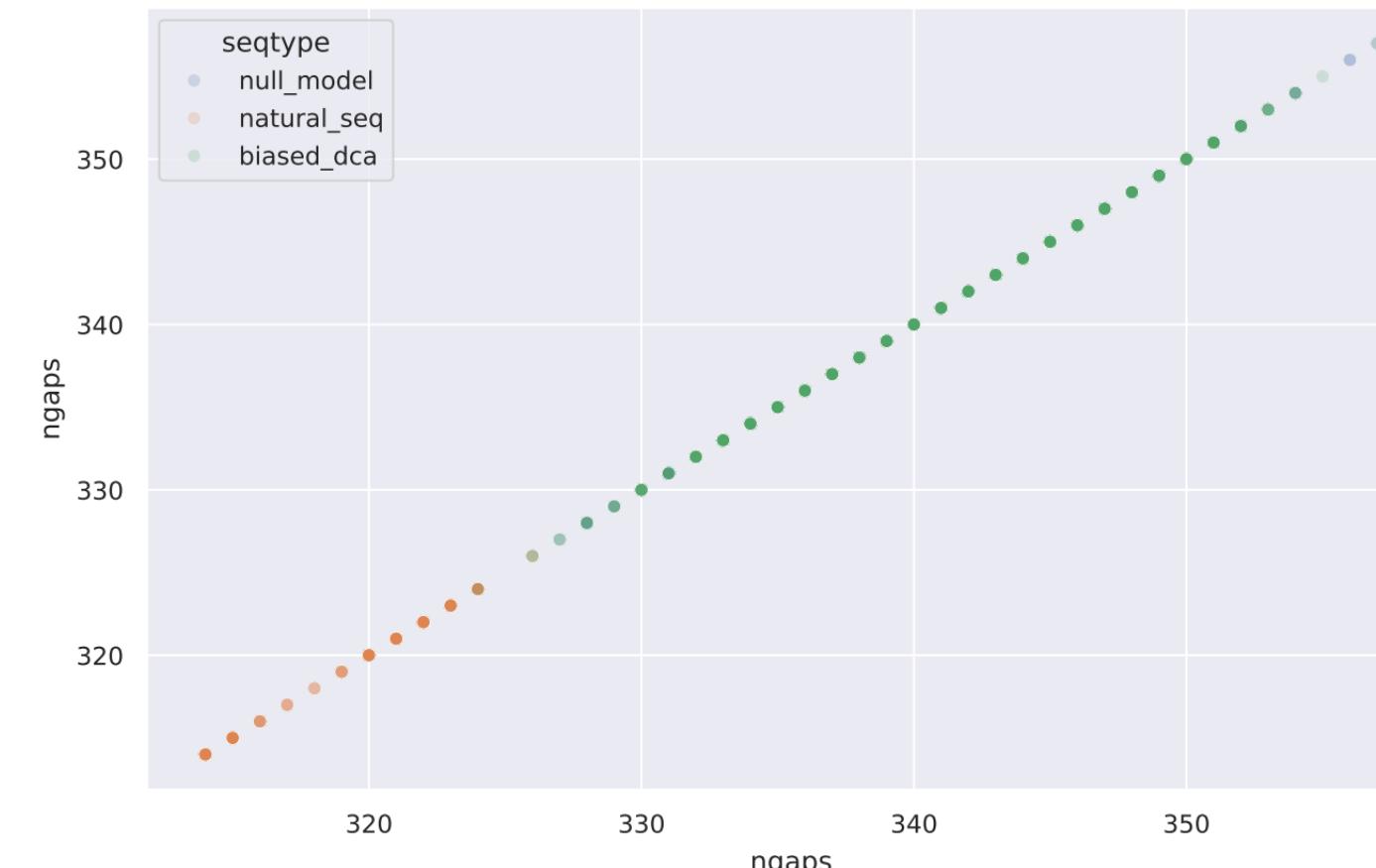
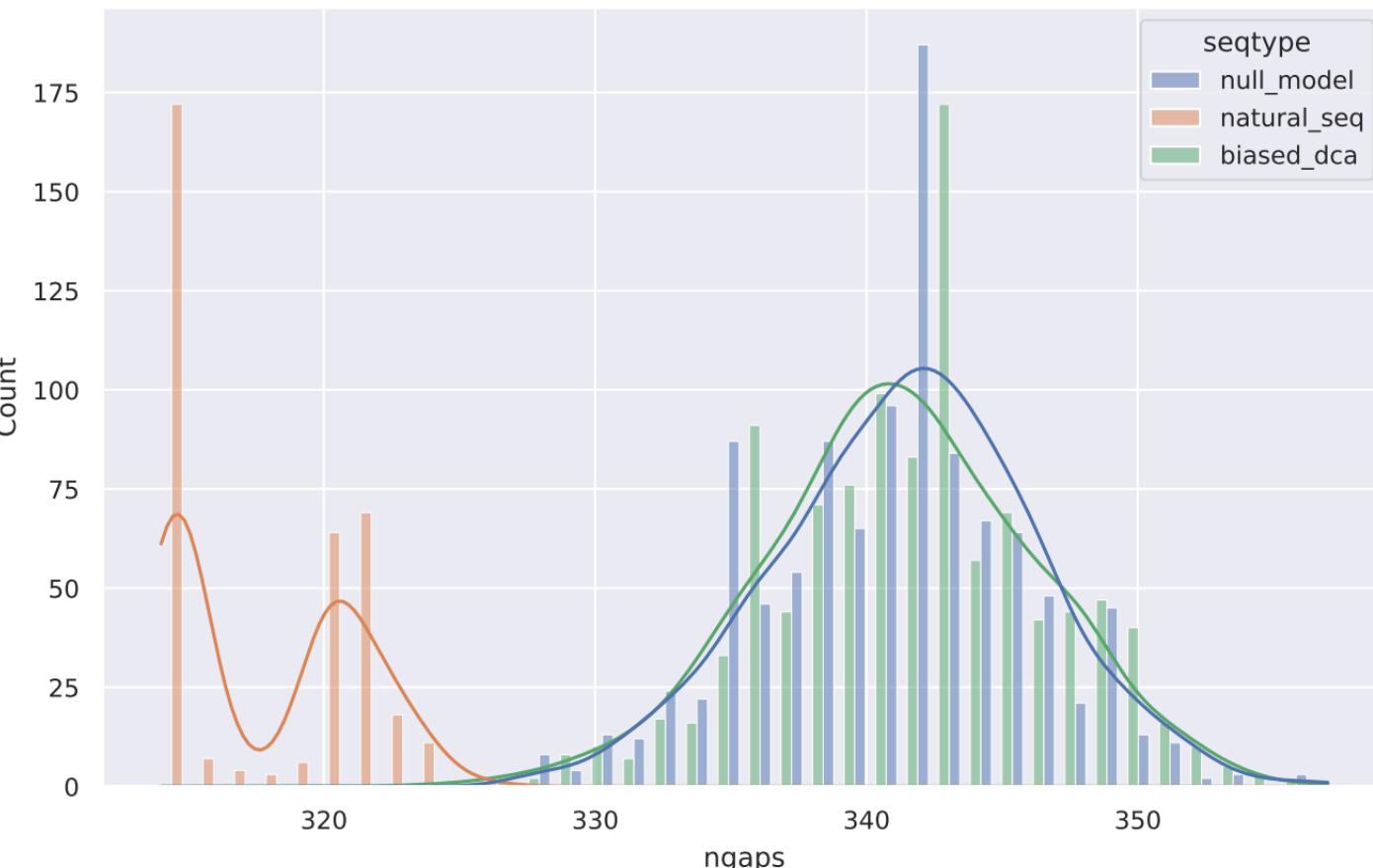
