

Predicting the length of stay and readmission probability of a patient in a hospital in Barcelona

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Introduction

Health service and hospitals are not topics that take so much time or importance on our day to day thoughts for most of us, but they are indeed crucial in guaranteeing our living standards. Public health care and public hospitals function on a complex balance between efficiency, budget restraints and the fulfill of moral and etic duties. The intricacy of how all this is achieved year after year is only known by the specialists; doctors, nurses, managers etc.

A common issue they have to deal with is the budget restrictions or limitations (a condition imposed by the government in order to conceed public funds) based on the number of readmissions that the health center has. On one hand a deeper study and prediction of the readmissions could be both interesting and key in the future development of hospitals, not only to reduce the budget sanctions but to improve the quality of the attention and health after a patient leaves the hospital.

On the other hand, the efficient allocation and use of their resources (human assets, lab tests and procedures...) can contribute to the profit of health services and hospitals. The availability of beds is an indicator of this efficient allocation of resources. Therefore, predicting the patient's length of stay in the hospital can be beneficial for the doctors and managers to act and react efficiently to each patient and circumstance.

All these challenges are particularly interesting for data science projects since their identification, model and prediction are not only important but they can also be rather complicated. Understanding this data and its challenges can lead to a powerful breakthrough, identifying waste in the system, which can allow the healthcare service to work more efficiently. The increase in efficiency could terefore potentially improve the allocation of resources and lead to an improved level of care for patients.

Data description

For this project we were provided with data from a hospital in Barcelona. The dataset provides us with the following variables:

-*Age*, *Gender*, *City* (where the patient lives), *Region* (where the patient lives), *Country* (where the patient lives), *Zip Code*, *Basic Health Area*, *Country of Birth* (where the patient was borned), *Insurance Company*, *Primary Diagnostic*, *Secondary diganostic/s*, *Number of comorbidities*, *Especial clinical condition/s*, *Mental illness* (binary whether the patient has one or not), *Alcohol and drugs* (binary whether the patient took some or not), *Neoplasm* (binary whether the patient has it or not), *Mentally handicapped* (binary whether the patient has it or not), *Respiratory disease* (binary whether the patient has one or not), *Diabetes* (binary whether the patient has it or not), *Heart failure* (binary whether the patient had it or not), *Loss of wieght* (binary whether the patient had it or not), *Depression* (binary whether the patient has one or not), *Anemia* (binary whether the patient has it or not), *PCC* (binary whether the patient is a chronic complex patient or not), *MACA* (binary whether the patient has a complex chronic disease or not), *Number of drugs prescribed at discharge*, *Date of admission*, *Time of admission*, *Day of the week of admission*, *Year of admission*, *Month of admission*, *Date of discharge*, *Time of discharge*, *Day of the week of discharge*, *Year of discharge*, *Month of discharge*, *Lenght of stay (LoS)*, *Medical specialty that handled that patient*, *Oncological area* (binary whether the patient was treated by the oncological area or not), *Admission origin*, *Number of procedures*, *Number of lab tests*, *Number of yearly admissions*, *Number of yearly emergencies*, *Number of yearly visits*, *Days until*

readmission, *Readmission of type 1* (whether a patient was admitted within the next 30 days after he was discharged and by a diagnostic that belongs to the same diagnostic group the original diagnostic he was admitted for belonged to), *Readmission of type 2* (same as readmission of type 1 but within the next 60 days after the patient was discharged instead of 30), *Readmission of type 3* (any patient that came back to the hospital within the next 15 days for any reason).

The response variables used in this project's analysis will be *LoS*, *Readmission of type 1*, *Readmission of type 2* and *Readmission on type 3*.

The number of observations of the original dataset are 18892.

##	EdatEnAlta	Sexe	Poblacio	Provincia	Pais	CodiPostal	AreaBasicaSalut	
## 1	67	D	BARCELONA	BARCELONA	ESPANYA	08036	BARCELONA 2-E	
## 2	67	D	BARCELONA	BARCELONA	ESPANYA	08017	BARCELONA 5-C	
## 3	55	D	BARCELONA	BARCELONA	ESPANYA	08003	BARCELONA 1-A	
## 4	83	D	BARCELONA	BARCELONA	ESPANYA	08022	BARCELONA 5-E	
## 5	85	D	BARCELONA	BARCELONA	ESPANYA	08032	BARCELONA 7-D	
##	PaisNaixement	Cobertura	GrupDiag	NumComorbiditats	CondClinEspec			
## 1	ESPANYA	SCS	07I	9	Y			
## 2	ESPANYA	SCS	02B	8	N			
## 3	REPUBLICA DOMINICANA	SCS	02B	8	Y			
## 4	ESPANYA	COL	VG	7	N			
## 5	ESPANYA	COL	VG	1	N			
##	MalaltiaMental	AlcoholDrogues	Neoplasia	RetardMental				
## 1	N	N	N	N				
## 2	N	N	Y	N				
## 3	N	N	Y	N				
## 4	N	N	N	N				
## 5	N	N	N	N				
##	MalaltiaRespiratoria	Diabetis	IC	PerduaPes	Depressio	Anemia	PCC	MACA
## 1	N	Y	N	N	N	N	N	N
## 2	N	N	N	N	N	N	N	N
## 3	N	Y	N	N	N	N	N	N
## 4	N	N	N	N	N	N	N	N
## 5	N	N	N	N	N	N	N	N
##	NumMedAlta	DataIngres	HoraIngres	HoraIngres2				
## 1	0	2012/11/22 00:00:00.000	1070	1200-1800				
## 2	0	2012/11/26 00:00:00.000	523	0600-1200				
## 3	0	2012/11/28 00:00:00.000	598	0600-1200				
## 4	0	2012/11/30 00:00:00.000	801	1200-1800				
## 5	0	2012/12/05 00:00:00.000	1041	1200-1800				
##	DiaSetmanaIngres	PeriodeIngres	MesIngres	DataAlta				
## 1	3	2012	10	2013/02/25 00:00:00.000				
## 2	0	2012	10	2013/01/04 00:00:00.000				
## 3	2	2012	10	2013/01/28 00:00:00.000				
## 4	4	2012	10	2013/01/04 00:00:00.000				
## 5	2	2012	11	2013/01/03 00:00:00.000				
##	HoraAlta	HoraAlta2	DiaSetmanaAlta	PeriodeAlta	MesAlta	DuradaEstada		
## 1	573	0600-1200	0	2013	1	95		
## 2	360	0000-0600	4	2013	0	39		
## 3	225	0000-0600	0	2013	0	61		
## 4	876	1200-1800	4	2013	0	35		
## 5	922	1200-1800	3	2013	0	29		
##	Servei	ServeiOnco	OrigenAdmissio	NumProc	NumLab			
## 1	DERM	N	Normal	0	0			

## 2	CGEN	N		Normal	0	0
## 3	ONCO	S		Normal	0	0
## 4	MI	N	OTROS CENTROS HOSPITALARIOS		0	0
## 5	MI	N	OTROS CENTROS HOSPITALARIOS		0	0
##	NumAdmissionsAny	NumUrgenciesAny	NumVisitesAny	DiesReadmissio		
## 1	0	0	0		0	
## 2	0	0	0		0	
## 3	0	0	0		0	
## 4	0	0	0		0	
## 5	0	0	0		0	
##	ReadmissionN1	ReadmissionN2	ReadmissionN3			
## 1	N	N	N			
## 2	N	N	N			
## 3	N	N	N			
## 4	N	N	N			
## 5	N	N	N			

Literature review and software

Literature review

A literature review revealed that little has been published with regard to this topic within Europe and that the vast majority of publications are from the USA ^{1, 3, 4, 5}. However, the main focus of this literature review will be on the following two papers due to their approach to the challenge of predicting hospital readmissions which could later be replicated by our study.

- Development and Implementation of a Real-Time 30-Day Readmission Predictive Model

Cronin et al., (2014) used the data of Massachusetts General Hospital (MGH) to explain the reasons behind the pressure on the hospitals to reduce readmission rates of patients. Cronin et al., (2014) explained that reliable predictions for patients rehospitalisation would allow for tailored interventions to be introduced for patients most at risk. This would require “the creation of a functional predictive model specifically designed to support real-time clinical operations” which comes with some challenges to solve and options to test.

- A comparison of models for predicting early hospital readmissions

Differently to the previous paper, Futuoma et al.², (2015) uses a dataset called “the New Zealand National Minimum Dataset, obtained from the New Zealand Ministry of Health”. The authors take into account that “there are a number of published risk models predicting 30 day readmissions for particular patient populations, however they often exhibit poor predictive performance and would be unsuitable for use in a clinical setting” and they “describe and compare several predictive models, some of which have never been applied to this task and which outperform the regression methods that are typically applied in the healthcare literature”.

Software

The software used in this thesis to clean, process, analyze, model and predict has been the open source statistical program R as well as the data mining with open source machine learning software in java Weka.

In the case of R, several packages have been used in order to maximize the efficiency and the quality of the outcome. Such packages are:

- For the data vizualization work: corrplot, ggbiplot, ggplot2, reshape2, caret and splines.

- For the analysis, modeling and predicting part: caret, e1071, plyr, parallel, gbm, ranger, caTools, glmnet and mlr.

In the case of Weka, the analysis conducted has used the costMatrix function (manually tuned) with a random forest of 100 trees and a data splitting of 75% training set and 25% test set.

Further explanation on how exactly the algorithms have been trained and tuned along with the challenges overcome and results of the process will be explained in subsequent sections of the project.

Case study

Aim

The goal is to predict the length of stay of a patient and the probability of readmission. These predictions (if made accurately) will both trigger the efficiency and quality of the hospital in treating their patients and reduce the sanctions by the government. This would therefore augment the budget of the hospital, increasing the capacity of the hospital to invest in R&D, new equipment, personal training etc.

Methodology

Models will be trained and tested to predict both the length of stay and the probability of readmissions. We will then compare the models and evaluate which model performs better, according to the specific criteria that this problem requires.

To train the models we will use almost all the variables listed in the data description section. The choice of the final variables along with the variable transformation will be explained in more detail in the analysis section.

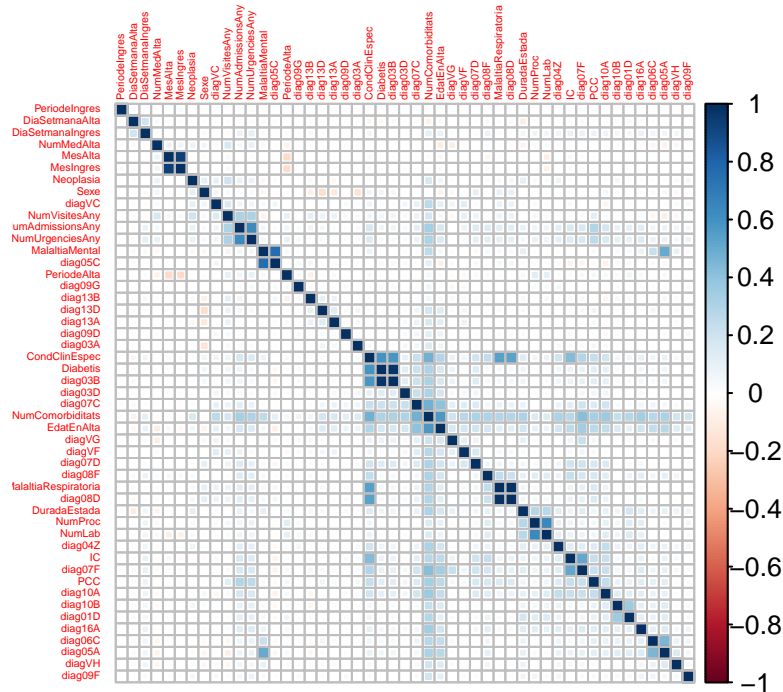
Visualization

To grasp a sense of what the dataset looks like, how the variables are related to each other and how the patients' characteristics are distributed we used some visualisation techniques that will be presented right after.

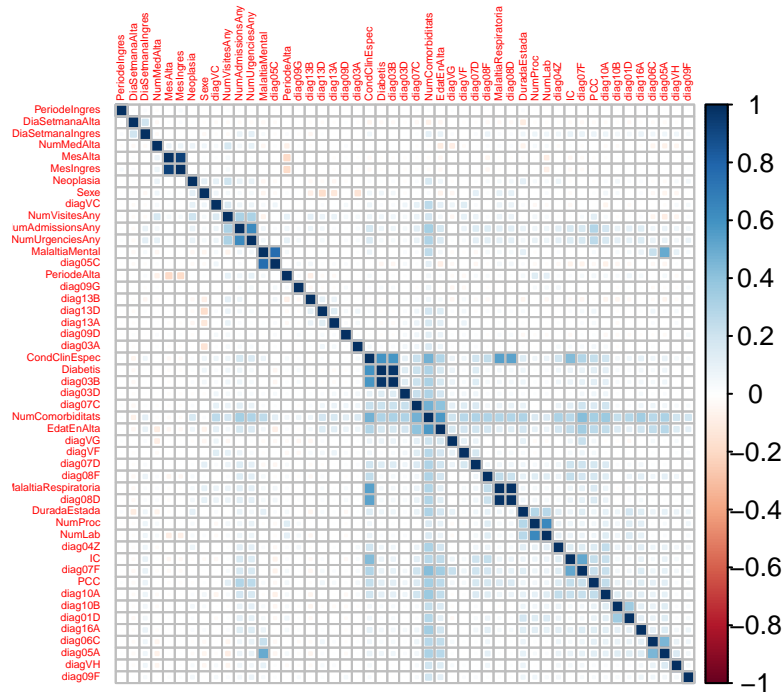
We first started by plotting some correlograms to see the possible correlation between the non-categorical variables (those that are numeric or binary and also doesn't have 0 or near zero variance).

There are three correlograms, the first one considers all the observations. The following two are subject to the readmission response variables, therefore one correlogram represents the correlation of the variables whose observations have a negative value in the response readmission variables. The other correlogram is considering those observations that have a positive response value in at least one of them.

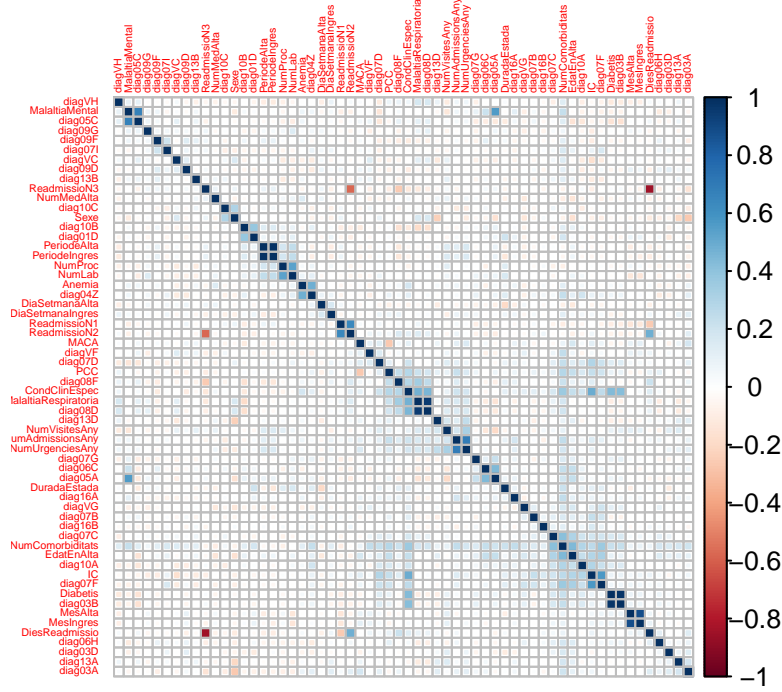
Correlogram with all observations



Correlogram with negative readmitted observations



Correlogram with positive readmitted observations



We can appreciate how we find some strong correlations like the correlation between *Month of admission* and *Month of discharge*. This is because the average length of stay in hospital is less than 30 days, therefore the month of admission and month of discharge are usually within the same month. We also see a strong correlation between some diagnostic codes and their strictly related diseases such as respiratory diseases, mental illness or diabetes. Finally the number of comorbidities has a strong correlation with some diagnostic codes (representing the most common secondary diagnostics).

