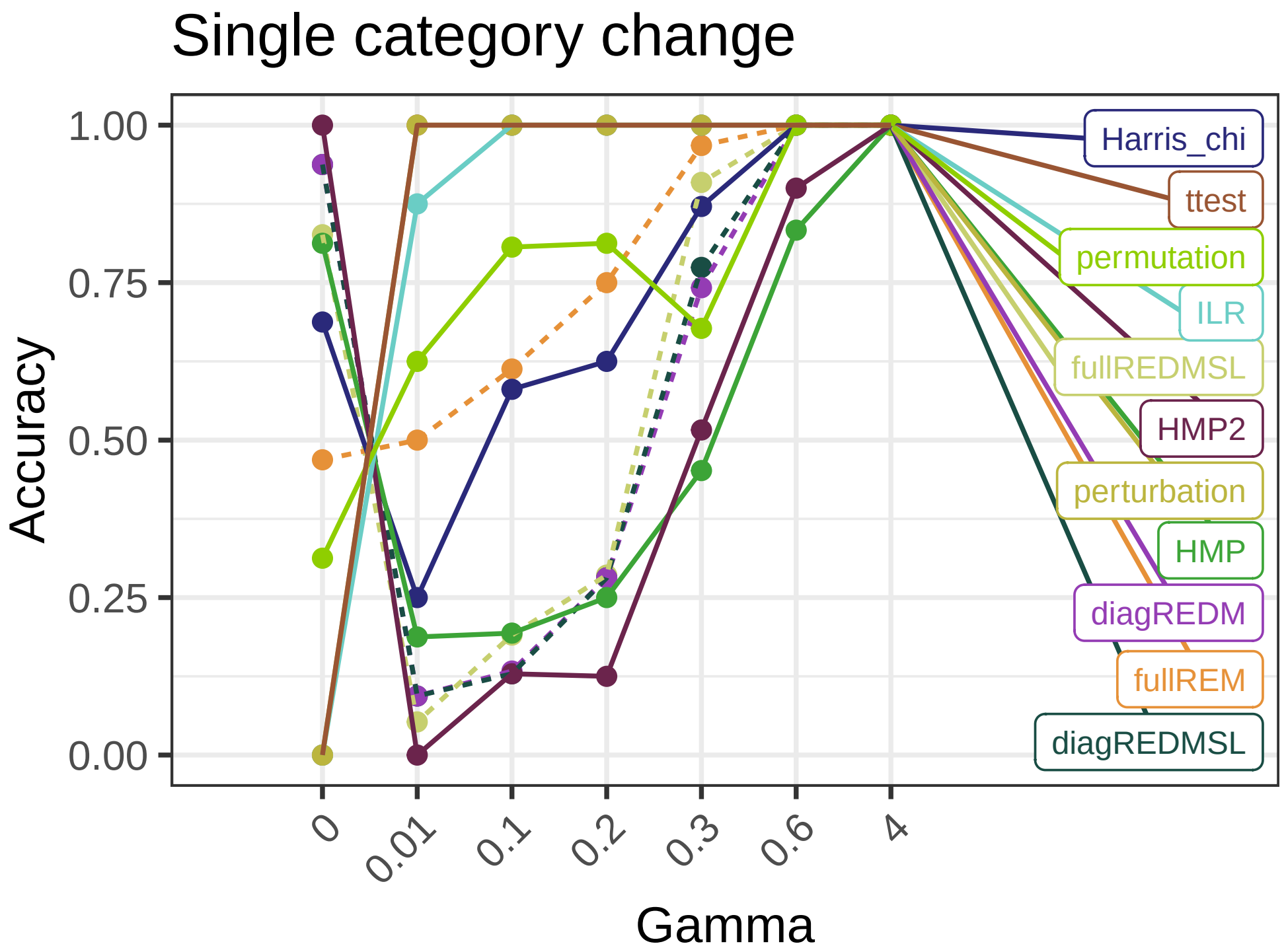
Single category change
Accuracy
1.00
0.75
0.50
0.25
0.00
0
0.01
0.1
0.2
0.3
0.6
4
Gamma
Harris_chi
ttest
permutation
ILR
fullREDMSL
HMP2
perturbation
HMP
diagREDM
fullREM
diagREDMSL