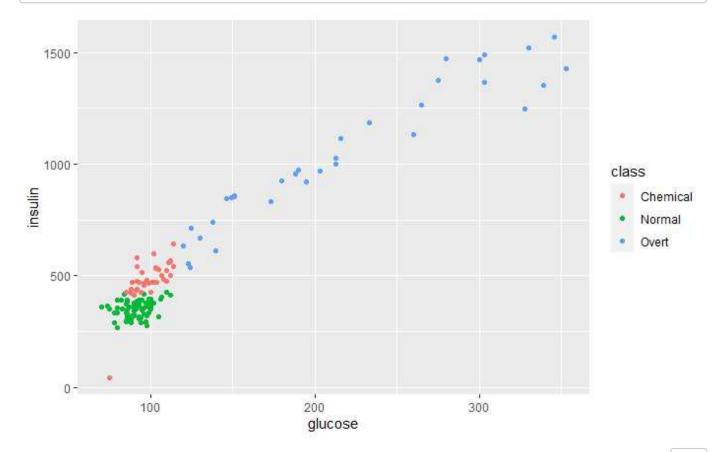
summary(diabetesTib)

```
class
                 glucose
                               insulin
Chemical:36
              Min.
                   : 70
                                 : 45.0
                            Min.
Normal :76
              1st Qu.: 90
                            1st Qu.: 352.0
Overt
        :33
              Median : 97
                            Median : 403.0
                                  : 540.8
              Mean
                    :122
                            Mean
              3rd Qu.:112
                            3rd Qu.: 558.0
              Max.
                    :353
                            Max.
                                   :1568.0
     sspg
Min.
      : 10.0
1st Qu.:118.0
Median :156.0
       :186.1
Mean
3rd Qu.:221.0
       :748.0
Max.
```

Plot the data around some of its classes:

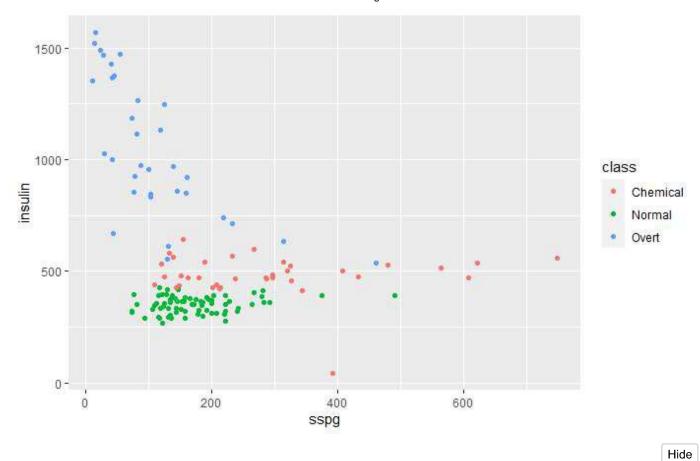
```
Hide
```

```
ggplot(diabetesTib, aes(glucose, insulin, col = class)) +
  geom_point()
```

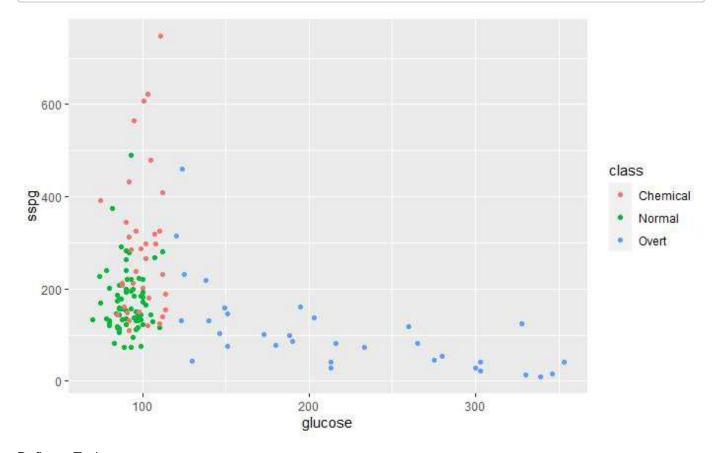


Hide

```
ggplot(diabetesTib, aes(sspg, insulin, col = class)) +
  geom_point()
```



ggplot(diabetesTib, aes(glucose, sspg, col = class)) +
 geom_point()



Define a Task:

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```
diabetesTask <- makeClassifTask(data = diabetesTib, target = "class")</pre>
```

Provided data is not a pure data.frame but from class tbl_df, hence it will be converted.

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diabetesTask

```
Supervised task: diabetesTib

Type: classif

Target: class

Observations: 145

Features:

numerics factors ordered functionals

3 0 0 0 0

Missings: FALSE

Has weights: FALSE
```

Classes: 3

Chemical Normal Overt 36 76 33

Positive class: NA

Has blocking: FALSE Has coordinates: FALSE

Define a learner: K Nearest Neighbours

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```
knn <- makeLearner("classif.knn", par.vals = list("k" = 2))</pre>
```

Train the model:

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```
knnModel <- train(knn, diabetesTask)</pre>
```

Use hold-out crossval:

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```
kfold <- makeResampleDesc("RepCV", folds = 10, reps = 50)</pre>
```