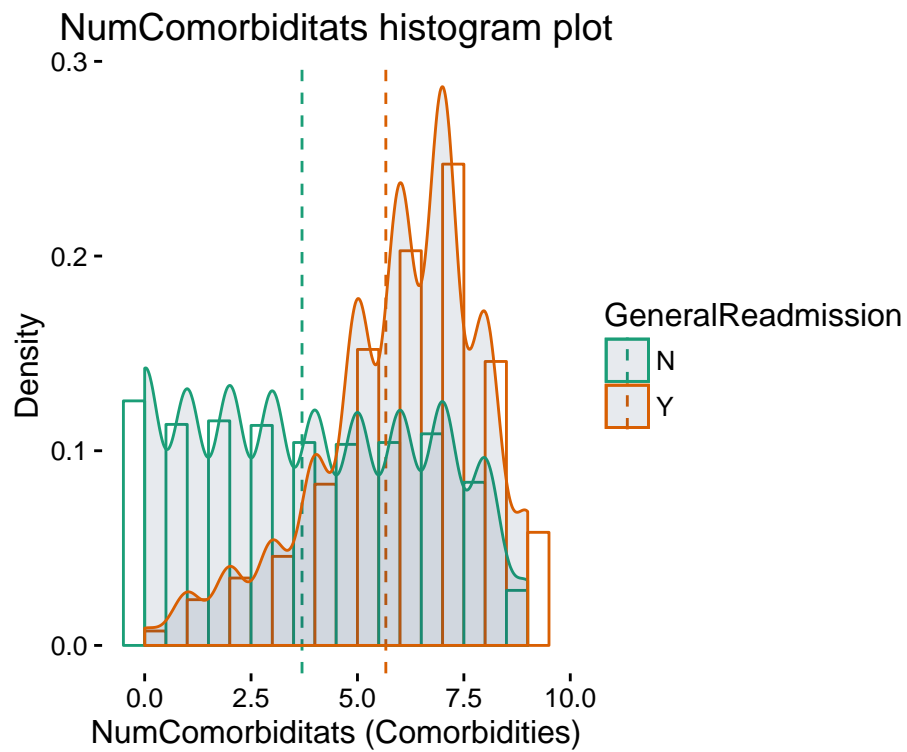
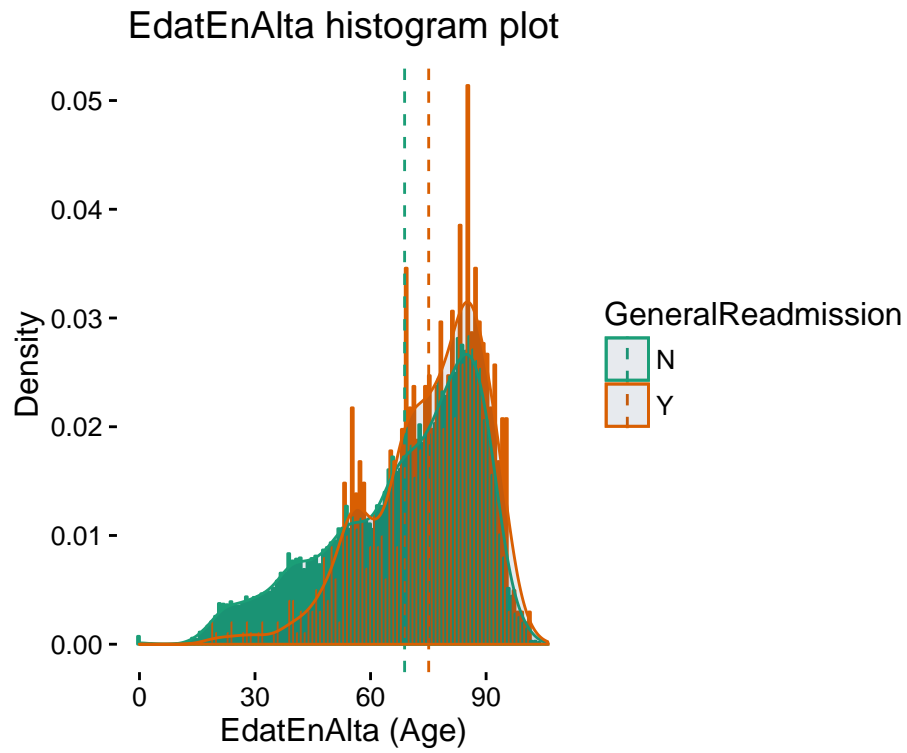
Most of the correlations we observe are positive (shown in blue while negative correlations will have a red tonality) and the pattern of correlation replicates through all three correlograms. One notable difference is in the one including only the observations with any kind of readmission where some strong negative correlations appear.

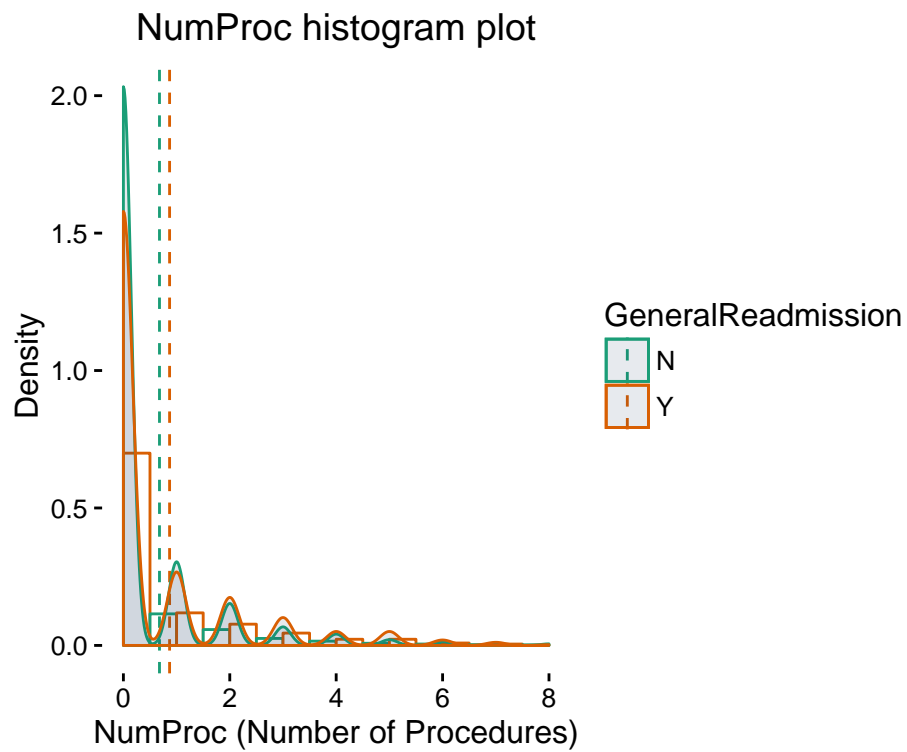
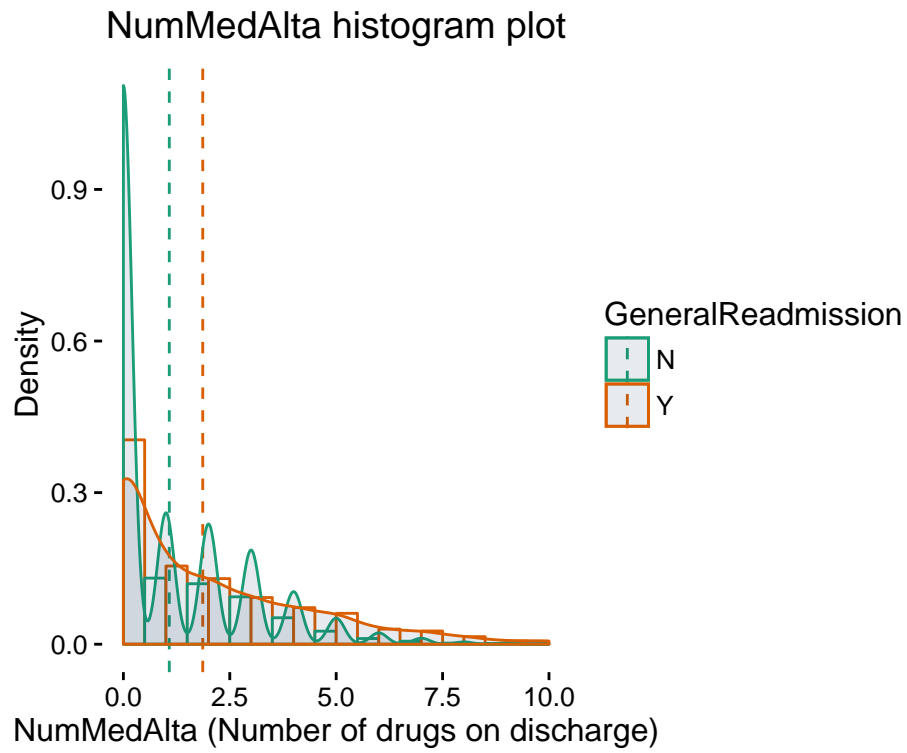
The strongest negative correlation appears between the *Days until readmission* variable and the *Readmission type 3* (R3) variable. This is due to the *R3* variable being defined as every readmission that happens for any reason within 15 days after discharge. As a result the positive responses of *R3* match exactly with the lower values of the *Days until readmission* variable.

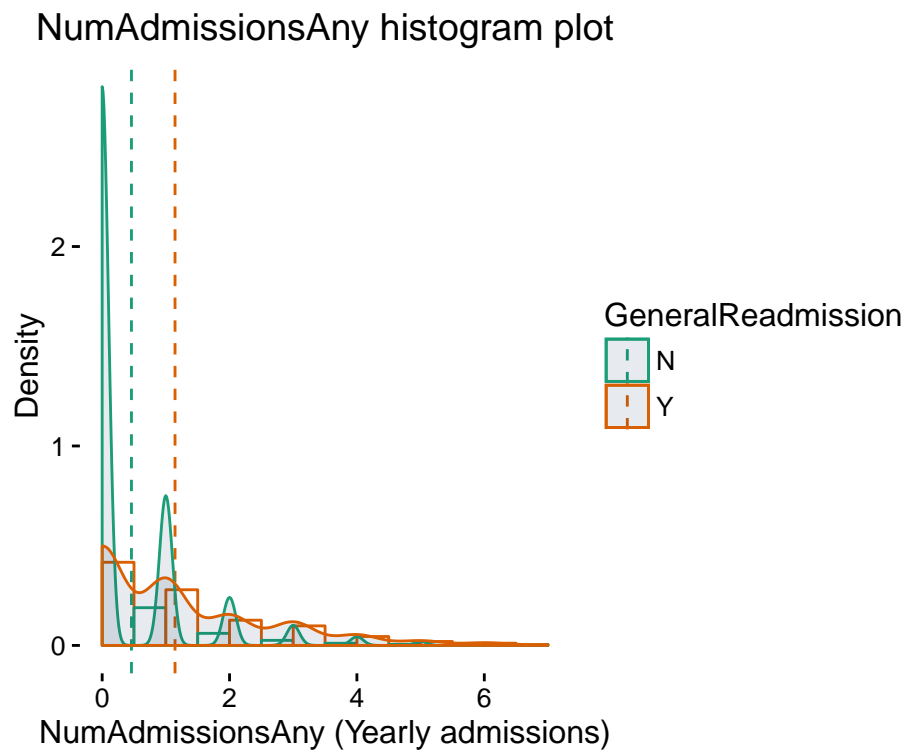
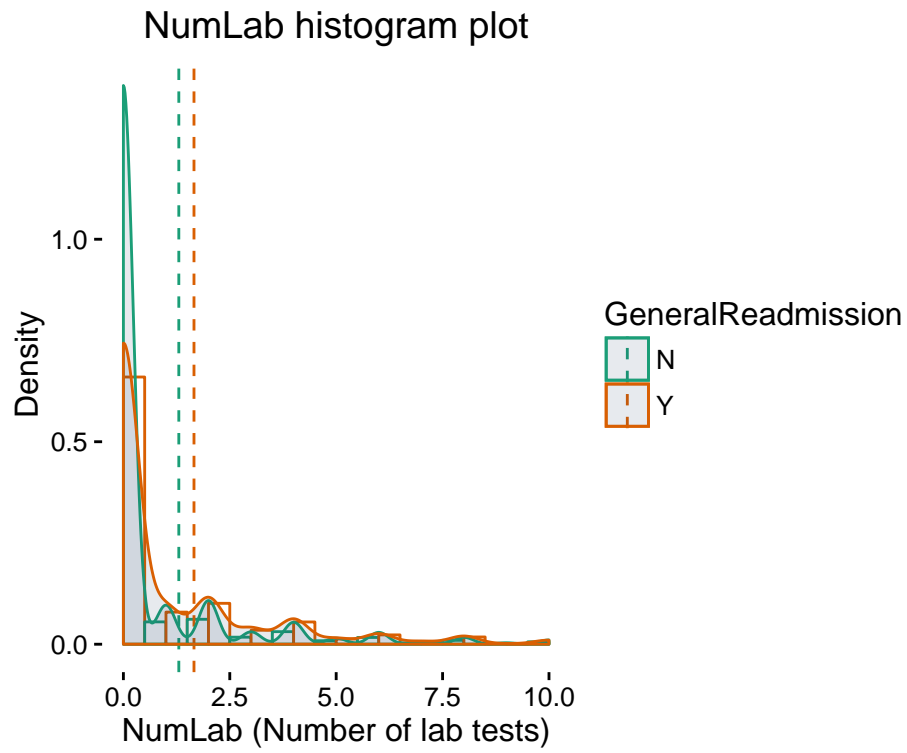
The other negative correlation appears between *R3* and the *Readmission type 2* (R2) variable. One reason for this could be because the R2 accepts readmissions up till 60 days after discharge and will capture those readmissions with the higher *Days until readmission* values (as opposed to *R3*). *R2* also have some medical restrictions: just returning to the hospital is enough for *R3* but not enough for *R1* and *R2*, as we explained above in the definition of the variables, so no all *R3* values will be at the same time *R2* or *R1*.

For some of the numeric variables it is also interesting to see how different the distributions are between the individuals with a positive response in, at least, one of the readmission variables and those with a negative response in all *R1*, *R2* and *R3*.

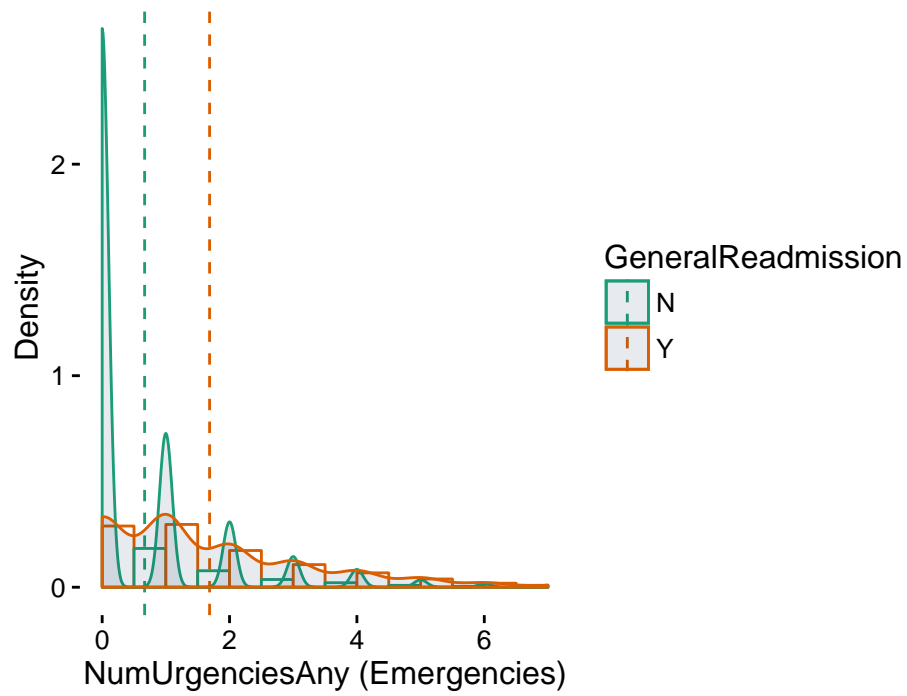
We built histograms to better display the information, plotting in different colors the different distributions and adding a dashed line that represents the mean of each group.