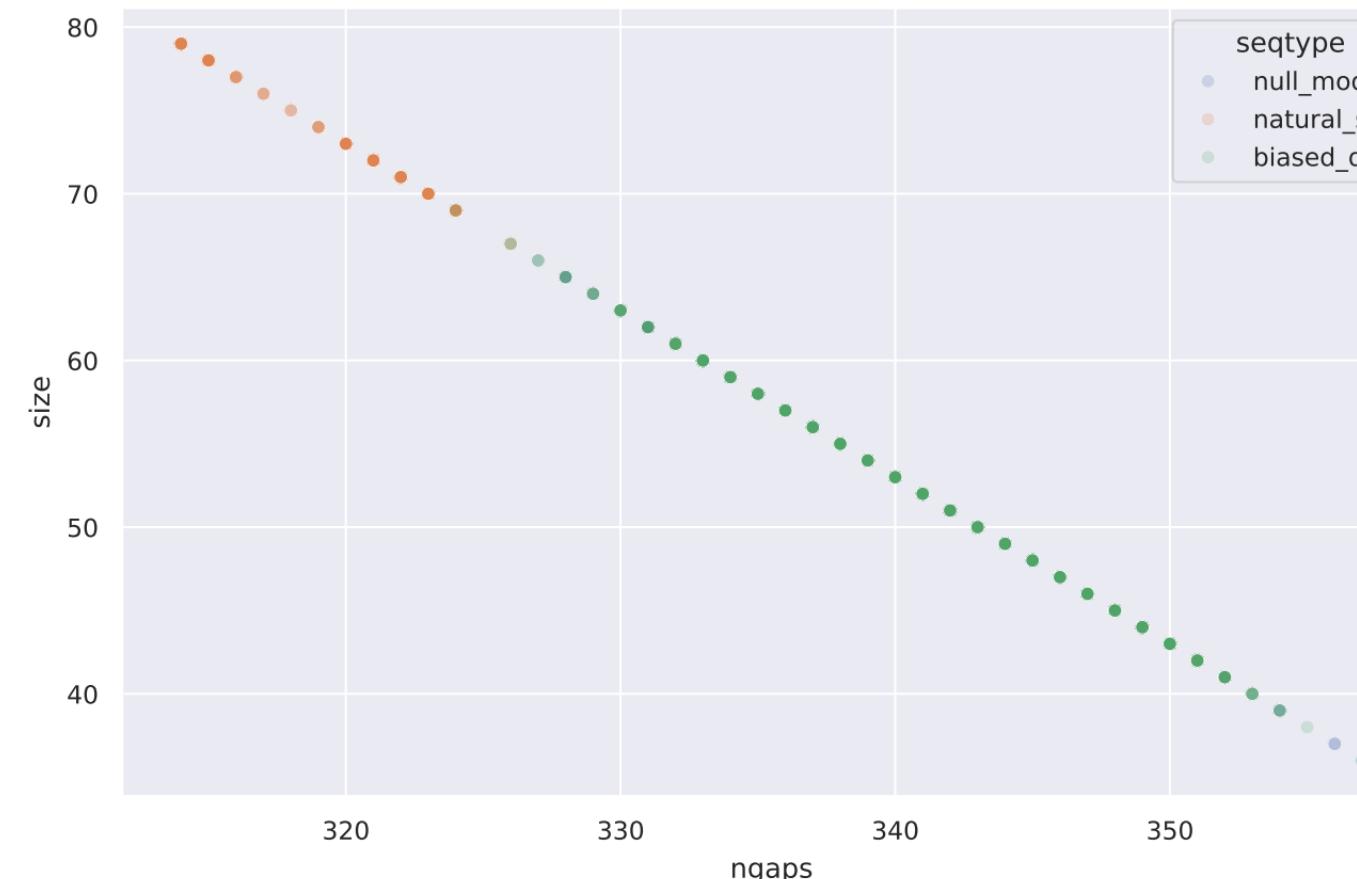
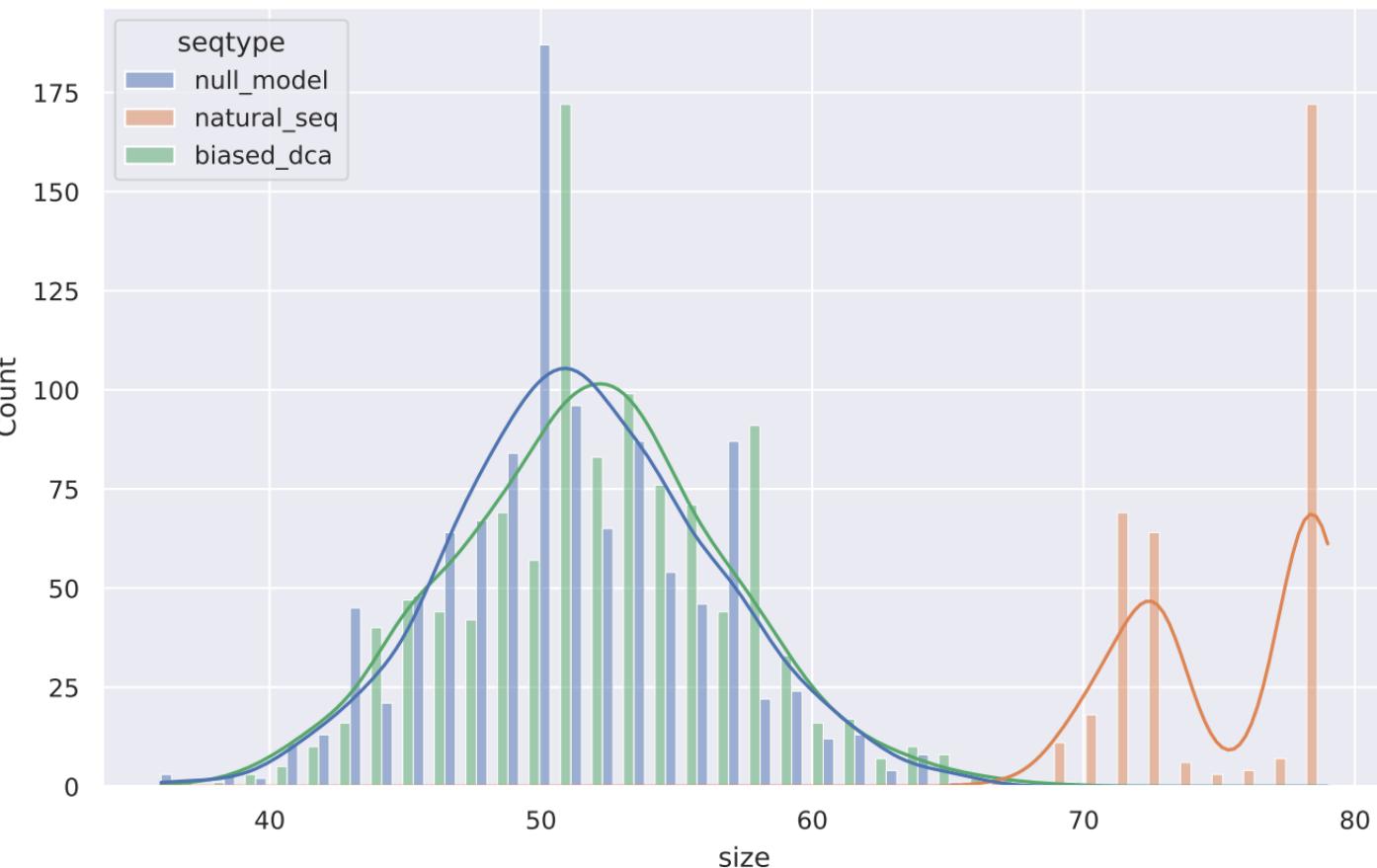


