## Ulcerative Colitis Metabolomics sPLSDA

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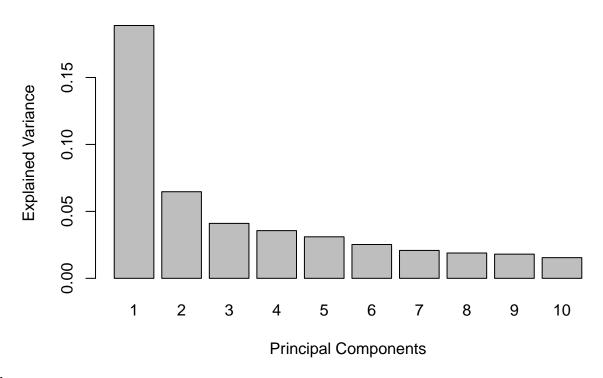
Data source: (The Inflammatory Bowel Disease Multi'omics Database)[https://ibdmdb.org/]

Data selection: n=120 (60 UC and 30 nonIBD samples selected with M/F split). Only C8-pos method data selected from metabolomic dataset.

Data preparation: Selected data was run through xmsPanda regression analysis to determine selected features. This analysis is performed on the regression selected features data.

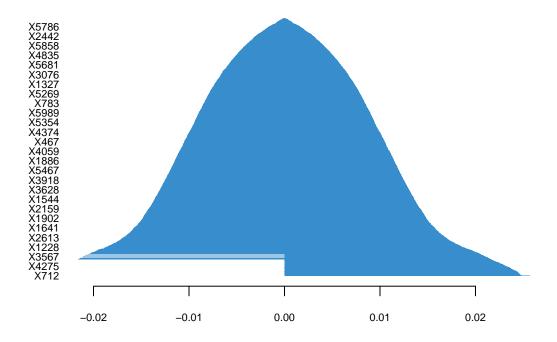
#### Initial sPLS-DA model

FIGURE 1: Barplot of the variance each principal component explains of the UC metabolomic



data.

# Loadings on comp 1



# **Loadings on comp 2**

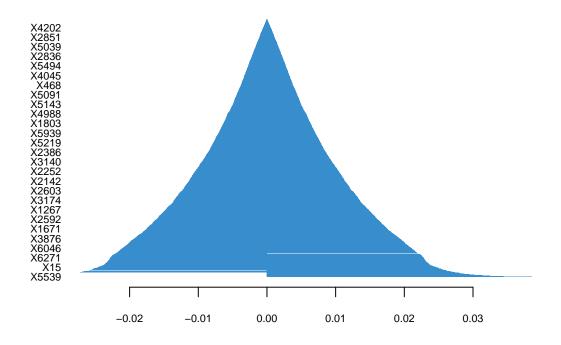
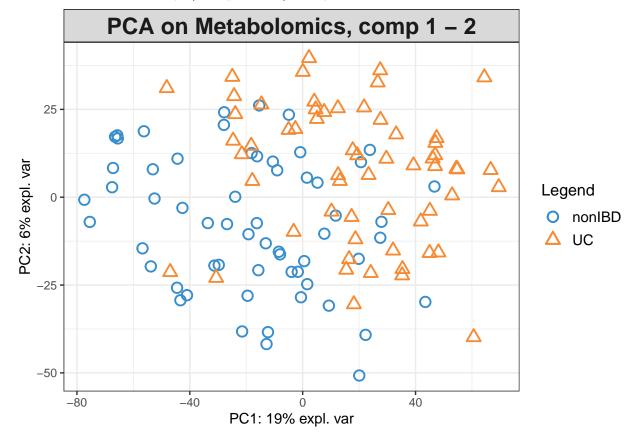


FIGURE 2: Preliminary (unsupervised) analysis with PCA on the UC metabolomic data



