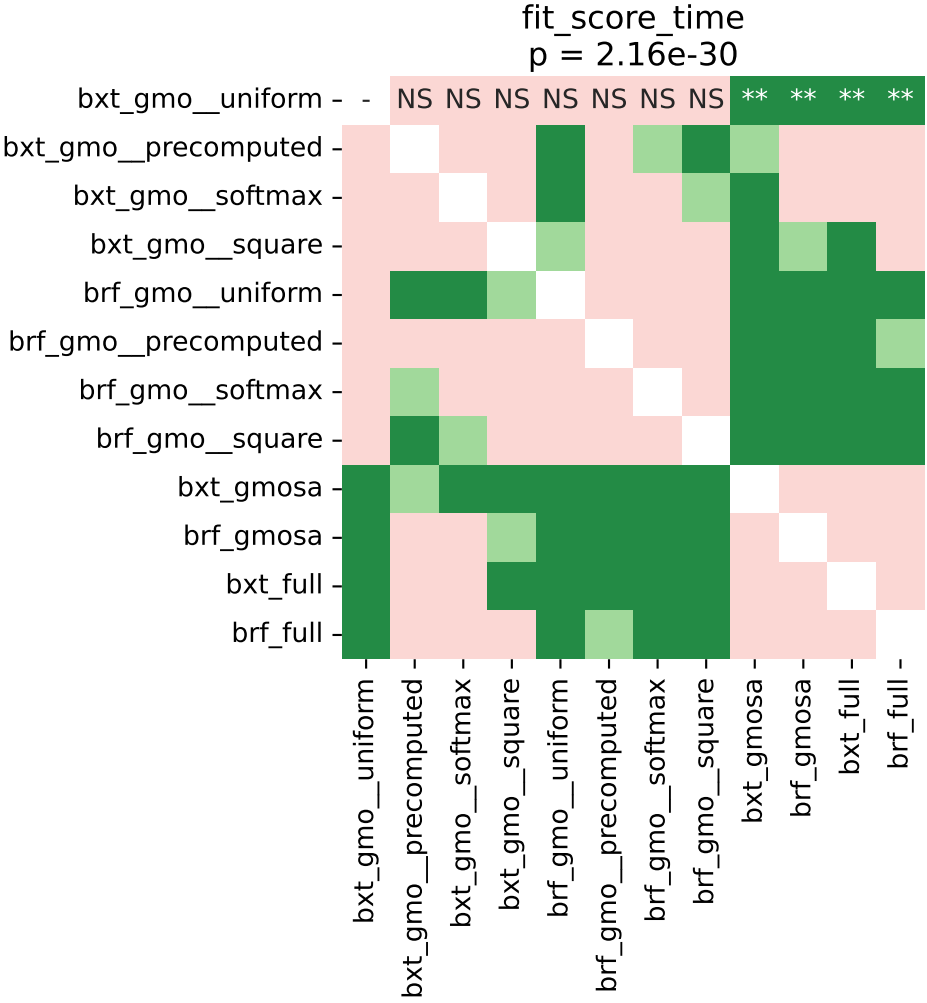
fit_score_time
p = 2.16e-30
bxt_gmo__uniform
bxt_gmo__precomputed
bxt_gmo__softmax
bxt_gmo__square
brf_gmo__uniform
brf_gmo__precomputed
brf_gmo__softmax
brf_gmo__square
bxt_gmosa
brf_gmosa
bxt_full
brf_full
- NS NS NS NS NS NS NS ** ** ** **