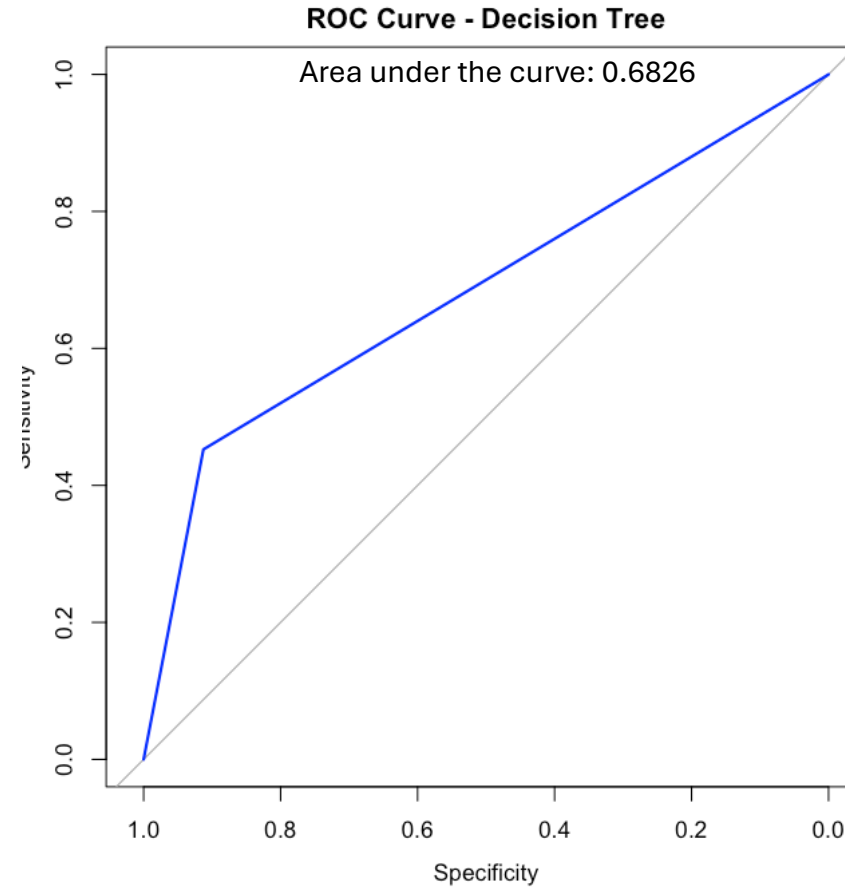


DECISION TREE 1

```
> print(importance)
```

	Overall
ap_hi	1521.4864
ap_lo	1036.4765
bp_category	1035.8156
bp_category_encoded	1035.8156
cholesterol	311.8565
gender	0.0000
height	0.0000
weight	0.0000
gluc	0.0000
smoke	0.0000
alco	0.0000
active	0.0000
age_years	0.0000
bmi	0.0000



Confusion Matrix and Statistics

	Reference	
Prediction	no	yes
no	10989	2168
yes	1051	1792

Accuracy : 0.7988
95% CI : (0.7925, 0.805)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.00000000000000022

Kappa : 0.4034

McNemar's Test P-Value : < 0.00000000000000022

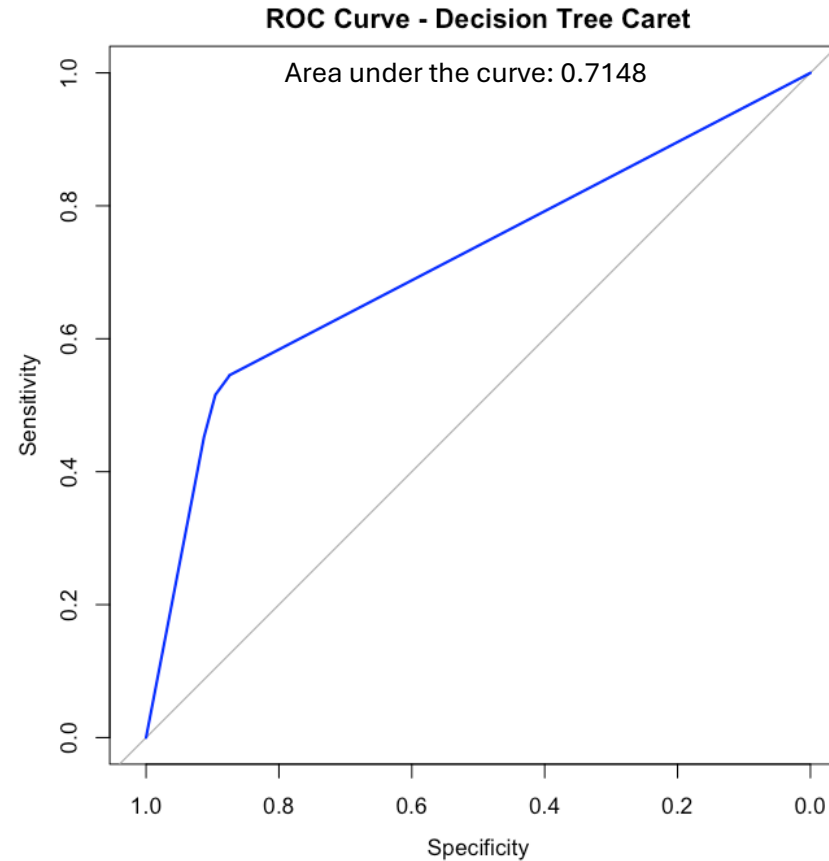
Sensitivity : 0.4525
Specificity : 0.9127
Pos Pred Value : 0.6303
Neg Pred Value : 0.8352
Prevalence : 0.2475
Detection Rate : 0.1120
Detection Prevalence : 0.1777
Balanced Accuracy : 0.6826

