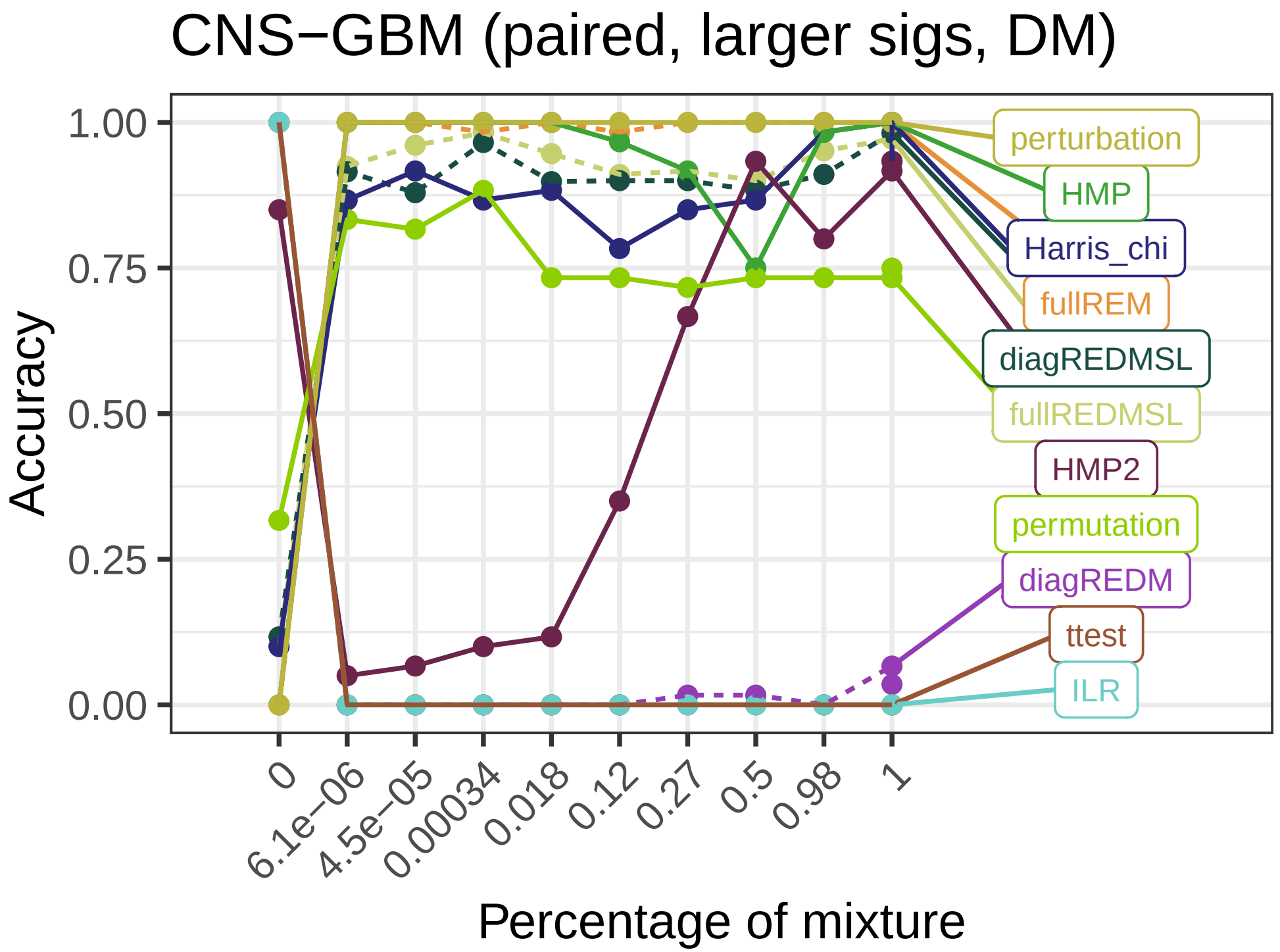
CNS−GBM (paired, larger sigs, DM)
Accuracy
1.00
0.75
0.50
0.25
0.00
0
6.1e−06
4.5e−05
0.00034
0.018
0.12
0.27
0.5
0.98
1
Percentage of mixture
perturbation
HMP
Harris_chi
fullREM
diagREDMSL
fullREDMSL
HMP2
permutation
diagREDM
ttest
ILR