fit score time p = 3.69e-34brf gmo -*** brf_gmosa -*** brf gso brf_lmo brf Iso -*** brf_sgso -*** NS *** *** *** *** *** brf sgso us bxt_gmosa -*** bxt gso -*** bxt_lmo bxt Iso -NS bxt_sgso -***

bxt sgso us brf_lmo . brf_Iso . brf_gmo brf_gso bxt_gso bxt_Imo brf_sgso orf_gmosa brf_sgso_us bxt_gmosa

bxt_sgso_us