

TT_balanced_accuracy
 $p = 1.81e-19$

0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9

brf_gmo_nrlmf (0.19)

brf_lmo_nrlmf (0.23)

brf_gso_nrlmf (0.34)

brf_gmosa_nrlmf (0.53)

(0.95) brf_gmo

(0.93) brf_lmo

(0.71) brf_gmosa

(0.62) brf_gso