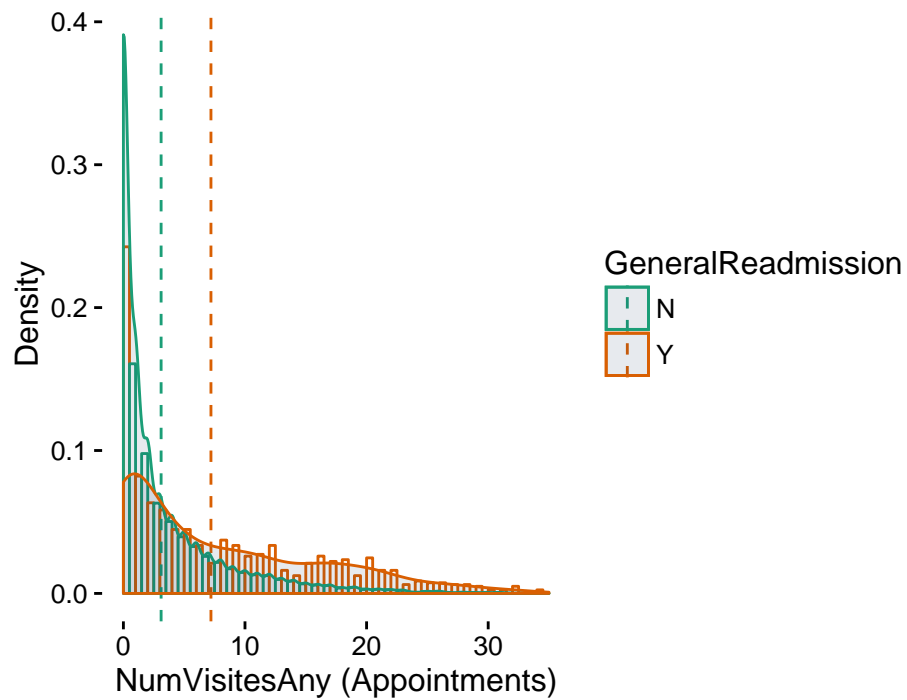


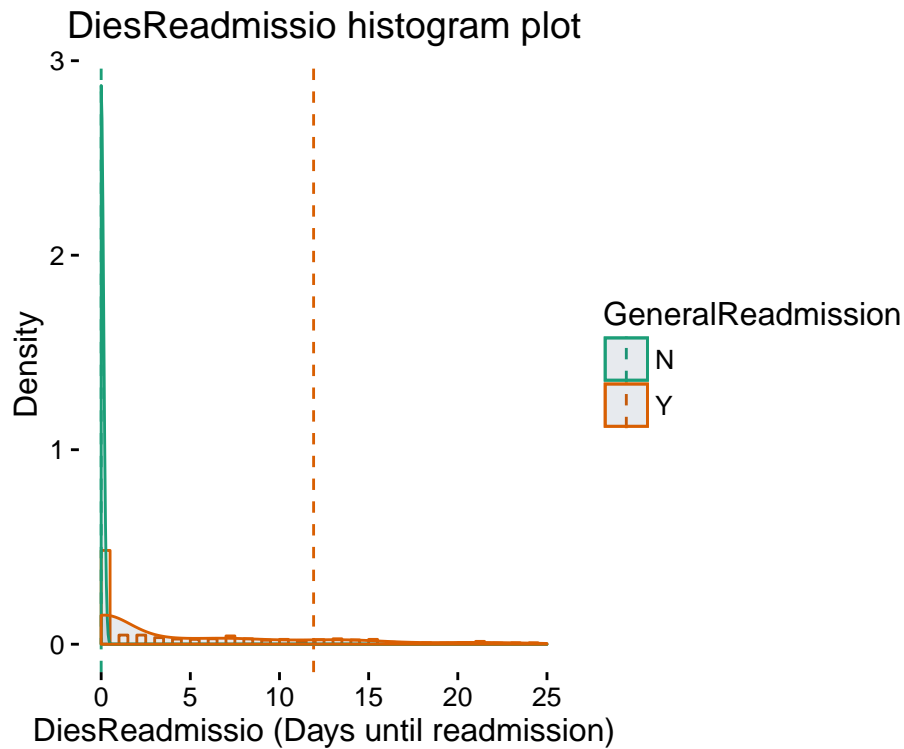


NumUrgenciesAny histogram plot



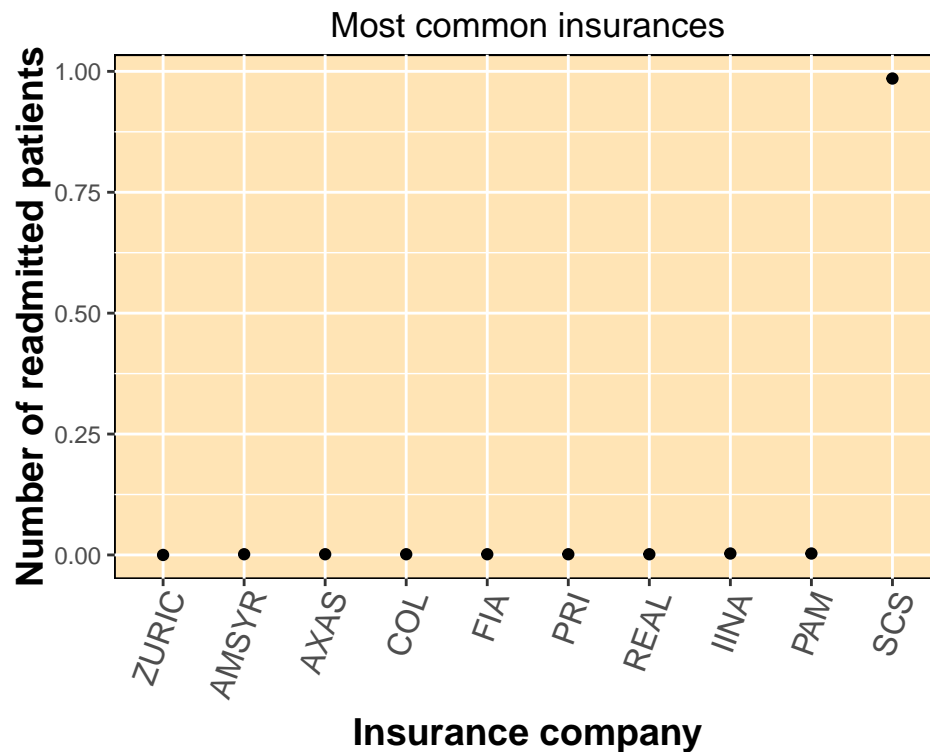
NumVisitesAny histogram plot

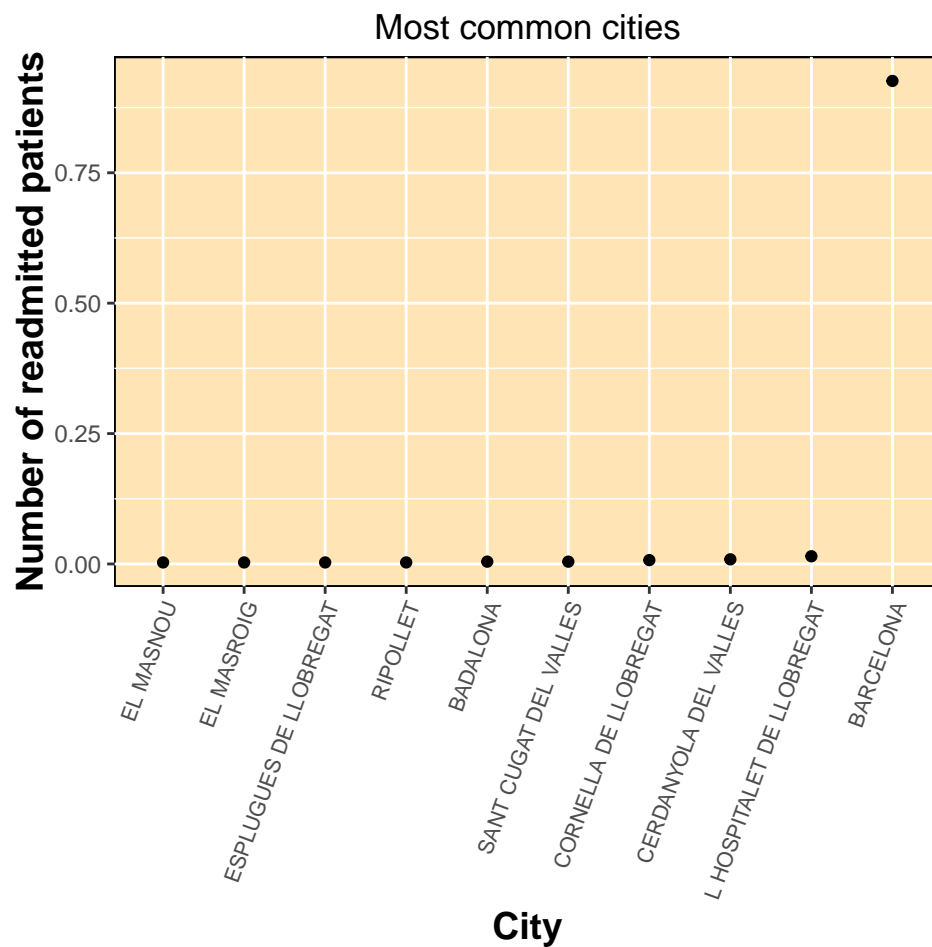
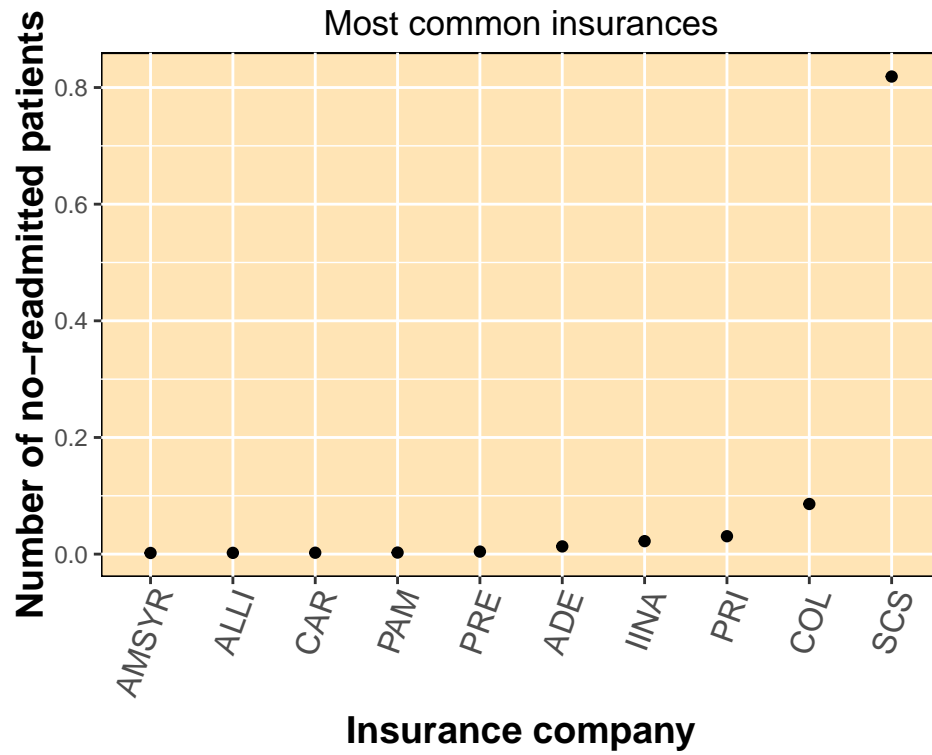


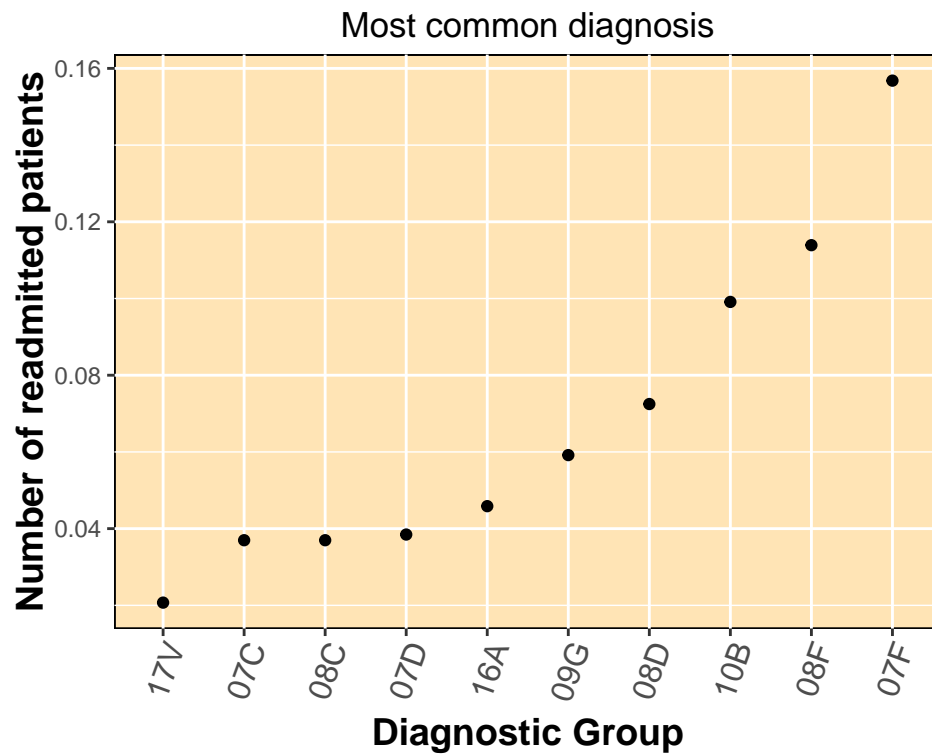
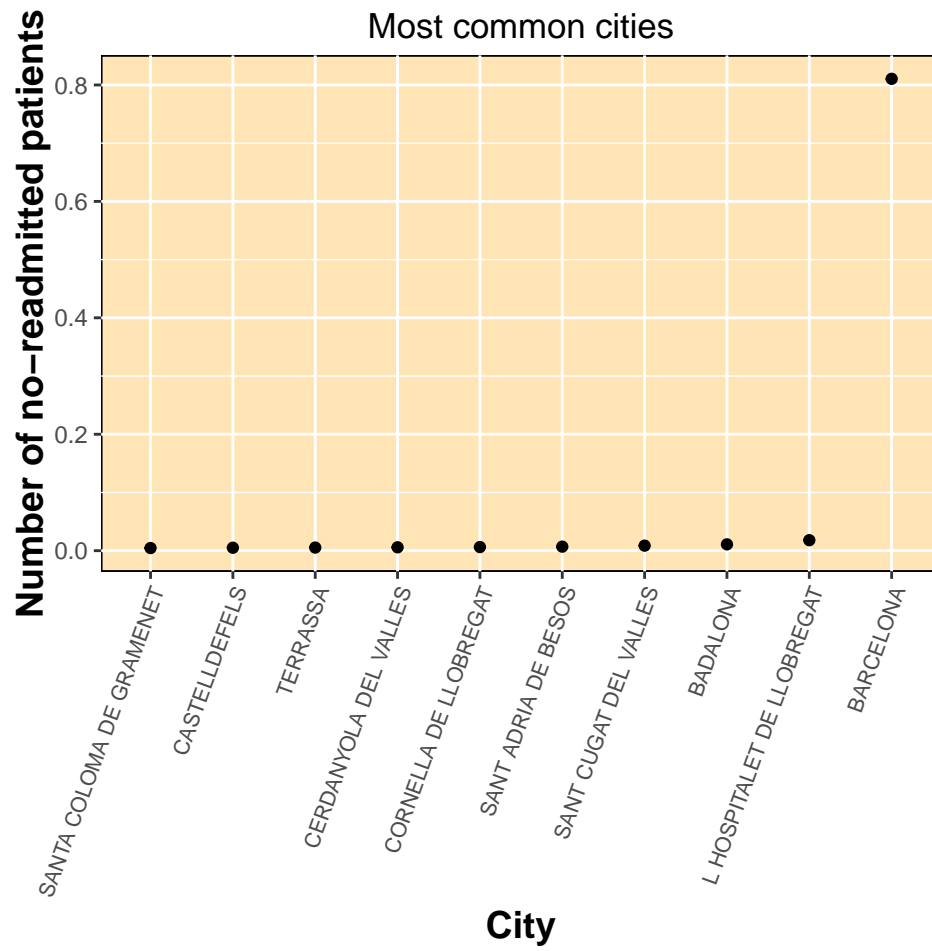


