

fit_score_time
p = nan

0.2

0.4

0.6

0.8

1.0

bxt_gso (0.12)

bxt_gmosa (0.26)

bxt_lmo (0.3)

bxt_lmo_nrlmf (0.34)

bxt_gmo (0.55)

(0.95) bxt_iso_nrlmf

(0.84) bxt_gmo_nrlmf

(0.78) bxt_iso

(0.76) bxt_gmosa_nrlmf

(0.6) bxt_gso_nrlmf

