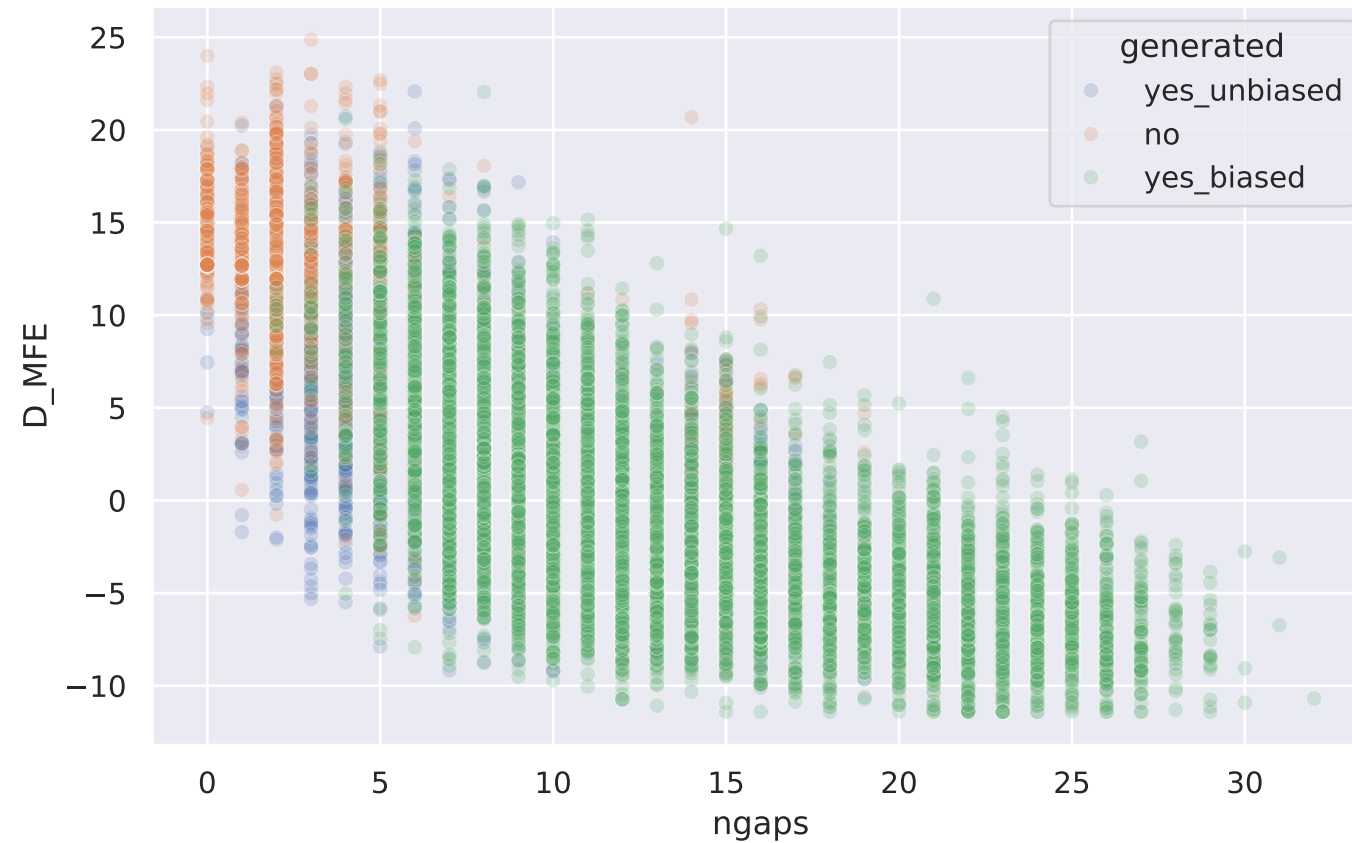
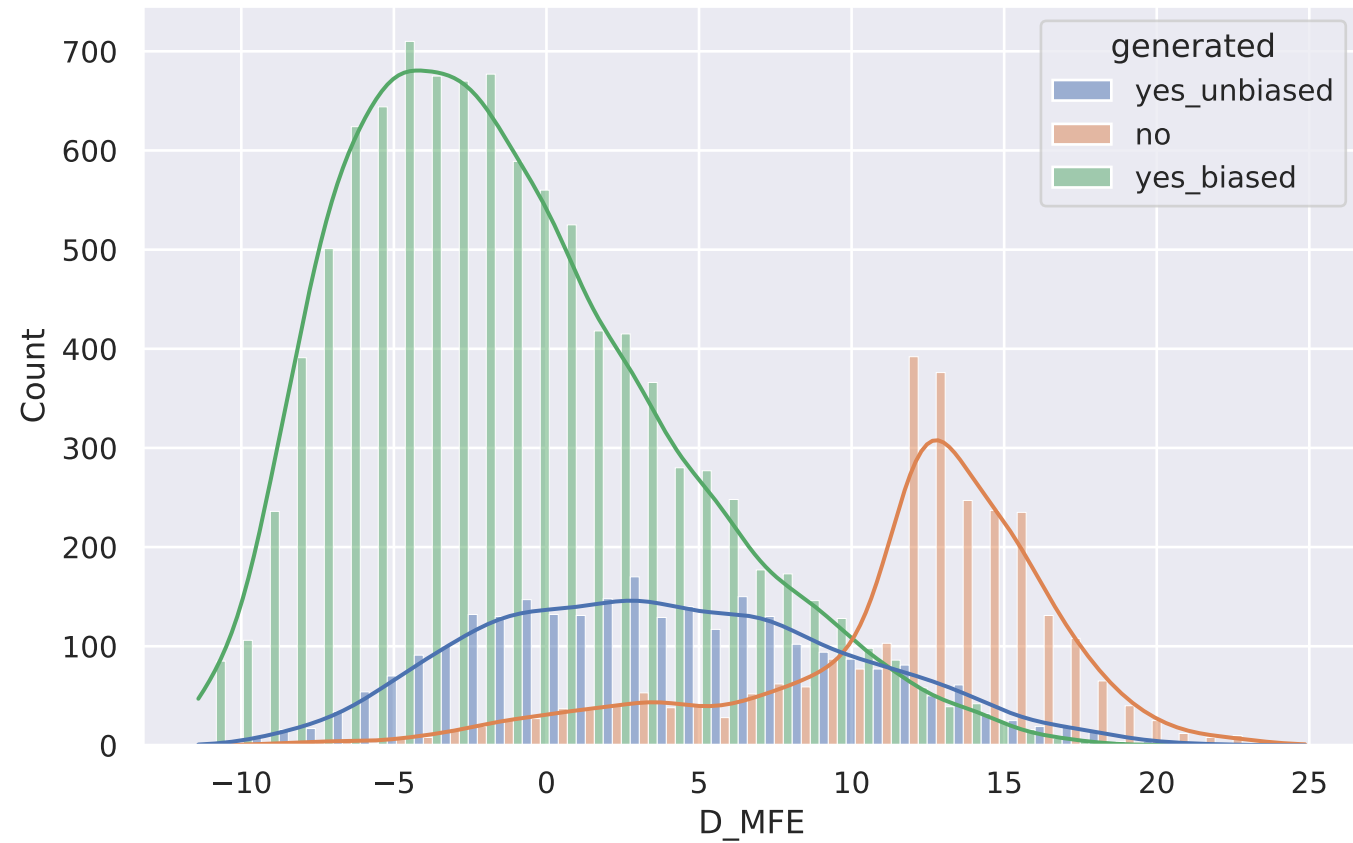
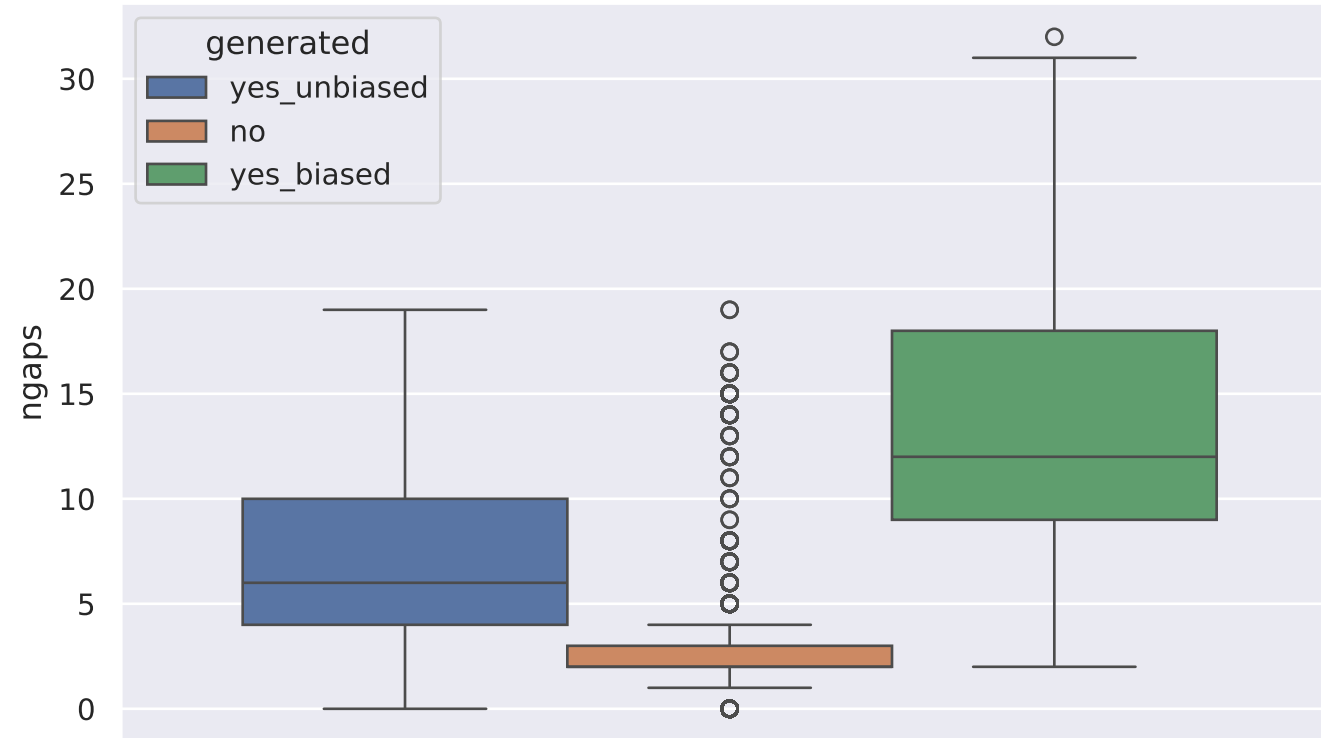
