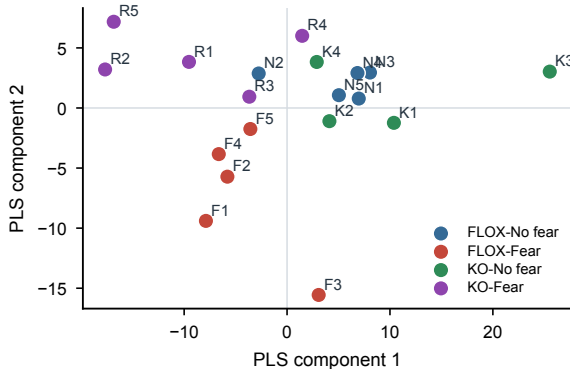


Lipidomics - positive ion mode - supervised overview

$R^2X=0.361$ $R^2Y=0.434$ $Q^2=-0.011$ CV accuracy=0.42 permutation P=0.060

A Four-group PLS-DA scores



B Response-label permutation

