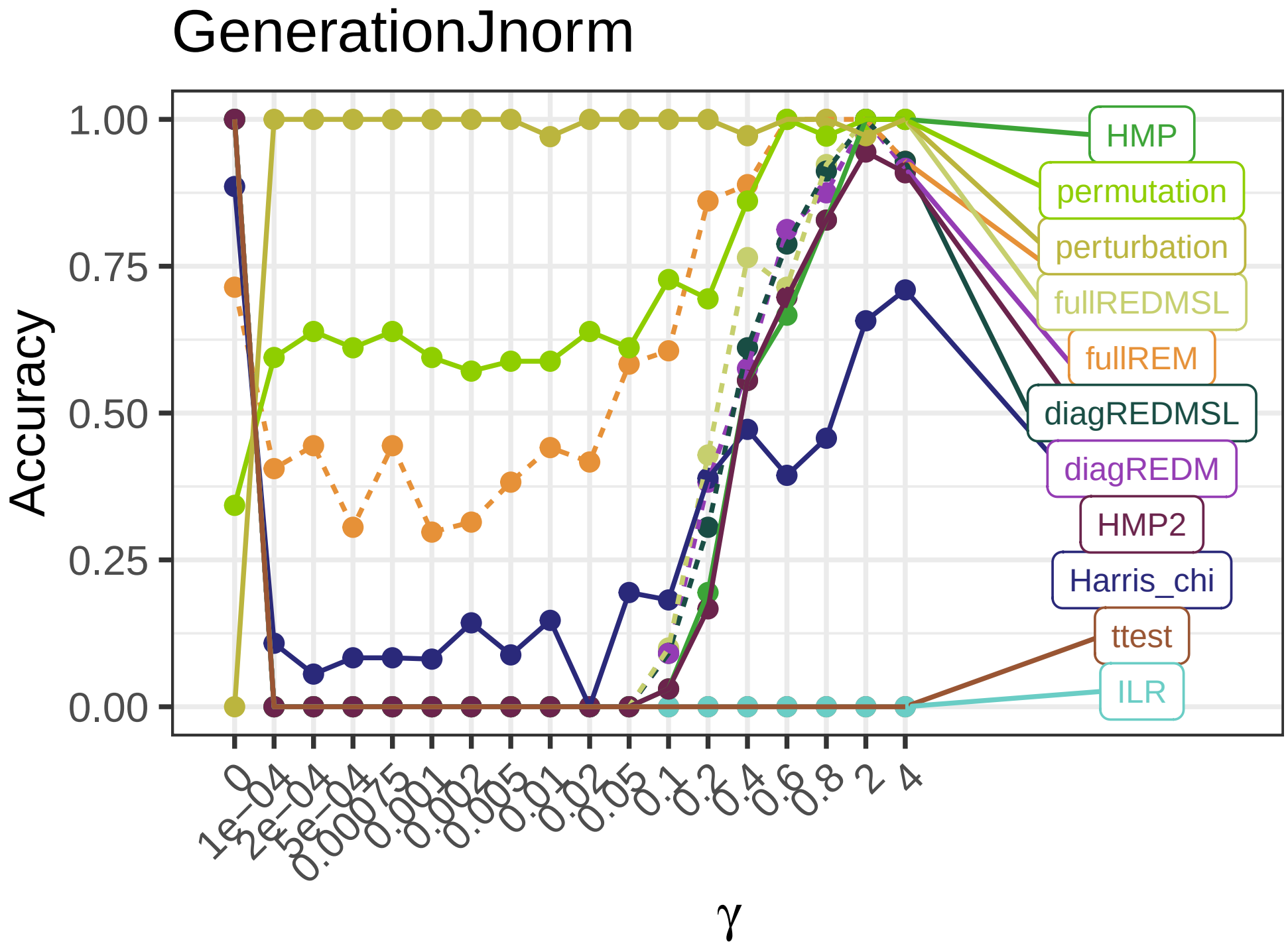
GenerationJnorm
Accuracy
1.00
0.75
0.50
0.25
0.00
0
1e−04
2e−04
5e−04
0.0001
0.0002
0.0005
0.001
0.002
0.005
0.01
0.02
0.05
0.1
0.2
0.4
0.6
0.8
2
4
γ
HMP
permutation
perturbation
fullREDMSL
fullREM
diagREDMSL
diagREDM
HMP2
Harris_chi
ttest
ILR