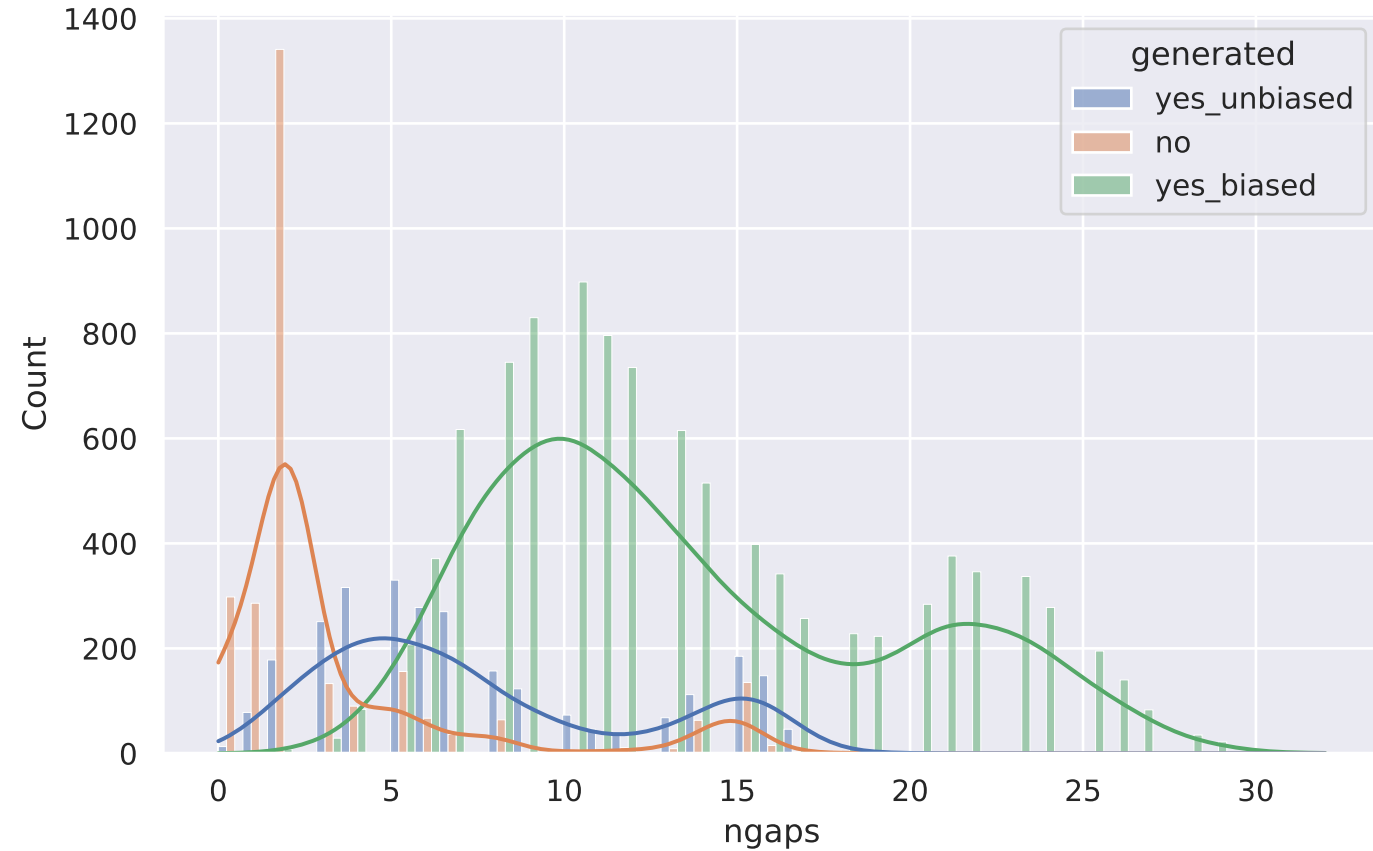


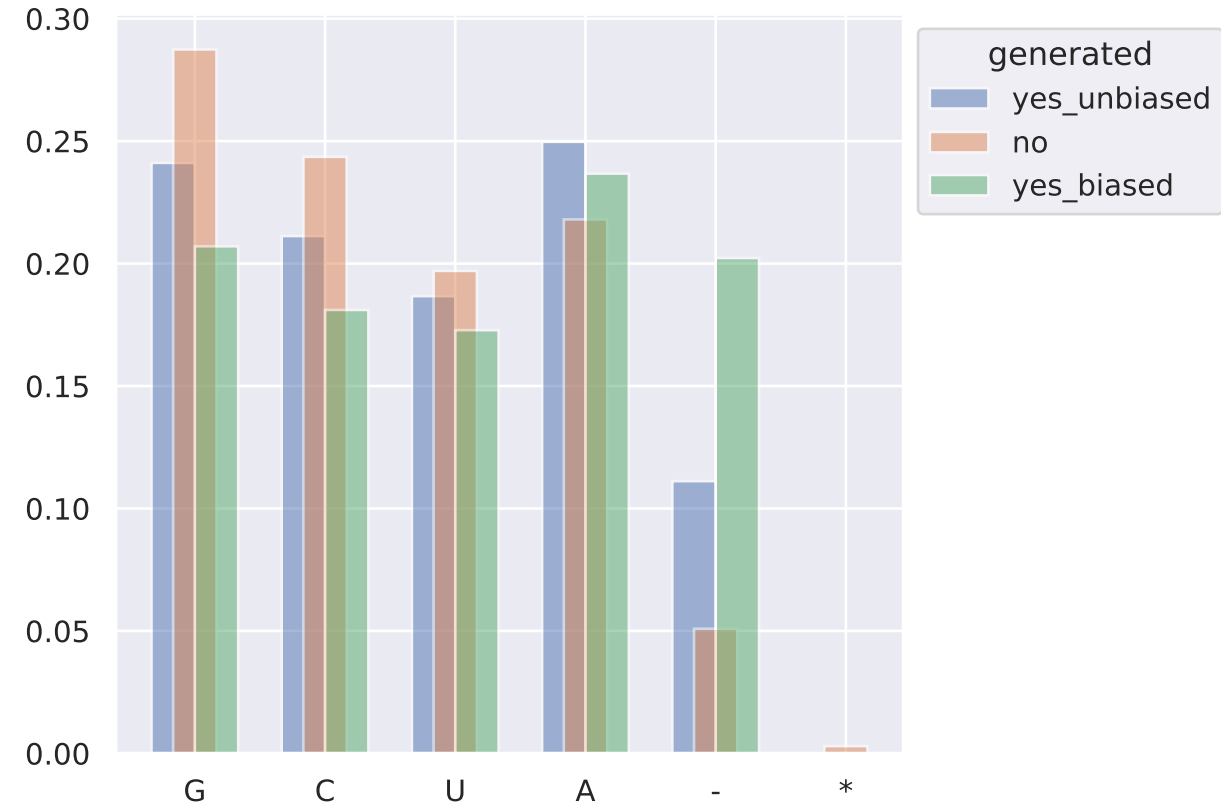
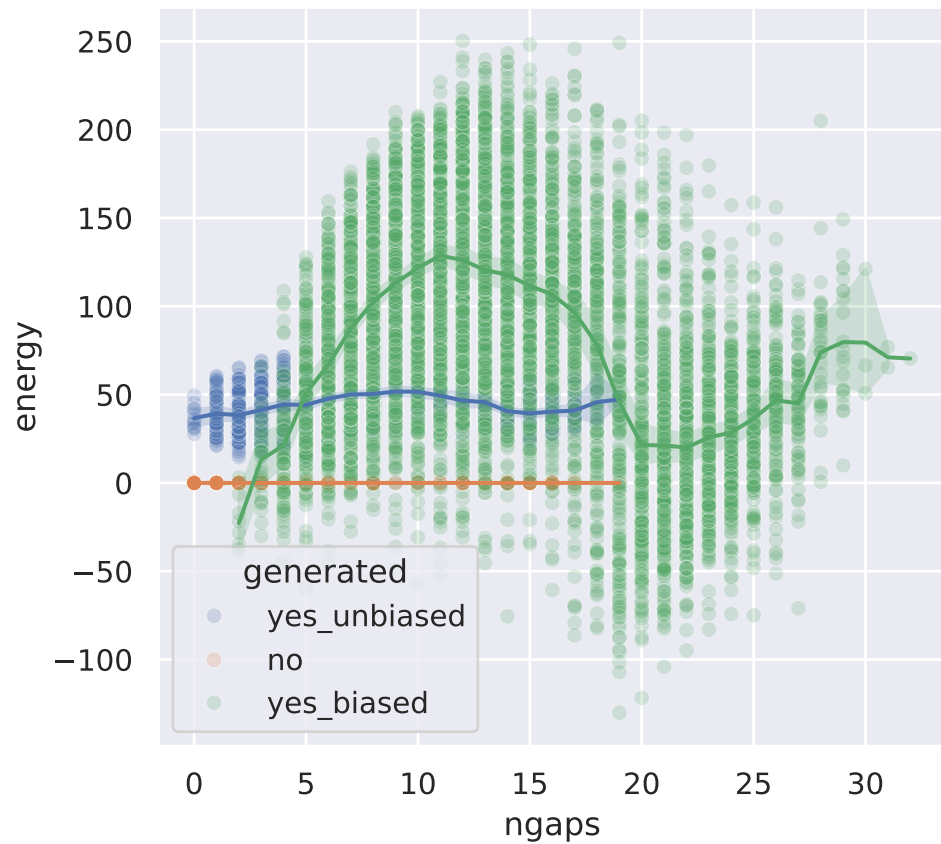
