Introduction to Machine Learning (NPFL054)

Homework 2

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Set up the project

```
rm(list = ls())
library(ISLR) # for the data
library(tidyverse) # convenient
library(rpart) # for decision trees
library(randomForest) # for ensemble learning
library(glmnet) # for regularized logistic regression
library(ROCR) # for ROC curves
```

```
## Reproduce the result
set.seed(123)

## Create the splitting vector

split <- sample(nrow(Caravan), 1000)

## Create the test dataset
d_test <- Caravan[split,]

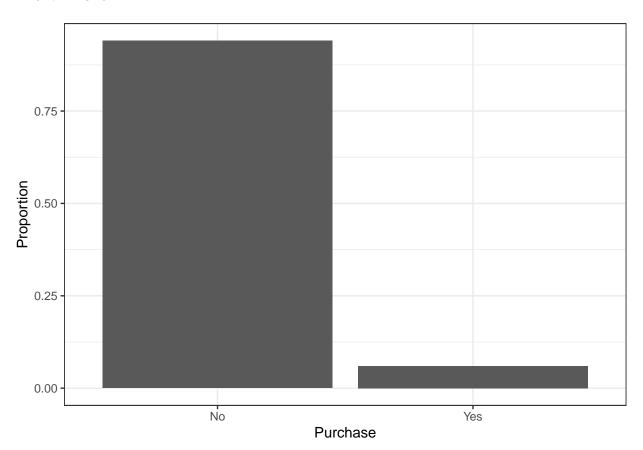
## Create the training dataset
d_train <- Caravan[-split,]</pre>
```

1. Task 1 - Data analysis

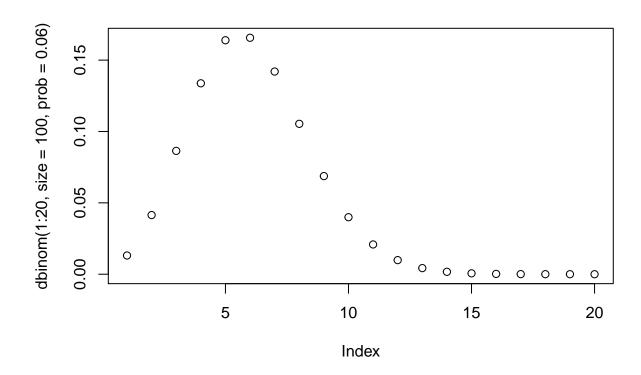
• First, check the distribution of the target attribute. What would be your precision if you select 100 examples by chance?

```
round(table(Caravan$Purchase), 2)
```

```
## No Yes ## 5474 348
```



```
plot(dbinom(1:20, size = 100, prob = .06))
```



We can see that there is 94% of customers who didn't purchase the insurance and that 6% who did. As the precision is the number of examples classified as *Yes* when the value is actually *Yes*, by chance, the precision should be 0.06.

• 1.a. Focus on the customer type MOSHOOFD: create a table with the number of customers that belong to each of 10 L2 groups and the percentage of customers that purchased a caravan insurance policy in each group. Comment the figures in the table. Then do the same for the customer subtype MOSTYPE (41 subgroups defined in L1).

MOSHOOFD type:

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group	size	purchase_prop
1	552	0.09
2	502	0.13
3	886	0.07
5	569	0.03
6	205	0.02
7	550	0.04
8	1563	0.06
9	667	0.06
10	276	0.02

From this first table, we can see that the customers that are more prone to purchase an insurance (13% of them) are the one belonging to the group 2, *i.e.* the *driven growers*. On the other hand, the customers belonging to the class 6 and 10, respectively the *cruising seniors* and the *farmers*, are less likely to subscribe to the insurance (only 2% in each group).

