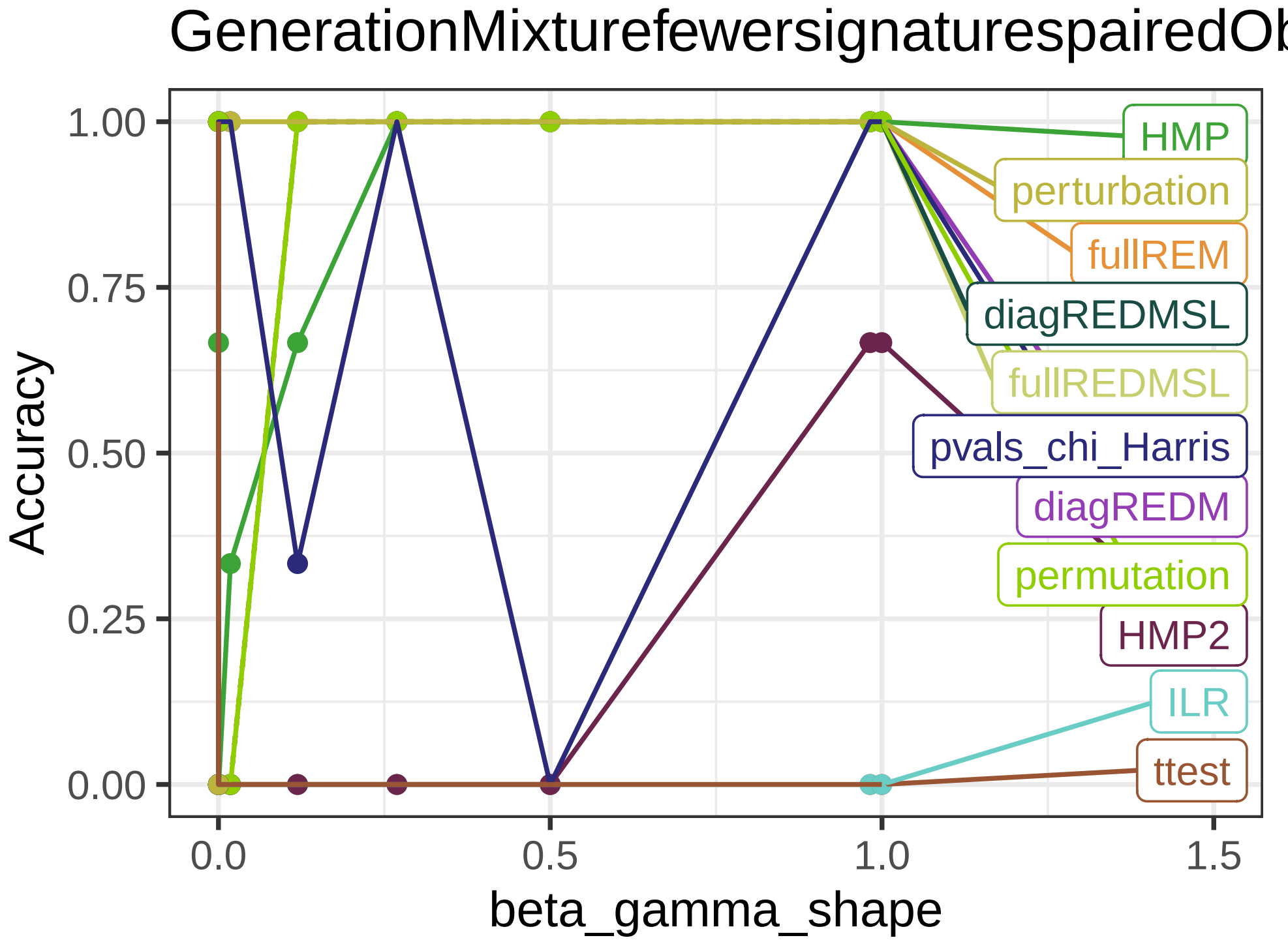
GenerationMixturefewersignaturespairedOb
Accuracy
1.00
0.75
0.50
0.25
0.00
0.0
0.5
1.0
1.5
beta_gamma_shape
HMP
perturbation
fullREM
diagREDMSL
fullREDMSL
pvals_chi_Harris
diagREDM
permutation
HMP2
ILR
ttest