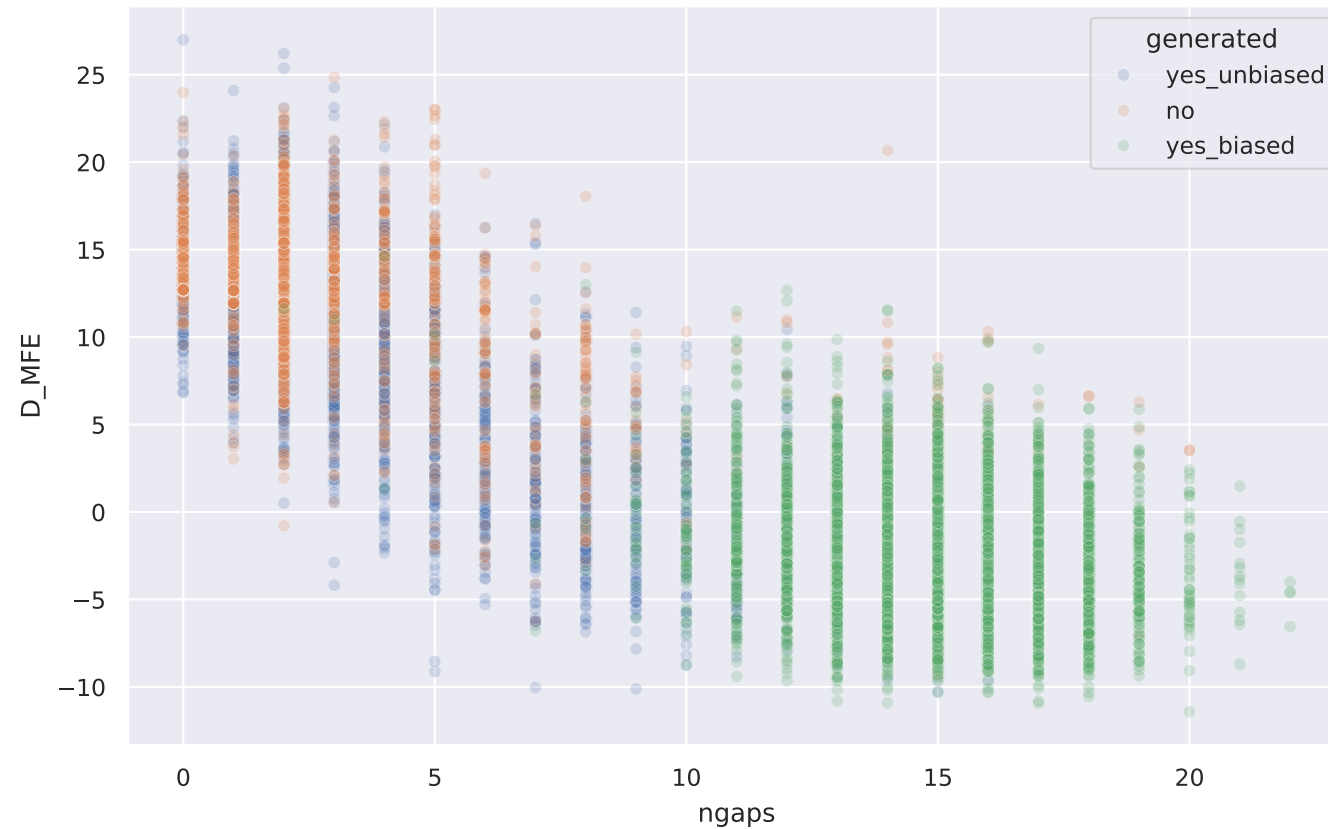
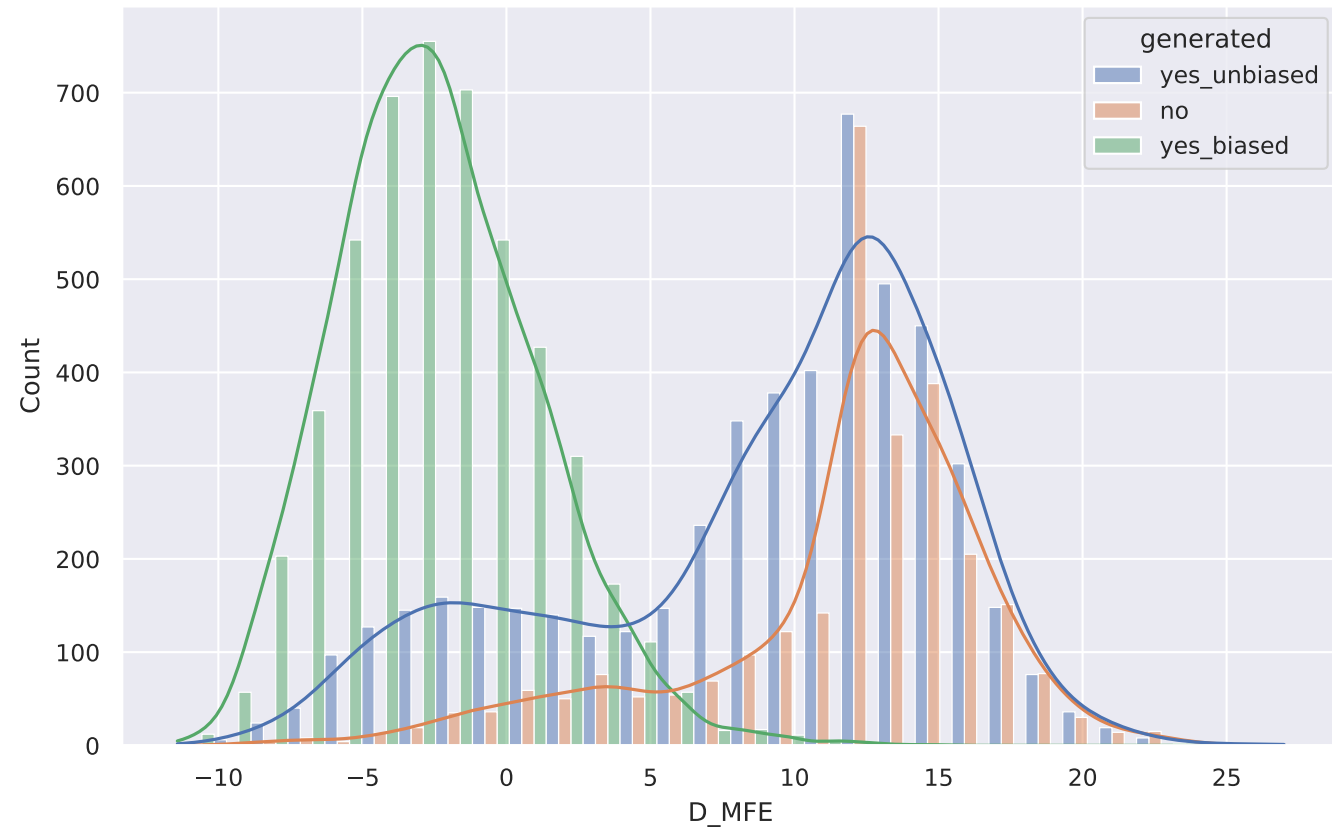