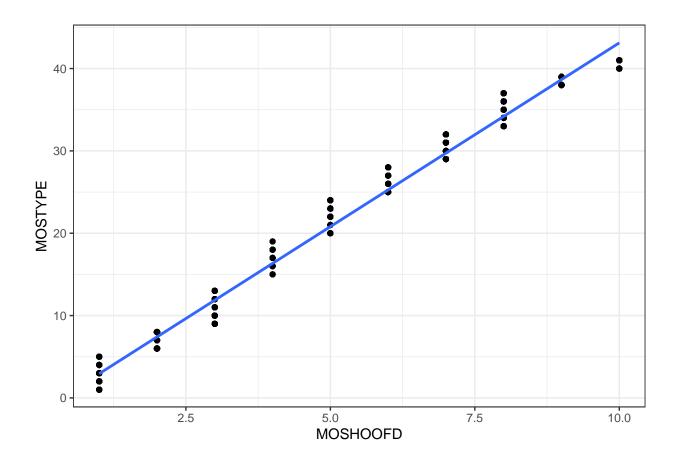
MOSHOOFD type:

group	size	purchase_prop	group	size	purchase_prop
8	339	0.15	10	165	0.05
12	111	0.14	34	182	0.05
1	124	0.10	4	52	0.04
3	249	0.10	5	45	0.04
6	119	0.10	9	278	0.04
20	25	0.08	22	98	0.04
37	132	0.08	35	214	0.04
2	82	0.07	24	180	0.03
7	44	0.07	30	118	0.03
13	179	0.07	31	205	0.03
36	225	0.07	23	251	0.02
38	339	0.07	25	82	0.02
11	153	0.06	26	48	0.02
32	141	0.06	27	50	0.02
33	810	0.06	29	86	0.02
39	328	0.06	41	205	0.02

The two groups more prone to buy an insurance are the group 8 and 12, which correspond respectively to *middle class families* and *affluent young families*. Thus, we can say that families are potential good targets to sell insurances. We can see that the class 25, 26, 27 and 29 all have a low proportion of individuals buying a insurance. They are all related to old people (*i.e.*, *Young seniors in the city*, *Own home elderly*, *Seniors in apartments*, *Porchless seniors: no front yard*). Thus, old people are not a good target to sell insurances.

1.b. Analyze the relationship between features MOSHOOFD and MOSTYPE.

```
Caravan %>%
  ggplot(aes(y = MOSTYPE, x = MOSHOOFD))+
  geom_point()+
  geom_smooth(method = "lm")+
  theme_bw()
```

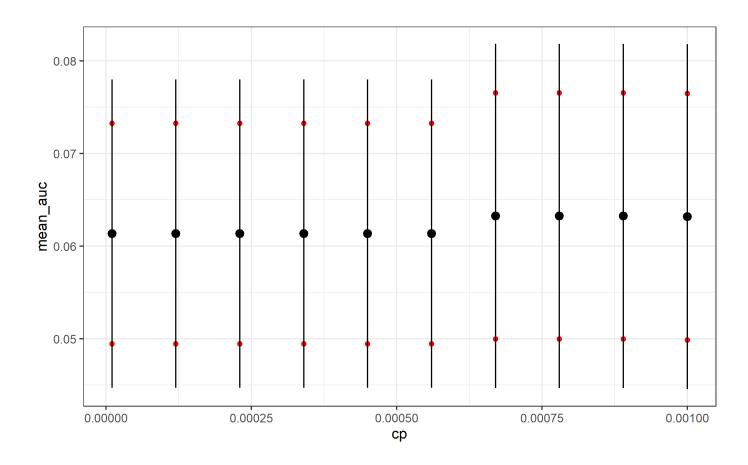


We can clearly see a relationship between these two features which are MOSHOOFD = Customer main type and MOSTYPE = Customer Subtype. This is expected because MOSTYPE is just a more precise social position. For instance, we can see that when MOSHOOFD=10, MOSTYPE=40|41. We can see that MOSHOOFD=10 correspond to Farmers and that MOSTYPE=40|41 are two subclasses of farmers: Large family farms and Mixed rurals, respectively.