# **Contribution on comp 1**

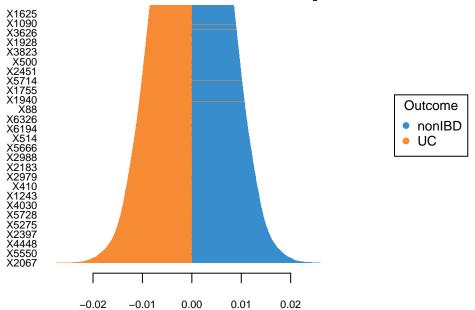
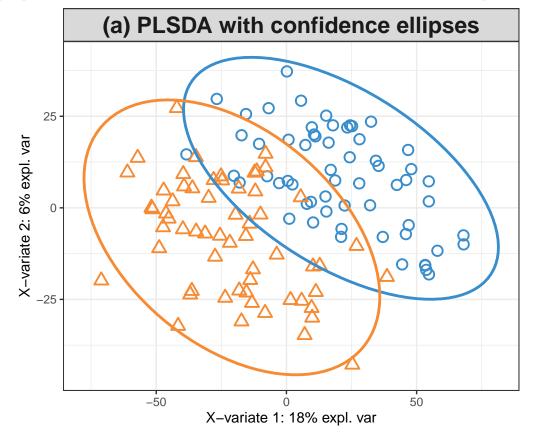


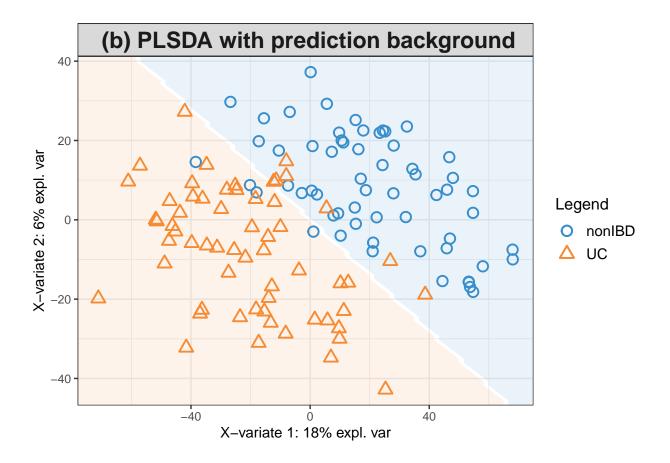
FIGURE 3: Sample plots of the UC metabolomic data after a basic PLS-DA model was oper-



Legend

○ nonII△ UC

ated on this data.



