

PH125_9 Choose Your Own Capstone Project

Mario De Toma

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Introduction

Objectives

Motivation) This project has been conducted as part of the Data Science Professional Certification path provided by HarvardX, an online learning initiative of Harvard University through edX. In particular this is the second data science project to submit for PH125.9x course denominated “Data Science: Capstone”. The name of the project, Choose Your Own, is due to the fact that the data set under analysis could be chosen by the learner from public available dataset. I choose the Fall Detection dataset from the curated list of datasets at the following link https://www.kaggle.com/annavictoria/ml-friendly-public-datasets?utm_medium=email&utm_source=intercom&utm_campaign=data+projects+onboarding as suggested by course staff.

Project objective) As per course project introduction the project aim is to apply machine learning techniques that go beyond standard linear regression. In particular the task of this project is multi class classification i.e. where the outcome variable is categorical with more than 2 classes. Specifically the problem statement is related to predict the type of activity among 6 different activities of daily living (ADLs) on the basis of monitored medical measures obtained through sensors worn by elder people.

Research question) The research question can be stated as: is it possible to predict 6 activities of daily living (ADLs) including Standing , Walking, Sitting, Falling, Cramps, Running from few predictors monitoring health status?

This project is not intended to solve the problem of Fall Detection which could be settled by binaryising the ADL information as Falling vs all the remaining activities.

Furthermore note that no causal inference claim can be raised after this study that focus only on supervised learning.

Dataset) Fall detection data set of Chinese hospitals of old age patients [1] is hosted by kaggle. It reports 16382 observations containing the ADL label and related 6 predictors.

Background

Starting point for conducting this study is the supervised learning process as described by Professor Rafael Irizarry in the PH125_8 edX course on Machine Learning and in his book: Introduction to Data Science [2]. In particular study is conducted using the ‘caret’ R package framework for machine learning [3].

Overview and outline

The study demonstrates that ADL can be predicted and that reachable accuracy of prediction depends on the model chosen.

This report is articulated in the following sections:

- *Methods:* where the dataset is explored in order to find some insight, then the design of the study is explained and different models are proposed. Finally the modeling will be described in details.

- *Results*: showing actual results achieved and models evaluation compared, furthermore a procedure is shown in order to explain the loca behavior of the best model
- *Conclusions*: summarizing achievement, discussing the project and indicate potential model improvement
- *Reproducibility*: providing information related to the reproducibility of the analysis including computation considerations, HW and SW stack used.

Methods

In order to answer the research question posed in the introcution section, the dataset Fall Detection has been analyzed and then the machine learning experiments consucted using different machine learning tecnicas adequate for the multi class classificaion task.

Exploratory Data Analysis

Fall detection dataset has been downloaded from kaggle, put on my github and then loaded into R and partitioned such that 70% of the observations belong to the training set and 30% the test set. The caret::createDataPartition function has been used in order to mantain the class distribution between train and test set.

Fall Detection dataset contains the following valriables:

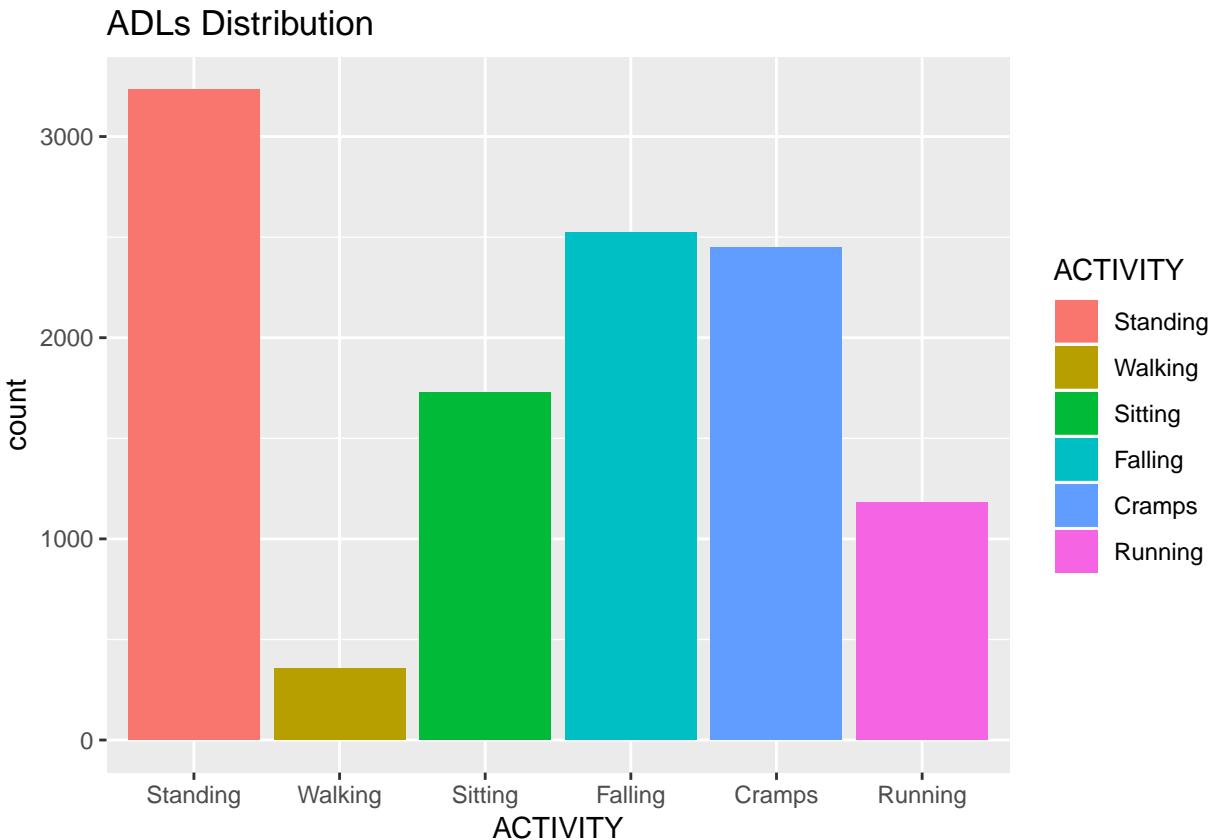
```
## Observations: 11,468
## Variables: 7
## $ ACTIVITY    <dbl> 3, 2, 2, 4, 4, 5, 3, 4, 5, 0, 4, 3, 3, 0, 0, 3, 2, ...
## $ TIME        <dbl> 4722.92, 4059.12, 4773.56, 8271.27, 7102.16, 7015....
## $ SL          <dbl> 4019.64, 2191.03, 2787.99, 9545.98, 14148.80, 7336...
## $ EEG          <dbl> -1600.000, -1146.080, -1263.380, -2848.930, -2381....
## $ BP           <dbl> 13, 20, 46, 26, 85, 22, 35, 82, 61, 59, 44, 16, 53...
## $ HR           <dbl> 79, 54, 67, 138, 120, 95, 157, 315, 214, 104, 156, ...
## $ CIRCLUATION <dbl> 317, 165, 224, 554, 809, 427, 1519, 5844, 1469, 65...
```

