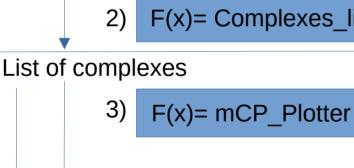


input mCP will take the data and prepare it as An imput to mCP, this one we can work later

Complexes list: It will take the data and make a list of all plots possible in the experiment in



mCP_Plotter: It will take the list and plot all protein Complexes & heatmaps that: 1) have more than 2 proteins 2) have more than 2 proteins and Pearson

reference to the Corum list.

Functions

F=0.93& F=0 mCP output 1 F(x) = mCP ROC

FDR output 1 (ROC)

correlation>0.93. mCP ROC gives FDR analysis by ROC method Take as true positive all complexes in Corum list. I have to work on that