'Positive' Class : yes

DECISION TREE TUNED FOR SENSITIVITY

```
> print(importance3)
```

	Overall
age_years	180.295709
ap_hi	1726.823074
ap_lo	1136.611726
bmi	25.055042
bp_category	1098.196674
bp_category_encoded	1038.108592
cholesterol	522.000594
gluc	25.782670
height	4.690020
weight	4.170049
gender	0.000000
smoke	0.000000
alco	0.000000
active	0.000000



Reference		
Prediction	no	yes
no	10780	1913
yes	1260	2047

Accuracy : 0.8017
95% CI : (0.7954, 0.8078)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.0000000000000022

Kappa : 0.4364

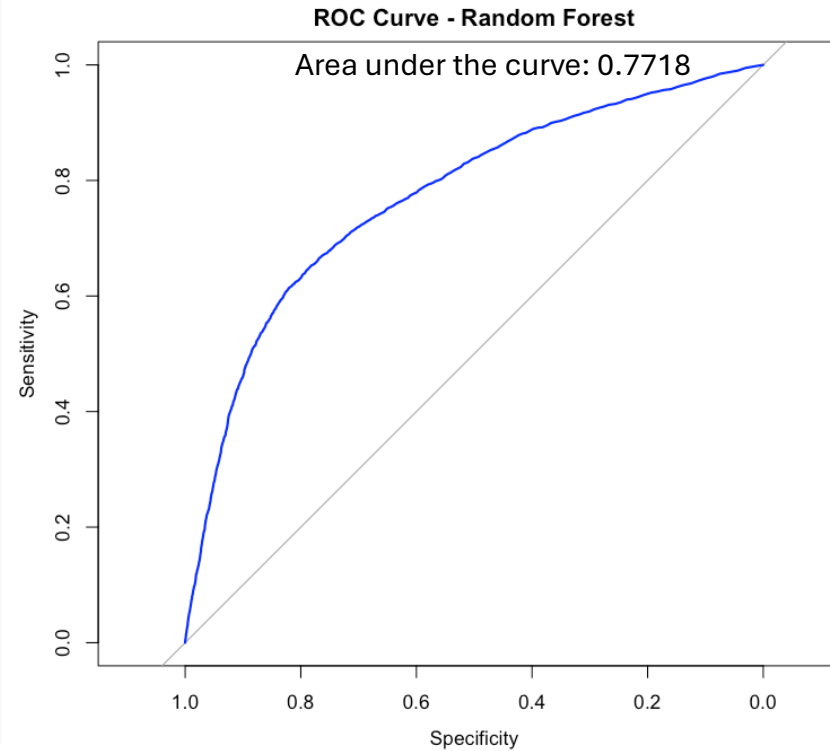
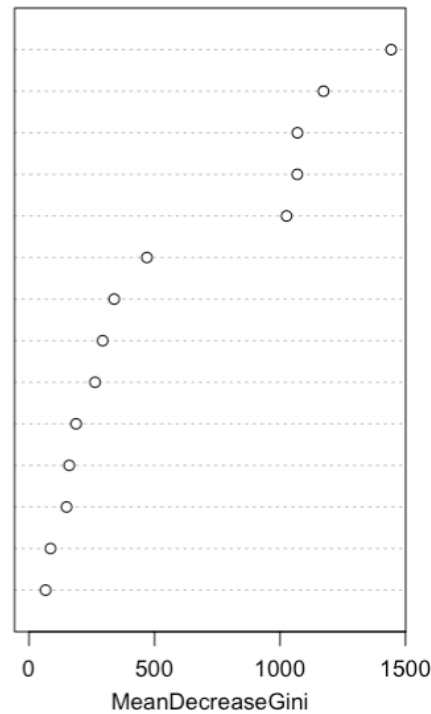
McNemar's Test P-Value : < 0.0000000000000022

Sensitivity : 0.5169
Specificity : 0.8953
Pos Pred Value : 0.6190
Neg Pred Value : 0.8493
Prevalence : 0.2475
Detection Rate : 0.1279
Detection Prevalence : 0.2067
Balanced Accuracy : 0.7061

'Positive' Class : yes

RANDOM FOREST (ntree = 500; No. of variables tried at each split: 4)

bmi
ap_hi
age_years
weight
height
ap_lo
cholesterol
bp_category_encoded
bp_category
gluc
gender
active
smoke
alco



Confusion Matrix and Statistics

Reference
Prediction no yes
no 10914 2191
yes 1126 1769

Accuracy : 0.7927
95% CI : (0.7863, 0.7989)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.00000000000000022

Kappa : 0.3882

Mcnemar's Test P-Value : < 0.00000000000000022

Sensitivity : 0.4467
Specificity : 0.9065
Pos Pred Value : 0.6111
Neg Pred Value : 0.8328
Prevalence : 0.2475
Detection Rate : 0.1106
Detection Prevalence : 0.1809
Balanced Accuracy : 0.6766

'Positive' Class : yes

NAIVE BAYES (normale e kernel density estimation)

```
> print(conf_matrix_norm)
Confusion Matrix and Statistics
```

	Prediction	1	0
Reference	1	1928	1504
	0	2032	10536

Accuracy : 0.779
95% CI : (0.7725, 0.7854)
No Information Rate : 0.7525
P-Value [Acc > NIR] : 2.003e-15

Kappa : 0.3789

McNemar's Test P-Value : < 2.2e-16

Sensitivity : 0.4869
Specificity : 0.8751
Pos Pred Value : 0.5618
Neg Pred Value : 0.8383
Prevalence : 0.2475
Detection Rate : 0.1205
Detection Prevalence : 0.2145
Balanced Accuracy : 0.6810

'Positive' Class : 1

```
> print(conf_matrix_kernel)
Confusion Matrix and Statistics
```