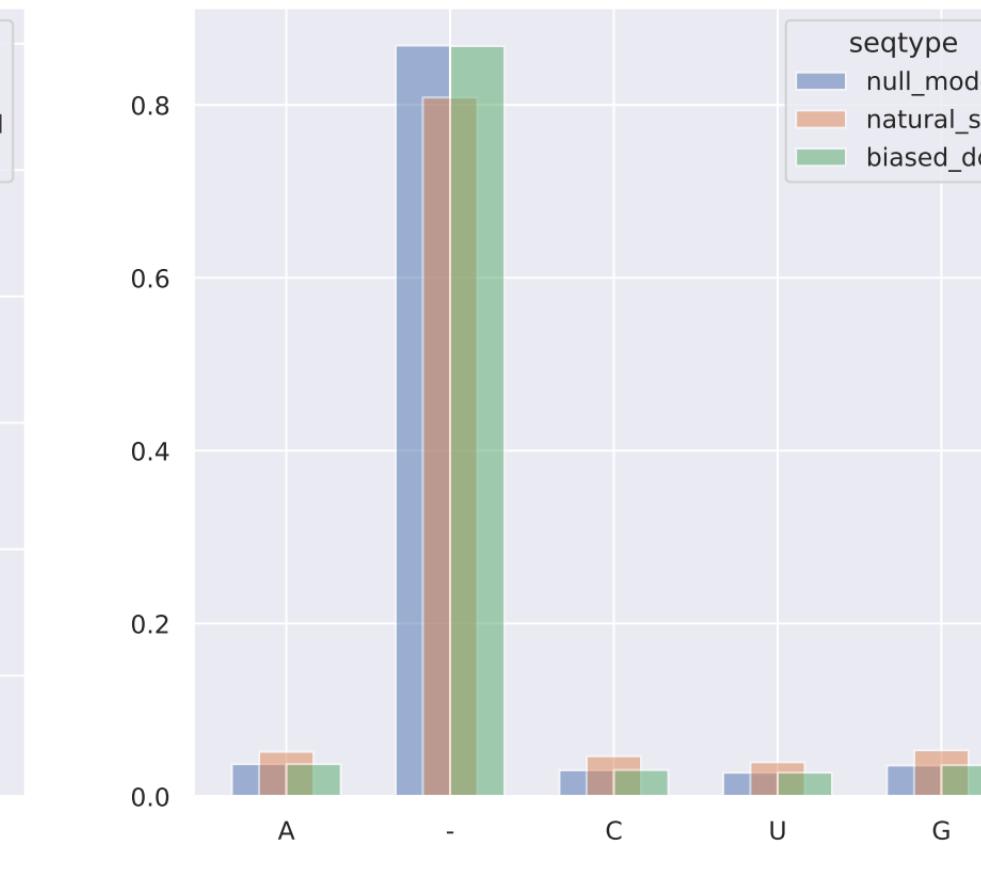
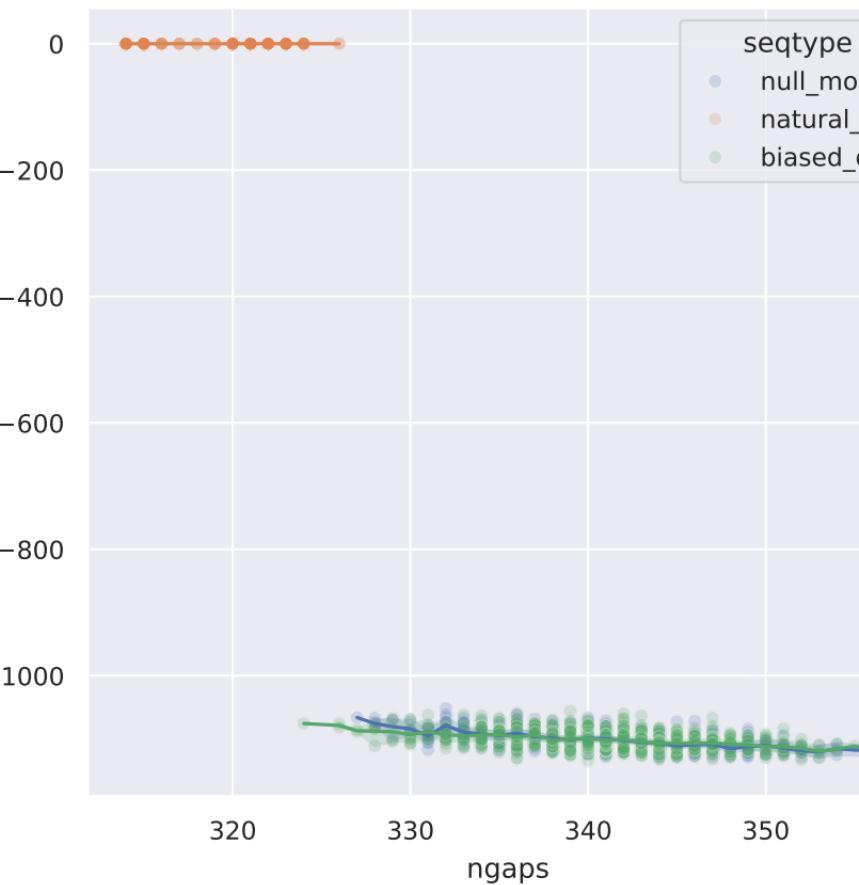