The class label ACTIVITY is a numeric variable. It had been converted to a factor with following levels: 0- Standing 1- Walking 2- Sitting 3- Falling 4- Cramps 5- Running

```
fall_train <- fall_train %>% mutate(ACTIVITY = factor(ACTIVITY,
                                                       labels = c('Standing', 'Walking', 'Sitting',
                                                       'Falling', 'Cramps', 'Running')))
fall_test <- fall_test %>% mutate(ACTIVITY = factor(ACTIVITY,
                                                       labels = c('Standing', 'Walking', 'Sitting',
                                                       'Falling', 'Cramps', 'Running')))
```

ADL Distribution

The ADL classes are not evenly distributed. In particular Walking ADL has few observations. This could make our multi class classification task harder.

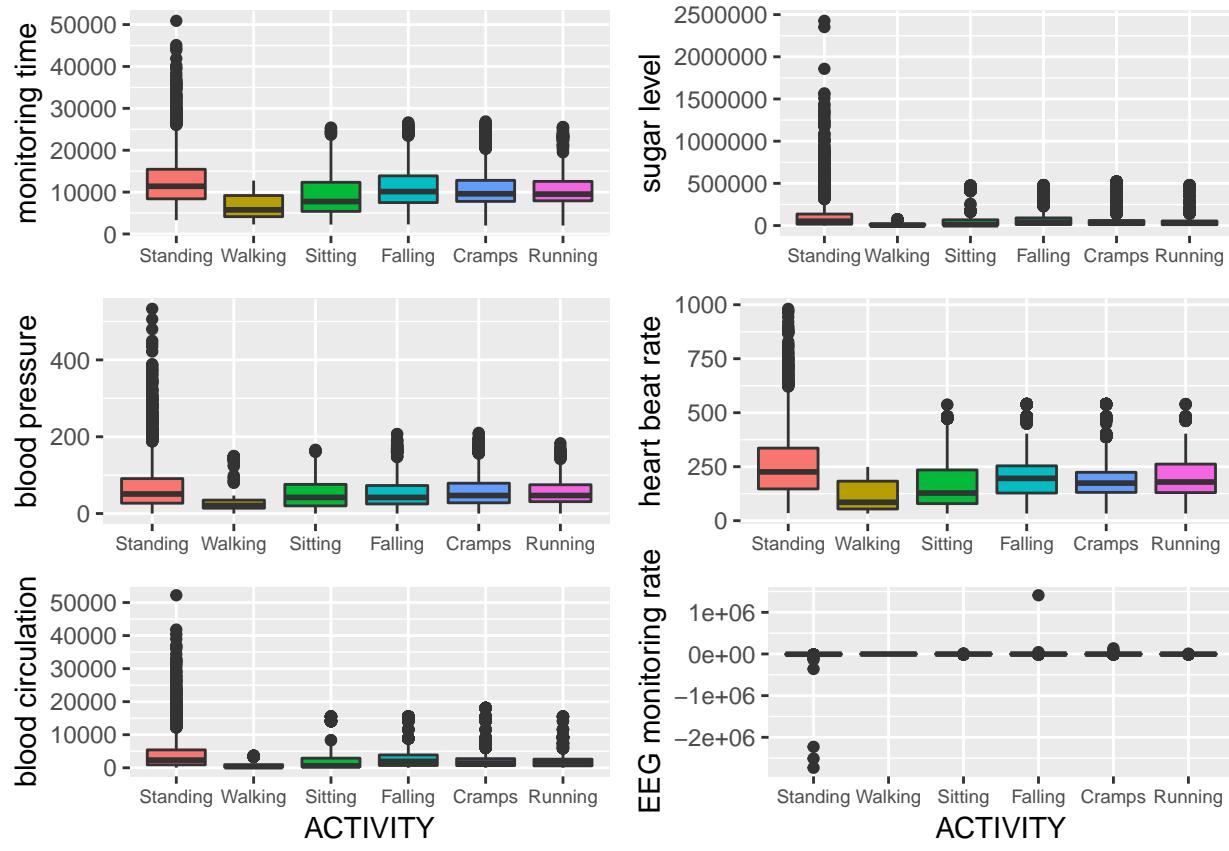


Predictors discriminative power

Predictors for ADL in Fall Detection dataset are:

- TIME monitoring time
- SL sugar level
- EEG monitoring rate
- BP blood pressure
- HR Heart beat rate
- CIRCLUATION Blood circulation

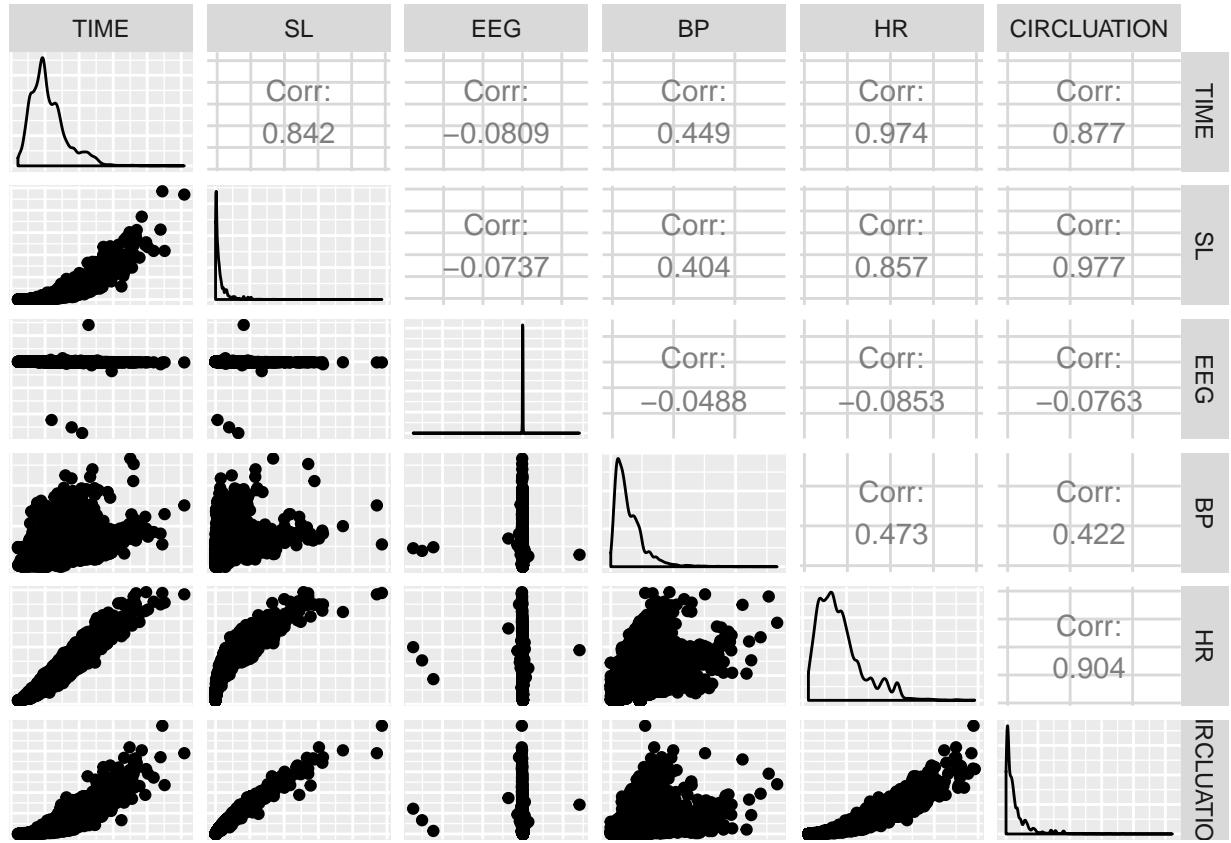
In order to check single predictor capacity to discriminate among classes, for each predictor the boxplot by class has been drawn.



The discriminative power of single predictors does not seem to be enough to classify ADL.

