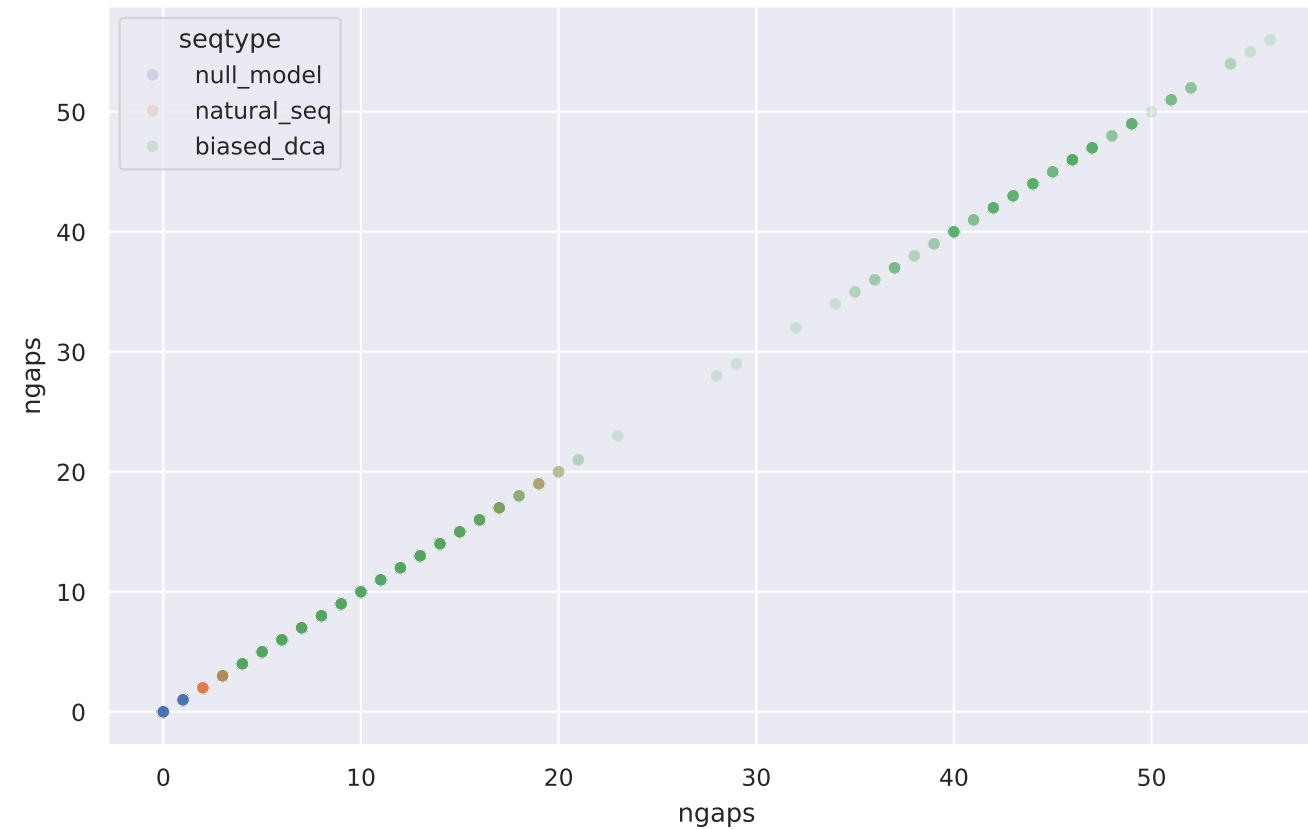
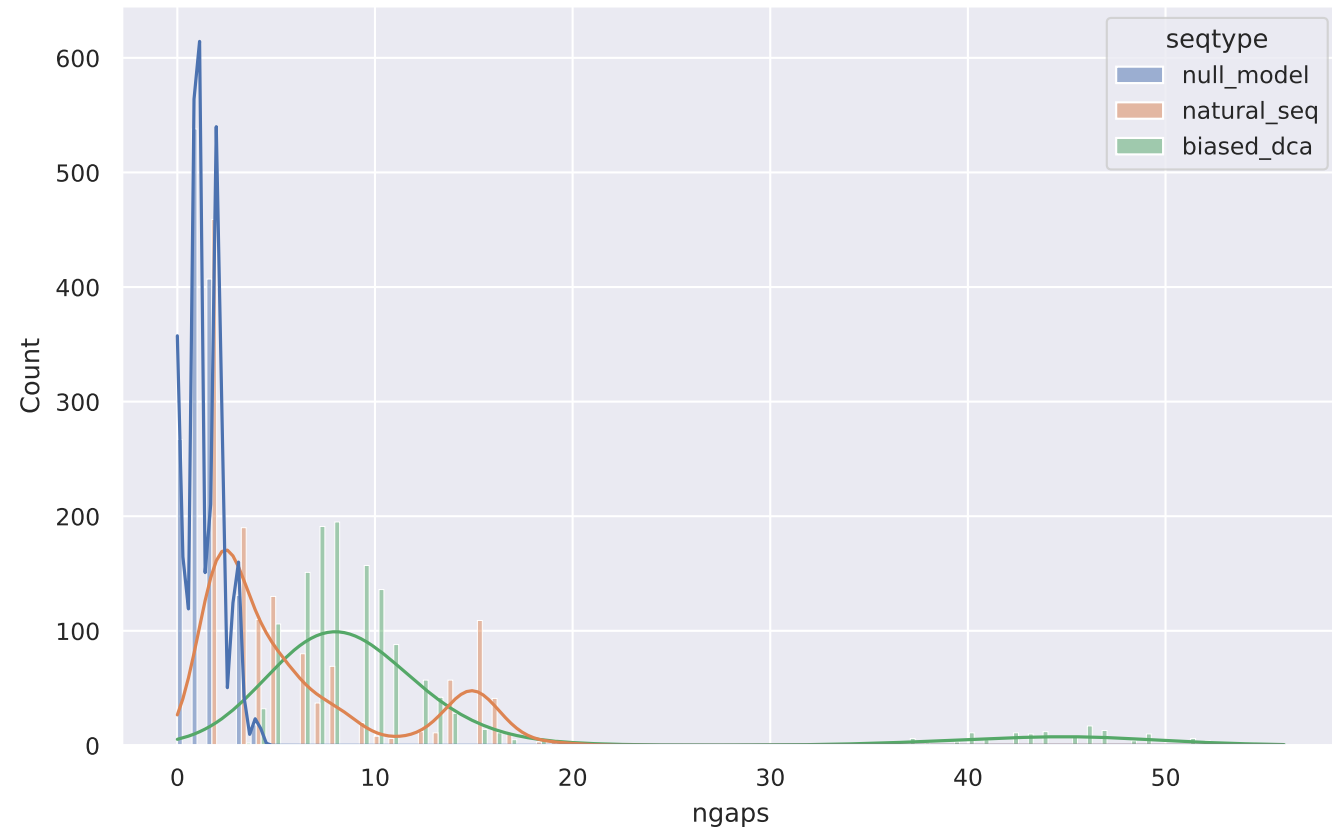
