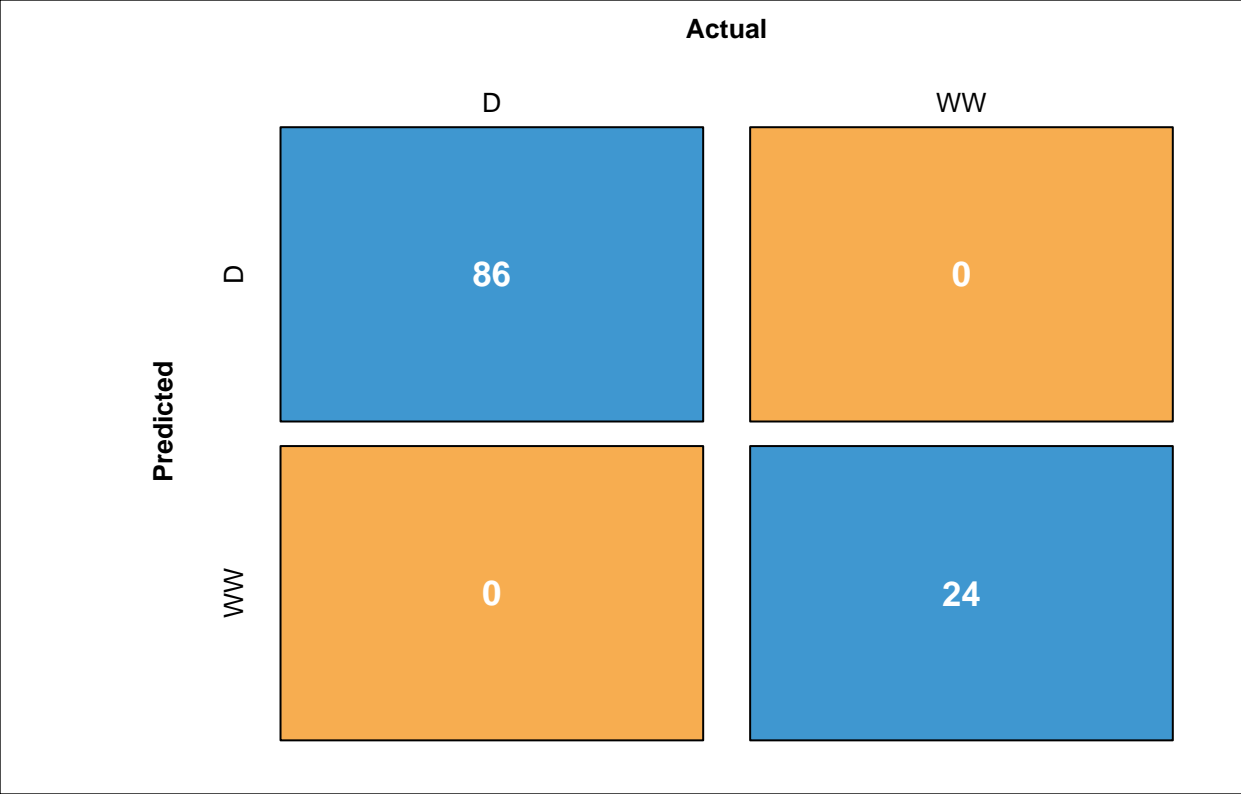


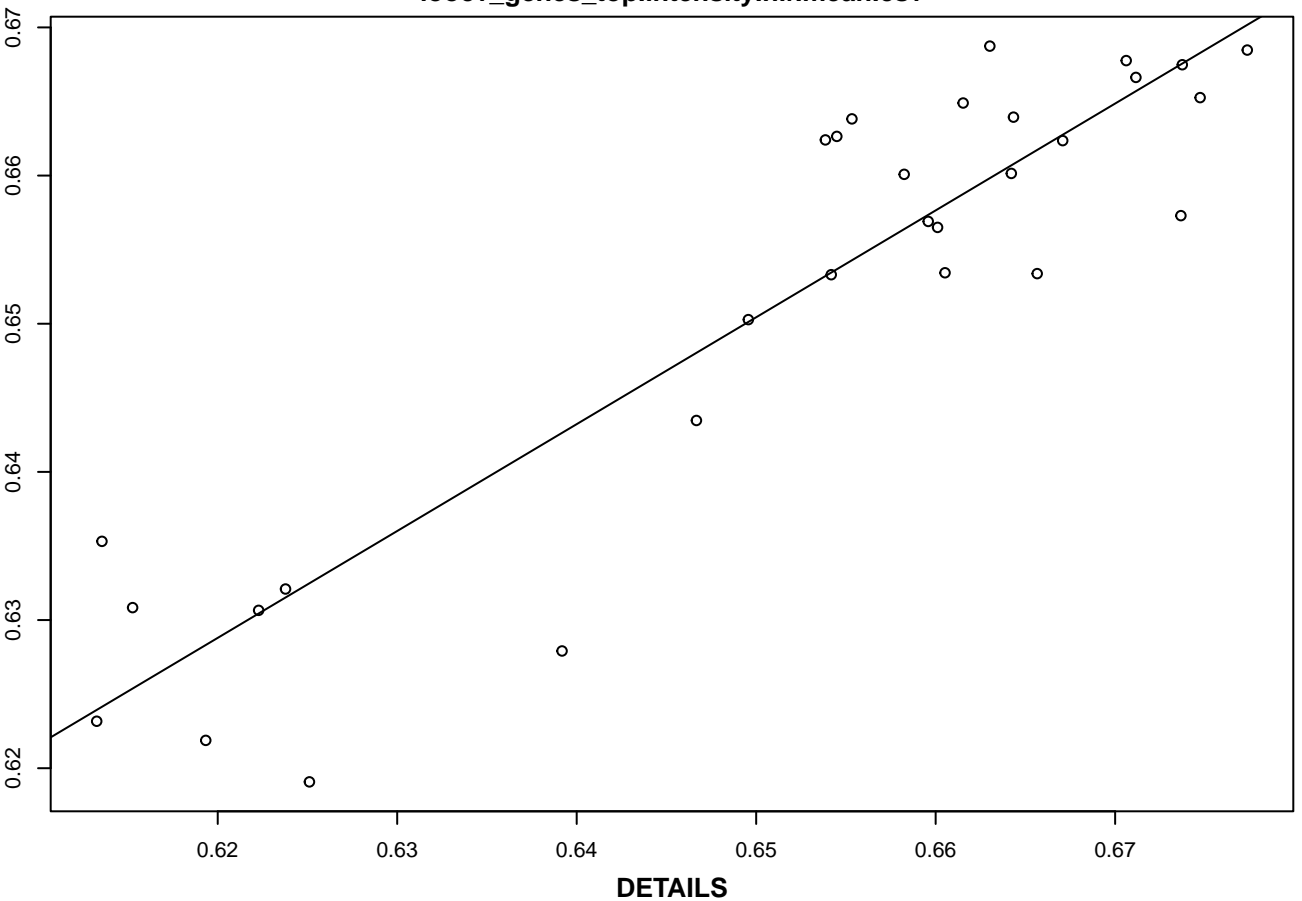
# CONFUSION MATRIX



## DETAILS

Sensitivity	Specificity	Precision	Recall	F1
1	1	1	1	1
