

TT_balanced_accuracy
 $p = 6.35e-03$

0.35 0.40 0.45 0.50 0.55 0.60 0.65 0.70

brf_gmo_full (0.33)

brf_gmo_uniform (0.52)

brf_gmo_square (0.61)

(0.72) brf_gmo_precomputed

(0.69) brf_gmo_softmax

(0.62) brf_gmosa

