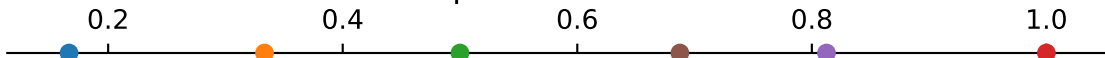


fit_score_time
 $p = 1.36e-15$



brf_gso (0.17)

(1) brf Iso

brf_lmo (0.33)

(0.81) brf_gmosa

brf_gmo (0.5)

(0.69) brf_sgso_us