

TT_balanced_accuracy
 $p = 1.06e-02$

0.5

0.6

0.7

0.8

brf_gmo_nrlmf (0.45)

brf_lmo_nrlmf (0.48)

brf_gmosa (0.5)

brf_gso (0.51)

(0.84) brf_gmosa_nrlmf

(0.6) brf_lmo

(0.59) brf_gmo

(0.53) brf_gso_nrlmf