Predictors correlation

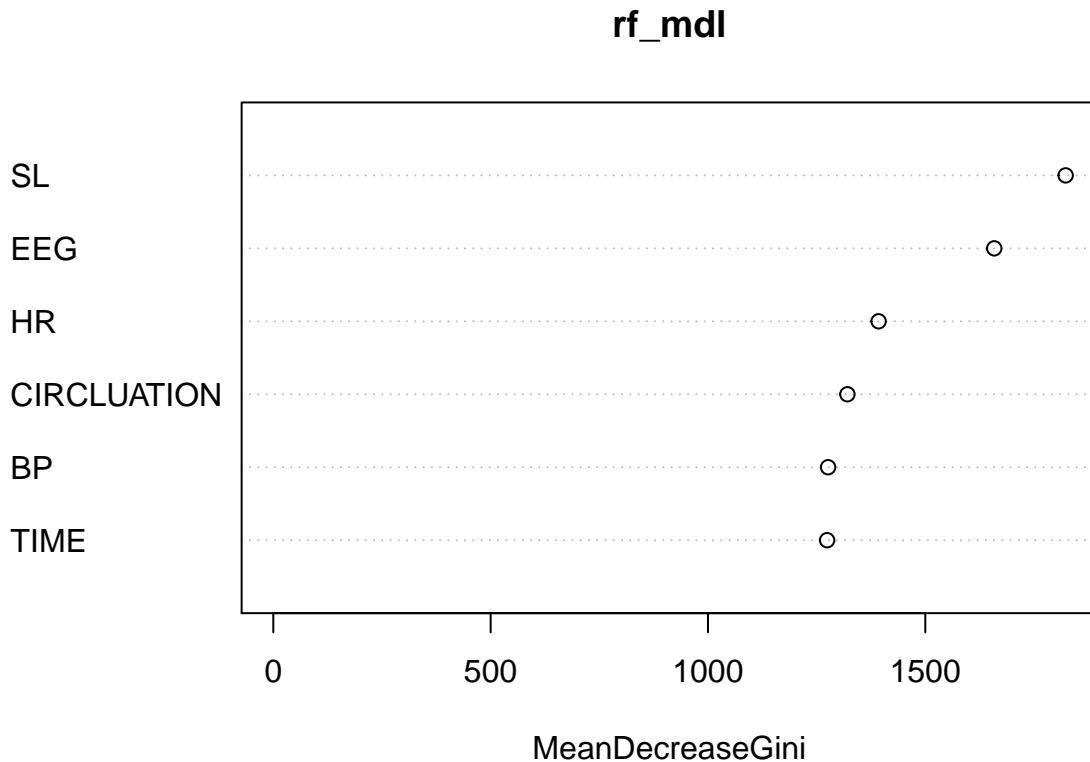
Further element of analysis is related to high level of correlation between predictors.



In particular SL (sugar level) and CIRCLUATION (blood circulation) with a correlation of 0.977 and TIME (monitoring time) and HR (heart beat rate) with a correlation of 0.974 are almost colinear. Also HR and CIRCULATION (0.904), HR with SL (0.857) and TIME with CIRCULATION (0.877) correlations are really high.

Feature importance

The feature importance has been therefore investigated through the random forest algorithm which provide as a side outcome the importance of a feature in discriminating one class from the other. In other words, for random forest algorithm is easy to compute how much each variable is contributing to the classification decision.



From the plot, SL is the most important predictor followed by EEG but all 6 are great help in the classification attempt.

Proposed models

In order to accomplish the multi class classification task the followin models has been tried:

- multinomial
- k Nearest Neighbors
- random forest

All the models are natural choice for multi class classification.

Study design

The study will be conducted using the training set, fall_train, for training and tuning of hyperparameters through cross validation while the final accuracy will be evaluated on the test set, fall_test.

The test dataset is not used in any former phase of the study and therefore it can simulate new data allowing to evaluate the capacity to generalize of the model.

Accuracy metric has been used to evaluate different models. For each model the confusion matrix will be produced in order to evaluate which class is harder to identify.

Cross validation has been configured in caret machine learning framework with 5 folds (80% for training, 20% for validation)

```
ctrl <- trainControl(method = 'cv', number = 5, p = .8)
```

Cross validation lead to long computation times because in the defined study design the model has to be trained and validated 5 times. Therefore cross validation computation has been parallelized making use of doParallel package and of the multithread architecture of the laptop used for this project.

Multinomial Logistic Regression

Multinomial logistic regression is a classification method that generalizes logistic regression to multiclass problems. The probability to belong to a particular class given the predictors is formulated as:

$$P(y = k|x^{(i)}, \theta) = \frac{\exp(\theta^{(k)T}x^{(i)})}{\sum_{j=1}^k \exp(\theta^{(j)T}x^{(i)})}$$

It is called also softmax regression.

