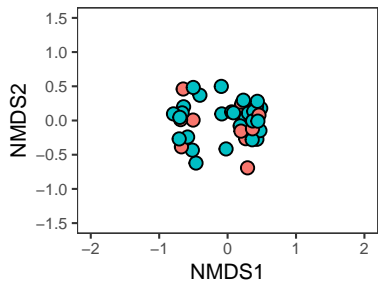
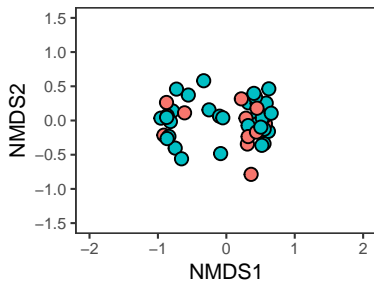


# noncdi\_abx\_total

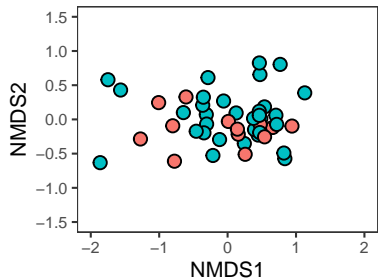
## Taxonomy



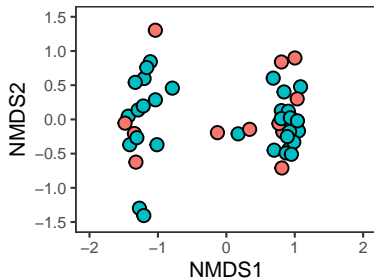
## Genes



## Metabolites



## MCBA



noncdi\_abx\_total

any

none