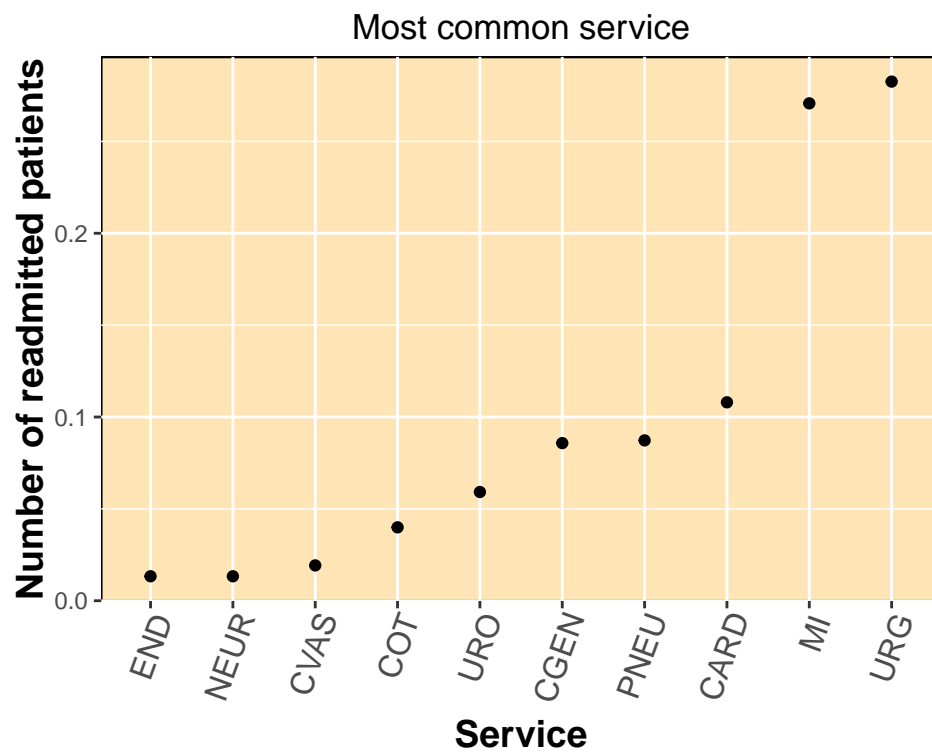
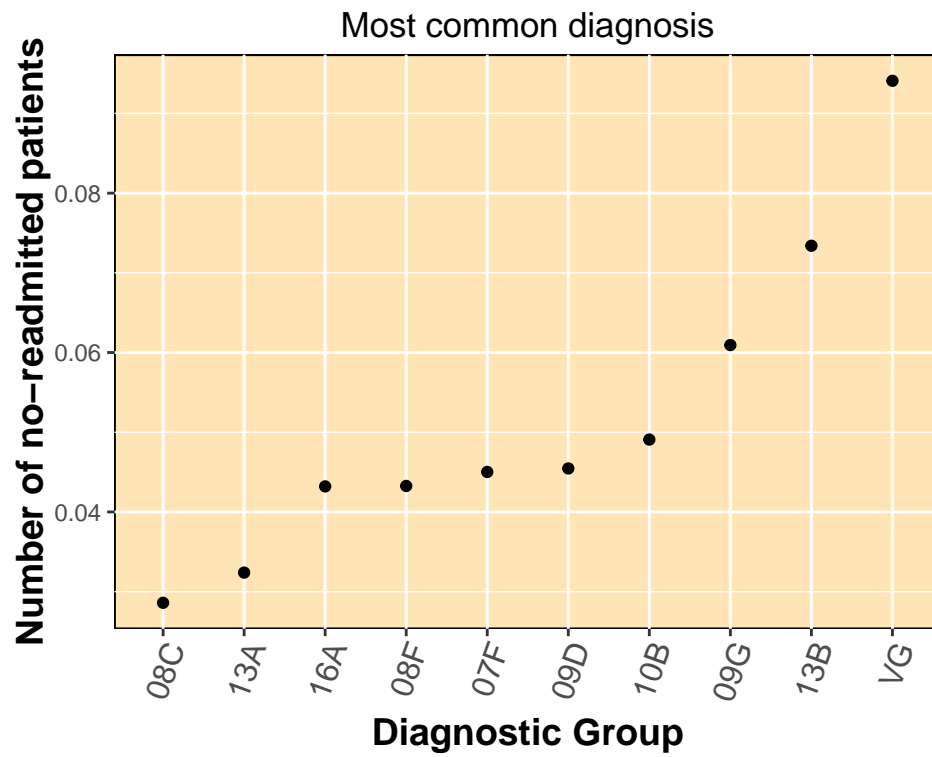
We can appreciate how the means are higher for the group with positive response in the readmission variables in all the histograms, this indicates that the patients that readmit are older, with more health problems, that required a longer and more complicated treatment and that attend the hospital more times along the year.

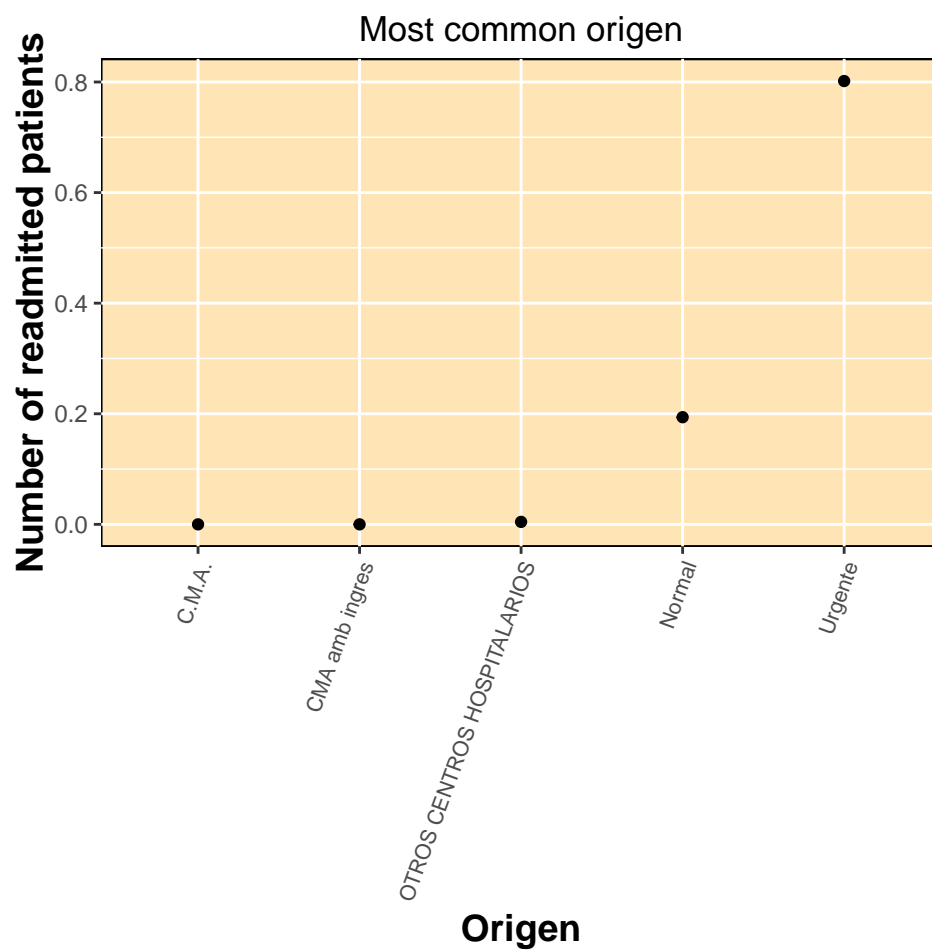
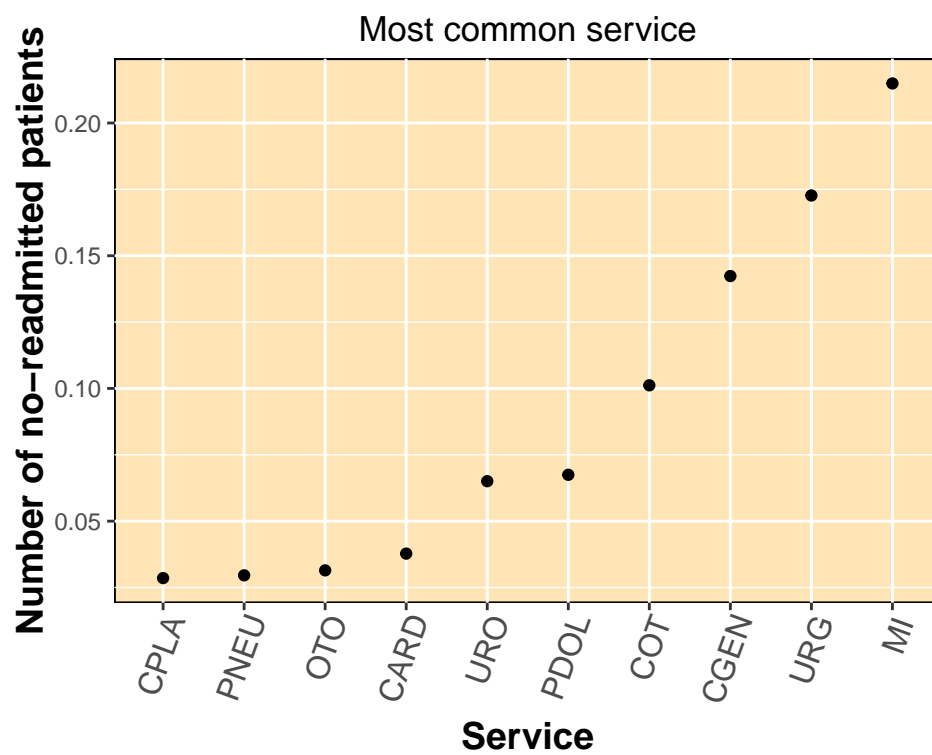
To visualize the most important categorical variables we made some plots where we can see the top 10 categories of each variable and the percentage of the number of observations belonging to each category.

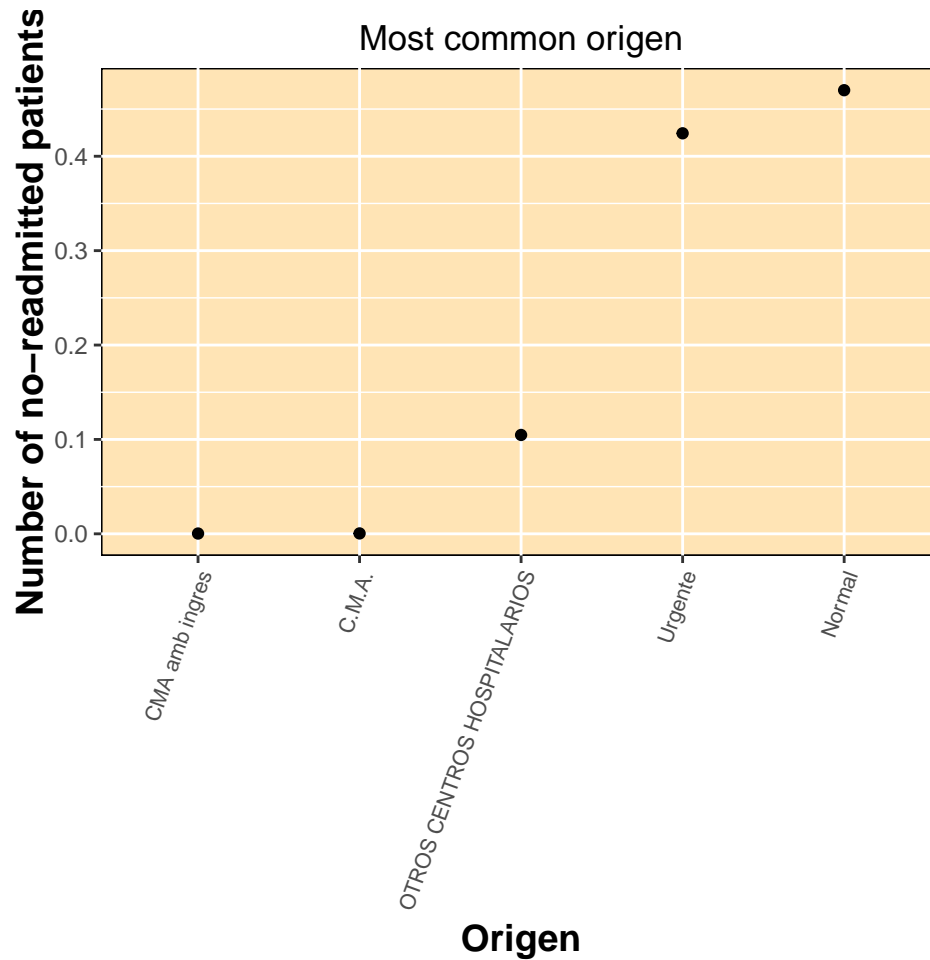












We can see that in almost all variables the vast amount of observations are concentrated in one category. The data is highly clustered in this sense because all of the data comes from the same hospital and most of the patients there share a common profile.

There are interesting differences to be noted between the two *Most common diagnostic* plots and *Most common services* plots. They are significantly different between the no-readmitted patients and the readmitted ones. Both plots are correlated to each other because different diagnostics were probably taken care in different health services. Taking into account this differences we can start assuming that maybe the diagnostic has something to do with the readmission.

Analysis

To analyze our dataset and predict our response variables we dropped some variables and needed to make a few changes in some other variables.

We dropped the *Region*, *Country* (both where the patient was born and lives), *Zip Code* and *Basic Health Area*, *Date of Admission*, *Days until readmission* and *Date of Discharge* for efficient purposes.

We generated the following variables:

-Population 2: this new variable breaks the *Population* variable into two categories: Barcelona and others. We did that because, as we saw in the visualization part we had the majority of observations grouped under the category Barcelona and therefore grouping all the other cities into one category gave us more predictive

power and made more sense in general since the hospital is indeed based in Barcelona so the difference would be if the patient lives in Barcelona or not.

- **Insurance company 2:** this new variable breaks the *Insurance company* variable in two categories: social security and others. We did this for similar reasons as the ones explained above in the *Population 2* variable.

- **Primary diagnostic:** what we did in this case was to group the primary diagnostics in groups that encapsule similar pathologies.

- **Secondary diagnostic:** what we did to this categorical variable was to transform each secondary diagnostic code into a binary variable and give it a 1 to all the observations that had that code in the old secondary diagnostic variable and a 0 otherwise. With this process we created 157 new binary variables.

- **Time of admission:** we transformed this variable and defined into the following blocks: from 00:00 to 06:00, from 6:00 to 12:00, from 12:00 to 18:00 and from 18:00 to 00:00, the intervals are with the first time included and the second one excluded. We justify this again for efficiency and predictive reasons, by grouping the times in these intervals we have more observations represented in each one than by considering the exact time for each observation.

- **Time of discharge:** we applied exactly the same procedure as we applied to the *Time of admission* variable.

Predictions

After this we moved on to the predictions. We first tried to predict the *LoS*. For that we tried gbm, glmnet and random forest (RF) methods and we realized that due to the distribution of the data (most of the values for LoS are 2) it was extremely hard to predict with accuracy the exact number of days. After some considerations on the actual value of predicting an exact number of days with a high probability of predicting them wrong we considered that for the information that the doctors are actually going to get from this prediction and for prediction accuracy purposes we should transform this numeric variable into a categorical one. The new *LoS* categorical variable had the following categories: less than 2 days, from 3 to 7 days and more than 7 days.