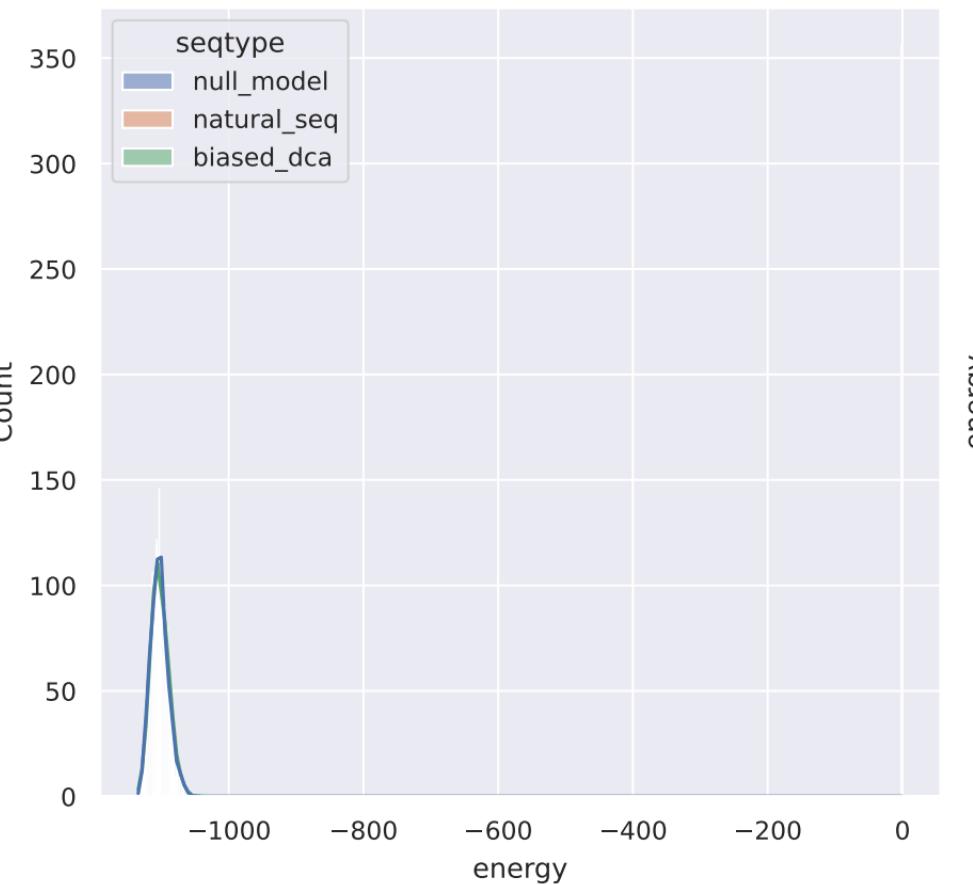
Numeric Feature : ngaps