	Prediction	1	0
Reference	1	1732	1303
	0	2228	10737

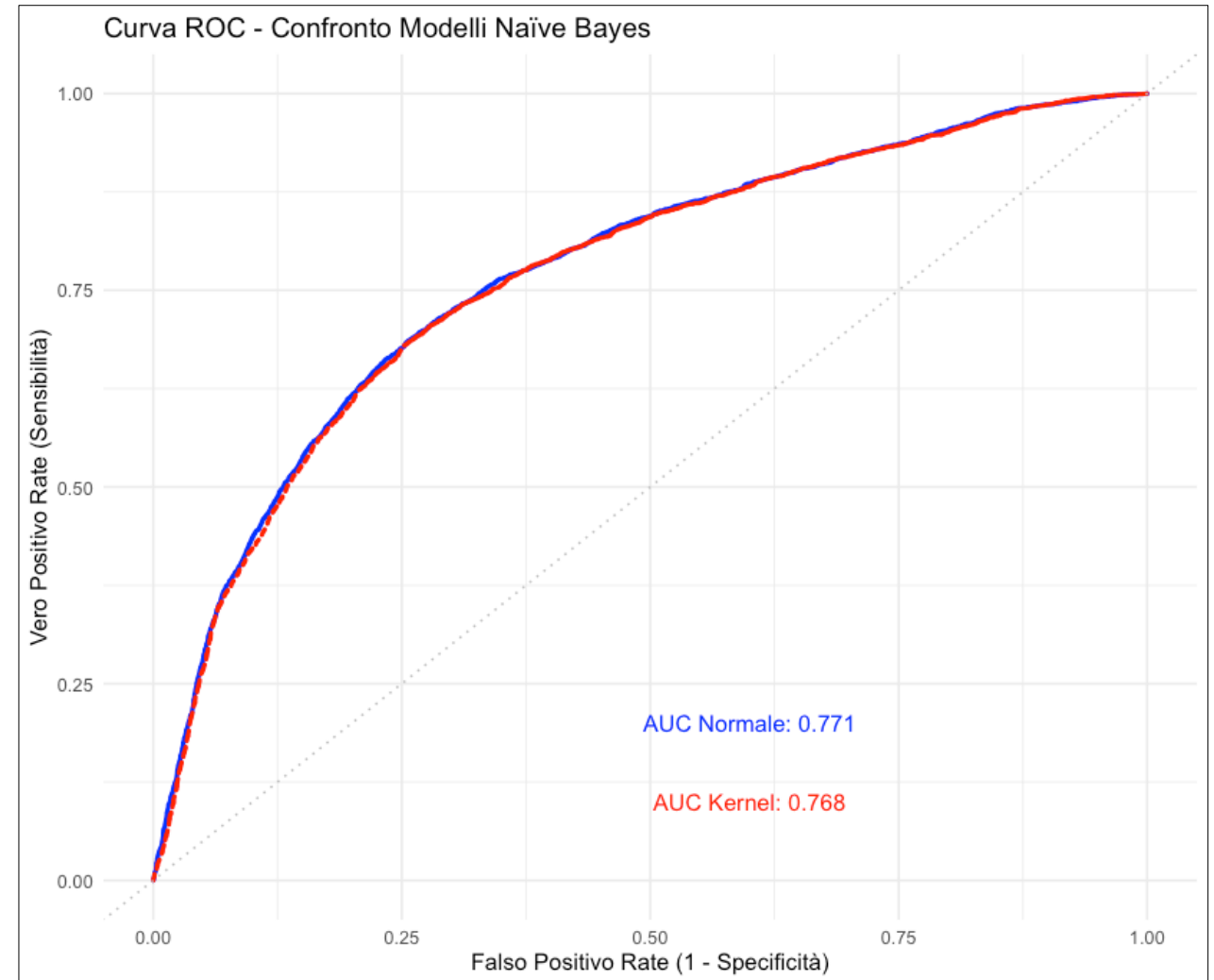
Accuracy : 0.7793
95% CI : (0.7728, 0.7857)
No Information Rate : 0.7525
P-Value [Acc > NIR] : 9.419e-16

Kappa : 0.3571

McNemar's Test P-Value : < 2.2e-16

Sensitivity : 0.4374
Specificity : 0.8918
Pos Pred Value : 0.5707
Neg Pred Value : 0.8282
Prevalence : 0.2475
Detection Rate : 0.1082
Detection Prevalence : 0.1897
Balanced Accuracy : 0.6646

'Positive' Class : 1



QDA

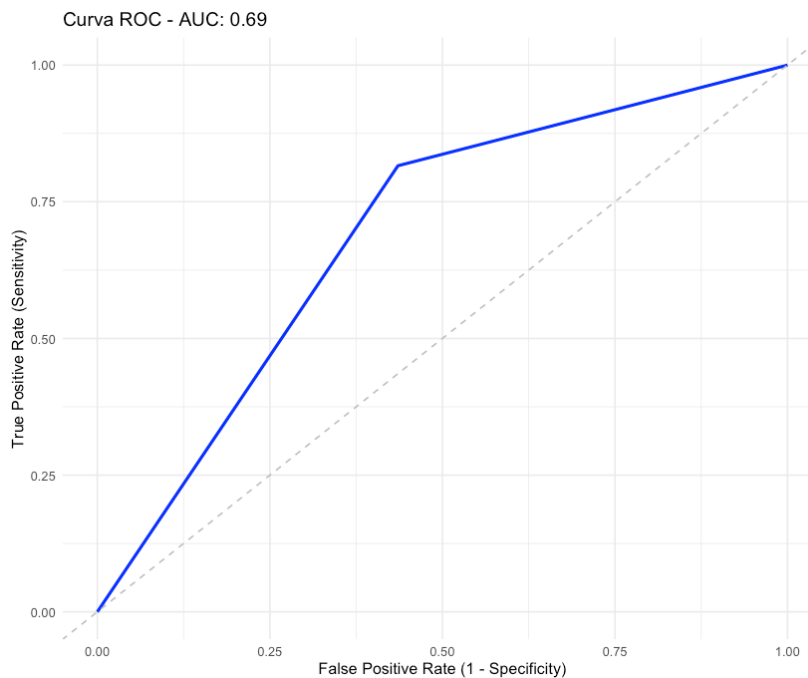
Modello logit usato per selezione variabili
per avere idea dell'importanza delle variabili