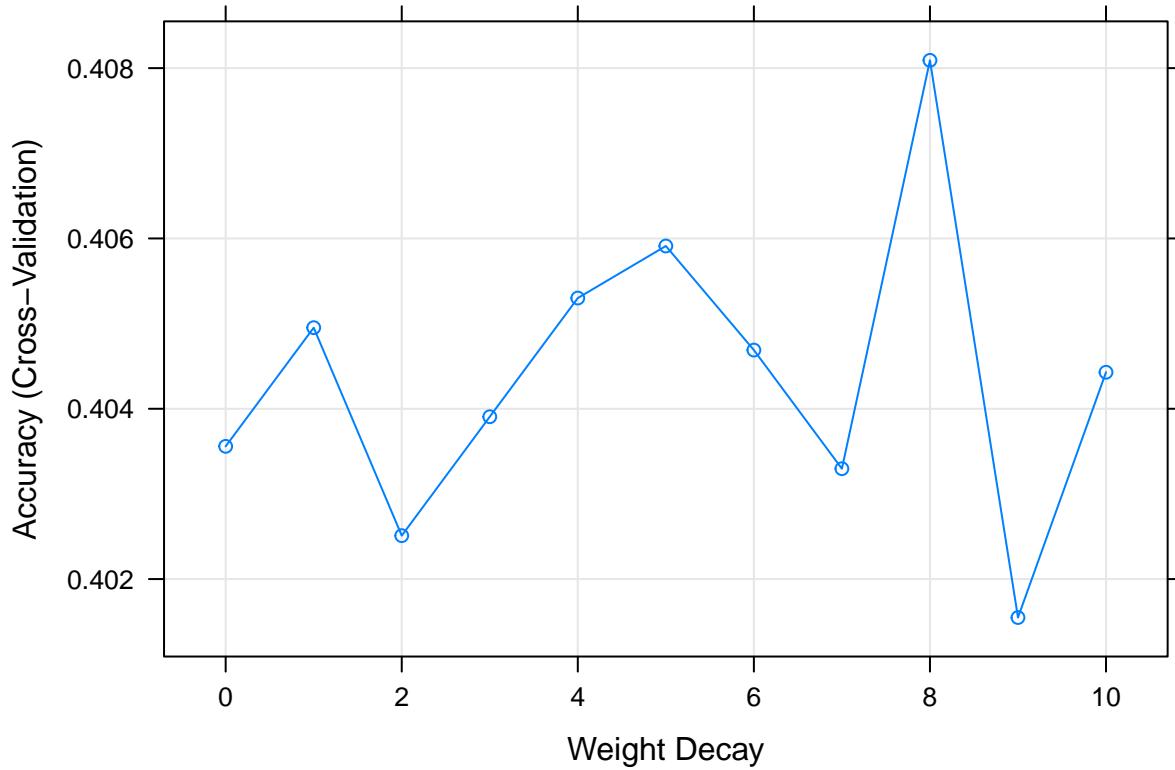
The multinomial model is implemented in r via neural networks by ‘nnet’ package. Under the ‘caret’ it is possible to tune the following parameter:

```
##      model parameter      label forReg forClass probModel
## 1 multinom      decay Weight Decay FALSE      TRUE      TRUE
```

The parameter Weight Decay is specific to neural networks and it helps the optimization process avoiding over-fitting. The training process will go through the following value for hyperparameters

```
tunegrid <- data.frame(decay = seq(from = 0, to = 10, by = 1))
```

The resulting cross-validation plot identify the best model.



After cross-validation training through the defined tune grid the best model found has the following parameter:

```
##     decay
## 9      8
```

After evaluating the accuracy of the model prediction to the test unseen data, the confusion matrix and the accuracy overall score is displayed.

```
##          Reference
## Prediction Standing Walking Sitting Falling Cramps Running
##   Standing      1008     45    199     297    337     169
##   Walking        0       0      0       0      0       0
##   Sitting       41     79    221     91     88      59
##   Falling       241     9    214    482    394     166
##   Cramps        85    13    141    194    229     112
##   Running        0       0      0       0      0       0

## [1] "model test accuracy:  0.39479"
```

As per this results, the multinomial model can be discarded. In some sense the bad accuracy results were expected since predictors are correlated and interconnected while multinomial as a type of general linear model has difficulty in getting the interactions. Furtermore we can see from the confusion matrix that Walking ADL is never predicted probably because it has few occurrence in the dataset. Also Running ADL is never predicted.

k Nearest Neighbour

k Nearest Neighbors is a non-parametric classification method that make use of distance (or similarity) measures. In particular for numerical predictors the euclidean distance is used. Euclidean distance is the length of the segment connecting 2 data points in the predictor space and it is defined as:

$$d(\vec{x}^{(i)}, \vec{x}^{(j)}) = \sqrt{(x_1^{(i)} - x_1^{(j)})^2 + (x_2^{(i)} - x_2^{(j)})^2 + \dots + (x_p^{(i)} - x_p^{(j)})^2}$$

The kNN algorithm stored all the data and classify new data points in relation of majority of k nearest (as per euclidean distance) points class.

The k Nearest Neighbors model is implemented in r by ‘e1071’ package. Under the ‘caret’ machine learning framework it is possible to tune the following parameter:

