

fit_time
 $p = 2.28e-11$

0.4

0.5

0.6

0.7

0.8

0.9

1.0

brf_gmo_uniform (0.35)

brf_gmo_precomputed (0.41)

brf_gmo_softmax (0.42)

(1) brf_gmo_full

(0.83) brf_gmosa

(0.49) brf_gmo_square