With this new variable we got the following results:

GBM	Less than 2 days	3 to 7 days	More than 7 days
Less than 2 days	93.59%	4.31%	6.1%
3 to 7 days	2.64%	80.7%	25.33%
More than 7 days	3.76%	15%	68.57%

GLMNET	Less than 2 days	3 to 7 days	More than 7 days
Less than 2 days	90.41%	17.31%	4.69%
3 to 7 days	7.94%	62.71%	27.57%
More than 7 days	1.64%	18.97%	67.73%

RF	Less than 2 days	3 to 7 days	More than 7 days
Less than 2 days	91.02%	8.53%	2.72%
3 to 7 days	6.82%	72.58%	25.33%
More than 7 days	2.15%	18.89%	71.95%

There are 2418 observations of the class *Less than 2 days* and the best prediction is made by GBM, 1207 of the class *3 to 7 days* and the best prediction is made by GBM and 1066 of the class *More than 7 days* and the best prediction is made by RF. Two out three of these predictions are made using GBM. Considering all this and the difference in the other predictions between models, if we had to pick a model as the model that performs better in predicting the response variable *LoS*, that would be GBM.

For the predictions of *R1*, *R2* and *R3* we went with the logistic regression and the glmnet and random forest.

The biggest challenge we had to face was the severe class imbalance that the dataset had. The positive response for *R1* represented only 1.54% of the observations, for *R2* represented 2.31% and for *R3* represented 2.22%. The approaches we took to solve this problem were different depending on the model.

For logistic regression and glmnet we optimized the threshold in order to maximize the F1 score, which is the harmonic mean of the precision and recall.

The following are the results for each model on each response variable:

- Logistic Regression (tuning the threshold and maximizing the F1 score)

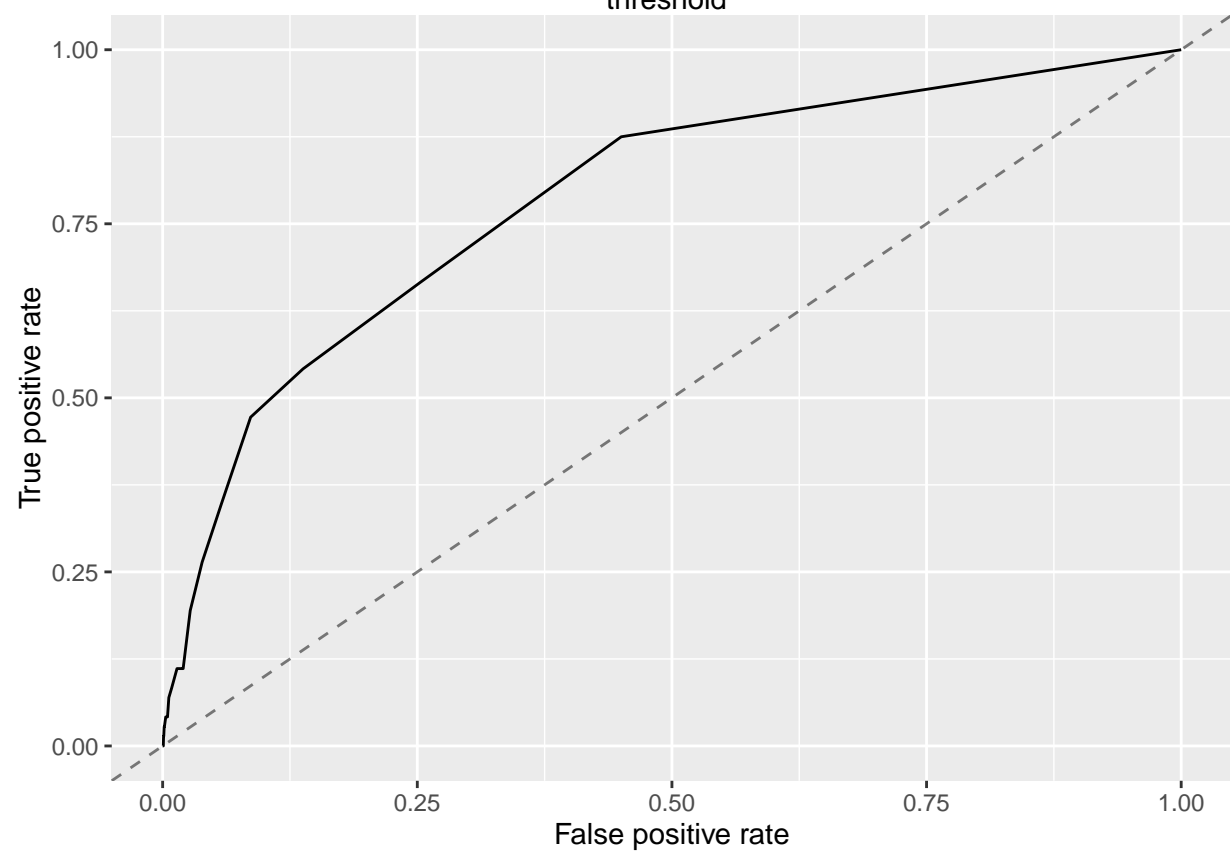
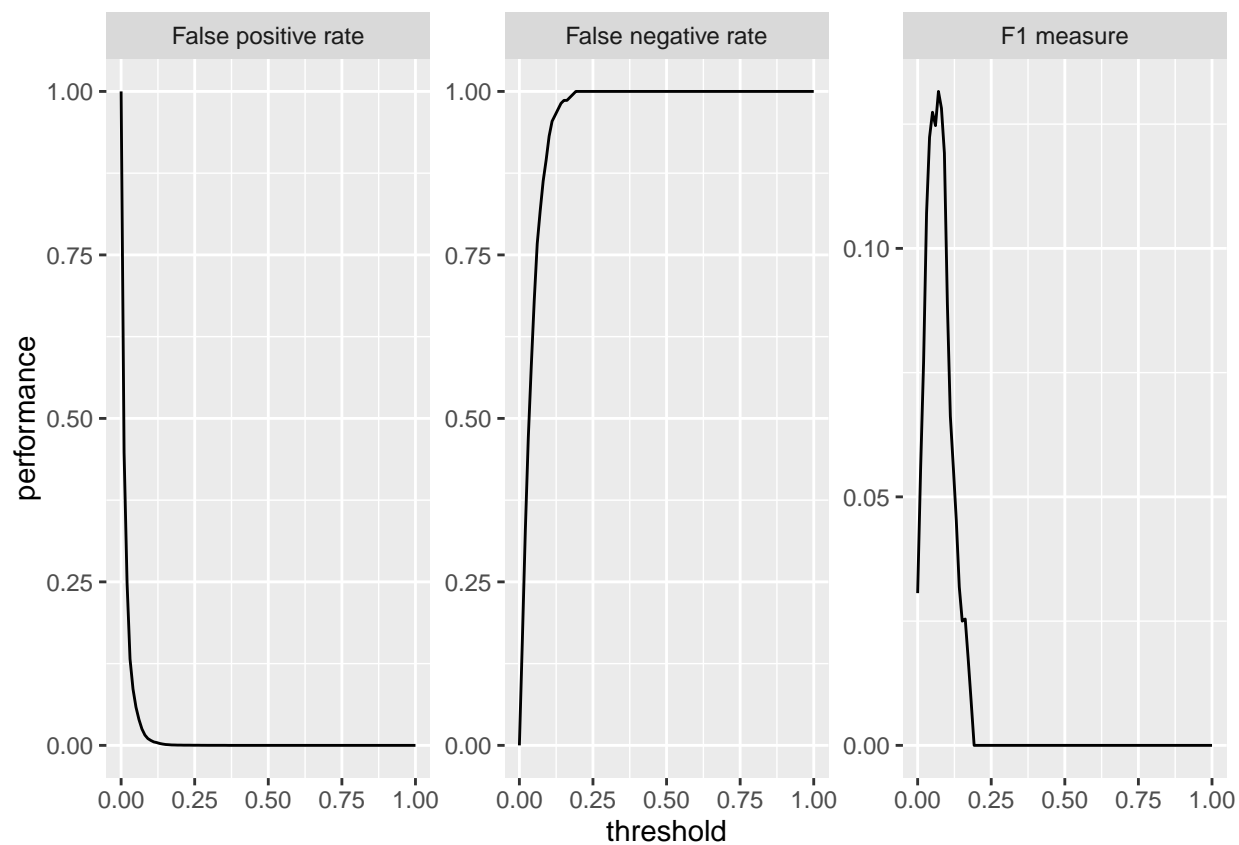
Readmission Type 1 results:

```
$train.metrics
      acc      auc      ppv      brier
0.98444492 0.79704448      NaN 0.01489564
```

```
$test.metrics
      acc      auc      ppv      brier
0.96099744 0.80295364 0.10071942 0.01467203
```

```
$train.matrix
      predicted
true      0 1 -SUM-
0      13860 0      0
1       219 0     219
-SUM-   219 0     219
```

```
$test.matrix
      predicted
true      0 1 -SUM-
0       4495 125    125
1        58 14     58
-SUM-    58 125    183
```



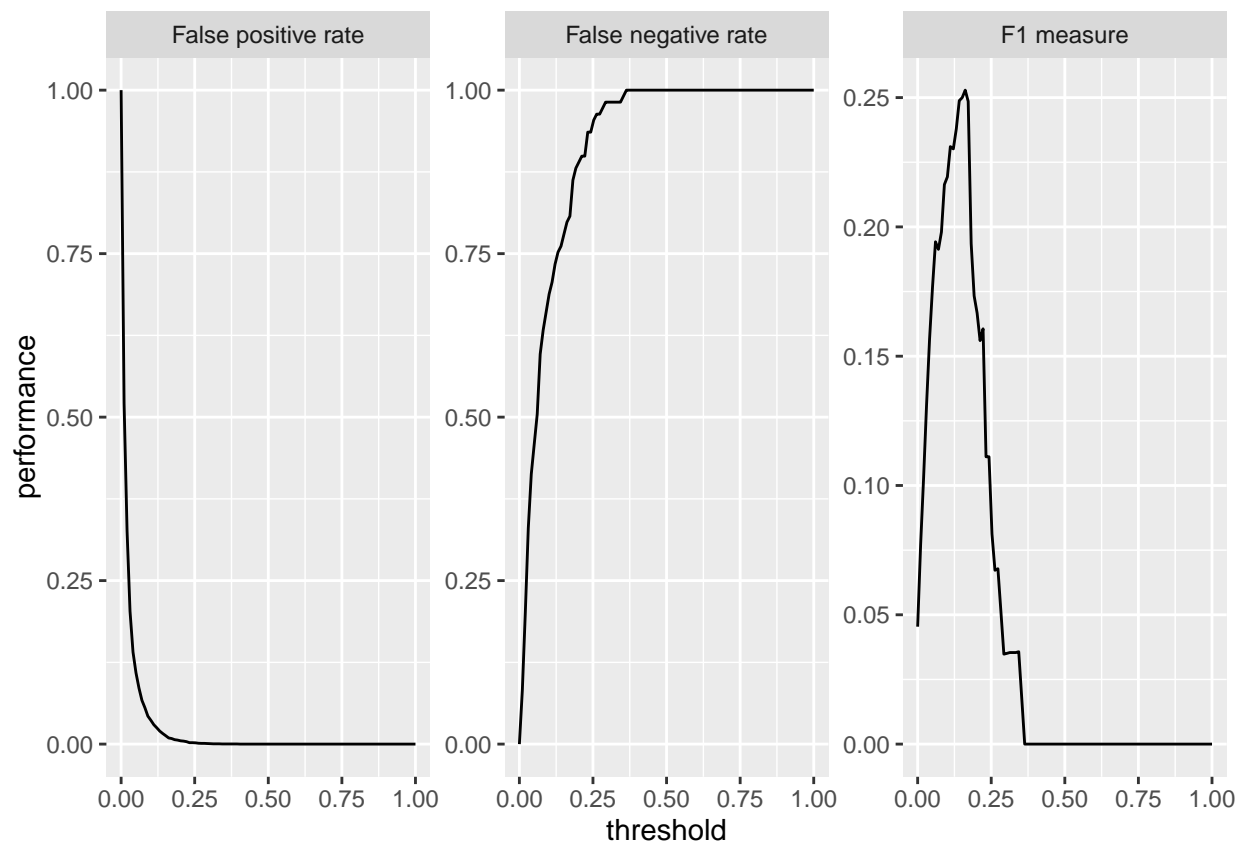
Readmission Type 2 results:

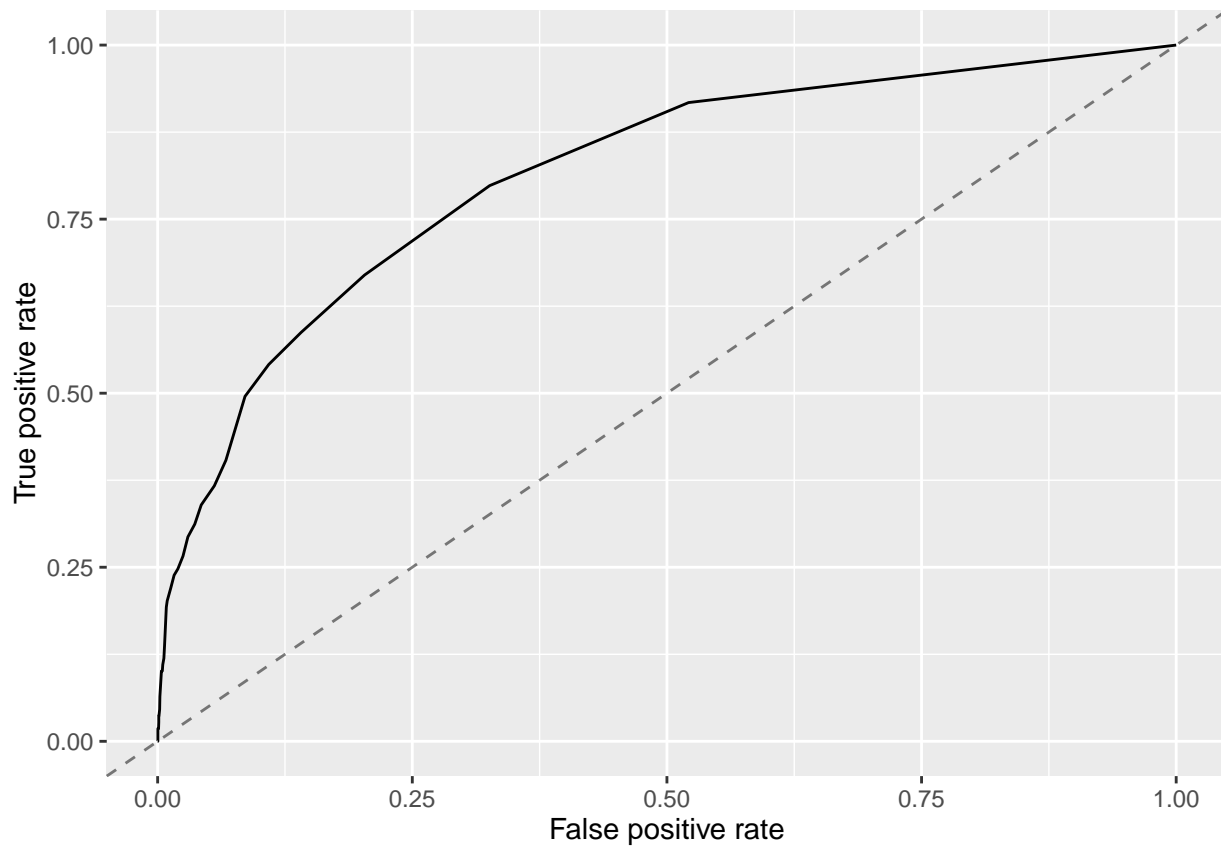
```
$train.metrics
      acc      auc      ppv      brier
0.97677392 0.82348734      NaN 0.02147668
```

```
$test.metrics
      acc      auc      ppv      brier
0.94842285 0.81646672 0.16915423 0.02108385
```

```
$train.matrix
      predicted
true      0 1 -SUM-
0      13752 0 0
1       327 0 327
-SUM-   327 0 327
```

```
$test.matrix
      predicted
true      0 1 -SUM-
0      4416 167 167
1       75 34 75
-SUM-   75 167 242
```