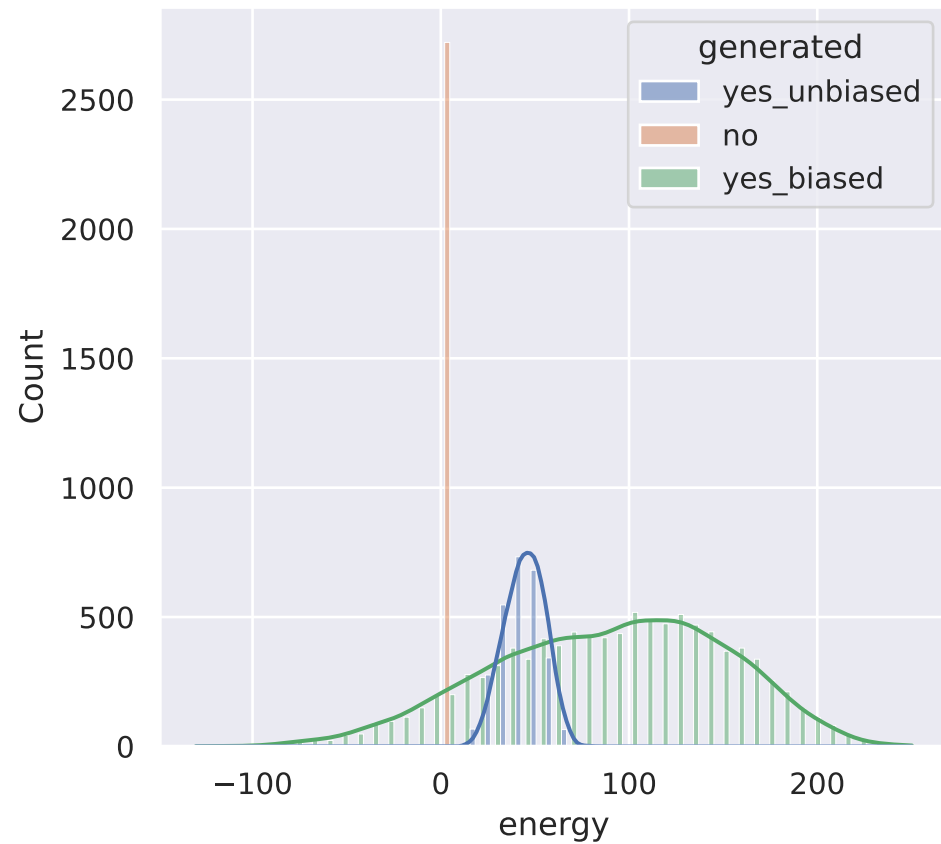
Numeric Feature : D_MFE



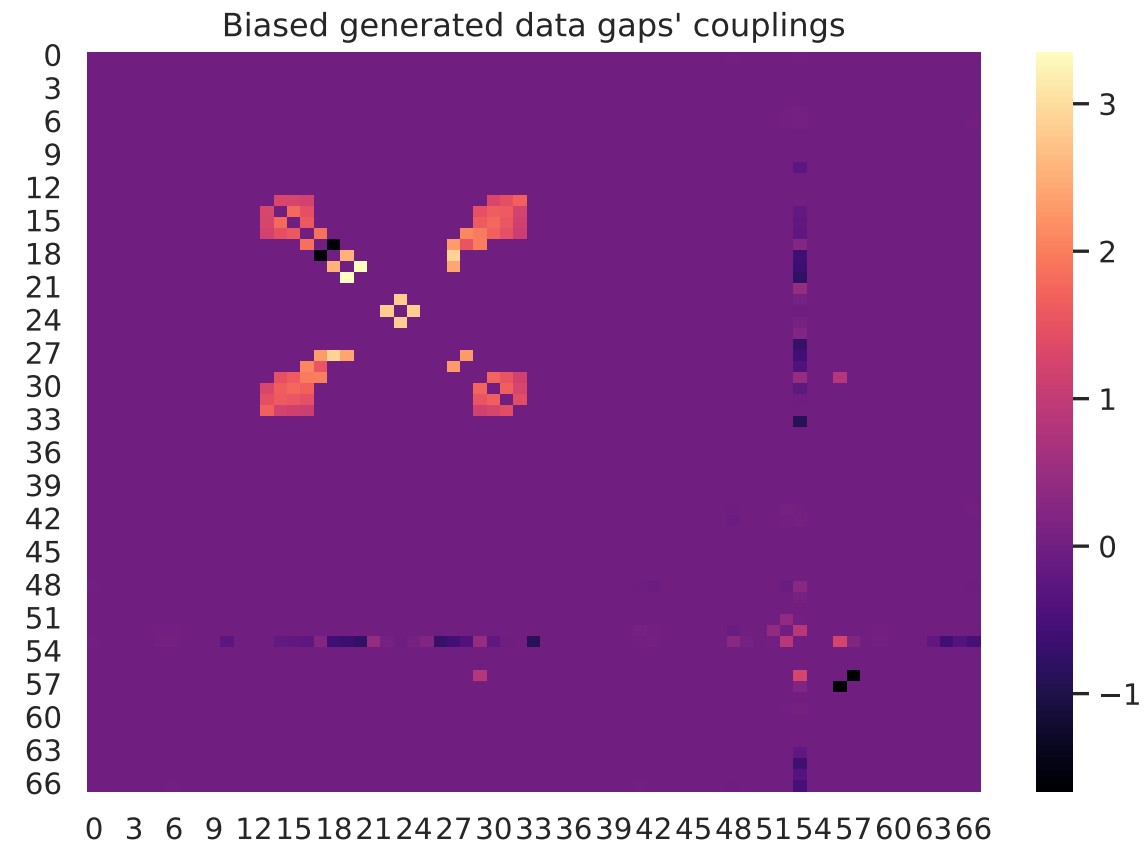
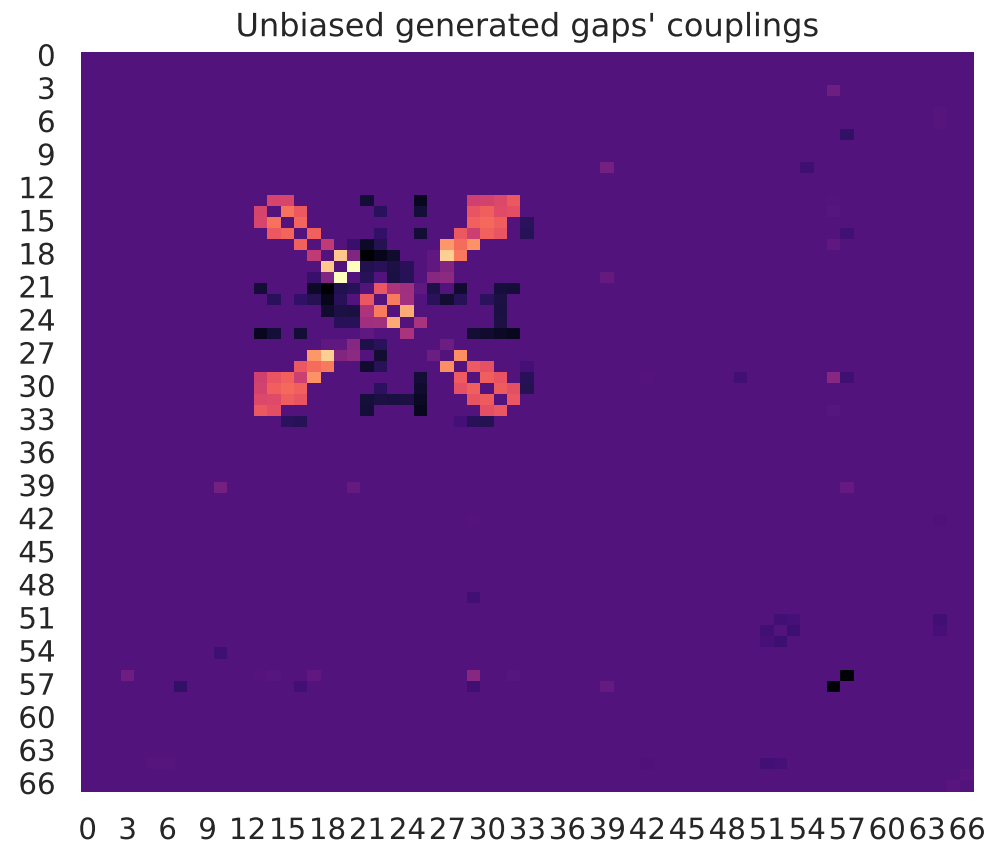