2. Task 2 - Model fitting, optimization, and selection

```
## Function to randomly extract the test dataset in d_train
## using always the same number of positive and negative
## values of Purchase
prepare_cv_folds <- function(k){</pre>
  # Create the subsets data containing Purchase == Yes
  # in one hand and Purchase == No in an other hand
  pos data <- d train[d train$Purchase == "Yes",]</pre>
  neg data <- d train[d train$Purchase == "No",]</pre>
  ## Compute the size of each fold
  fold.size.pos <- nrow(pos_data)%/%k
  fold.size.neg <- nrow(neg_data)%/%k
  ## Randomly rearrange the indexes
  set.seed(12); s_pos <- sample(nrow(pos_data))</pre>
  set.seed(12); s_neg <- sample(nrow(neg_data))</pre>
  ## create the list that will contain the test folds
  f.idx <- list()
  ## For each fold, extract the dataset that will be used as test
  for(i in 1:k){
      f.idx[[i]] <-
        rbind(pos_data[s_pos[(1 + (i-1)*fold.size.pos):(i*fold.size.pos)],],
              neg_data[s_neg[(1 + (i-1)*fold.size.neg):(i*fold.size.neg)],])
  }
  return(f.idx)
## Use the function to create the 10 test datasets
split data <- prepare_cv_folds(10)</pre>
```

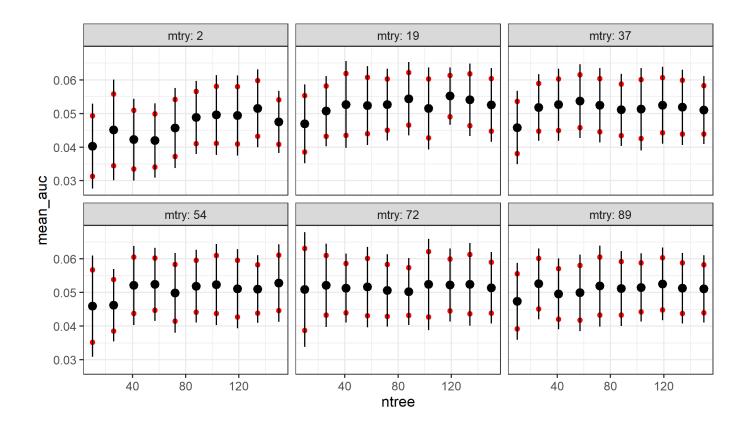
2.1 Decision tree



The graphique above shows the mean AUC as a function of different values of cp. The black lines represent the standard deviation and the red dots the Confidence Intervals. As we can see on this plot, reducing the complexity parameter below cp=0.001 doesn't change the mean $AUC_{0.2}$. This means that cp=0.001 is already sufficiently low. As a low cp means a more complex model, we are looking for the highest value of cp maximizing the mean AUC. Thus, we can select cp=0.001 to learn the decision tree.

However, as we can see, the mean AUC is always equal to 0.027, which is rather disappointing.

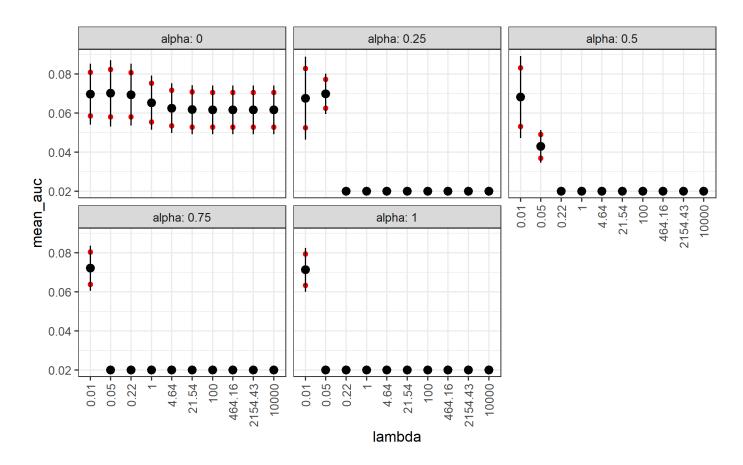
2.2 Random Forest



This plot shows the mean auc as a function of the number of trees. Each square correspond to a value of mtry indicated in the grey square. As we can see, the highest value of $AUC_{0.2}$ is for mtry=72 and ntree=10.

