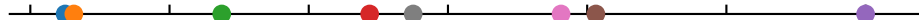


TT_balanced_accuracy
 $p = 1.22e-04$

0.3 0.4 0.5 0.6 0.7 0.8 0.9



brf_gmosa (0.33)

brf_gmo_nrlmf (0.33)

brf_lmo_nrlmf (0.44)

brf_gso (0.54)

(0.9) brf_gmo

(0.71) brf_gmosa_nrlmf

(0.68) brf_lmo

(0.57) brf_gso_nrlmf