Accuracy		Kappa		
1		1		

loocv\_genes\_top.intensity.nir.mean.csv



**RMSE**

0.008

**NRMSE**

1.3 %

**MAE**

0.007

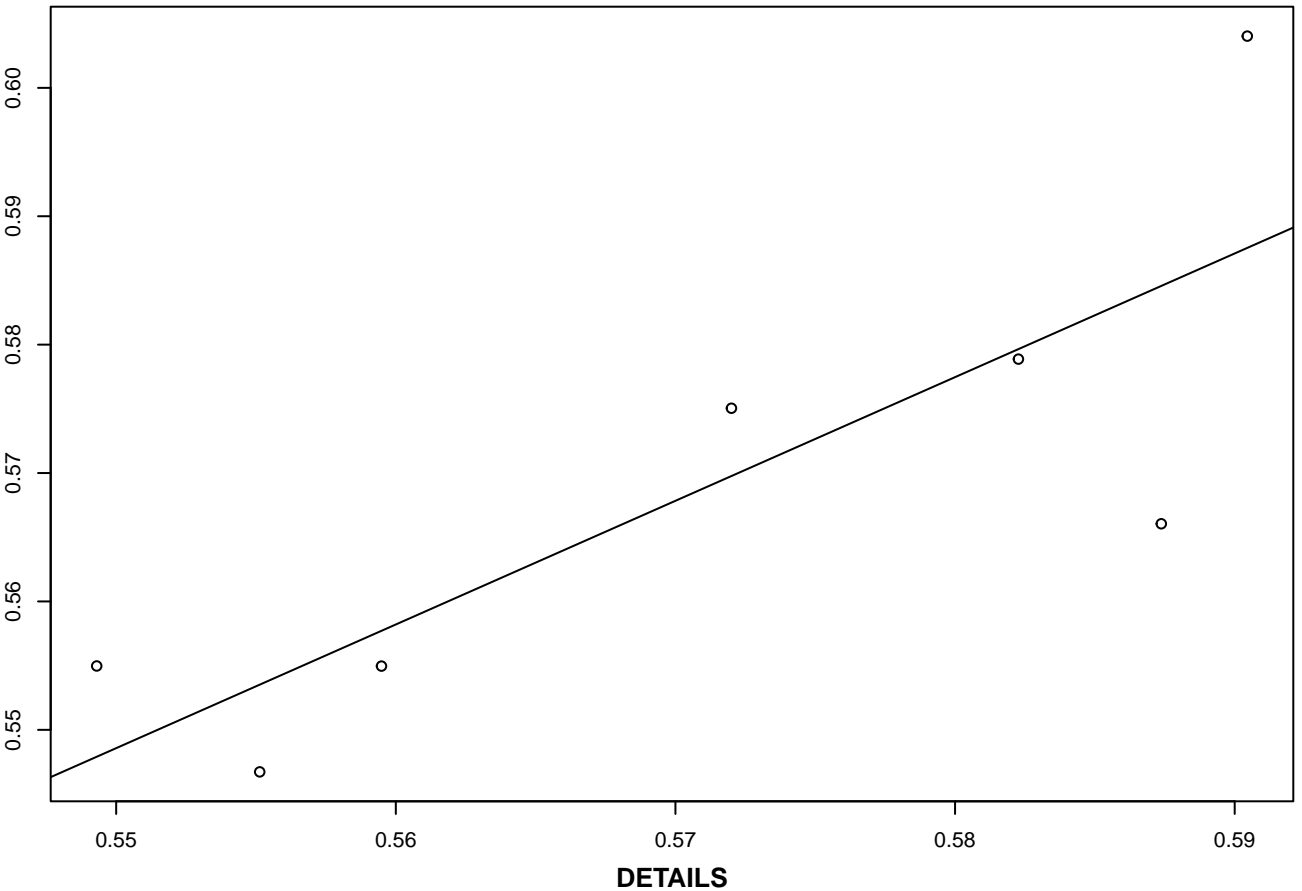
**NMAE**

1.1 %

**R<sup>2</sup>**

0.836

loocv\_genes\_top.intensity.vis.hsv.s.mean.csv



**RMSE**

0.011

**NRMSE**

1.9 %

**MAE**

0.009

**NMAE**

1.5 %

**R^2**

0.669