However, we must keep in mind that the Confidence Intervals always overlap, which means that the difference between the mean $AUC_{0.2}$ are not significant.

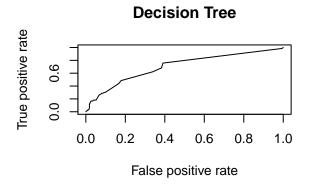
2.3 Regularized logistic regression

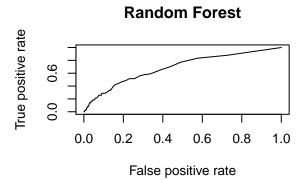


2.4 Evaluation on the test dataset

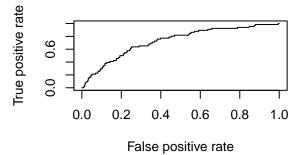
Thanks to the previous steps, we have chosen:

- 1. A decision tree with $\ensuremath{cp} = 0.00067$
- 2. A random forest with $ntree=120\ \mathrm{and}\ mtry=19$
- 3. A regularized logistic regression with $\lambda=0.01$ and $\alpha=0.75$



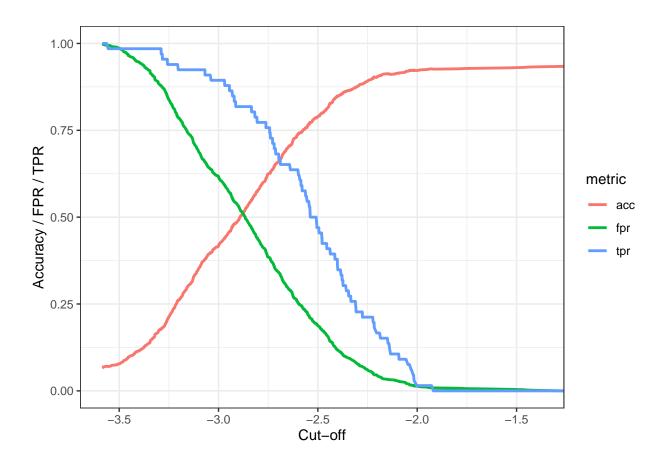


Regularized Logistic Regression



model	auc
Decision Tree	0.060
Random Forest	0.055
Regularized Logistic Regression	0.057

2.5 Setting cutoff threshold



The optimal cut-off seems to be 0.15 1) because we can see that this is the value where the accuracy is reaching a threshold and 2) because this is where the difference between the True Positive Rate and the False Positive Rate is the greatest. This second remark is important because here, we are looking for maximizing the number of true positive while minimizing the number of False Positive.

Now, we can compute the confusion matrix with a threshold of 0.15:

3. Task 3 - Model interpretation and feature selection

Table 3.1: Variable importance for the decision tree

feature	MeanDecreaseGini		
		feature	MeanDecreaseGini
PPERSAUT	16.39	MBERMIDD	3.81
PBRAND	12.61	MGODPR	3.63
APERSAUT	12.06	MINKGEM	3.53
MKOOPKLA	10.73	APLEZIER	3.19
PWAPART	9.52	PPLEZIER	3.19
MOSTYPE	9.41	МНКООР	3.14
MFGEKIND	9.25	MSKD	3.12
MBERHOOG	8.81	MFALLEEN	3.03
AWAPART	8.71	MSKC	2.90
MOSHOOFD	7.16	MGODOV	2.85
MOPLMIDD	6.77	MAUT0	2.73
MOPLLAAG	6.68	MSKB2	2.63
MRELGE	6.64	MBERARBO	2.52
ABRAND	6.16	MINK7512	2.35
MFWEKIND	5.93	MAUT1	2.10
MINK4575	5.69	MINKM30	1.83
MGODGE	5.58	MAUT2	1.77
MRELOV	5.25	MRELSA	1.13
MZPART	4.99	MAANTHUI	0.83
MOPLHOOG	4.83	MBERZELF	0.79
MZFONDS	4.68	MGODRK	0.78
MBERARBG	4.66	ALEVEN	0.70
MSKA	4.65	PLEVEN	0.70
MGEMLEEF	4.55	AWALAND	0.34
MHHUUR	4.53	PWALAND	0.34
MSKB1	4.24	ABYSTAND	0.31
MINK3045	4.14	PBYSTAND	0.31
MGEMOMV	3.85	PTRACTOR	0.30

