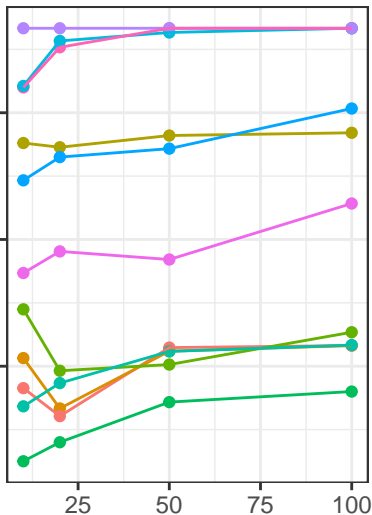


Accuracy

0.8  
0.6  
0.4

n



diagREDM

diagREDMSL

fullREDMSL

fullREM

HMP

HMP2

ILR

permutation

perturbation

pvals\_chi\_Harris

ttest