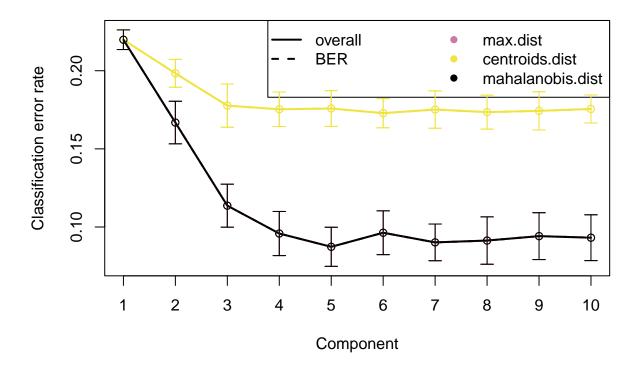
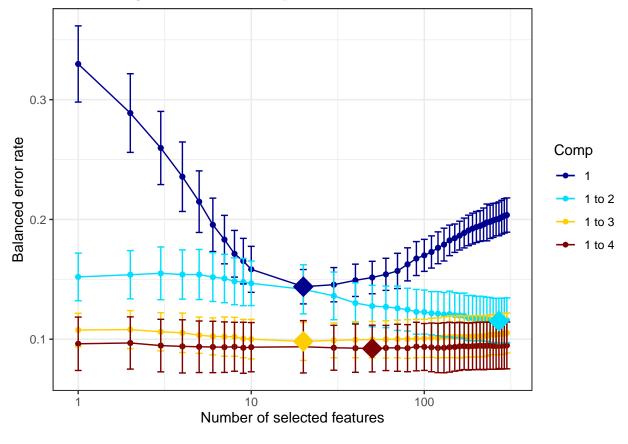
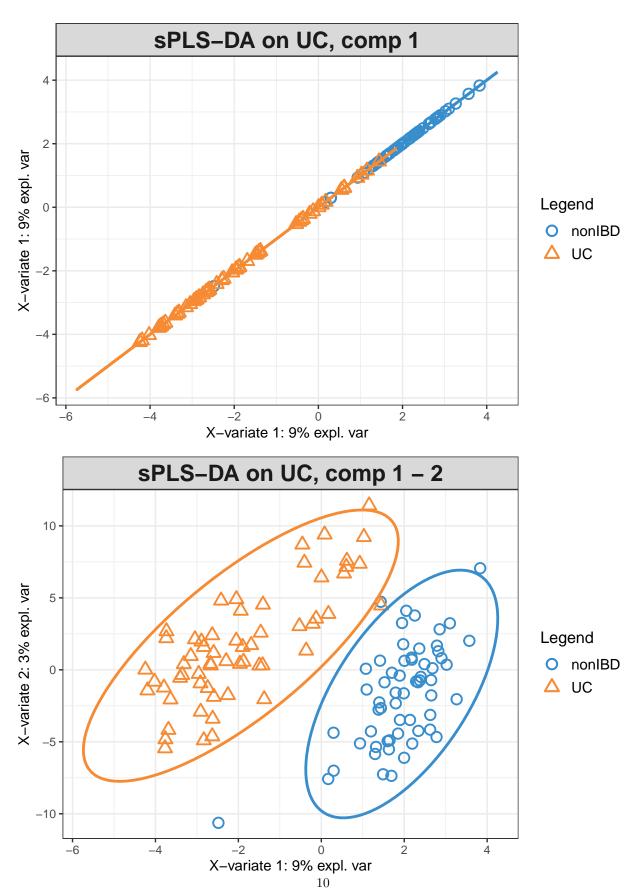
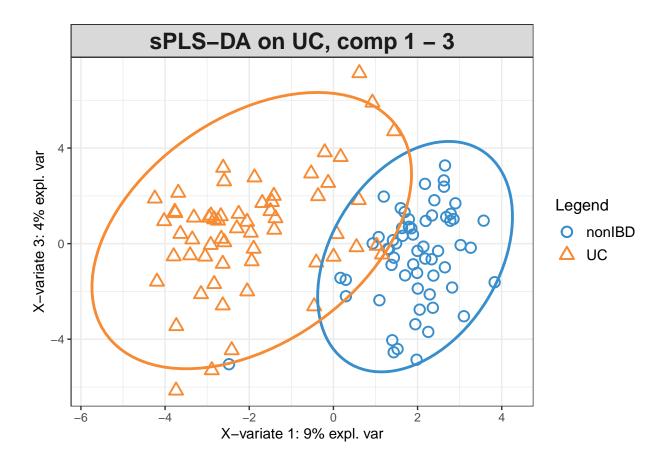