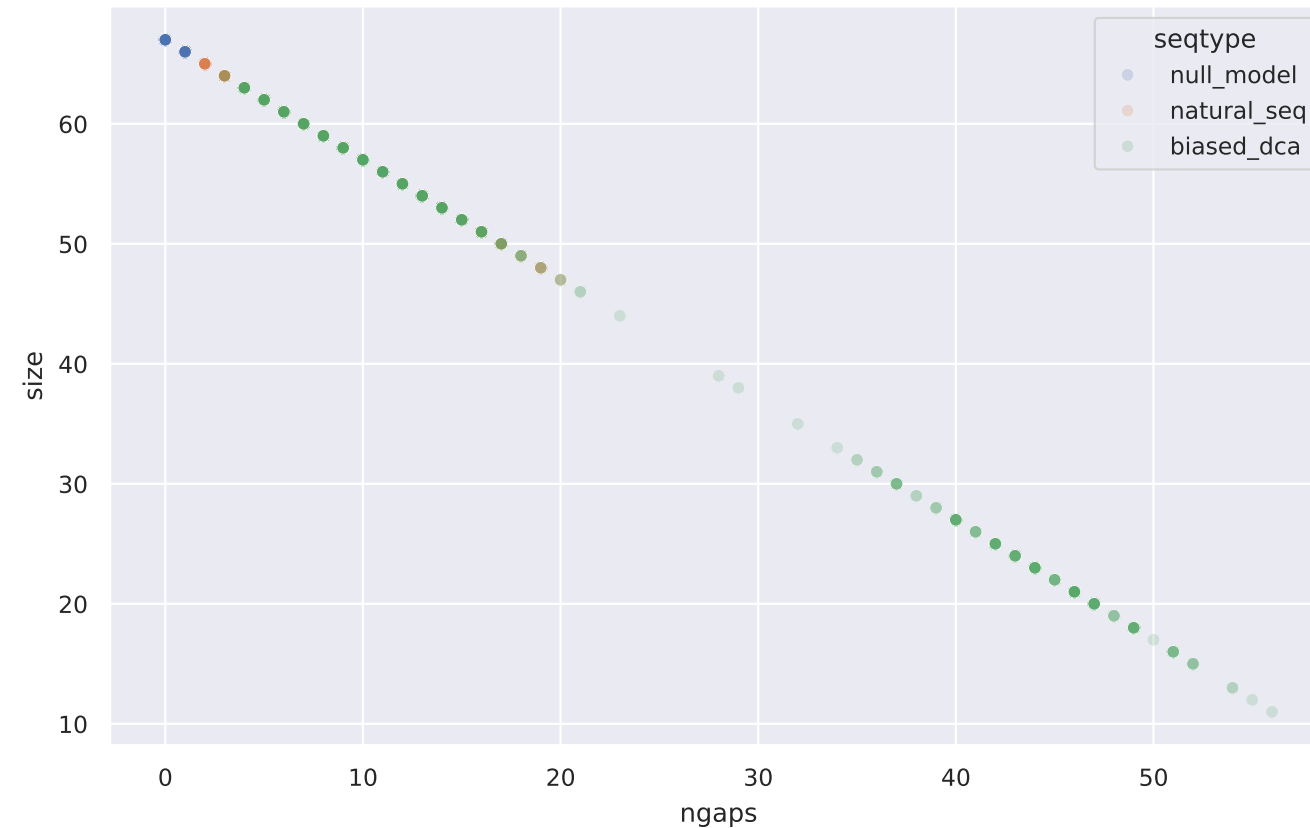
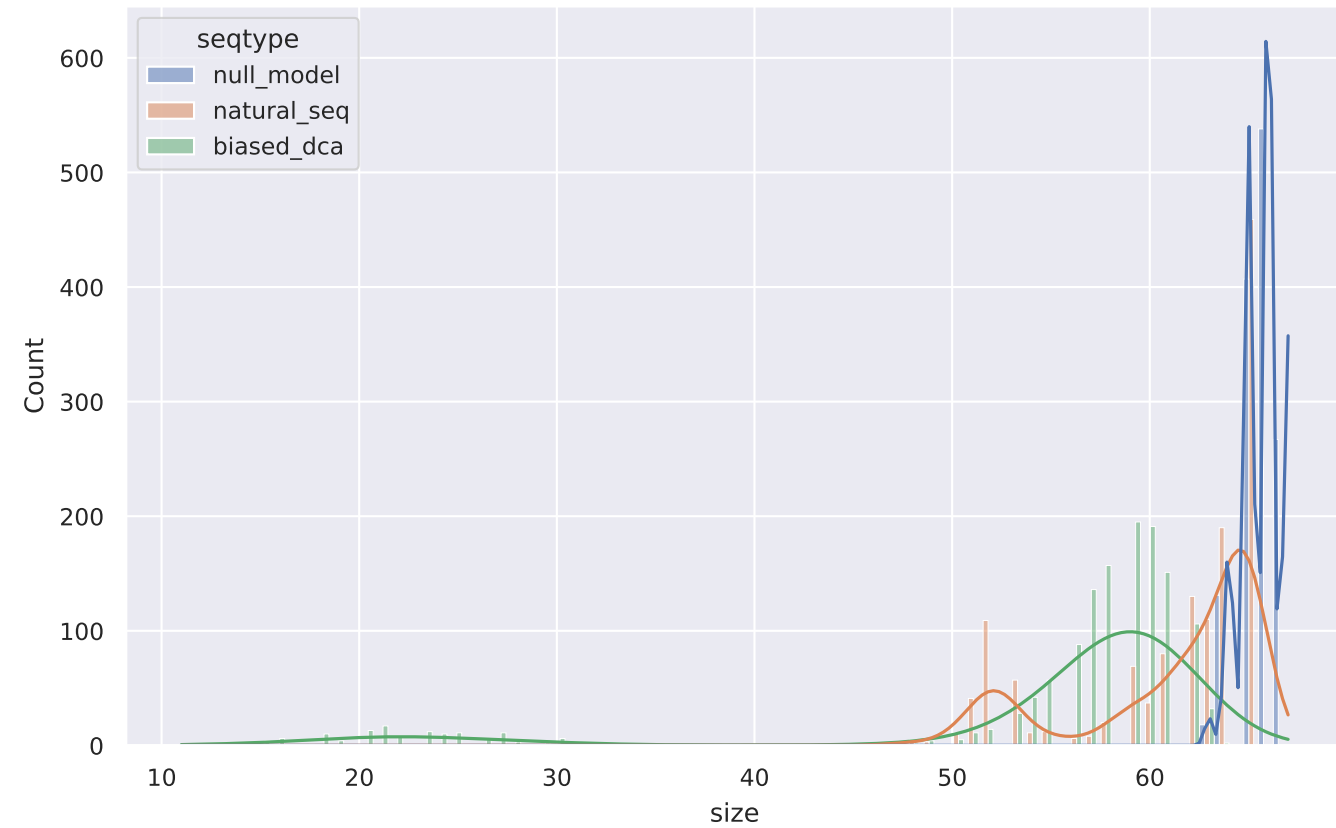


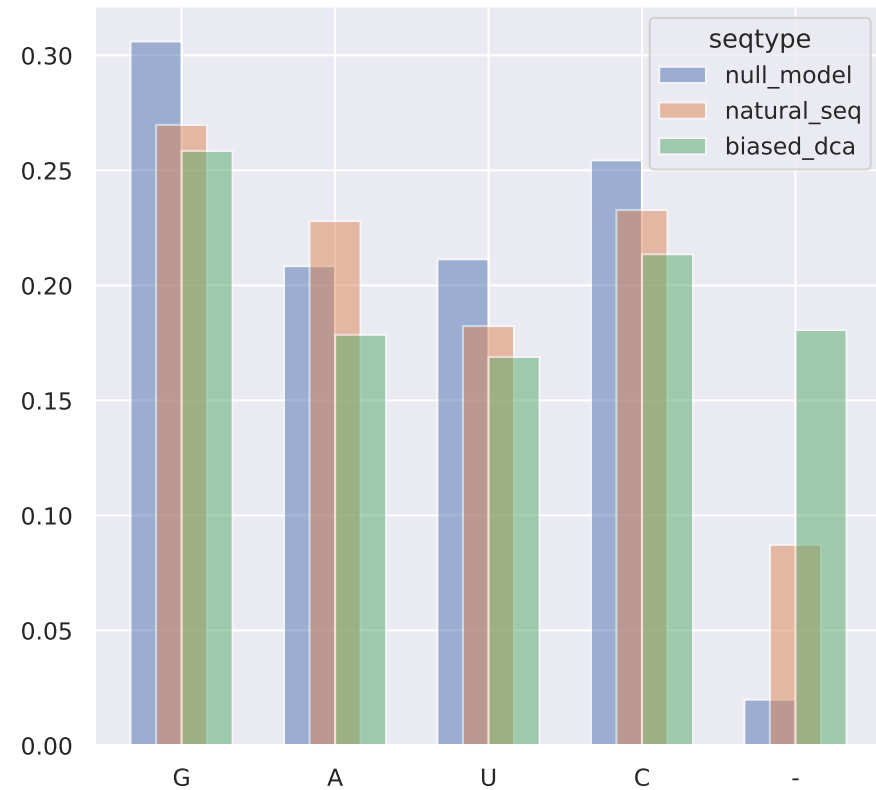
