

fit_time
 $p = 3.15e-11$

0.4

0.5

0.6

0.7

0.8

0.9

1.0

brf_gmo_uniform (0.35)

(1) brf_gmo_full

brf_gmo_precomputed (0.44)

(0.83) brf_gmosa

brf_gmo_softmax (0.44)

(0.44) brf_gmo_square

