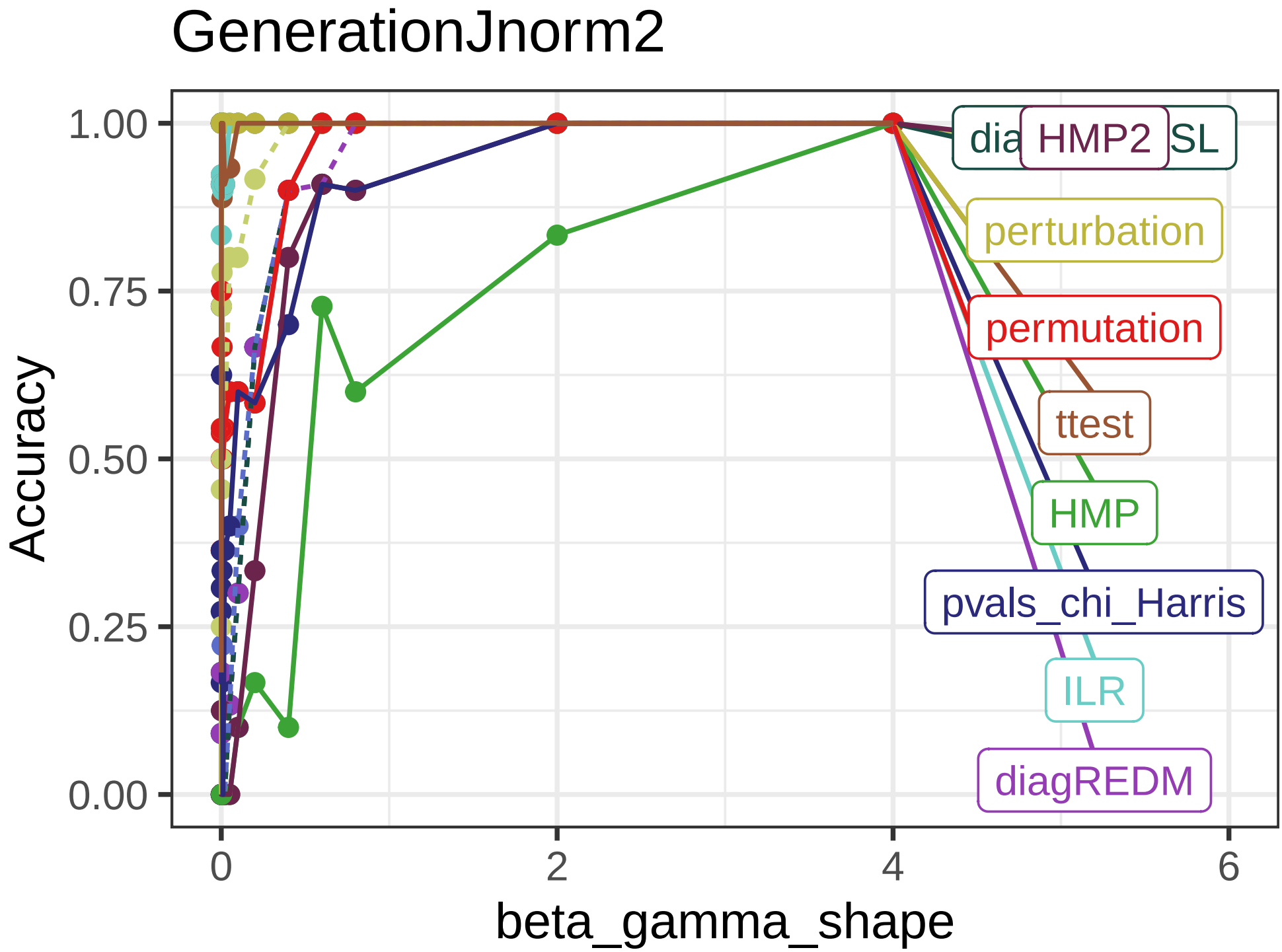
GenerationJnorm2
Accuracy
1.00
0.75
0.50
0.25
0.00
0
2
4
6
beta_gamma_shape
dia
HMP2
SL
perturbation
permutation
ttest
HMP
pvals_chi_Harris
ILR
diagREDM