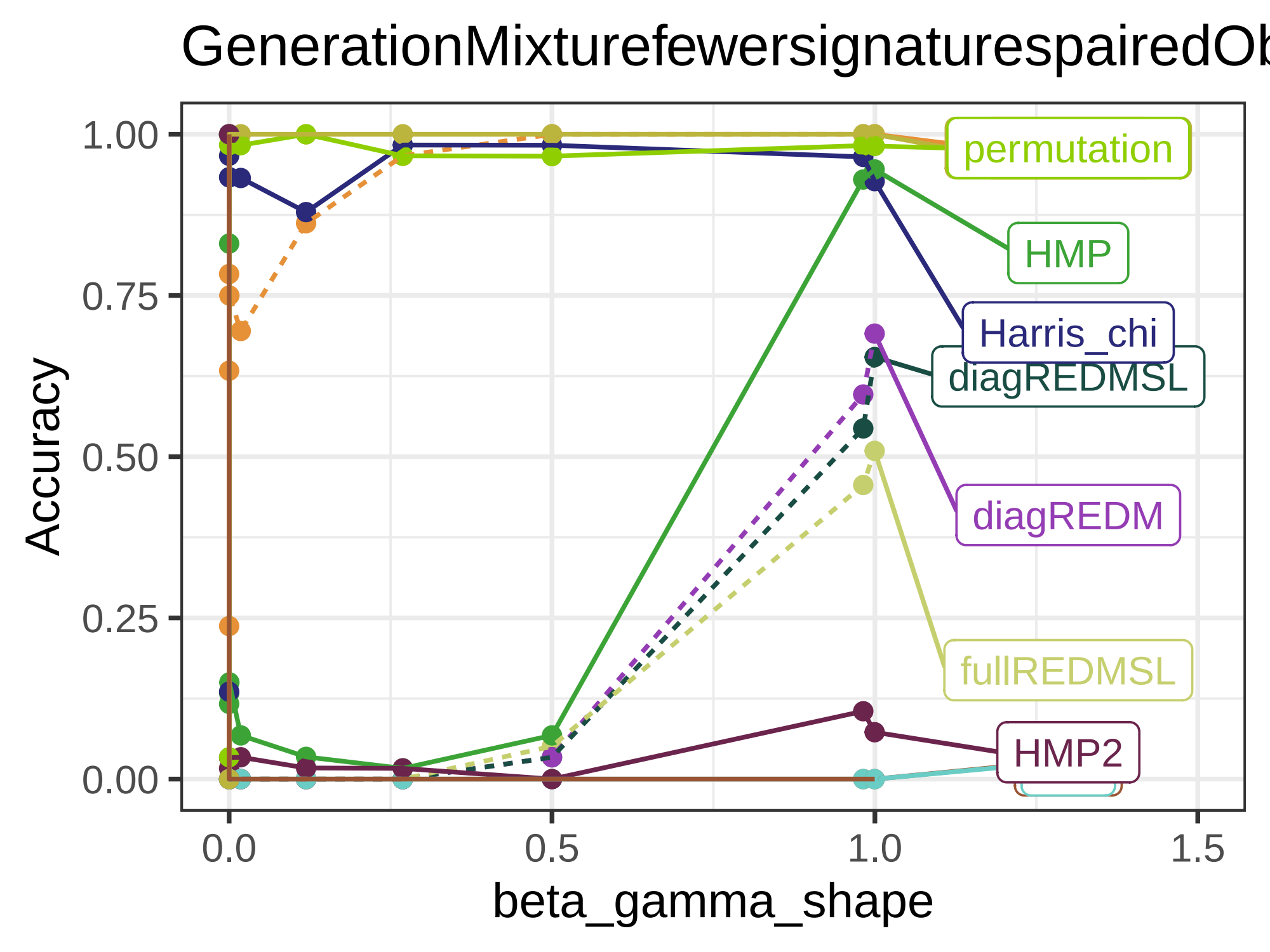
GenerationMixturefewersignaturespairedOb
Accuracy
1.00
0.75
0.50
0.25
0.00
0.0
0.5
1.0
1.5
beta_gamma_shape
permutation
HMP
Harris_chi
diagREDMSL
diagREDM
fullREDMSL
HMP2