Single term deletions

Model:

cardio ~ weight + ap_hi + ap_lo + cholesterol + gluc + smoke +
alco + active + age_years + bp_category

	Df	Deviance	F value	Pr(>F)	
<none>		22278			
weight	1	22317	42.0540	9.052e-11	***
ap_hi	1	23038	818.3020	< 2.2e-16	***
ap_lo	1	22282	4.1071	0.0427141	*
cholesterol	1	22522	262.9126	< 2.2e-16	***
gluc	1	22290	13.3802	0.0002548	***
smoke	1	22287	10.2024	0.0014044	**
alco	1	22290	13.4441	0.0002463	***
active	1	22306	30.6818	3.072e-08	***
age_years	1	22604	350.7295	< 2.2e-16	***
bp_category	3	22343	23.3584	4.323e-15	***



Confusion Matrix and Statistics

	Reference	
Prediction	1	0
1	2235	2217
0	1725	9823

Accuracy : 0.7536
95% CI : (0.7469, 0.7603)
No Information Rate : 0.7525
P-Value [Acc > NIR] : 0.3748

Kappa : 0.365

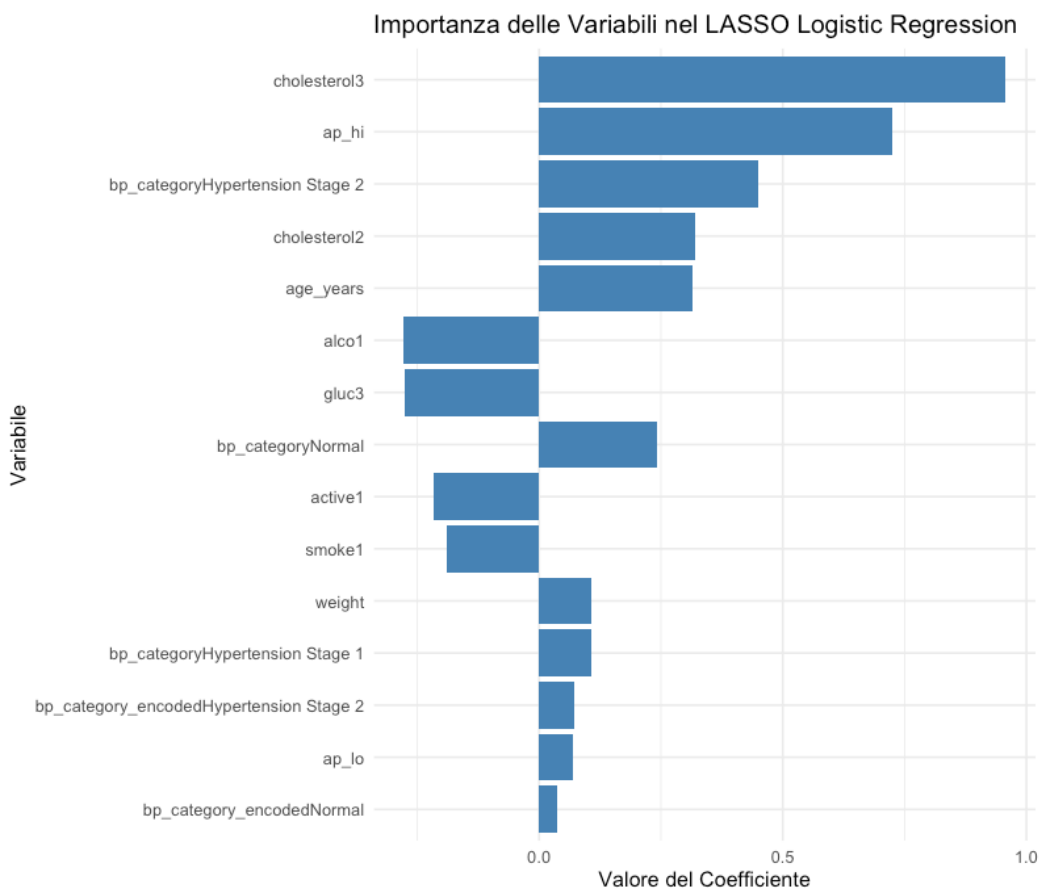
Mcnemar's Test P-Value : 5.27e-15

Sensitivity : 0.5644
Specificity : 0.8159
Pos Pred Value : 0.5020
Neg Pred Value : 0.8506
Prevalence : 0.2475
Detection Rate : 0.1397
Detection Prevalence : 0.2782
Balanced Accuracy : 0.6901

