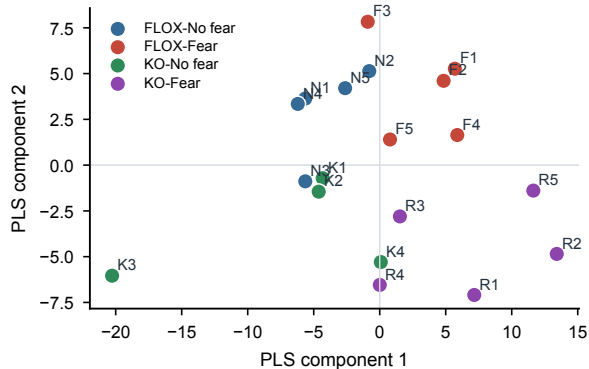


# Lipidomics - negative ion mode - supervised overview

$R^2X=0.319$   $R^2Y=0.440$   $Q^2=-0.208$  CV accuracy=0.21 permutation P=0.433

## A Four-group PLS-DA scores



## B Response-label permutation

