

TT_balanced_accuracy

$p = 1.21e-11$

0.2 0.4 0.6 0.8

brf_full (0.22)

bxt_full (0.31)

brf_gmo_uniform (0.38)

brf_gmo_square (0.46)

brf_gmosa (0.47)

brf_gmo_softmax (0.48)

(0.8) bxt_gmo_softmax

(0.79) bxt_gmo_square

(0.78) bxt_gmo_uniform

(0.75) bxt_gmo_precomputed

(0.54) brf_gmo_precomputed

(0.53) bxt_gmosa

