fit score time p = 2.32e-31brf gmo precomputed - -NS brf gmo softmax - NS brf gmo square brf_gmo__uniform brf gmo full -*** *** brf gmosa bxt_gmo__precomputed bxt_gmo softmax bxt gmo square bxt_gmo uniform bxt gmo full bxt gmosa brf_gmo__precomputed brf_gmo__softmax brf_gmo__square brf_gmo__uniform brf_gmo_full brf_gmosa oxt_gmo__precomputed bxt_gmo__softmax bxt_gmo_square bxt_gmo_uniform bxt_gmo_ful