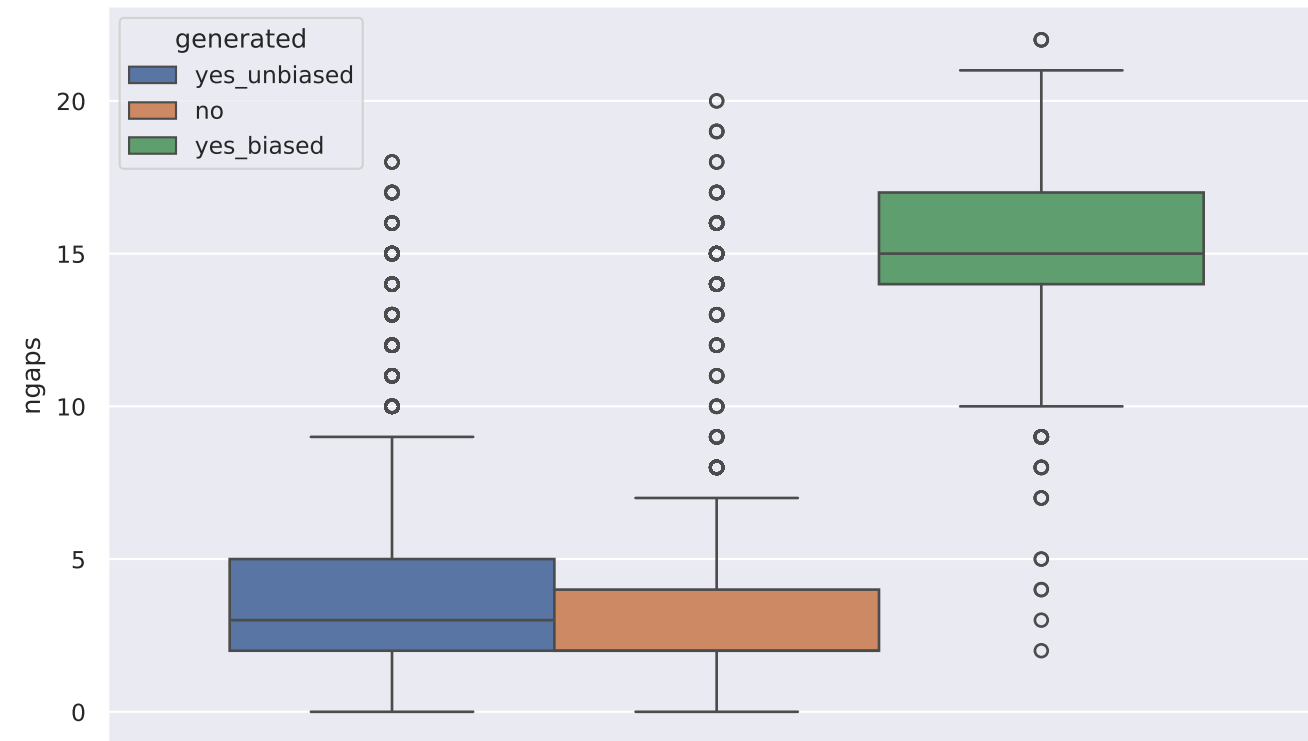
