

fit_score_time
 $p = 2.12e-15$

0.2 0.4 0.6 0.8 1.0

brf_gmosa (0.17)

(1) brf_gmo_full

brf_gmo_precomputed (0.36)

(0.82) brf_gmo_uniform

brf_gmo_softmax (0.47)

(0.68) brf_gmo_square

