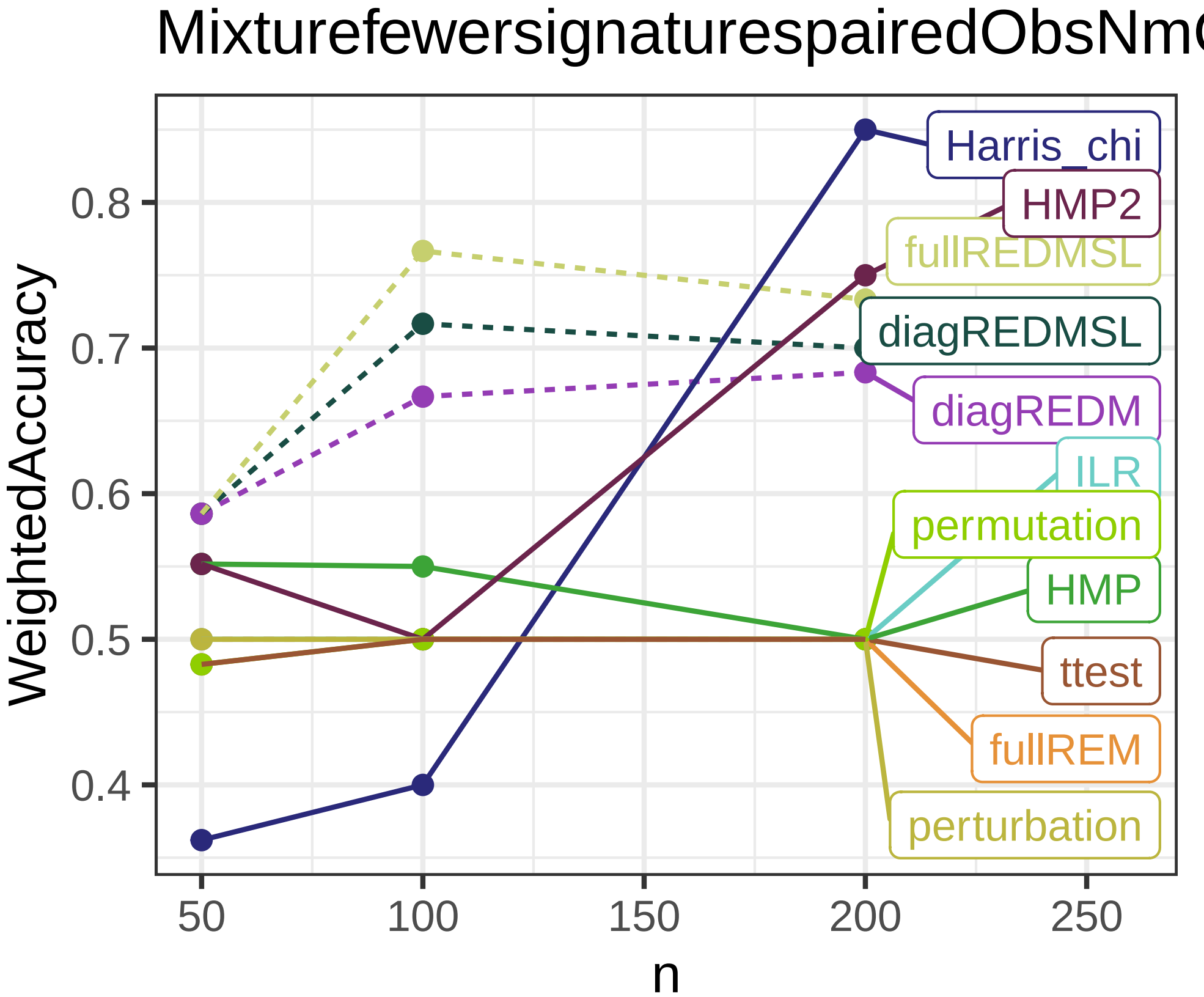
MixturefewersignaturespairedObsNmC
WeightedAccuracy
0.8
0.7
0.6
0.5
0.4
50
100
150
200
250
n
Harris_chi
HMP2
fullREDMSL
diagREDMSL
diagREDM
ILR
permutation
HMP
ttest
fullREM
perturbation