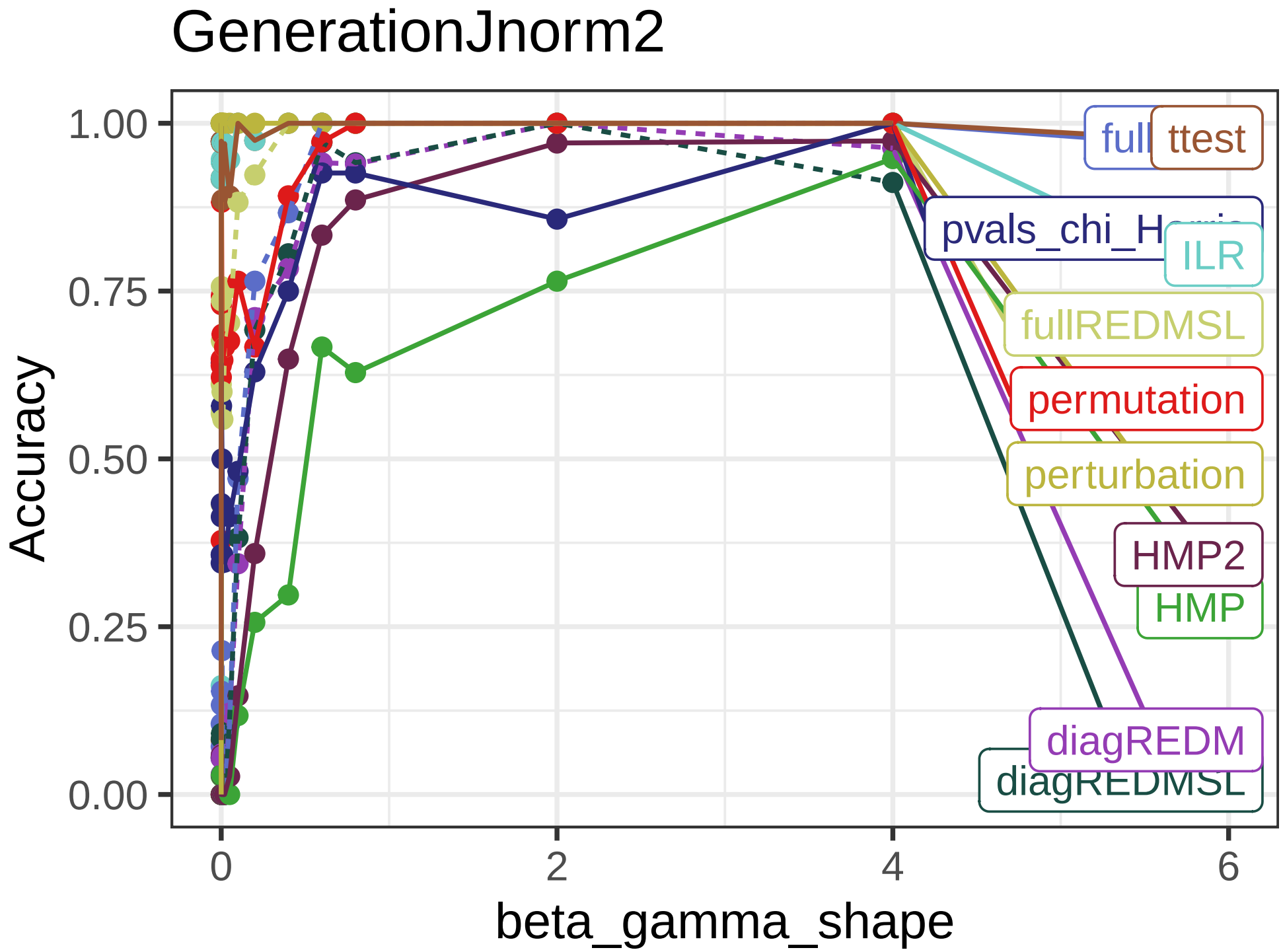
GenerationJnorm2
Accuracy
1.00
0.75
0.50
0.25
0.00
0
2
4
6
beta_gamma_shape
full
ttest
pvals_chi_H
ILR
fullREDMSL
permutation
perturbation
HMP2
HMP
diagREDM
diagREDMSL