fit score time p = 2.28e-34brf gmo -*** brf_gmosa -*** brf gso -*** brf_lmo brf Iso -*** *** brf_sgso -*** *** *** brf sgso us -*** *** bxt_gmosa -*** bxt gso -*** bxt_lmo bxt Iso bxt_sgso -***

bxt sgso us brf_lso brf_gso . brf_lmo . brf_sgso_us brf_sgso bxt_gso bxt_lmo brf_gmo orf_gmosa bxt_gmosa oxt_sgso_us