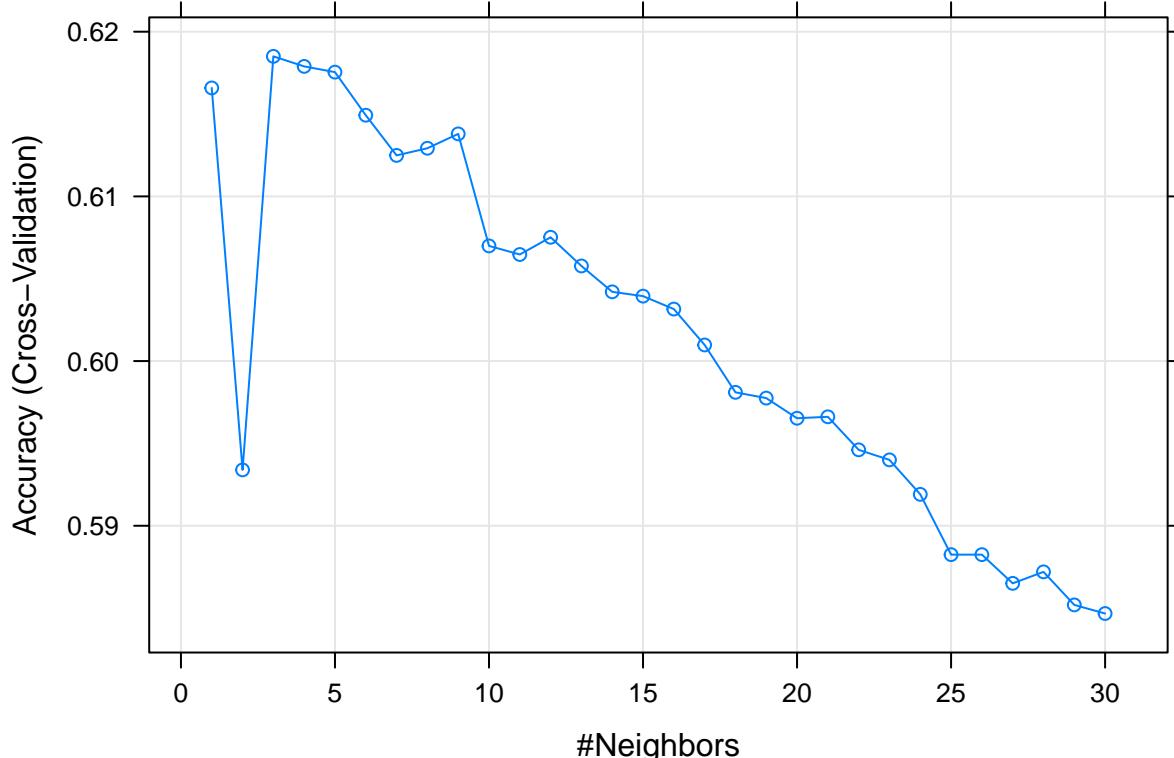
```
##   model parameter      label forReg forClass probModel
## 1   knn                 k #Neighbors    TRUE      TRUE      TRUE
```

As parameter k increase the decision boundary get more smooth. It can be thought as a mean of regularization. The training process will go through the following value for hyperparameters

```
tunegrid <- data.frame(k= seq(1, 30, 1))
```

Given that kNN algorithm is based on distance/similarity measures, data needs to be scaled (by dividing by respective standard deviation) and centered (by subtracting the mean) before training in order to avoid that predictors with largest numerical range mask the effect of other predictors.

The resulting cross-validation plot identify the best model.



After cross-validation training through the defined tune grid the best model found has the following parameter:

```
##   k  
## 3 3
```

After evaluating the accuracy of the model prediction to the test unseen data, the confusion matrix and the accuracy overall score is displayed.

```
##          Reference  
## Prediction Standing Walking Sitting Falling Cramps Running  
##   Standing      911      4     12     19     48     48  
##   Walking       13    107     28      8      7      2  
##   Sitting       52     31    525    162     57     24  
##   Falling      136      2    149    663    194     54  
##   Cramps       165      0     45    179    615    123  
##   Running       98      2     16     33    127    255  
  
## [1] "model test accuracy: 0.62597"
```

kNN model succeed in classifying all 6 ADLs and it gets a good accuracy score considering that we have 6 class. Even Walking despite of class small numerosity is predicted with good accuracy. The best model is obtained with a quite small k, a complex model, as expected because of the interactions among predictors.

Random Forest

Random Forest algorithm builds multiple decision trees and merges them together to get a more accurate and stable prediction reducing variance and avoiding overfitting in respect of the single decision tree. Random forest improves the predictive performance of decision tree through bagging, averaging models learned on multiple bootstrapped samples from the original dataset, and randomly selecting the predictors among which identify the one for partitioning data so that purest node are created at each split.

The Random Forest model is implemented in r by ‘randomForest’ package. Under the ‘caret’ it is possible to tune the following parameter:

