Number of binding positions in top 10 predicted – dnabackbone method hits fraction in top predicted
0
.0 aa_ref_overlap_individuals_change_ratio instances_individuals_change_ratio jsd_100way_aa_not_used_ratio jsd_100way_instances_major_ratio jsd_mul_aa_ref_SE jsd_SE_diff_ratio jsds_ratio jsds_subtraction SE_jsd_diff_ratio 0.0 measurment