Numeric Feature : ngaps



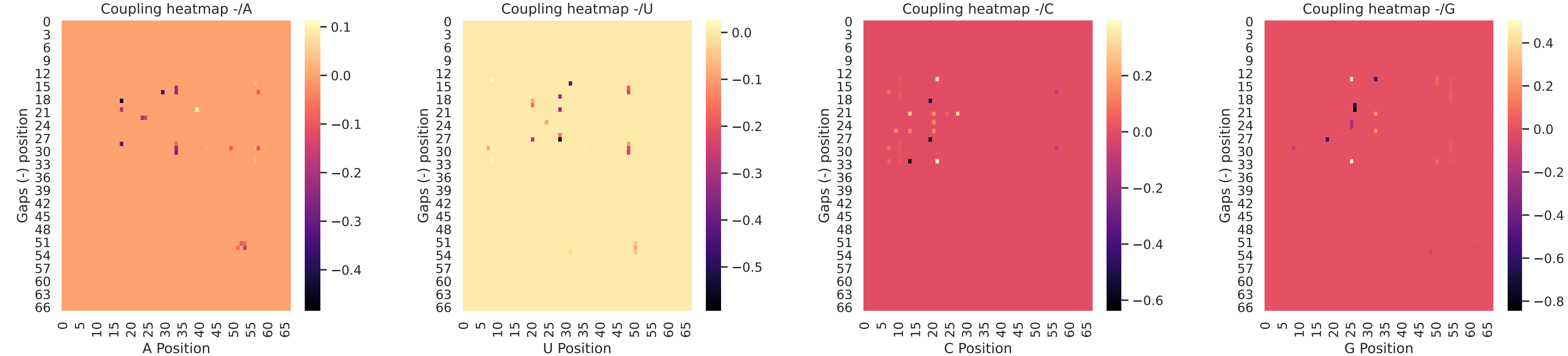
Numeric Feature : Sequences Energy and Nucleic acid