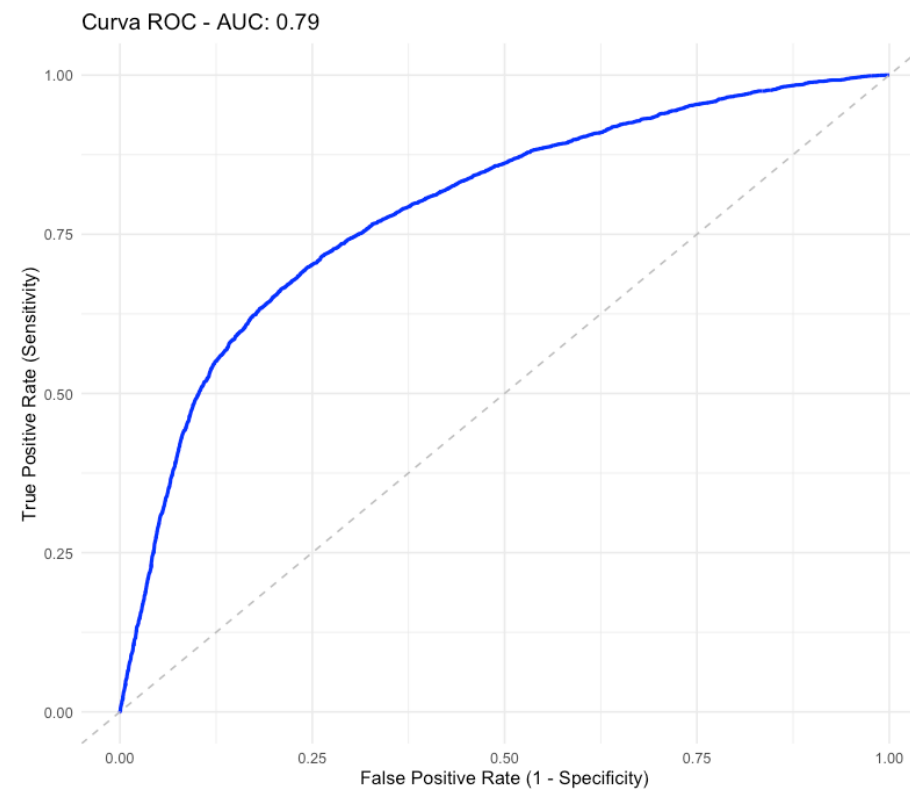
'Positive' Class : 1

LOGISTIC LASSO

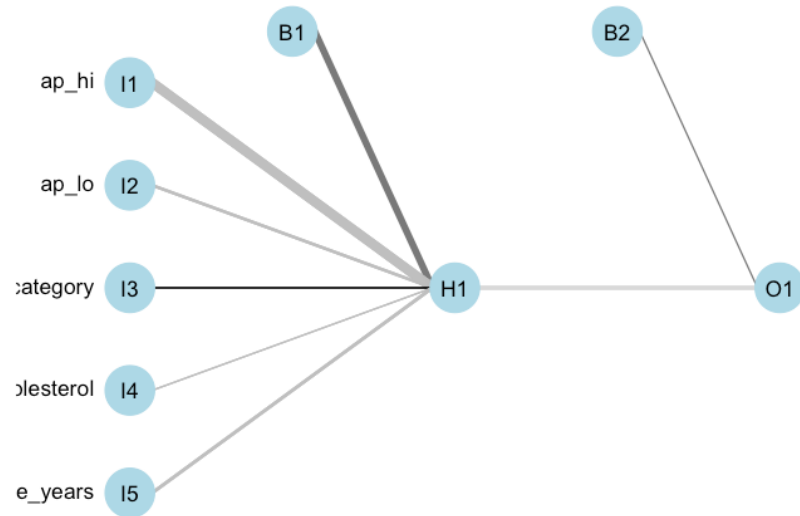
(Miglior valore di lambda: 0.0003789426)



Accuracy: 0.7919
Precision: 0.646
Sensitivity: 0.352
F1-score: 0.4557



PERCEPTRON



Confusion Matrix and Statistics

Prediction \ Reference	Reference	
	0	1
0	10949	2165
1	1091	1795

Accuracy : 0.7965
95% CI : (0.7902, 0.8027)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.00000000000000022

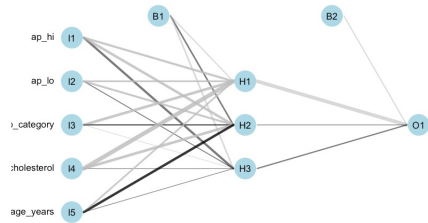
Kappa : 0.399

McNemar's Test P-Value : < 0.00000000000000022

Sensitivity : 0.4533
Specificity : 0.9094
Pos Pred Value : 0.6220
Neg Pred Value : 0.8349
Prevalence : 0.2475
Detection Rate : 0.1122
Detection Prevalence : 0.1804
Balanced Accuracy : 0.6813

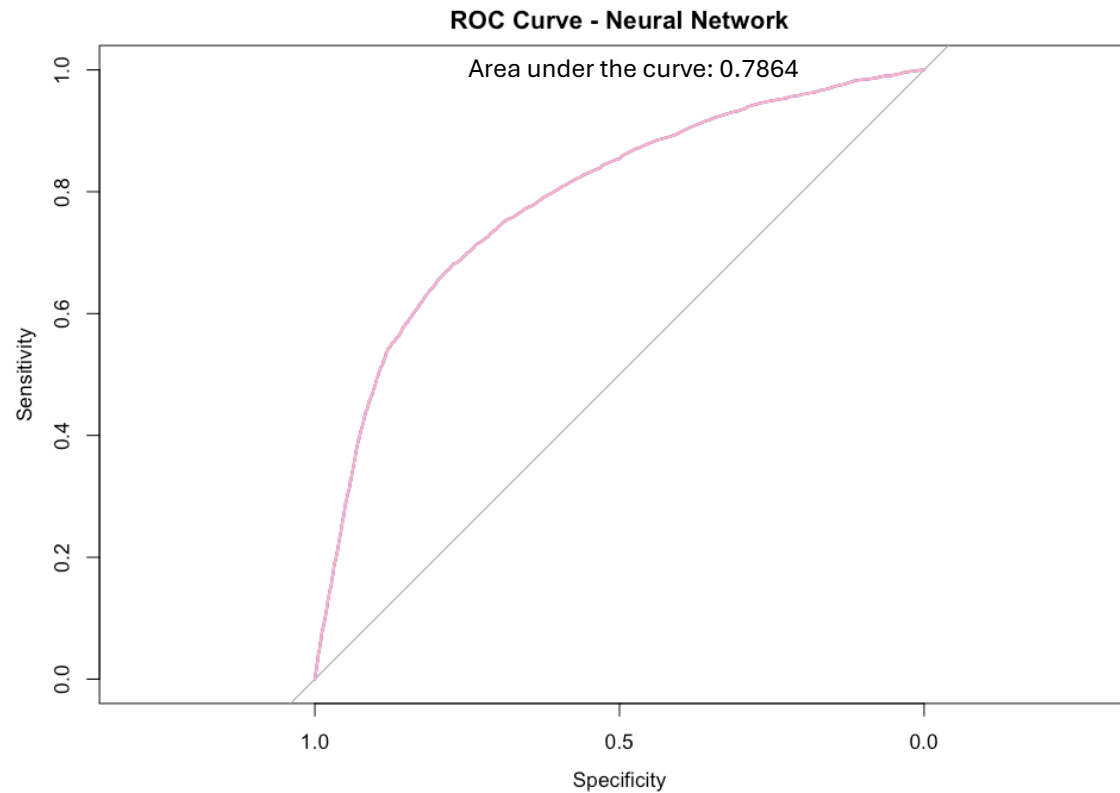
'Positive' Class : 1

RETE NEURALE MLP



```
> print(importance)
```

```
Overall
X1 34.96618
X2 16.03156
X3 11.32777
X4 23.11674
X5 14.55775
```



