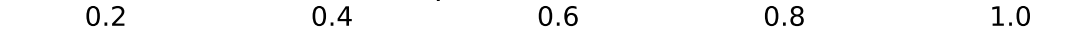


fit_time
 $p = 1.36e-15$



brf_lmo (0.17)

(1) brf_iso

brf_sgso_us (0.33)

(0.83) brf_gmosa

brf_gso (0.52)

(0.65) brf_gmo