average frequency



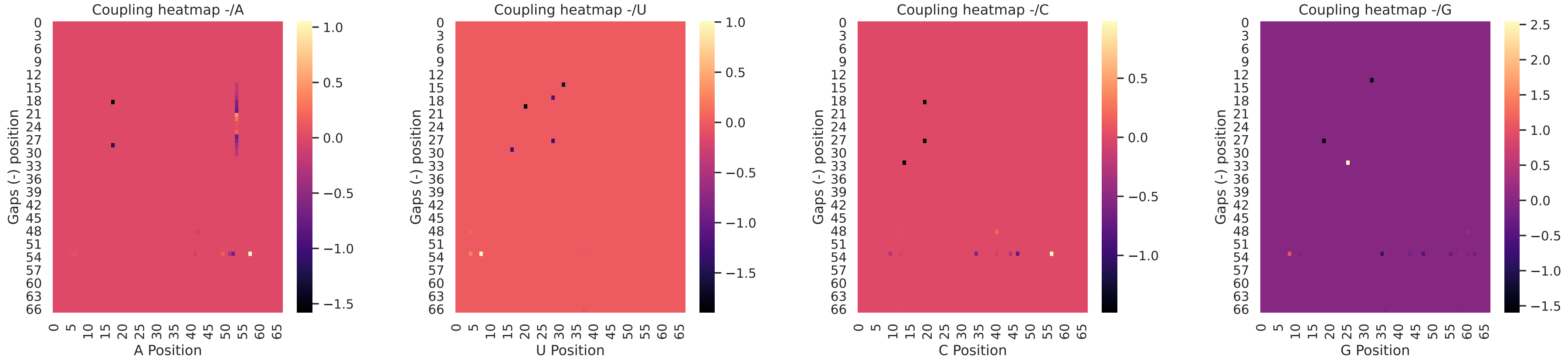
Couplings heatmap for -/-



Couplings heatmap for Unbiased