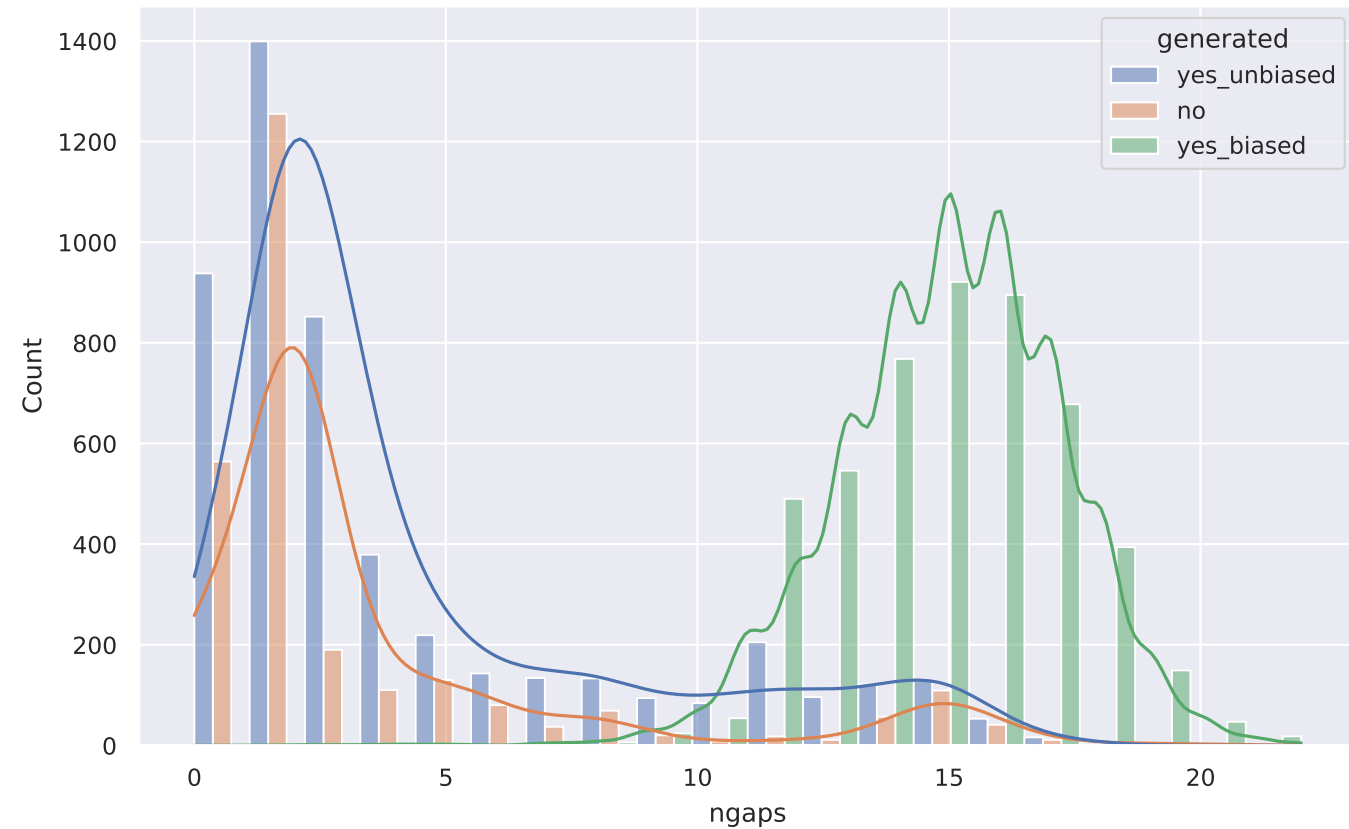


