

TT_balanced_accuracy
 $p = 1.83e-02$

0.40 0.45 0.50 0.55 0.60 0.65 0.70 0.75

brf_gmo_uniform (0.42)

brf_gmosa (0.5)

brf_gmo_full (0.54)

(0.74) brf_gmo_square

(0.73) brf_gmo_precomputed

(0.56) brf_gmo_softmax