Hide

```
kFoldCV <- resample(learner = knn, task = diabetesTask, resampling = kfold)</pre>
```

```
Resampling: repeated cross-validation
Measures:
                      mmce
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[Resample] iter 401:
                      0.1428571
[Resample] iter 402:
                      0.1333333
[Resample] iter 403:
                      0.0714286
```

```
[Resample] iter 404:
                      0.0000000
[Resample] iter 405:
                      0.0714286
[Resample] iter 406:
                      0.0666667
[Resample] iter 407:
                      0.0666667
[Resample] iter 408:
                      0.1333333
[Resample] iter 409:
                      0.1333333
[Resample] iter 410:
                      0.1428571
[Resample] iter 411:
                      0.0666667
[Resample] iter 412:
                      0.0666667
[Resample] iter 413:
                      0.1428571
[Resample] iter 414:
                      0.0000000
[Resample] iter 415:
                      0.1428571
[Resample] iter 416:
                      0.0666667
[Resample] iter 417:
                      0.0000000
[Resample] iter 418:
                      0.1333333
[Resample] iter 419:
                      0.1428571
[Resample] iter 420:
                      0.0000000
[Resample] iter 421:
                      0.1333333
[Resample] iter 422:
                      0.1428571
[Resample] iter 423:
                      0.0000000
[Resample] iter 424:
                      0.0666667
[Resample] iter 425:
                      0.0000000
[Resample] iter 426:
                      0.1333333
[Resample] iter 427:
                      0.2000000
[Resample] iter 428:
                      0.2000000
[Resample] iter 429:
                      0.0714286
[Resample] iter 430:
                      0.1428571
[Resample] iter 431:
                      0.1333333
[Resample] iter 432:
                      0.0000000
[Resample] iter 433:
                      0.0666667
[Resample] iter 434:
                      0.1333333
[Resample] iter 435:
                      0.0714286
[Resample] iter 436:
                      0.2000000
[Resample] iter 437:
                      0.0000000
[Resample] iter 438:
                      0.1428571
[Resample] iter 439:
                      0.1428571
[Resample] iter 440:
                      0.2142857
[Resample] iter 441:
                      0.0000000
[Resample] iter 442:
                      0.0000000
[Resample] iter 443:
                      0.0666667
[Resample] iter 444:
                      0.2000000
[Resample] iter 445:
                      0.2666667
[Resample] iter 446:
                      0.1428571
[Resample] iter 447:
                      0.0000000
[Resample] iter 448:
                      0.2000000
[Resample] iter 449:
                      0.1428571
                      0.0000000
[Resample] iter 450:
[Resample] iter 451:
                      0.0714286
[Resample] iter 452:
                      0.0000000
[Resample] iter 453:
                      0.2142857
[Resample] iter 454:
                      0.0666667
[Resample] iter 455:
                      0.2000000
[Resample] iter 456:
                      0.1428571
[Resample] iter 457:
                      0.0666667
[Resample] iter 458:
                      0.1333333
[Resample] iter 459:
                      0.1428571
[Resample] iter 460:
                      0.0000000
[Resample] iter 461:
                      0.2666667
```

```
[Resample] iter 462: 0.0000000
[Resample] iter 463:
                     0.1428571
[Resample] iter 464: 0.0000000
[Resample] iter 465: 0.0000000
[Resample] iter 466: 0.0000000
[Resample] iter 467: 0.1428571
[Resample] iter 468: 0.1333333
[Resample] iter 469: 0.1428571
[Resample] iter 470: 0.1333333
[Resample] iter 471: 0.2000000
[Resample] iter 472: 0.0000000
[Resample] iter 473:
                     0.2857143
[Resample] iter 474: 0.0714286
[Resample] iter 475: 0.0714286
[Resample] iter 476: 0.1333333
[Resample] iter 477: 0.0000000
[Resample] iter 478:
                     0.0000000
[Resample] iter 479: 0.2142857
[Resample] iter 480: 0.0666667
[Resample] iter 481: 0.0666667
[Resample] iter 482: 0.0666667
[Resample] iter 483:
                     0.0714286
[Resample] iter 484: 0.0666667
[Resample] iter 485: 0.0714286
[Resample] iter 486: 0.0666667
[Resample] iter 487: 0.0714286
[Resample] iter 488:
                     0.1428571
[Resample] iter 489: 0.0666667
[Resample] iter 490: 0.1428571
[Resample] iter 491: 0.0666667
[Resample] iter 492: 0.1333333
[Resample] iter 493: 0.0666667
[Resample] iter 494: 0.2142857
[Resample] iter 495: 0.1333333
[Resample] iter 496: 0.0666667
[Resample] iter 497: 0.0714286
[Resample] iter 498: 0.0714286
[Resample] iter 499: 0.0000000
[Resample] iter 500:
                     0.1428571
Aggregated Result: mmce.test.mean=0.1033143
```

Display the confusion matrix:

Hide

calculateConfusionMatrix(kFoldCV\$pred)

```
predicted
true
           Chemical Normal Overt -err.-
  Chemical
                1473
                        164
                               163
                                      327
  Normal
                 154
                       3646
                                 0
                                      154
                 267
  0vert
                           0
                             1383
                                      267
  -err.-
                 421
                        164
                               163
                                      748
```

We can plot the Knn model:

Hide

plotLearnerPrediction(learner = knn, task = diabetesTask)