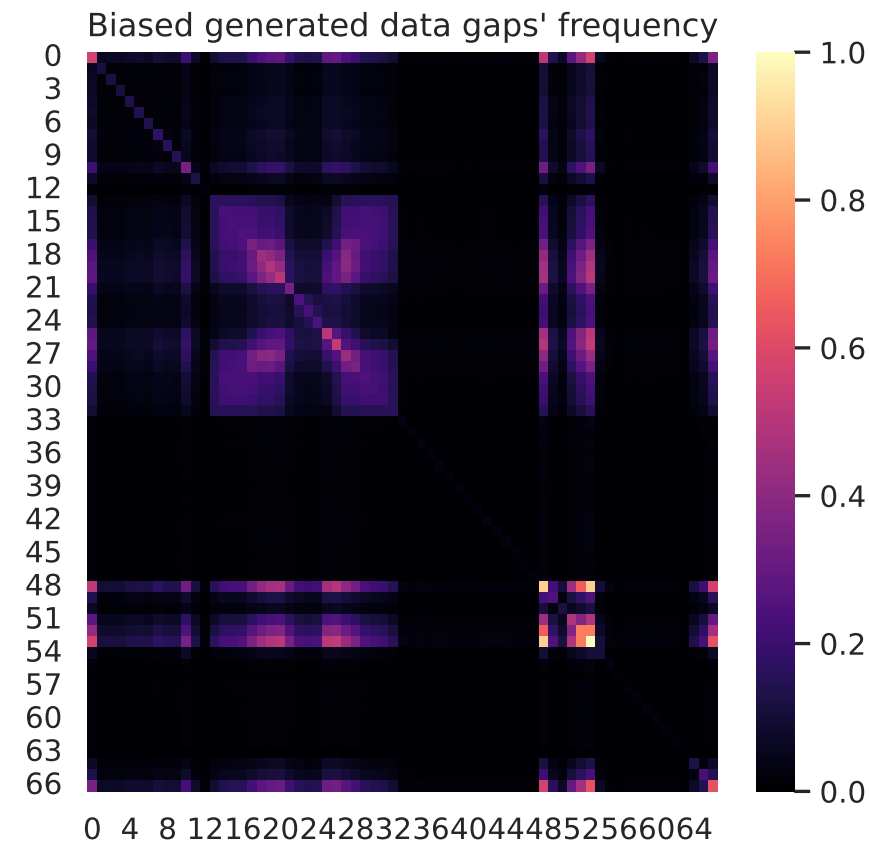
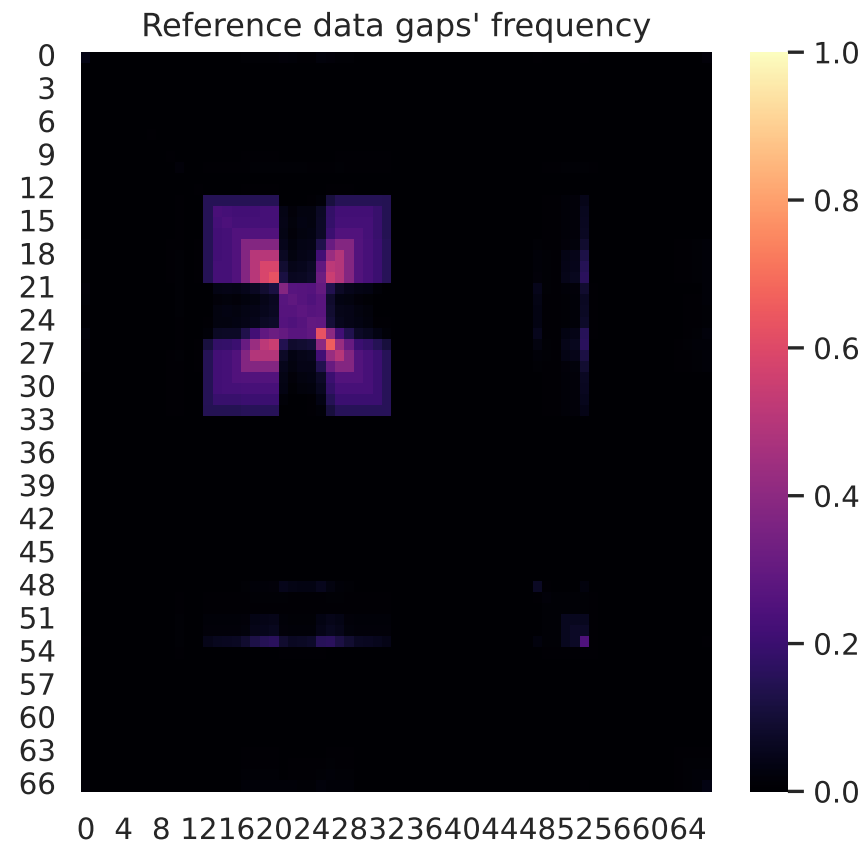
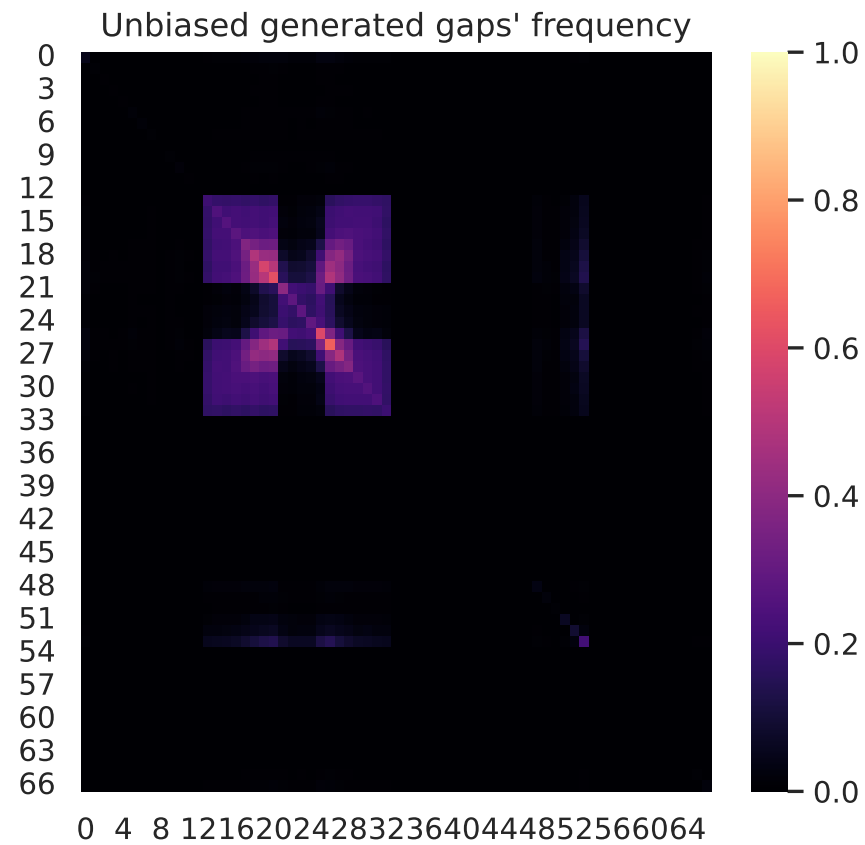


