

TT_balanced_accuracy
 $p = 5.70e-05$

0.3 0.4 0.5 0.6 0.7 0.8

brf_gmo_uniform (0.31)

brf_gmo_softmax (0.43)

brf_gmosa (0.61)

(0.78) brf_gmo_square

(0.74) brf_gmo_full

(0.62) brf_gmo_precomputed