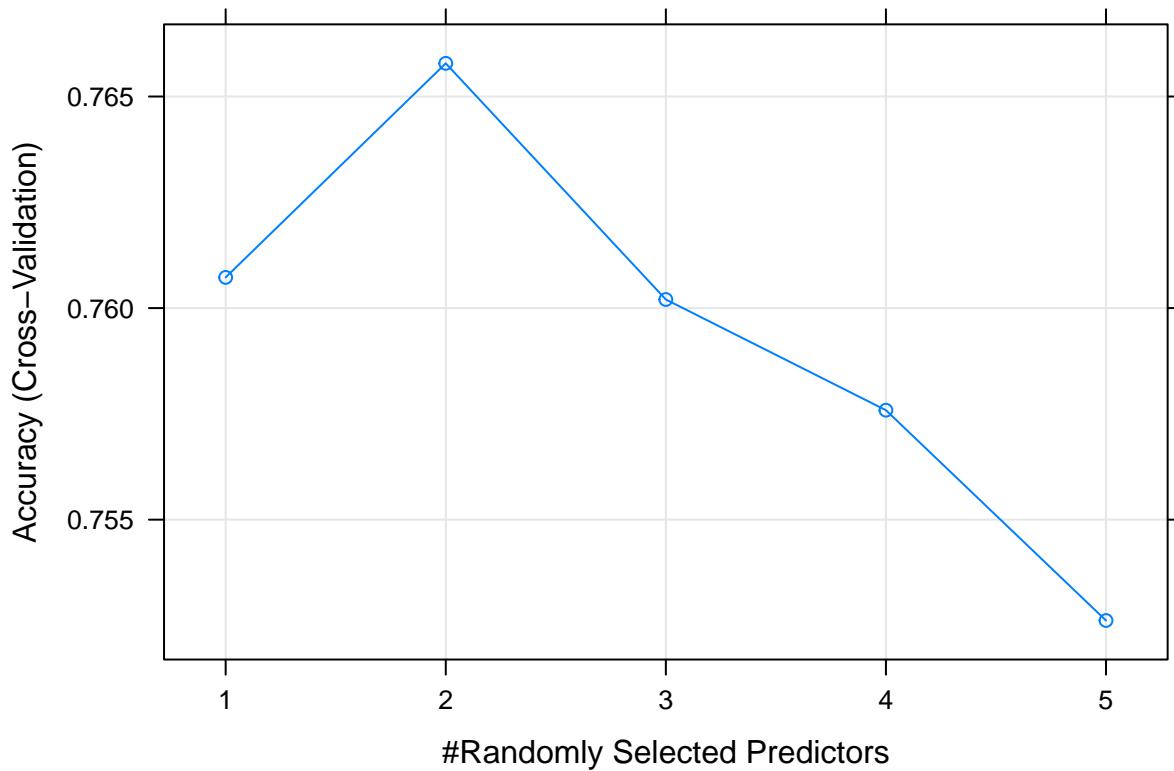
```
##   model parameter           label forReg forClass probModel  
## 1   rf      mtry #Randomly Selected Predictors   TRUE    TRUE    TRUE
```

that set the number of variables randomly sampled as candidates at each split.

The training process will go through the following value for hyperparameters

```
tunegrid <- data.frame(mtry= seq(1, 5, 1))
```

The resulting cross-validation plot identify the best model.



After cross-validation training through the defined tune grid the best model found has the following parameter:

```
##     mtry
## 2      2
```

After evaluating the accuracy of the model prediction to the test unseen data, the confusion matrix and the accuracy overall score is displayed.

```
##          Reference
## Prediction Standing Walking Sitting Falling Cramps Running
##   Standing      1330      0       7       6      22      28
##   Walking        0     120      21       3       3       4
##   Sitting        4      26     588     121      46      12
##   Falling       15      0    139     777     177      42
##   Cramps        20      0      19     143     700     129
##   Running        6      0       1      14     100     291

## [1] "model test accuracy:  0.77452"
```

Random Forest model succeed in classifying all 6 ADLs and it gets a more than good accuracy score considering that we have 6 class. Even Walking despite of class small numerosity is predicted with good accuracy. The best model is obtained with a small mtry.

Results

The following table showed the results achieved for all models.

method	accuracy
multinom	0.3947904
knn	0.6259666
rf	0.7745218

Multinomial model tends to perform badly on the Fall Detection dataset because predictors are heavily correlated. Better accuracy performance can be achieved with kNN a memory based algorithm but with a small number k of neighbors revealing an intrinsic complexity. Random Forest model predicts with a good accuracy all 6 ADLs because trees understand interactions between predictors.

Interpretation

RandomForest model achieved a good accuracy result but it remains a black-box model with low interpretability even if considering the variable importance side result. This consideration about interpretability seems to be almost generalizable in respect to machine learning black-box methods adoption: among relevant literature it is possible to mention the provoking titled paper “Why should I trust you? . . .” [7] and the as much provoking book “Weapons of Math Destruction . . .” [8]. So in order to provide explanation about the model result that can be validated by domain expert, a “ceteris paribus” analysis has been performed. The concept behind this kind of analysis is very simple: take one or more single predictions of interest for domain expert and analyze what the model have learned varying only one predictor at a time keeping all other variable unchanged (this is the meaning of latin word ‘ceteris paribus’: all else unchanged). The prediction selected is:

ACTIVITY	TIME	SL	EEG	BP	HR	
Falling	20368.8	193489	-4900	93	403	
The analysis for multiclass classification	one_report	assification	follow	the procedure documented at https://pbiecek.github.io/ceteris-paribus/		
In order to understand how the results change	one_report	ing plot showing				
		lesions/figure				

It is a local analysis. At the selected prediction the probability of “Falling” is way higher than for other ADLs. There is a range varying HR in which predicted probabilities remain the same. Decreasing a little HR Heart Beat Rate does not change the prediction while below about 300 the most probable ADLs become “Standing”. Also increasing HR till about 350 the model mantain the prediction at “Falling”. Without any domain expertise it is impossible to say if physiology theory can support the result learned by our model, but this is why generally Data Science Team includes domain experts together with Data Scientists.

Conclusions

Going back to our research question, it is possible to state that ADLs can be predicted from basic health measuree with an accuracy over 77%. This means that the best model guesses the right ADL among 6 more than 3 times over 4. It is a remarkable result.

Validity

Results can be considered valid because of this 3 main reasons:

- a consistent training / validation / test study design has been followed consistently for all models
- test and training/validation set contains thousands of observations
- all used machine learning techniques are consolidated
- ceteris paribus analysis presented is a way to validate the model consistency with domain theory

Limitations

This project is a data science project in the context of supervised learning focused on the study of prediction.