Confusion Matrix and Statistics

	Reference	
Prediction	0	1
0	10984	2177
1	1056	1783

Accuracy : 0.7979
95% CI : (0.7916, 0.8041)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.00000000000000022

Kappa : 0.4006

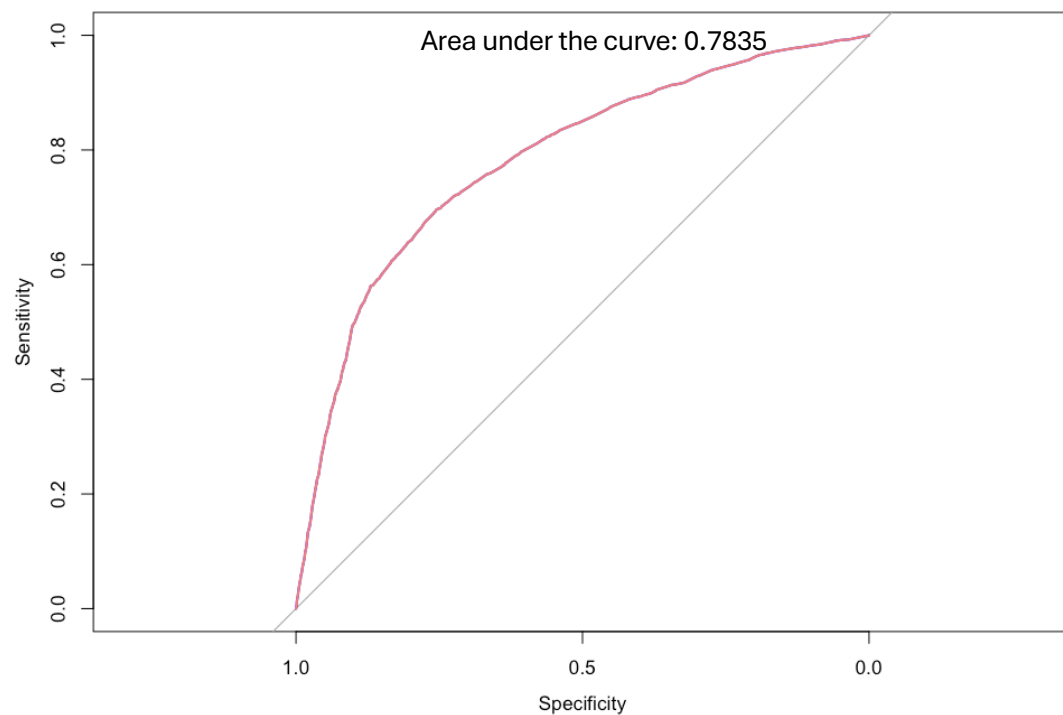
McNemar's Test P-Value : < 0.00000000000000022

Sensitivity : 0.4503
Specificity : 0.9123
Pos Pred Value : 0.6280
Neg Pred Value : 0.8346
Prevalence : 0.2475
Detection Rate : 0.1114
Detection Prevalence : 0.1774
Balanced Accuracy : 0.6813

'Positive' Class : 1

KNN

ROC Curve - KNN



Confusion Matrix and Statistics

	Reference	
Prediction	no	yes
no	10901	2087
yes	1139	1873

Accuracy : 0.7984
95% CI : (0.7921, 0.8046)
No Information Rate : 0.7525
P-Value [Acc > NIR] : < 0.0000000000000022