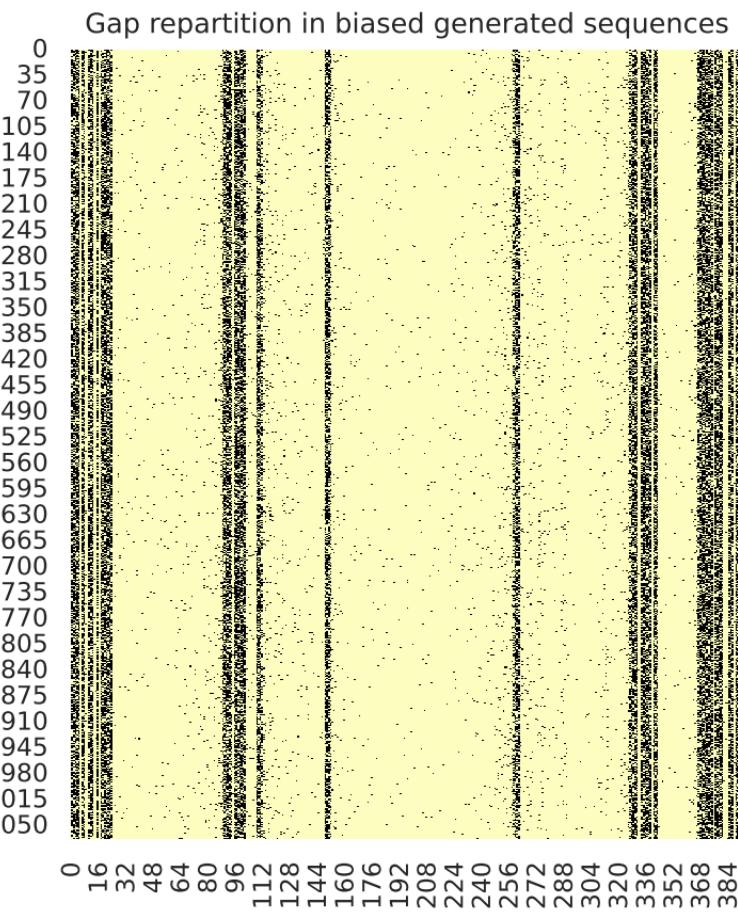
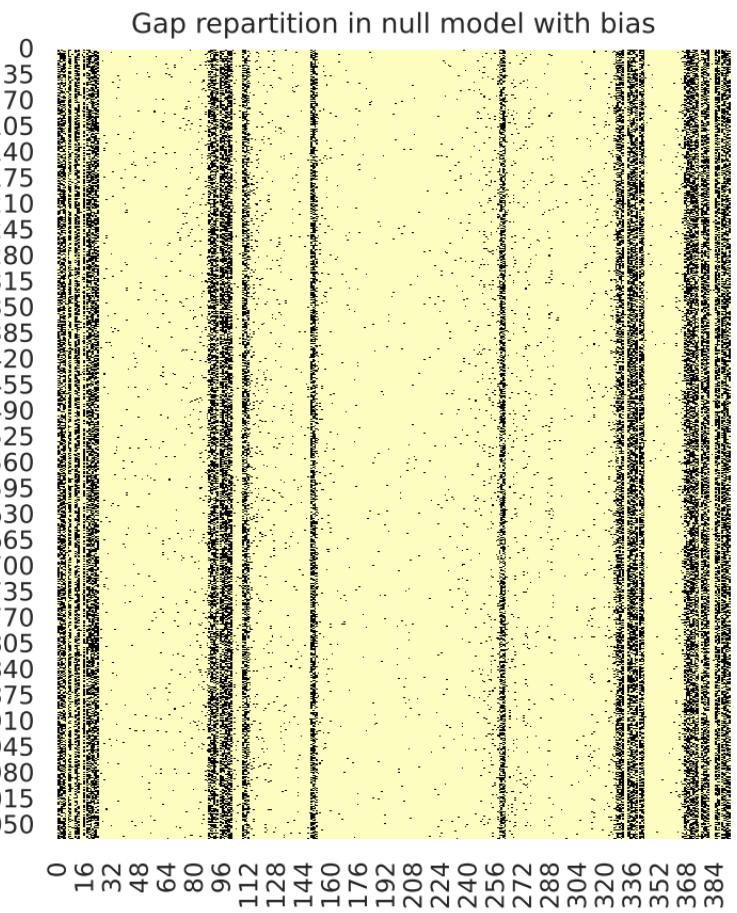