Therefore the following 2 general limitations apply:

- from the results cannot be inferred anything about causation;
- the results validity depends on the accuracy of the data collected and contained in the dataset under study. Any sampling or measurement bias could be reflected in the results.

Model improvements

Future research should look at evaluating different machine learning techniques such as implementing a deep neural network with enough hidden layers for getting all the interactions between predictors. Another possibility for increasing the accuracy of the prediction could be stacking: an ensemble method that builds a classification model at an upper level in regards of the studied models using prediction of the lower levels model as predictors for the upper level model.

Implications

This project helped me in consolidating

- my understanding of the data science research methodology;
- the ability to communicate data science results in a reproducible report;
- and the expert use of statistical computation tools.

Reproducibility

R script and rmarkdown file are available for review on public github repository:

👉 https://github.com/mdt-ds/PH125.9_cyo_fallDetection .

R scripts are intended to be reproducible.

- All package loading is checked for package installation.
- Directories are all indicated in relative fashion.
- Seed for random number generation has been set to guarantee reproducible results wherever it is needed
- Furthermore in order to facilitate reproducibility, HW and SW used for this project have been reported below.

HW

The computation has been performed on my laptop:

```
## [1] "Machine:      AMD Ryzen 5 PRO 2500U w/ Radeon Vega Mobile Gfx"  
## [1] "Num cores:    4"  
## [1] "Num threads: 8"  
## [1] "RAM:          8GB"
```

SW

The software stack is shown below launching sessionInfo() R function.

```
R version 3.5.3 (2019-03-11)  
Platform: x86_64-w64-mingw32/x64 (64-bit)  
Running under: Windows 10 x64 (build 16299)  
  
Matrix products: default  
  
locale:  
[1] LC_COLLATE=Italian_Italy.1252  LC_CTYPE=Italian_Italy.1252  
[3] LC_MONETARY=Italian_Italy.1252 LC_NUMERIC=C  
[5] LC_TIME=Italian_Italy.1252  
  
attached base packages:  
[1] parallel  stats      graphics   grDevices  utils      datasets  methods  
[8] base  
  
other attached packages:  
[1] benchmarkme_1.0.0     ceterisParibus_0.3.1  gower_0.2.0  
[4] DALEX_0.4            doParallel_1.0.14    iterators_1.0.10  
[7] foreach_1.4.4        randomForest_4.6-14  GGally_1.4.0  
[10] gridExtra_2.3       caret_6.0-81       lattice_0.20-38  
[13]forcats_0.4.0       stringr_1.4.0      dplyr_0.8.0.1  
[16] purrr_0.3.2        readr_1.3.1       tidyverse_1.2.1  
[19] tibble_2.1.1        ggplot2_3.1.0      tidyverse_1.2.1  
  
loaded via a namespace (and not attached):  
[1] httr_1.4.0           jsonlite_1.6       splines_3.5.3  
[4] prodlm_2018.04.18   modelr_0.1.4      assertthat_0.2.1  
[7] highr_0.8            stats4_3.5.3      cellranger_1.1.0  
[10] yaml_2.2.0          ipred_0.9-8       pillar_1.3.1  
[13] backports_1.1.3    glue_1.3.1        digest_0.6.18  
[16] RColorBrewer_1.1-2 rvest_0.3.2      colorspace_1.4-1  
[19] recipes_0.1.5     htmltools_0.3.6   Matrix_1.2-15  
[22] plyr_1.8.4          timeDate_3043.102 pkgconfig_2.0.2  
[25] broom_0.5.1         haven_2.1.0       scales_1.0.0  
[28] lava_1.6.5          generics_0.0.2    withr_2.1.2  
[31] nnet_7.3-12         lazyeval_0.2.2    cli_1.1.0
```

```

[34] survival_2.43-3      magrittr_1.5          crayon_1.3.4
[37] readxl_1.3.1        evaluate_0.13       fansi_0.4.0
[40] nlme_3.1-137        MASS_7.3-51.1       xml2_1.2.0
[43] class_7.3-15         tools_3.5.3         data.table_1.12.0
[46] hms_0.4.2           munsell_0.5.0       compiler_3.5.3
[49] e1071_1.7-1          rlang_0.3.2         grid_3.5.3
[52] rstudioapi_0.10     labeling_0.3        rmarkdown_1.12
[55] gtable_0.3.0         ModelMetrics_1.2.2 codetools_0.2-16
[58] reshape_0.8.8         curl_3.3           benchmarkmeData_1.0.1
[61] reshape2_1.4.3        R6_2.4.0            lubridate_1.7.4
[64] knitr_1.22           utf8_1.1.4          stringi_1.4.3
[67] Rcpp_1.0.1            rpart_4.1-13        tidyselect_0.2.5
[70] xfun_0.5

```

Computation time

Sourcing the script containing all the analysis on my laptop configured as above took less than 3 minutes to complete.

Acknowledgments

I gratefully aknowledge the efforts of Professor Rafael Irizarry and all HarvardX Course staff for teaching this learning path towards a deeper understanding of Data Science.

References

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- [8] O’NEIL, C. (2016). Weapons of math destruction: how big data increases inequality and threatens democracy.
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