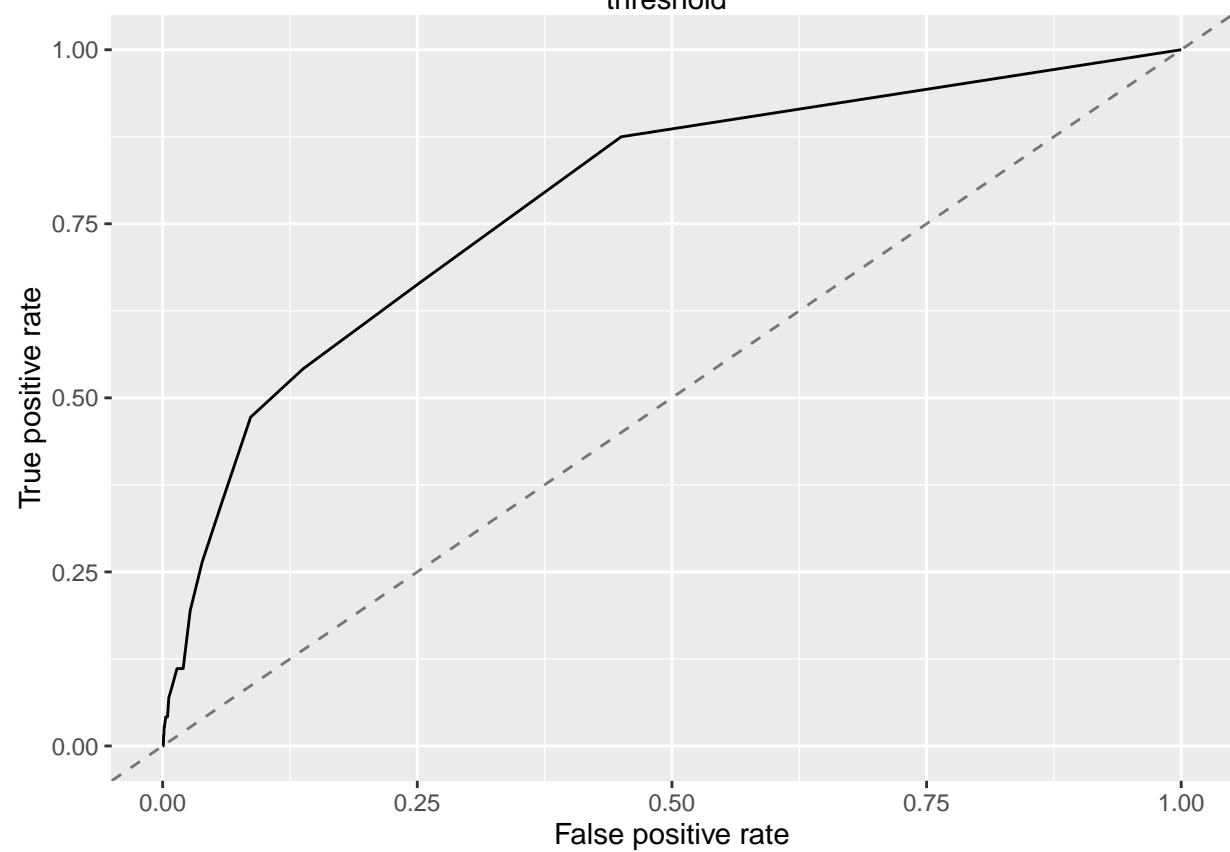
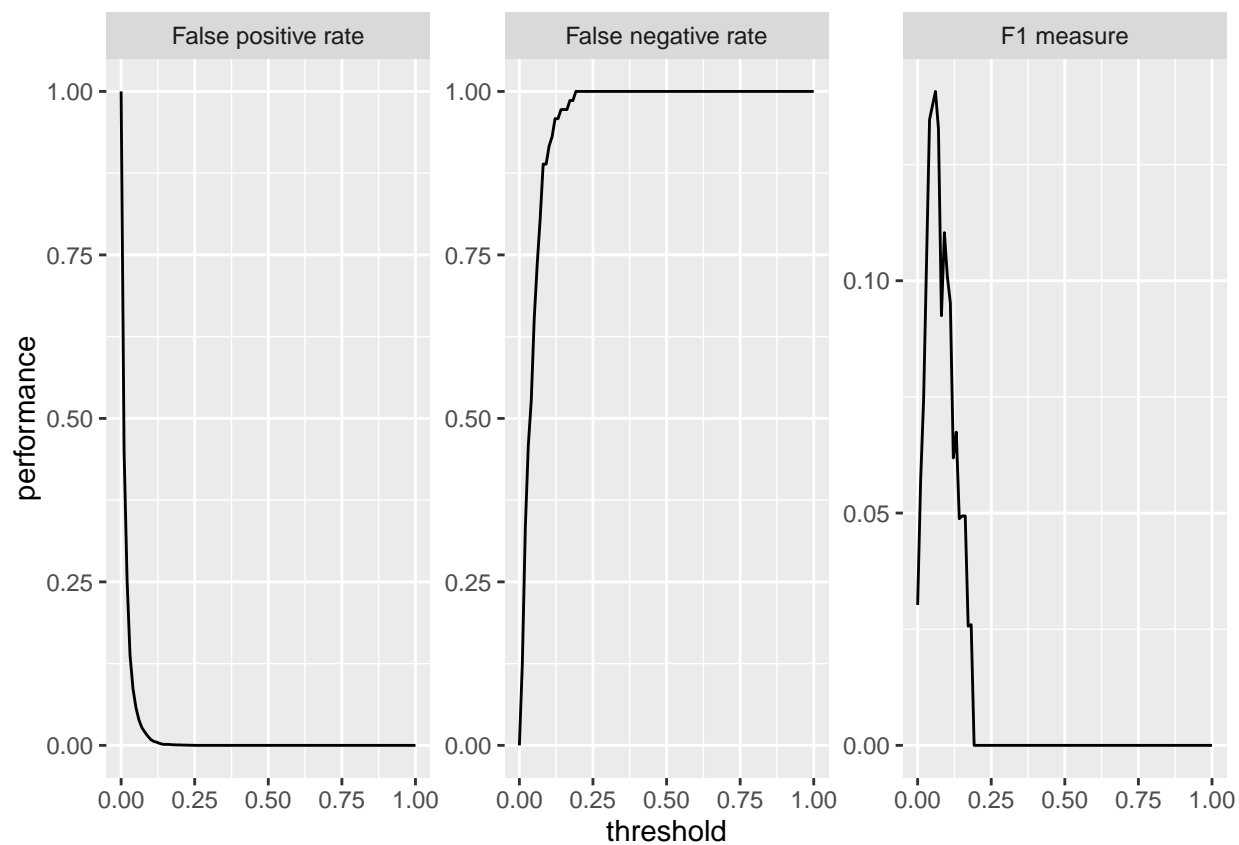
Readmission Type 3 results:

```
$train.metrics
      acc      auc      ppv      brier
0.97755522 0.78373454 0.00000000 0.02131409
```

```
$test.metrics
      acc      auc      ppv      brier
0.90920716 0.79600631 0.08740360 0.02143638
```

```
$train.matrix
      predicted
true    0 1 -SUM-
0      13763 1      1
1       315 0     315
-SUM-   315 1     316
```

```
$test.matrix
      predicted
true    0 1 -SUM-
0      4232 355    355
1       71 34     71
-SUM-   71 355    426
```



- Glmnet (with 10 folds of cross validation and grid tuning the alpha parameter)

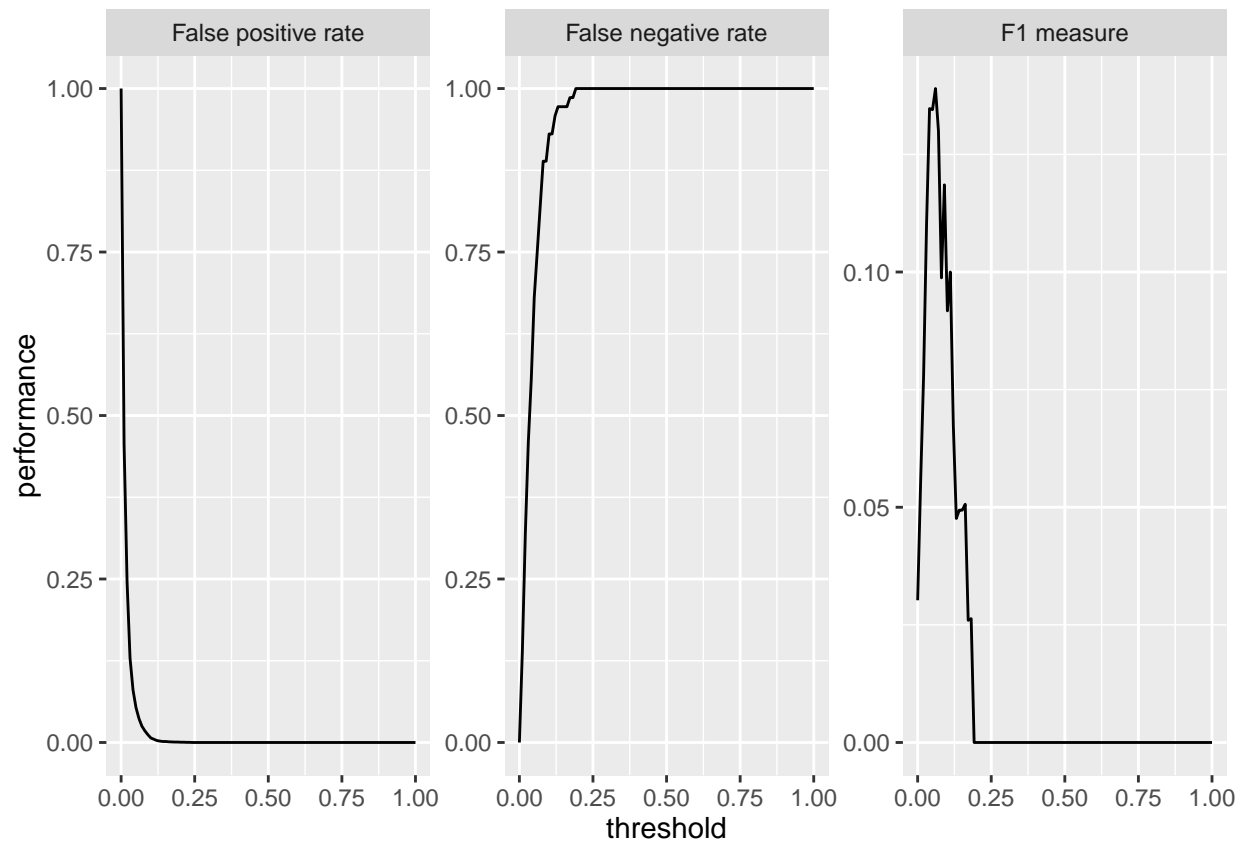
Readmission Type 1 results:

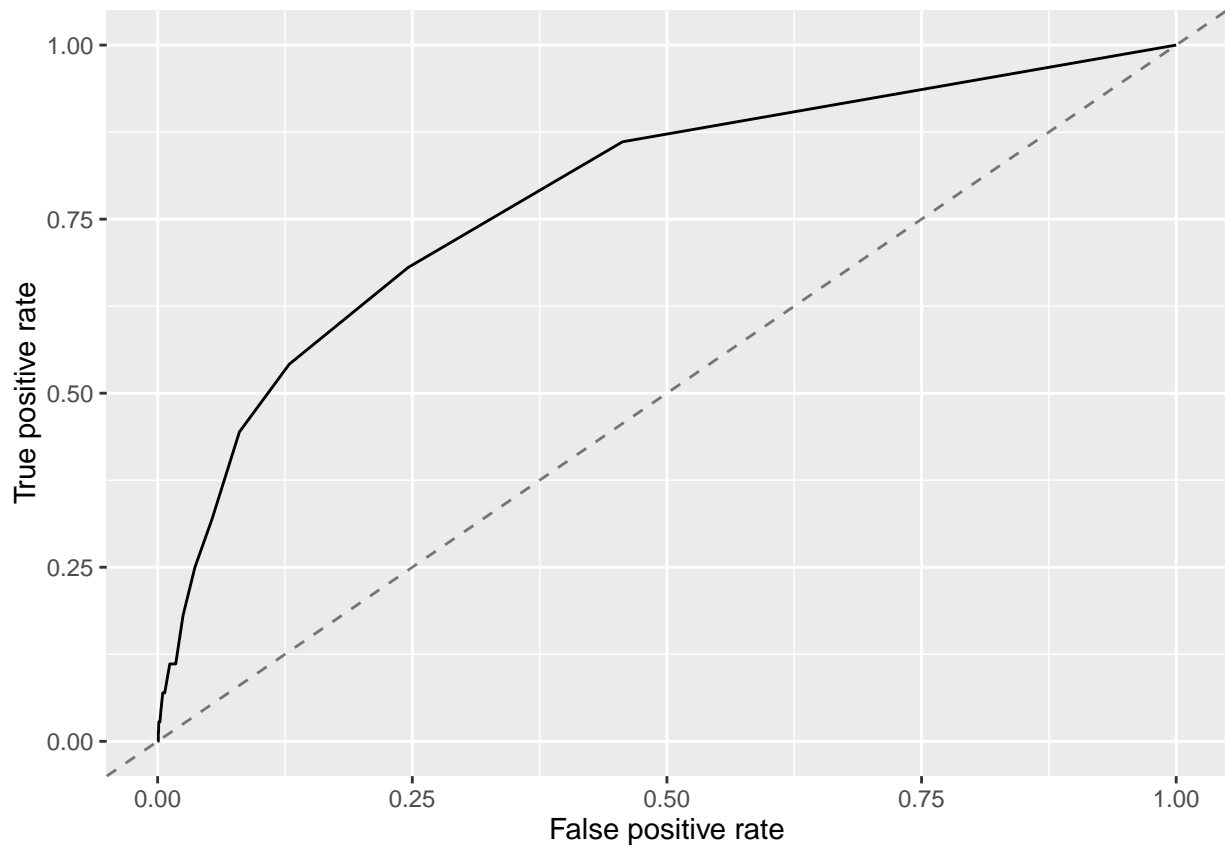
```
$train.metrics
      acc      auc      ppv      brier
0.98444492 0.79560823      NaN 0.01489673
```

```
$test.metrics
      acc      auc      ppv      brier
0.96291560 0.80520984 0.10156250 0.01466212
```

```
$train.matrix
      predicted
true    0 1 -SUM-
0      13860 0      0
1        219 0     219
-SUM-    219 0     219
```

```
$test.matrix
      predicted
true    0 1 -SUM-
0      4505 115    115
1        59 13     59
-SUM-    59 115    174
```