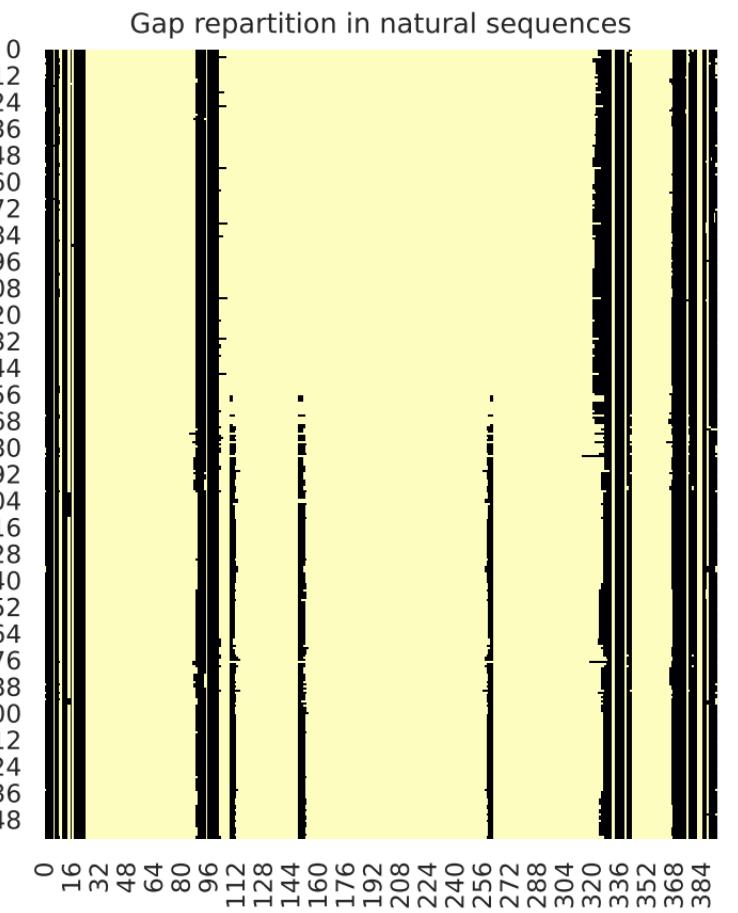
Numeric Feature : size



Numeric Feature : Sequences Energy and Nucleic acid average frequency



Comparison of gap reparation in generated MSA respect to the null model and natural short sequences



1.0
0.8
0.6
0.4
0.2
0.0