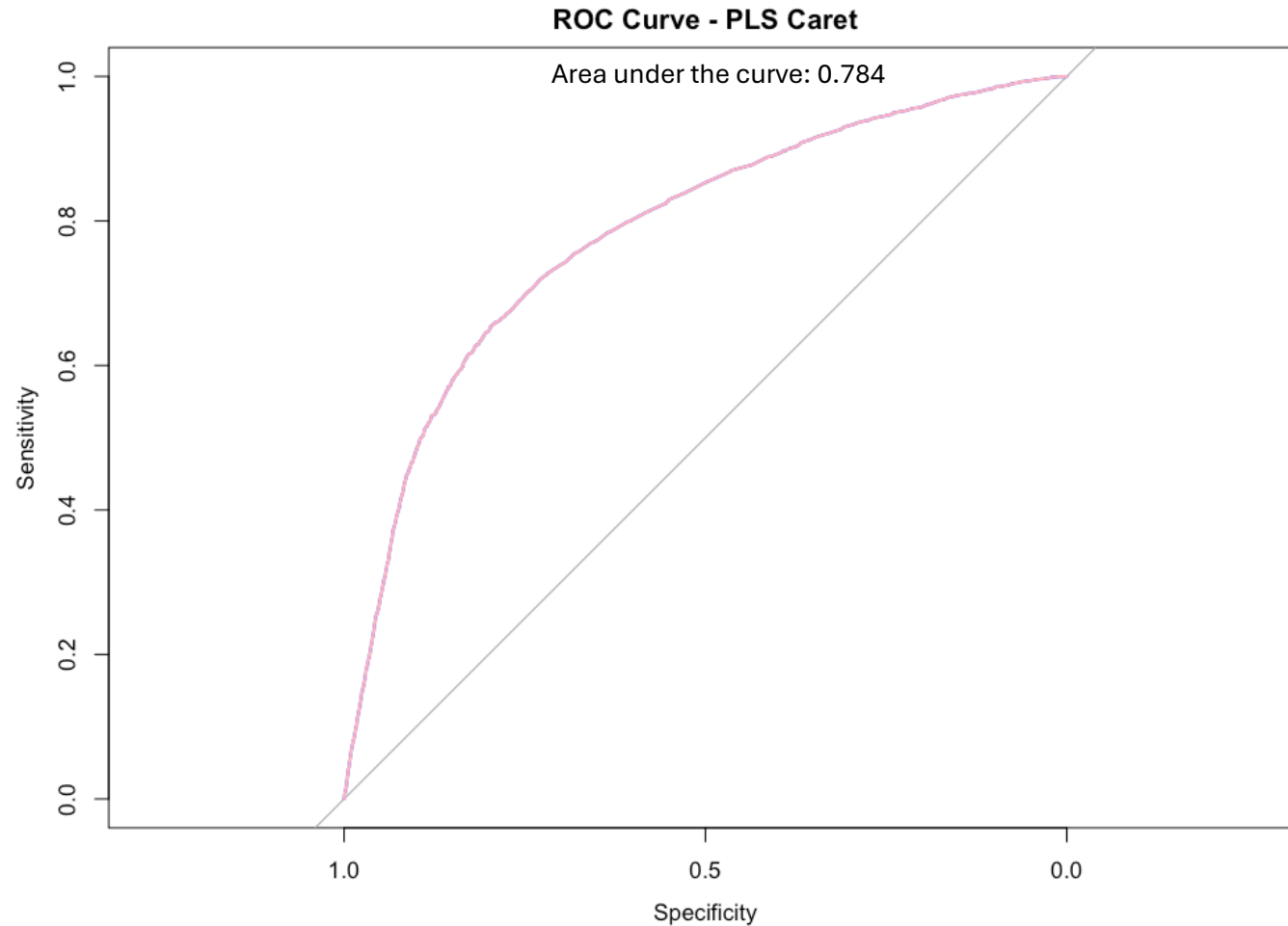
Kappa : 0.4114

McNemar's Test P-Value : < 0.0000000000000022

Sensitivity : 0.4730
Specificity : 0.9054
Pos Pred Value : 0.6218
Neg Pred Value : 0.8393
Prevalence : 0.2475
Detection Rate : 0.1171
Detection Prevalence : 0.1883
Balanced Accuracy : 0.6892

'Positive' Class : yes

PLS



Confusion Matrix and Statistics

Prediction	Reference	
	1	0
1	1145	632
0	2815	11408

Accuracy : 0.7846

95% CI : (0.7781, 0.7909)

No Information Rate : 0.7525

P-Value [Acc > NIR] : < 0.00000000000000022

Kappa : 0.2904

Mcnemar's Test P-Value : < 0.00000000000000022

Sensitivity : 0.28914

Specificity : 0.94751

Pos Pred Value : 0.64434

Neg Pred Value : 0.80208

Prevalence : 0.24750

Detection Rate : 0.07156

Detection Prevalence : 0.11106

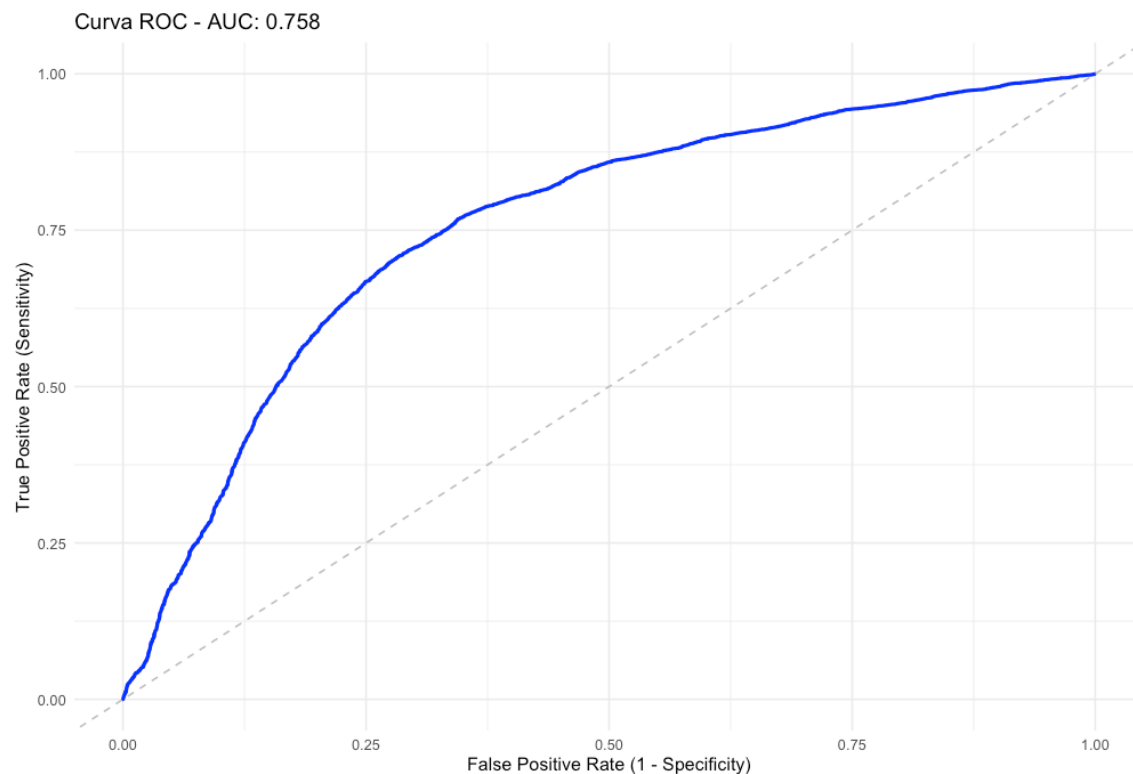
Balanced Accuracy : 0.61832

'Positive' Class : 1

MIGLIOR MODELLO PER SENSITIVITY: QDA

I RISULTATI SEGUENTI SEGUONO CROSS VALIDATION PER MIGLIOR SOGLIA PER MASSIMIZZARE F1_SCORE

"Miglior soglia per massimizzare F1-score: 0.1"