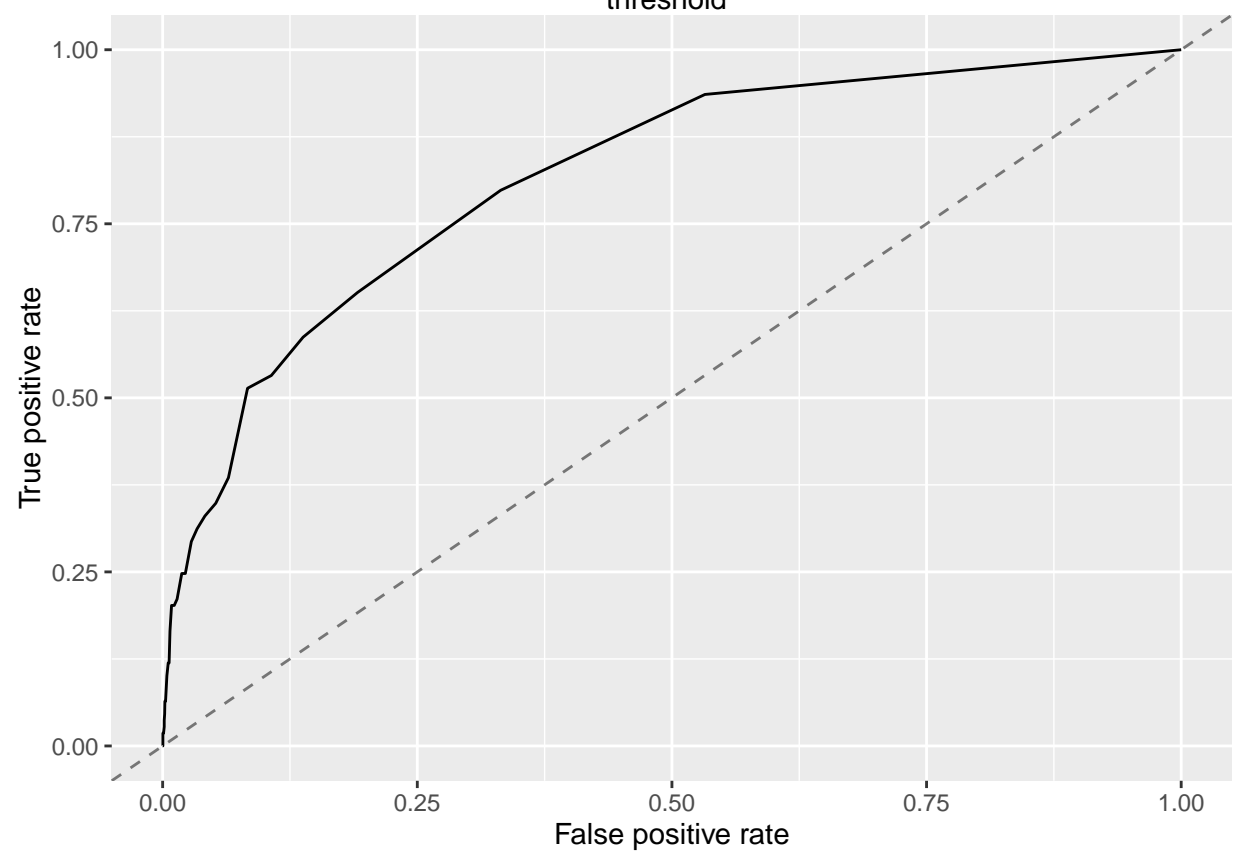
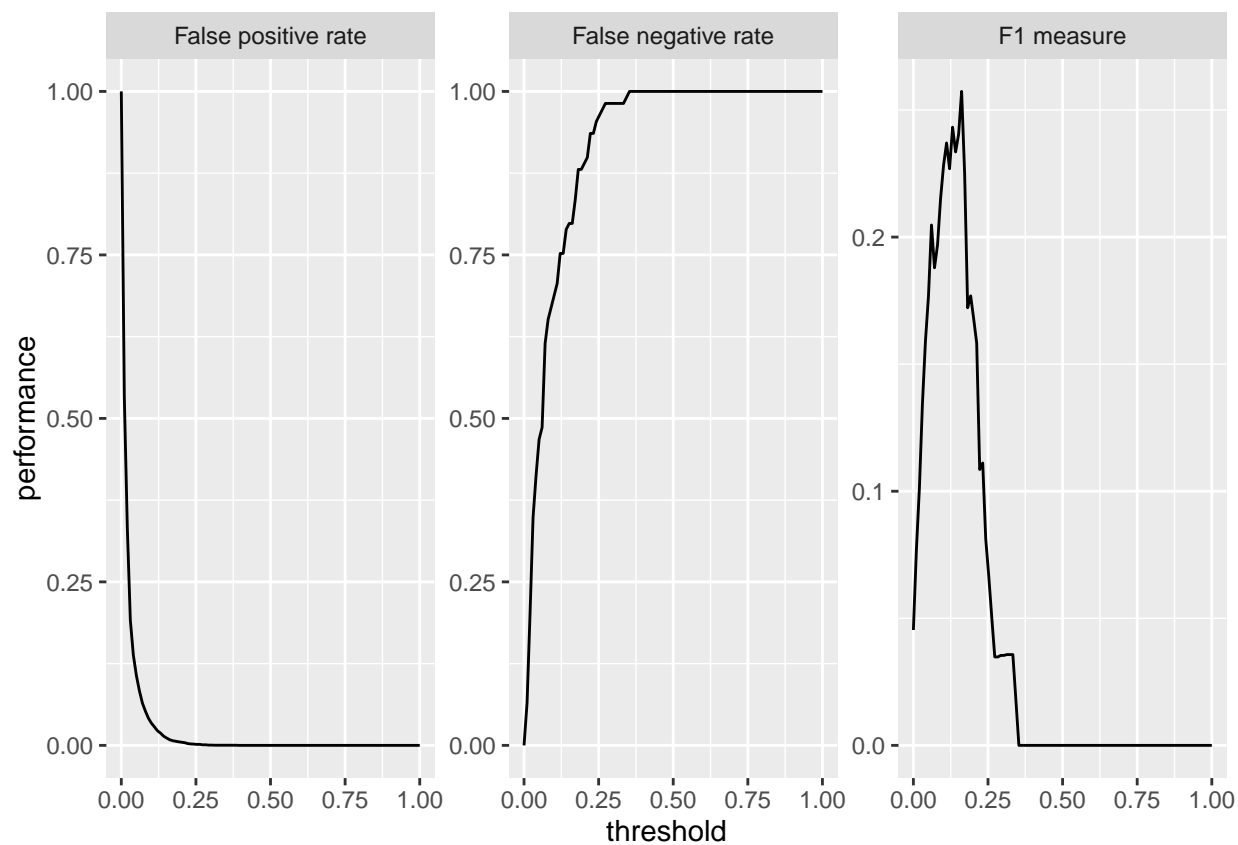
Readmission Type 2 results:

```
$train.metrics
      acc      auc      ppv      brier
0.97677392 0.82371616      NaN 0.02148113
```

```
$test.metrics
      acc      auc      ppv      brier
0.94394714 0.82219591 0.15929204 0.02109058
```

```
$train.matrix
      predicted
true    0 1 -SUM-
0      13752 0    0
1       327 0   327
-SUM-   327 0   327
```

```
$test.matrix
      predicted
true    0 1 -SUM-
0      4393 190   190
1       73 36    73
-SUM-   73 190   263
```

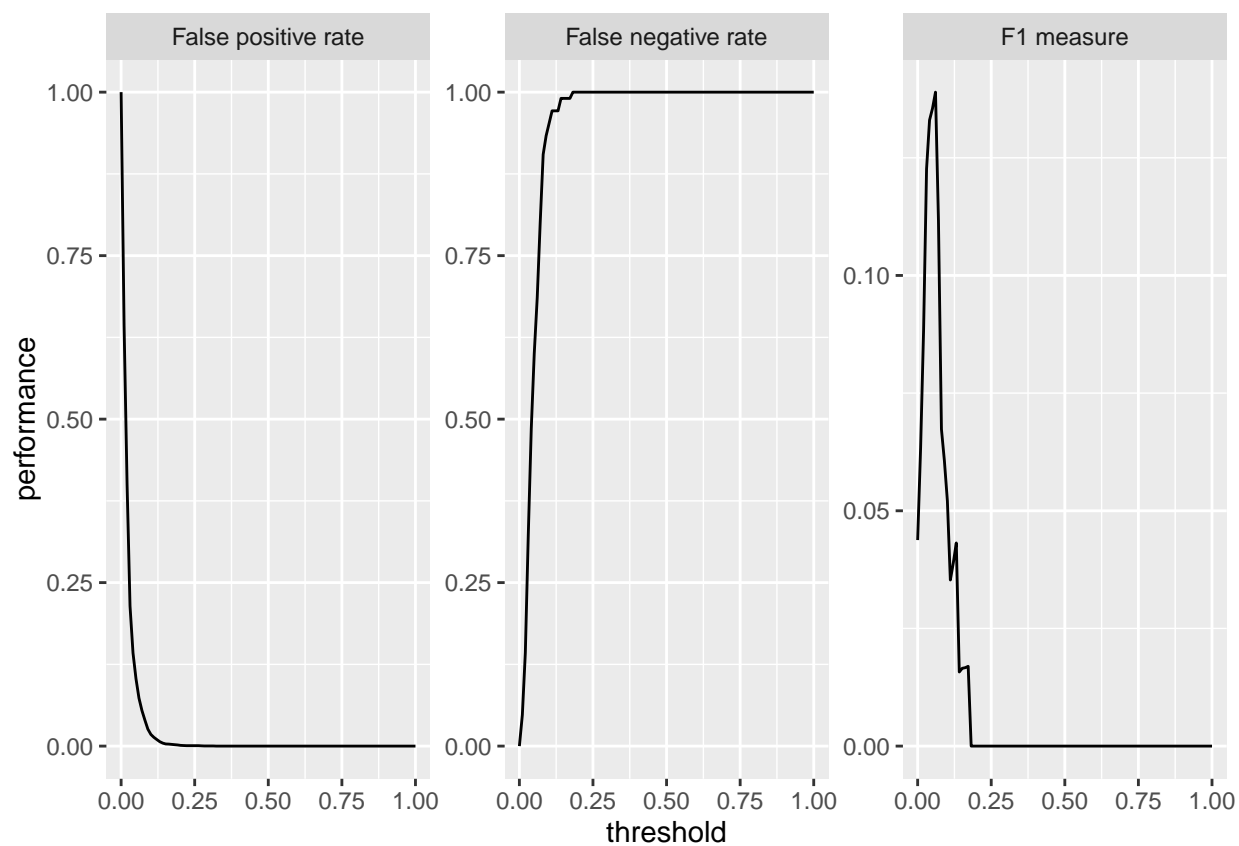
Readmission Type 3 results:

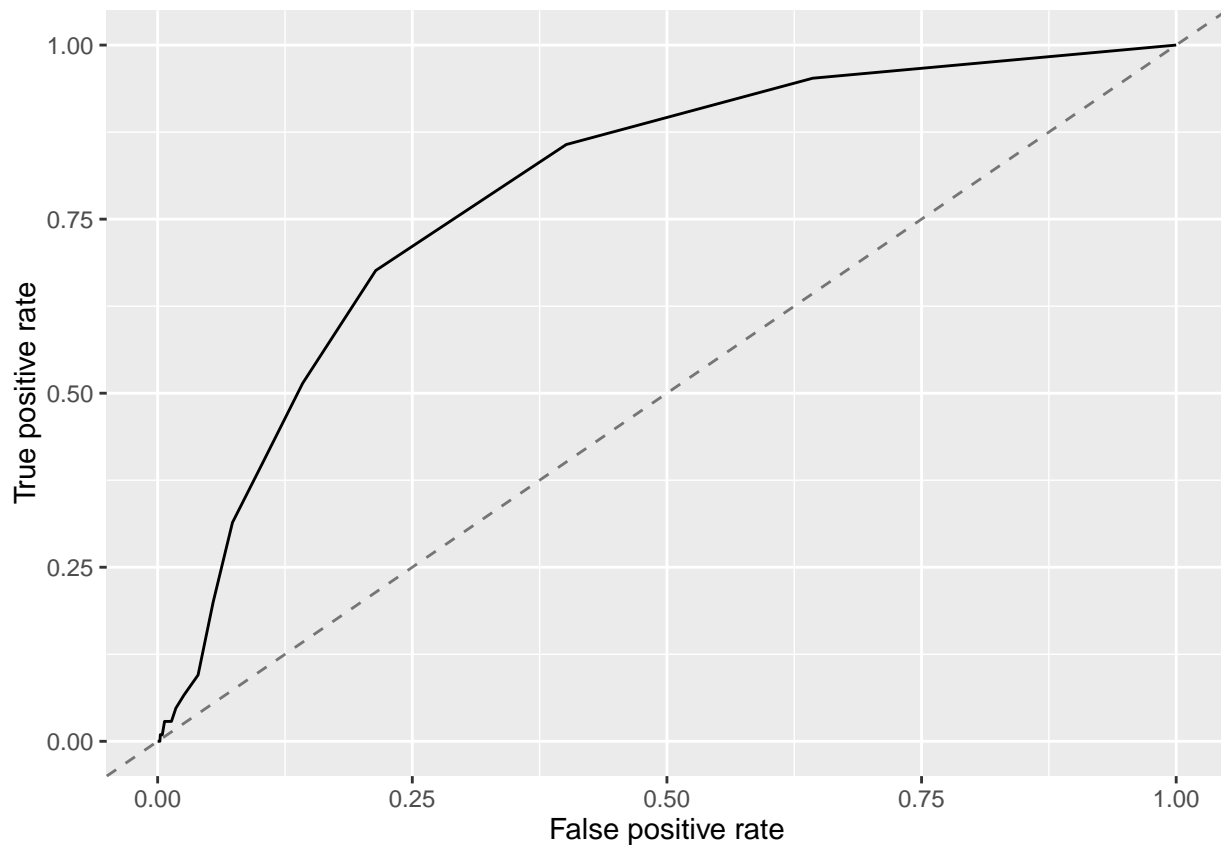
```
$train.metrics
      acc      auc      ppv      brier
0.97755522 0.78381400 0.00000000 0.02131442
```

```
$test.metrics
      acc      auc      ppv      brier
0.91283035 0.79830889 0.08918919 0.02141667
```

```
$train.matrix
      predicted
true    0 1 -SUM-
0      13763 1      1
1       315 0     315
-SUM-   315 1     316
```

```
$test.matrix
      predicted
true    0 1 -SUM-
0      4250 337    337
1       72 33     72
-SUM-   72 337   409
```





For the random forest we introduced different cost matrices to try to optimize our results. Of course in our case of study the false negatives (FN) are more important than the false positives (FP) so that is why the cost matrix penalizes the FN more than the FP. This also gave us a way to compare the results with the previous models.

- Weka results of random forest:

Readmission Type 1 results:

Readmissio N1, cost matrix 20-1, random forest, split 75%

=== Evaluation on test split ===
 === Summary ===

Correctly Classified Instances	4642	98.285 %
Incorrectly Classified Instances	81	1.715 %
Kappa statistic	0.0445	
Mean absolute error	0.0421	
Root mean squared error	0.1307	
Relative absolute error	135.4414 %	
Root relative squared error	102.5708 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.999	0.974	0.984	0.999	0.991	0.76	N
	0.026	0.001	0.286	0.026	0.047	0.76	Y
Weighted Avg.	0.983	0.958	0.972	0.983	0.976	0.76	

=== Confusion Matrix ===

```

a    b  <-- classified as
4640  5 |    a = N
   76  2 |    b = Y

```

Readmissio N1, cost matrix 100-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	3802	80.4997 %
Incorrectly Classified Instances	921	19.5003 %
Kappa statistic	0.0622	
Mean absolute error	0.2459	
Root mean squared error	0.3576	
Relative absolute error	790.2177 %	
Root relative squared error	280.5712 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.809	0.41	0.992	0.809	0.891	0.807	N
	0.59	0.191	0.049	0.59	0.091	0.807	Y
Weighted Avg.	0.805	0.407	0.976	0.805	0.878	0.807	

=== Confusion Matrix ===

```

a    b  <-- classified as
3756 889 |    a = N
   32  46 |    b = Y

```

Readmissio N1, cost matrix 60-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	4328	91.6367 %
Incorrectly Classified Instances	395	8.3633 %
Kappa statistic	0.1074	

```

Mean absolute error          0.1482
Root mean squared error      0.2486
Relative absolute error      476.3702 %
Root relative squared error   195.0567 %
Total Number of Instances    4723

```

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.925	0.615	0.989	0.925	0.956	0.813	N
	0.385	0.075	0.08	0.385	0.132	0.813	Y
Weighted Avg.	0.916	0.606	0.974	0.916	0.942	0.813	

=== Confusion Matrix ===

```

  a    b  <-- classified as
4298 347 |    a = N
  48   30 |    b = Y

```

Readmissio N1, cost matrix 50-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

```

Correctly Classified Instances    4435          93.9022 %
Incorrectly Classified Instances   288          6.0978 %
Kappa statistic                   0.1257
Mean absolute error               0.1211
Root mean squared error           0.2176
Relative absolute error           389.1505 %
Root relative squared error       170.7035 %
Total Number of Instances        4723

```

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.949	0.679	0.988	0.949	0.968	0.794	N
	0.321	0.051	0.096	0.321	0.148	0.792	Y
Weighted Avg.	0.939	0.669	0.973	0.939	0.955	0.794	

=== Confusion Matrix ===

```

  a    b  <-- classified as
4410 235 |    a = N
  53   25 |    b = Y

```

Readmission Type 2 results:

Readmissio N2, cost matrix 20-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	4561	96.57	%
Incorrectly Classified Instances	162	3.43	%
Kappa statistic	0.1727		
Mean absolute error	0.0777		
Root mean squared error	0.1749		
Relative absolute error	171.4773	%	
Root relative squared error	115.973	%	
Total Number of Instances	4723		