FIGURE 4: Tuning the number of components in PLS-DA on the UC metabolomic data



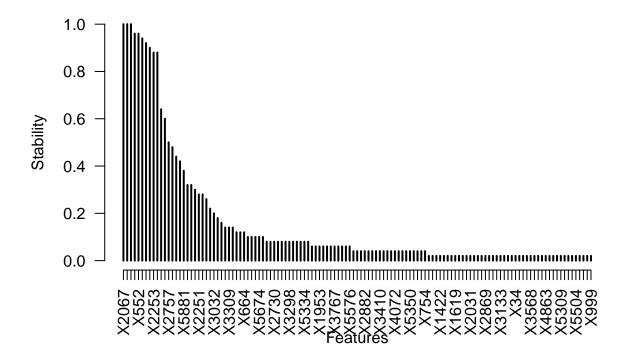
### Creating the final model



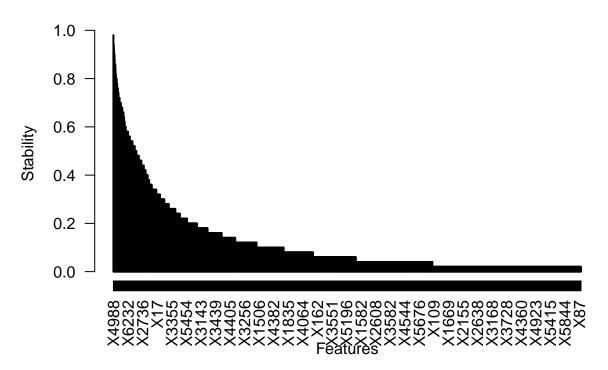


## Error in cim plot: figure margins too large. See ?cim for help.

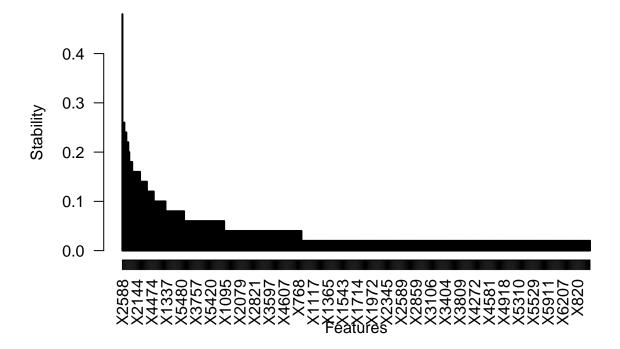
# (a) Comp 1



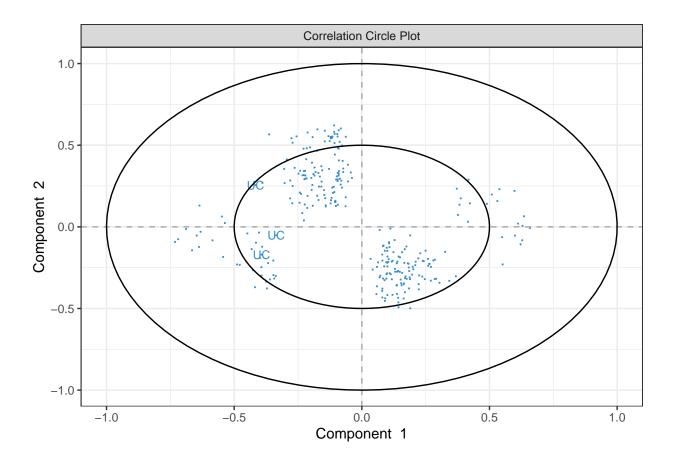
# (b) Comp 2



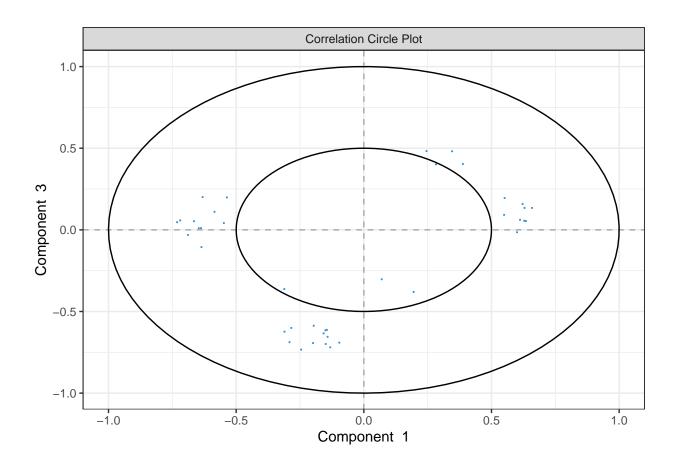
# (c) Comp 3



## Warning: Removed 287 rows containing missing values ('geom\_text()').



## Warning: Removed 40 rows containing missing values ('geom\_text()').



### Training the model

### Confusion matrix

Table 1: Confusion matrix

	nonIBD	UC
nonIBD	29	4
UC	0	24

```
## Attaching package: 'caret'

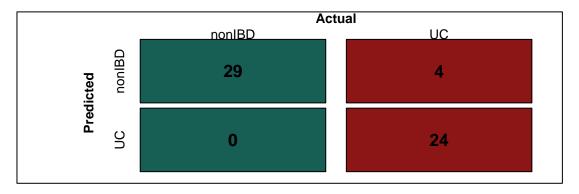
## The following objects are masked from 'package:mixOmics':
##
## nearZeroVar, plsda, splsda

## Confusion Matrix and Statistics
##
## Reference
## Prediction nonIBD UC
```