Confusion Matrix and Statistics

	Reference	
Prediction	1	0
1	2295	10663
0	1665	1377

Accuracy : 0.2295
95% CI : (0.223, 0.2361)
No Information Rate : 0.7525
P-Value [Acc > NIR] : 1

Kappa : -0.1737

Mcnemar's Test P-Value : <2e-16

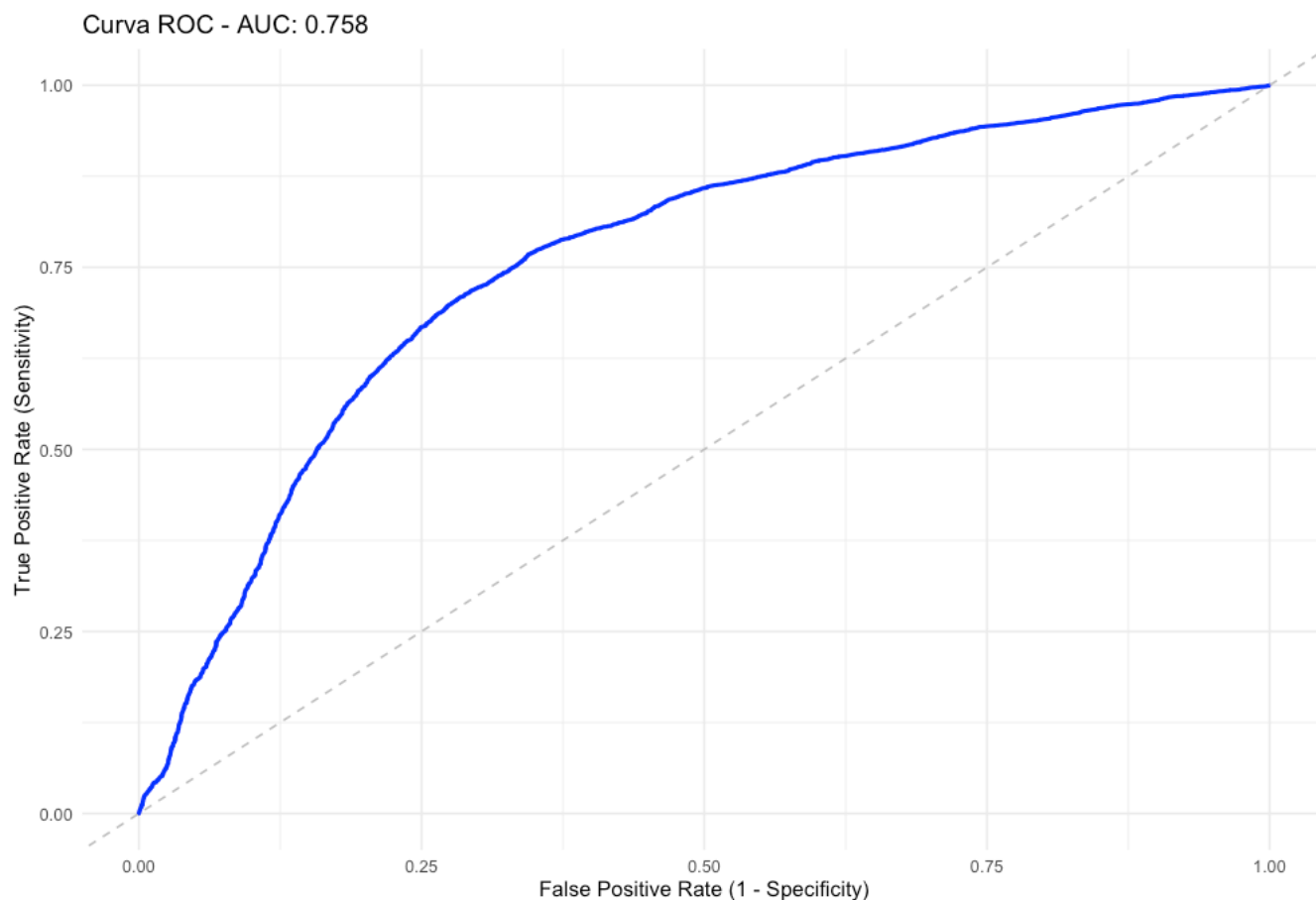
Sensitivity : 0.5795
Specificity : 0.1144
Pos Pred Value : 0.1771
Neg Pred Value : 0.4527
Prevalence : 0.2475
Detection Rate : 0.1434
Detection Prevalence : 0.8099
Balanced Accuracy : 0.3470

'Positive' Class : 1

MIGLIOR MODELLO PER SENSITIVITY: QDA

I RISULTATI SEGUENTI SEGUONO CROSS VALIDATION PER MIGLIOR SOGLIA PER MASSIMIZZARE Area Under the Curve: TP/FN

"Miglior soglia per massimizzare AUC: 0.75"



Confusion Matrix and Statistics

Reference

Prediction 1 0

1 1358 9207

0 2602 2833

Accuracy : 0.2619

95% CI : (0.2551, 0.2688)

No Information Rate : 0.7525

P-Value [Acc > NIR] : 1

Kappa : -0.2704

McNemar's Test P-Value : <2e-16

Sensitivity : 0.34293

Specificity : 0.23530

Pos Pred Value : 0.12854

Neg Pred Value : 0.52125

Prevalence : 0.24750

Detection Rate : 0.08488

Detection Prevalence : 0.66031

Balanced Accuracy : 0.28911

'Positive' Class : 1