Couplings heatmap for biased

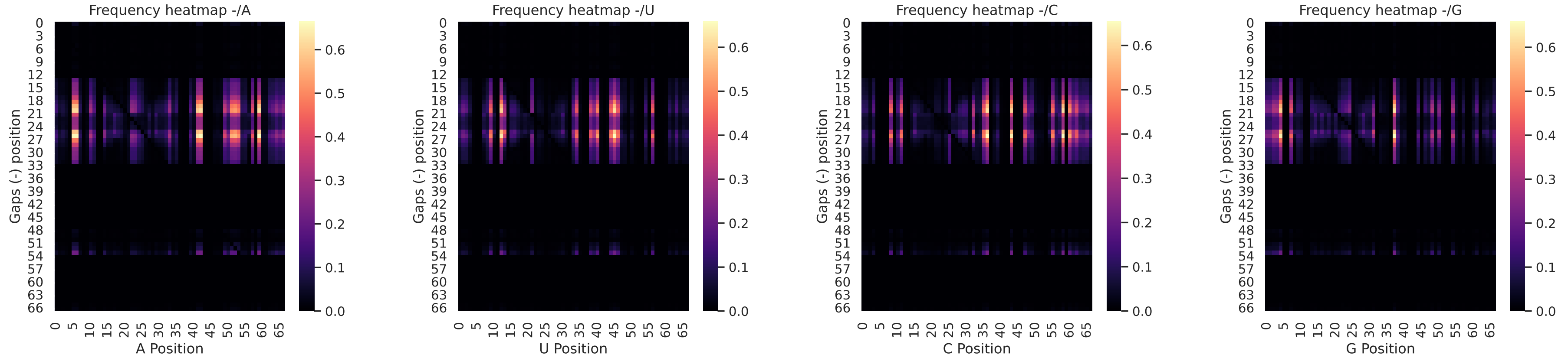


Gaps' frequency depending on position in the MSA's sequences