Table 3.2: Variable importance for the random forest

feature	MeanDecreaseGini	feature	MeanDecreaseGini	feature	MeanDecreaseGini
PBRAND	20.09	MINKM30	7.55	AMOTSCO	2.73
MOSTYPE	17.20	MSKA	7.47	PBROM	2.70
PPERSAUT	16.73	MINK7512	7.41	PFIETS	2.62
APERSAUT	14.74	MZFONDS	7.40	ABROM	2.44
MKOOPKLA	11.94	MGODOV	7.33	MAANTHUI	2.43
PWAPART	11.54	MFALLEEN	7.29	PWAOREG	2.36
MGODGE	11.08	MRELGE	7.20	PGEZONG	1.65
MBERMIDD	11.03	MAUT1	6.95	AWAOREG	1.59
MGODPR	10.97	MZPART	6.82	PTRACTOR	1.20
MOPLMIDD	10.90	MRELOV	6.26	PINBOED	1.15
MOSHOOFD	10.27	MAUT0	6.18	AGEZONG	1.11
MBERARBG	10.12	MAUT2	5.99	AINBOED	1.02
MINK3045	9.88	MGEMLEEF	5.83	PAANHANG	0.97
MOPLLAAG	9.86	ALEVEN	5.55	PWABEDR	0.88
MFGEKIND	9.49	MSKD	5.39	ATRACTOR	0.78
MFWEKIND	9.46	PLEVEN	5.07	AWABEDR	0.73
MOPLHOOG	9.39	MGEMOMV	4.77	AAANHANG	0.69
MBERARBO	9.11	MGODRK	4.72	PBESAUT	0.49
MHHUUR	8.98	AFIETS	4.71	ABESAUT	0.47
MINK4575	8.82	MRELSA	4.61	AZEILPL	0.27
MSKB1	8.74	MBERZELF	4.37	APERSONG	0.25
MBERHOOG	8.71	PBYSTAND	3.99	PZEILPL	0.22
MSKC	8.31	ABYSTAND	3.73	AWALAND	0.19
МНКООР	8.27	PPLEZIER	3.70	PPERSONG	0.17
MINKGEM	7.92	APLEZIER	3.53	PWALAND	0.15
ABRAND	7.72	MBERBOER	3.38	PWERKT	0.04
MSKB2	7.66	PMOTSCO	3.24	PVRAAUT	0.01
AWAPART	7.65	MINK123M	2.76	AWERKT	0.01
	1			AVRAAUT	0.00

Table 3.3: Variable importance for the lasso regression

feature	s0	feature	s0
APLEZIER	1.2945711	PWABEDR	-0.0584269
AZEILPL	1.1414119	MOPLHOOG	0.0515774
ABYSTAND	0.7523564	MGODRK	-0.0435352
AFIETS	0.4165938	MAUT1	0.0356634
PWAOREG	0.2678450	MHHUUR	-0.0344877
PPERSAUT	0.2038757	MBERMIDD	0.0309560
MBERBOER	-0.1672857	PGEZONG	0.0288318
PWAPART	0.1551758	MGEMLEEF	0.0271291
MINK123M	-0.1506243	PWERKT	-0.0263719
PWALAND	-0.1470406	MGODGE	-0.0245771
ATRACTOR	-0.1432662	PBESAUT	-0.0218699
PBRAND	0.1026457	MKOOPKLA	0.0191547
MINKGEM	0.0765117	MRELSA	-0.0169228
MRELGE	0.0675735	MINK7512	0.0124976
MOPLLAAG	-0.0633978	MSKD	-0.0099615