

fit_score_time
p = 4.64e-20

0.2

0.4

0.6

0.8

1.0

bxt_gmo_nrlmf (0.12)

brf_gmo_nrlmf (0.33)

brf_gso_nrlmf (0.34)

bxt_gmo (0.47)

(1) brf_gmo

(0.85) bxt_gmosa_nrlmf

(0.77) brf_lmo

(0.61) bxt_gso_nrlmf

