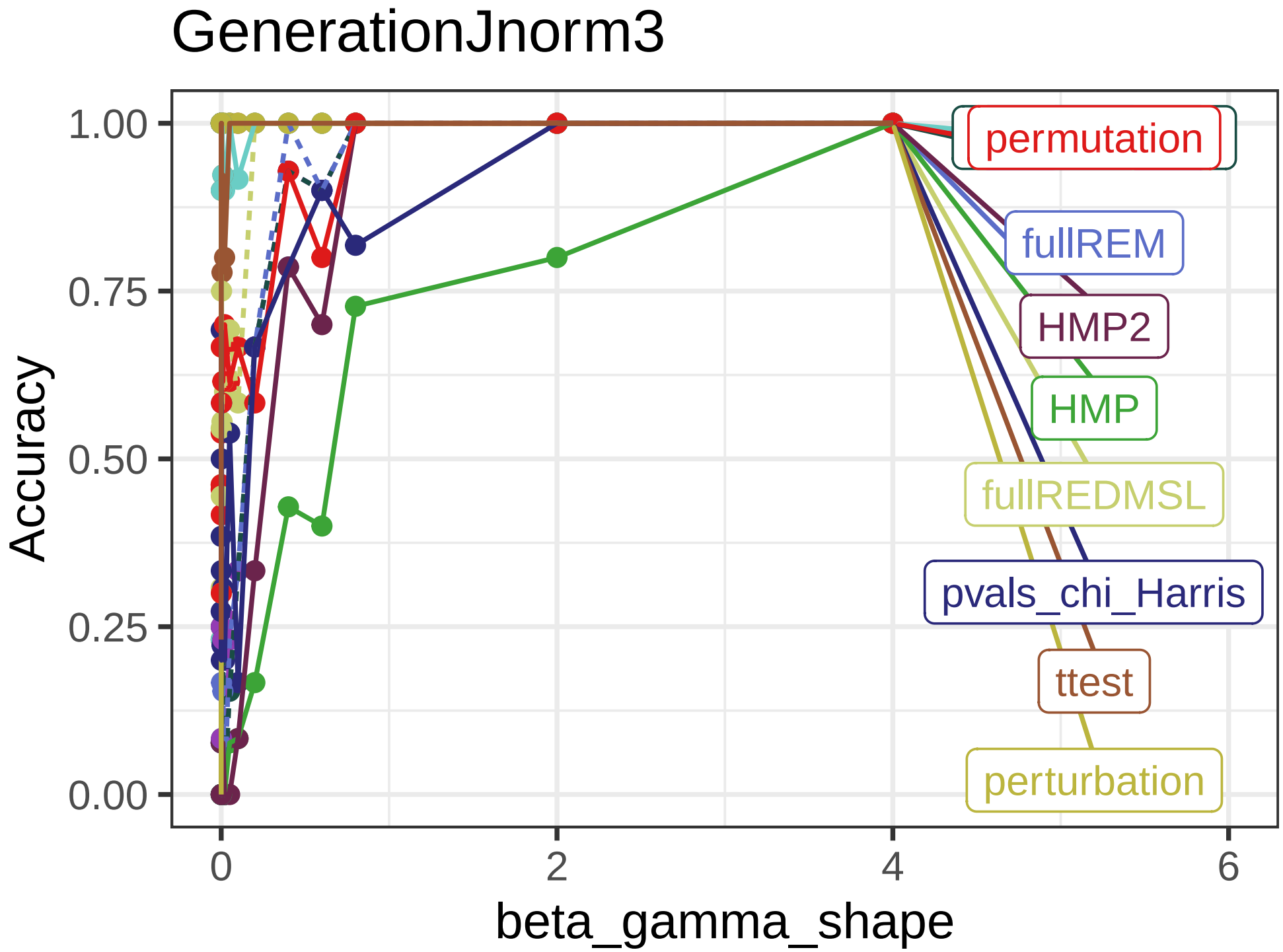
GenerationJnorm3
Accuracy
1.00
0.75
0.50
0.25
0.00
0
2
4
6
beta_gamma_shape
permutation
fullREM
HMP2
HMP
fullREDMSL
pvals_chi_Harris
ttest
perturbation