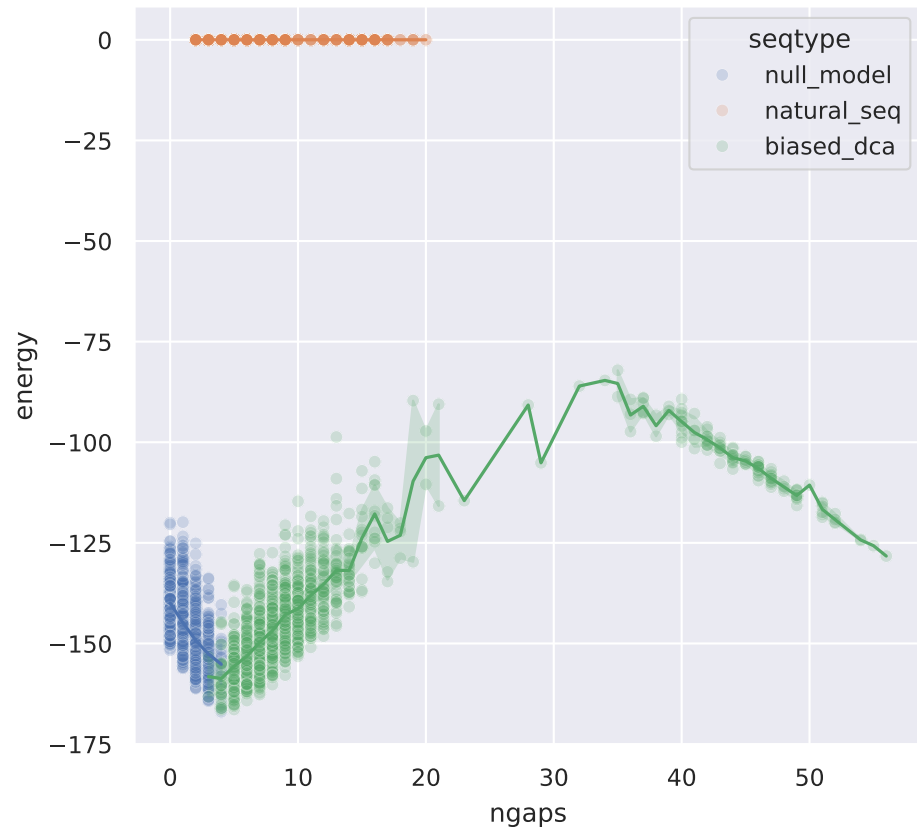
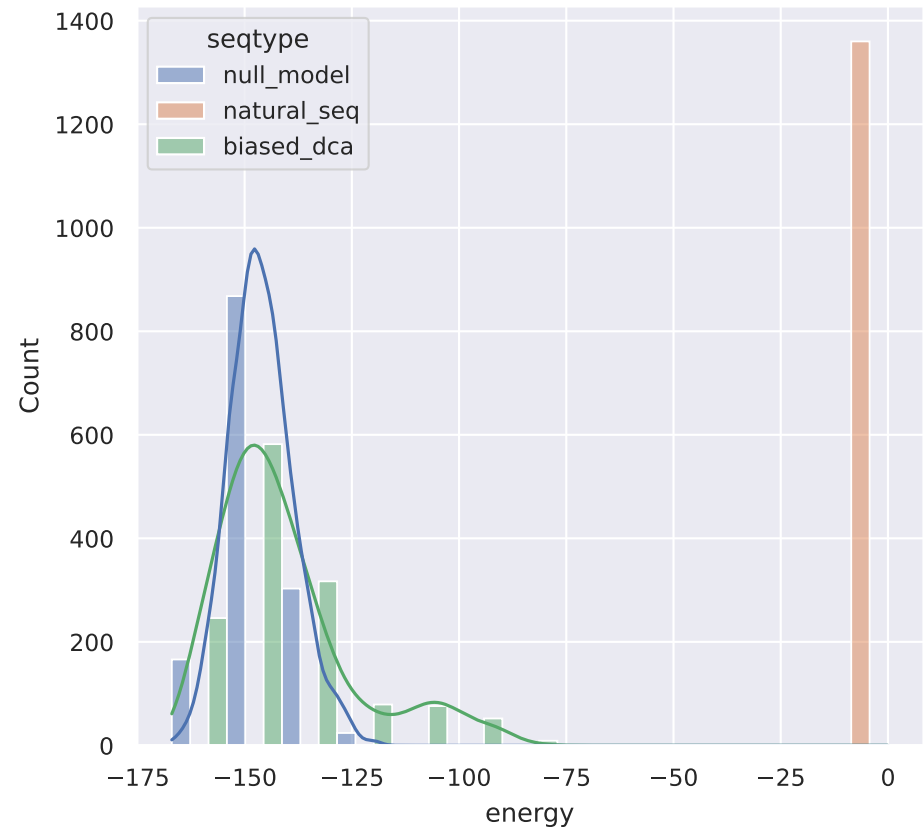
Numeric Feature : ngaps



Numeric Feature : size



Numeric Feature : Sequences Energy and Nucleic acid average frequency



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