## nonIBD 29 4 ## UC 0 24 ## ## Accuracy : 0.9298 95% CI: (0.83, 0.9805) ## No Information Rate : 0.5088 ## P-Value [Acc > NIR] : 6.94e-12 ## ## ## Kappa: 0.8593 ## ## Mcnemar's Test P-Value : 0.1336 ## ## Sensitivity: 0.8571 Specificity: 1.0000 ## ## Pos Pred Value : 1.0000 Neg Pred Value: 0.8788 ## ## Prevalence: 0.4912 Detection Rate: 0.4211 ## ## Detection Prevalence : 0.4211 Balanced Accuracy: 0.9286 ## ## ## 'Positive' Class : UC ##

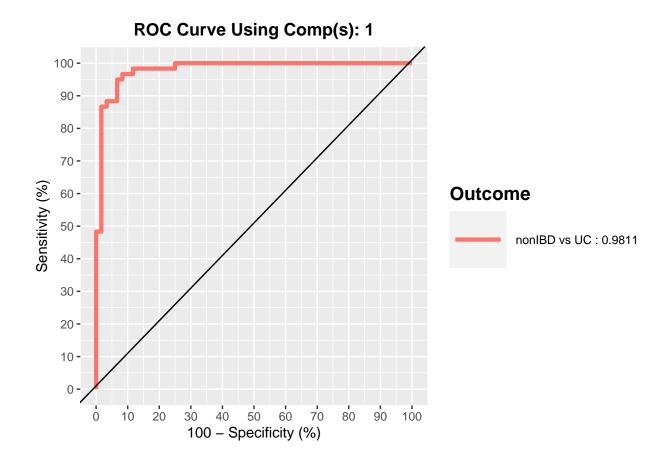
### Performance plots

## **Confusion Matrix**

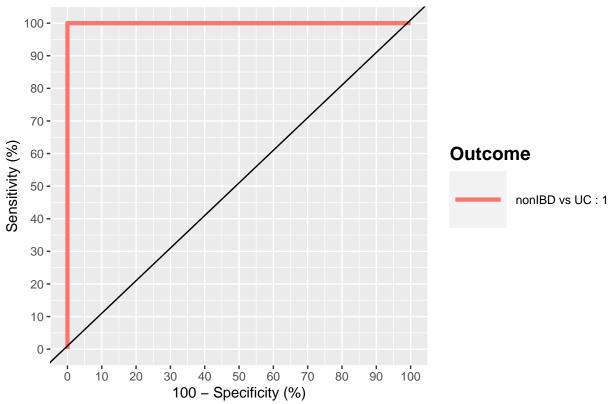


### **Confusion matrix statistics**

Sensitivity	Specyficity	Precision	Recall Balancepotaccuracy
, ,	Acgugacy	0.00	Kanga (1.33)







### Odds ratio for possible confounding variables

```
Odds ratio
##
                                   2.5 %
                                           97.5 %
## (Intercept)
                             1 0.6028659 1.658744
## samples$sexMale
                             1 0.4888607 2.045572
##
                       Odds ratio
                                       2.5 %
                                               97.5 %
## (Intercept)
                         0.893486 0.4715531 1.692953
## samples$consent_age
                         1.004058 0.9851150 1.023366
##
                                     Odds ratio 2.5 % 97.5 %
## (Intercept)
                                   4.254481e+07
                                                         Inf
## samples$raceMore than one race 1.762847e-08
                                                          Inf
## samples$raceWhite
                                   2.056655e-08
                                                          Inf
```