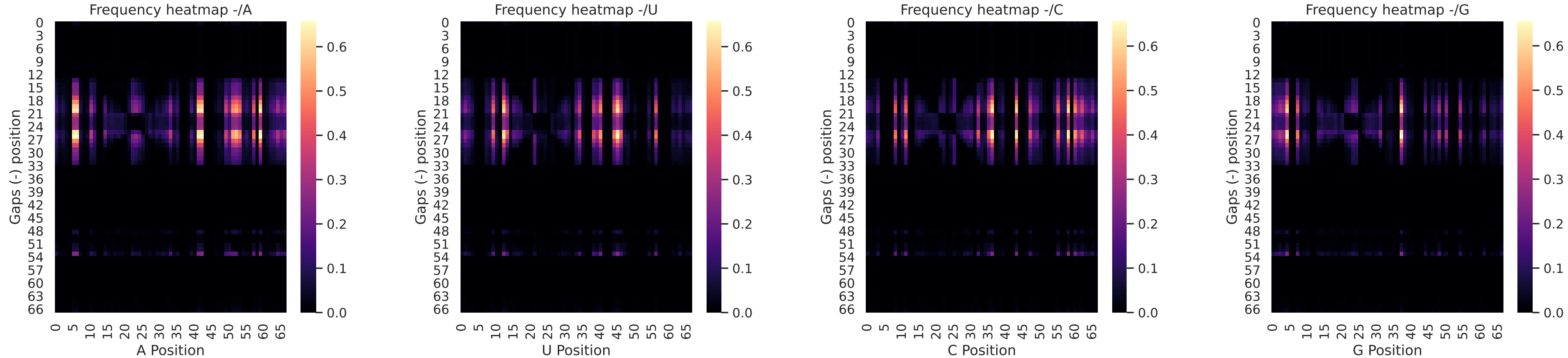




Frequency heatmap for Unbiased



Frequency heatmap for ref



Frequency heatmap for biased

