

TL_balanced_accuracy
 $p = 1.01e-13$

0.2

0.4

0.6

0.8

1.0

brf_gmo_nrlmf (0.12)

brf_gso_nrlmf (0.36)

brf_lmo_nrlmf (0.52)

brf_gmo (0.54)

(0.95) brf_lmo

(0.79) brf_gso

(0.62) brf_gmosa_nrlmf

(0.6) brf_gmosa