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.985	0.827	0.98	0.985	0.982	0.792	N
	0.173	0.015	0.211	0.173	0.19	0.792	Y
Weighted Avg.	0.966	0.808	0.962	0.966	0.964	0.792	

=== Confusion Matrix ===

a	b	<-- classified as
4542	71	a = N
91	19	b = Y

Readmissio N2, cost matrix 100-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	3179	67.3089	%
Incorrectly Classified Instances	1544	32.6911	%
Kappa statistic	0.0603		
Mean absolute error	0.3429		
Root mean squared error	0.4671		
Relative absolute error	757.0491	%	
Root relative squared error	309.6936	%	
Total Number of Instances	4723		

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.67	0.209	0.993	0.67	0.8	0.818	N
	0.791	0.33	0.054	0.791	0.101	0.818	Y
Weighted Avg.	0.673	0.212	0.971	0.673	0.784	0.818	

=== Confusion Matrix ===

```
      a      b  <-- classified as
3092 1521 |      a = N
    23   87 |      b = Y
```

Readmissio N2, cost matrix 60-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	3823	80.9443 %
Incorrectly Classified Instances	900	19.0557 %
Kappa statistic	0.0865	
Mean absolute error	0.237	
Root mean squared error	0.3588	
Relative absolute error	523.2731 %	
Root relative squared error	237.8628 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.815	0.418	0.988	0.815	0.893	0.814	N
	0.582	0.185	0.07	0.582	0.125	0.814	Y
Weighted Avg.	0.809	0.413	0.967	0.809	0.875	0.814	

=== Confusion Matrix ===

```
      a      b  <-- classified as
3759  854 |      a = N
    46   64 |      b = Y
```

Readmissio N2, cost matrix 50-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	4028	85.2848 %
Incorrectly Classified Instances	695	14.7152 %
Kappa statistic	0.1049	
Mean absolute error	0.203	
Root mean squared error	0.3208	
Relative absolute error	448.1796 %	
Root relative squared error	212.6706 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.861	0.482	0.987	0.861	0.92	0.815	N
	0.518	0.139	0.082	0.518	0.141	0.815	Y
Weighted Avg.	0.853	0.474	0.966	0.853	0.901	0.815	

=== Confusion Matrix ===

```

  a    b  <-- classified as
3971 642 |    a = N
  53   57 |    b = Y

```

Readmission Type 3 results:

Readmissio N3, cost matrix 20-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	4606	97.5228 %
Incorrectly Classified Instances	117	2.4772 %
Kappa statistic	0.0133	
Mean absolute error	0.0658	
Root mean squared error	0.1596	
Relative absolute error	148.9269 %	
Root relative squared error	105.811 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.998	0.991	0.977	0.998	0.987	0.762	N
	0.009	0.002	0.111	0.009	0.017	0.762	Y
Weighted Avg.	0.975	0.968	0.957	0.975	0.965	0.762	

=== Confusion Matrix ===

```

  a    b  <-- classified as
4605   8 |    a = N
 109   1 |    b = Y

```

Readmissio N3, cost matrix 100-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	2990	63.3072 %
Incorrectly Classified Instances	1733	36.6928 %


```

Kappa statistic          0.0495
Mean absolute error      0.3669
Root mean squared error  0.4878
Relative absolute error   829.8135 %
Root relative squared error 323.4365 %
Total Number of Instances 4723

```

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.629	0.209	0.992	0.629	0.77	0.782	N
	0.791	0.371	0.048	0.791	0.091	0.782	Y
Weighted Avg.	0.633	0.213	0.97	0.633	0.754	0.782	

=== Confusion Matrix ===

```

      a      b  <-- classified as
2903 1710 |      a = N
      23     87 |      b = Y

```

Readmissio N3, cost matrix 60-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

```

Correctly Classified Instances 3744          79.2716 %
Incorrectly Classified Instances 979          20.7284 %
Kappa statistic              0.0702
Mean absolute error          0.2512
Root mean squared error      0.3645
Relative absolute error      568.0387 %
Root relative squared error   241.6528 %
Total Number of Instances    4723

```

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.799	0.455	0.987	0.799	0.883	0.774	N
	0.545	0.201	0.061	0.545	0.109	0.774	Y
Weighted Avg.	0.793	0.449	0.965	0.793	0.865	0.774	

=== Confusion Matrix ===

```

      a      b  <-- classified as
3684  929 |      a = N
      50     60 |      b = Y

```

 Readmissio N3, cost matrix 50-1, random forest, split 75%

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	4013	84.9672 %
Incorrectly Classified Instances	710	15.0328 %
Kappa statistic	0.0799	
Mean absolute error	0.2107	
Root mean squared error	0.3179	
Relative absolute error	476.4429 %	
Root relative squared error	210.7556 %	
Total Number of Instances	4723	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	0.86	0.573	0.984	0.86	0.918	0.775	N
	0.427	0.14	0.068	0.427	0.117	0.776	Y
Weighted Avg.	0.85	0.563	0.963	0.85	0.899	0.775	

=== Confusion Matrix ===

a	b	<-- classified as
3966	647	a = N
63	47	b = Y

After considering our analysis, we have established that in order to achieve the perfect balance between the FP and the FN and optimize the results for each model, careful consideration should be given to the real cost for the hospital to have a FN than a FP. In the case of the logistic regression and the glmnet we should therefore design a measure that could capture accurately the cost we are trying to minimize. In the case of the random forest with cost matrix we should also optimize the cost matrix subject to the real cost of allowing more FP to reduce the FN. That is an important, complicated and precise challenge that falls out of the scope of this thesis but that we will clearly encourage anyone working on the subject to spend time on it.

To increase the reliability of our analysis an outlier detection analysis was performed in order to remove the outliers from our data. This analysis did not lead to an increase in the quality of our results. This could be because those “outliers” are extreme values that appear precisely in those “readmitted” observations and most likely are part of the reason those patients were readmitted. Therefore all extreme values were included in our final analysis.

Conclusion

A readmission predictive model can be successfully implemented in a academic hospital using existing data.

However it is not something easy to predict, developers of those predictive models need to consider more than the statistics when developing models. An implementable model balances clinical priorities, statistical requirements, availability of data and technical requirements.

The following are the conclusions we can get from the models predicting each response variable:

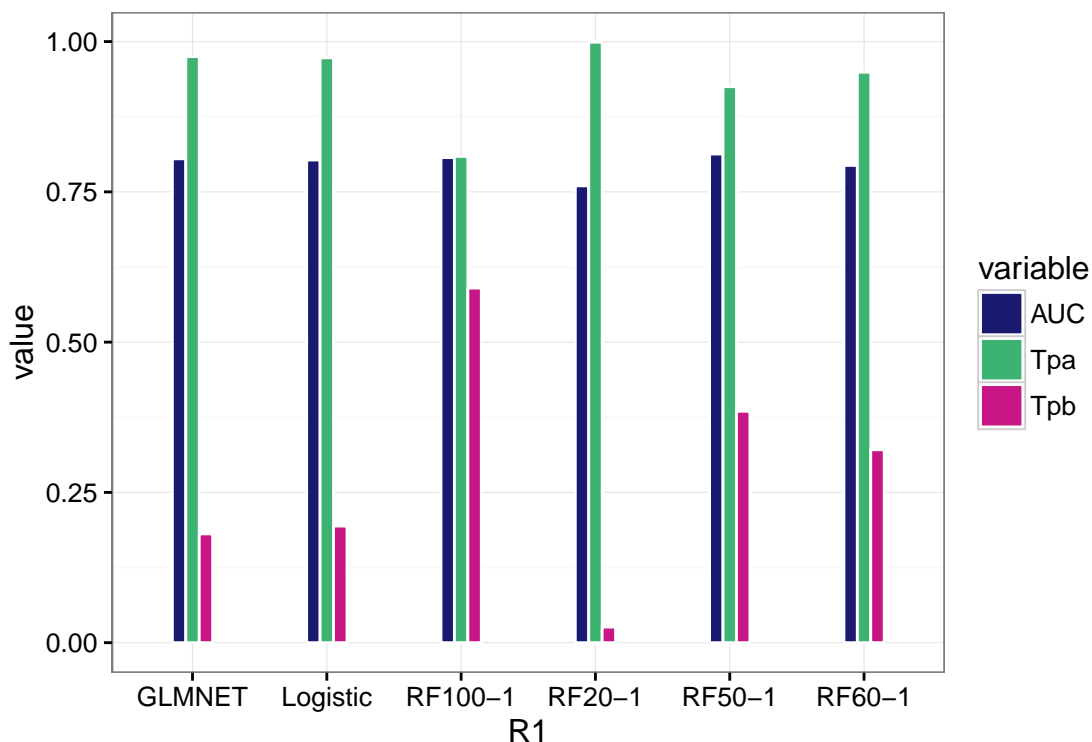
- Length of stay

The best model for predicting the *LoS* in our case study is the GBM for the *Less than 2 days* and *3 to 7 days* categories and the Random Forest for the *More than 7 days* category. If we had to pick just one model we would choose GBM because the difference between the GBM and the RF in the first two categories is significantly larger than the difference between the RF and the GBM in the third category.

- Readmissions

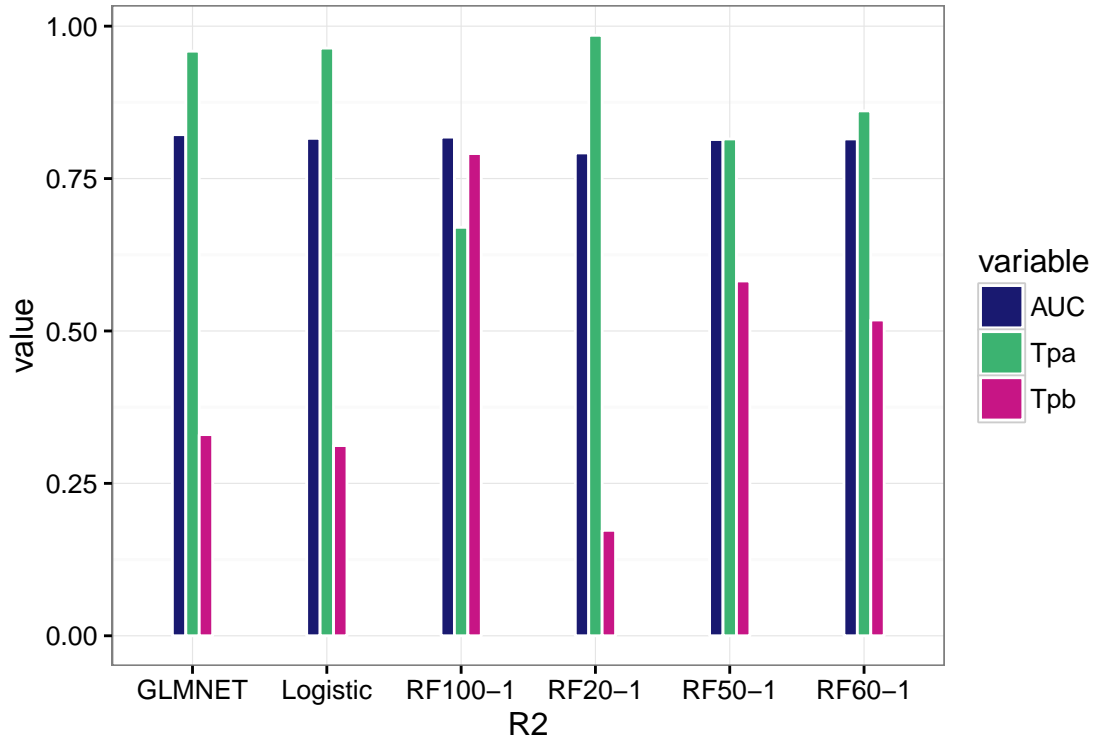
It is not that apparent through model comparison which one is the best model in the readmissions. We plotted all the models and 3 different measures to try to pick one. The three different measures are the AUC (Area under the curve), the Tpa (true positive values for class a, the not readmitted patients) and the Tpb (true positive values for class b, the readmitted patients). The three of this measures should be maximized but sometimes from one model to another there is a tradeoff where one measure increases and another decreases. In this case study we picked the higher AUC in order to decide on those cases.

- Readmission type 1



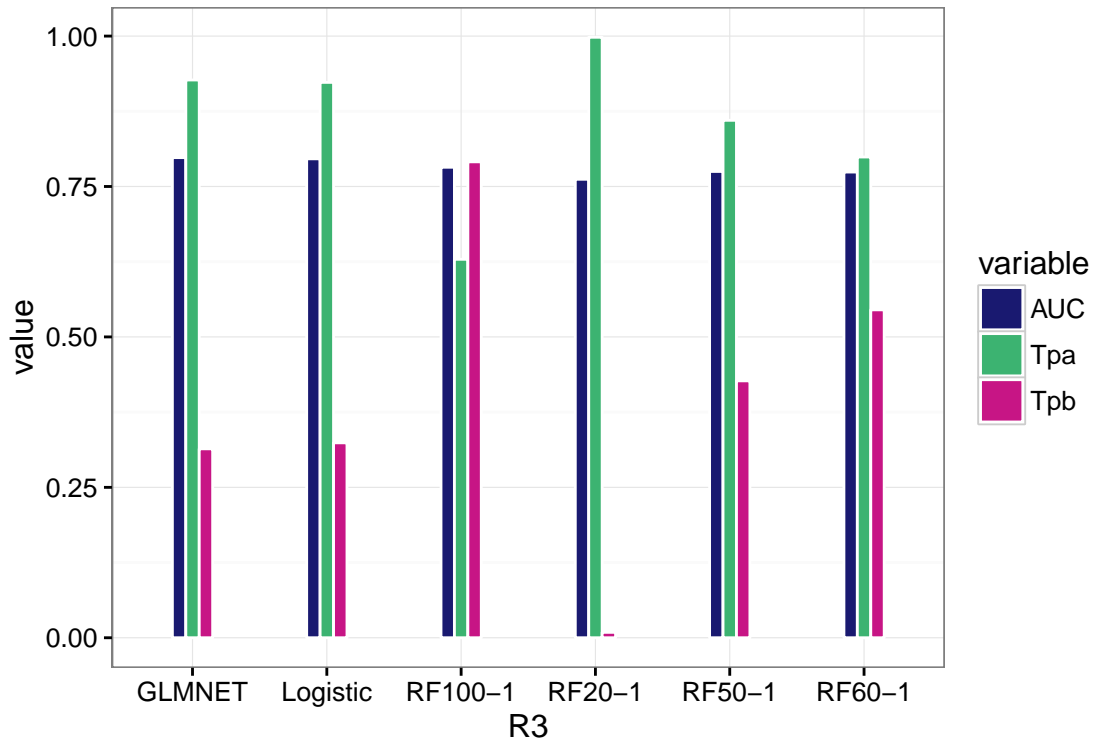
In this case the best model is the RF100-1.

- *Readmission type 2*



In this case the best model is the GLMNET.

- *Readmission type 3*



In this case the best model is the GLMNET.

At a first look the RF100-1 might appear to be the best model in predicting $R2$ and $R3$ because it predicts better the class b, but when checking the AUC we pick GLMNET. It is true that the AUC of the RF100-1 is close to GLMNET in both cases. It is also true that we said above in the report that we care more about predicting the class b better even at expenses of predicting the class a worse. But as we stated at the very end of the *Predictions* section, to correctly determine which is the exact tradeoff between the increase in predictions of class b and the decrease in predictions of class a that we can assume, a better cost measure should be designed. For that we would need more information on the economic value that the hospital gives to each wrongly predicted observation.

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- ³Kiyana Zolfaghar, Naren Meadem, Ankur Teredesai, Senjuti Basu Roy, Si-Chi Chin, Brian Muckian, 2013. “Big Data Solutions for Predicting Risk-of-Readmission for Congestive Heart Failure Patients”. IEEE International Conference on Big Data. <https://cwds.uw.edu/sites/default/files/publications/Big%20Data%20Solutions%20for%20Predicting%20Risk-of-Readmission%20for%20Congestive%20Heart%20Failure.pdf>
- ⁴Phillips Healthcare Transformation Services. “Reducing avoidable readmissions using predictive analytics” URL: http://www.philips.fi/b-dam/b2bhc/us/hts/population-health/Reducing_avoidable_readmissions_using_predictive_analytics.pdf (visited in 15/04/2016)
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