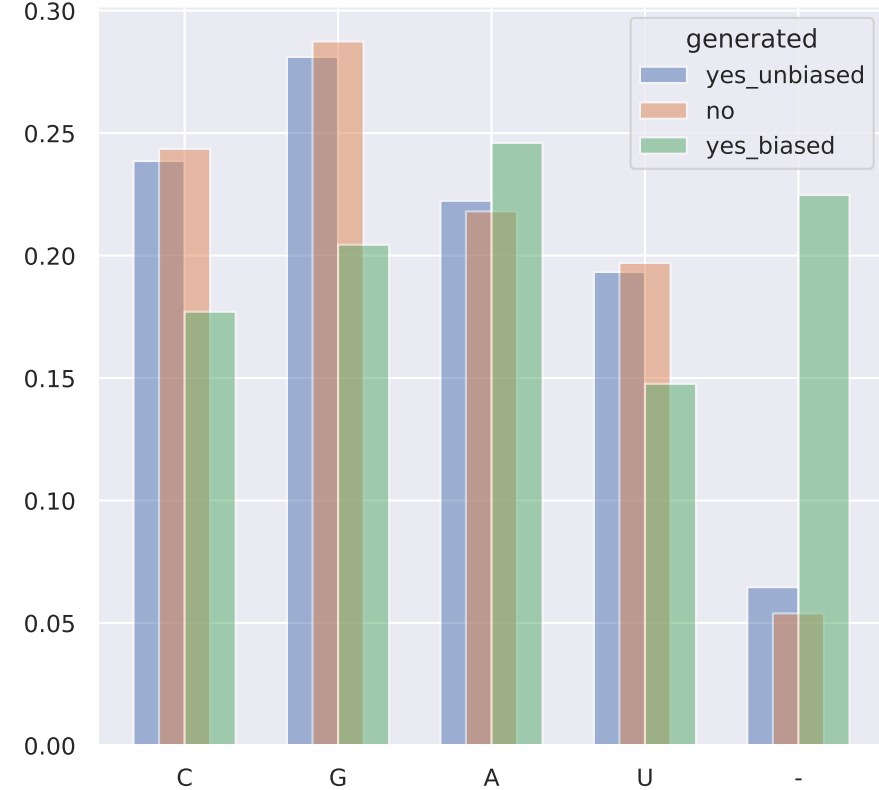
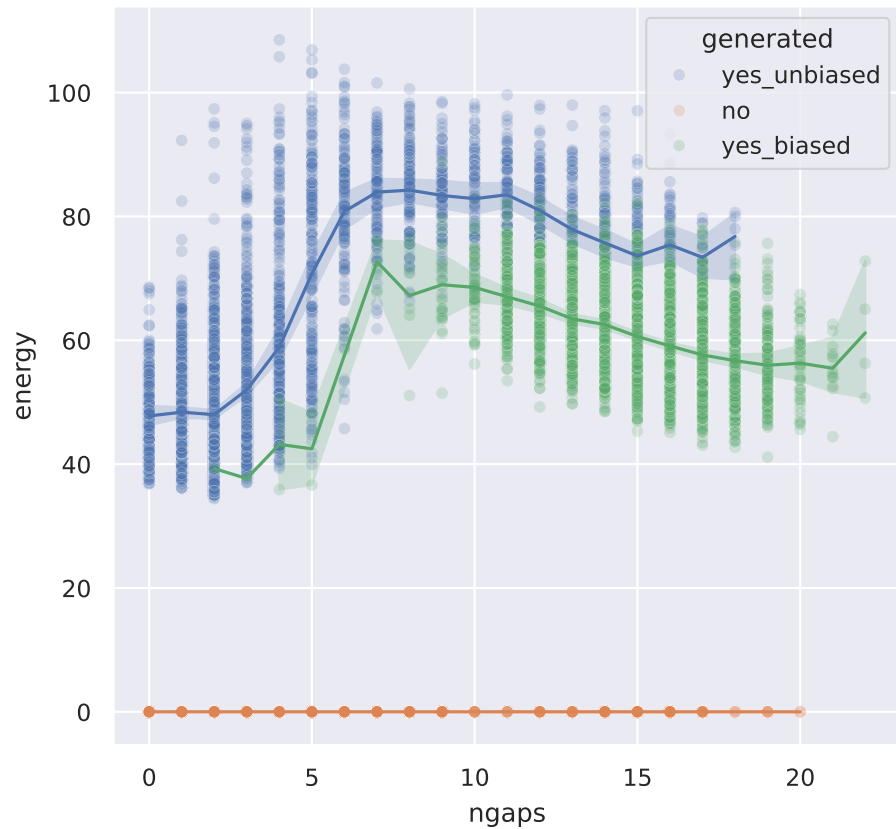
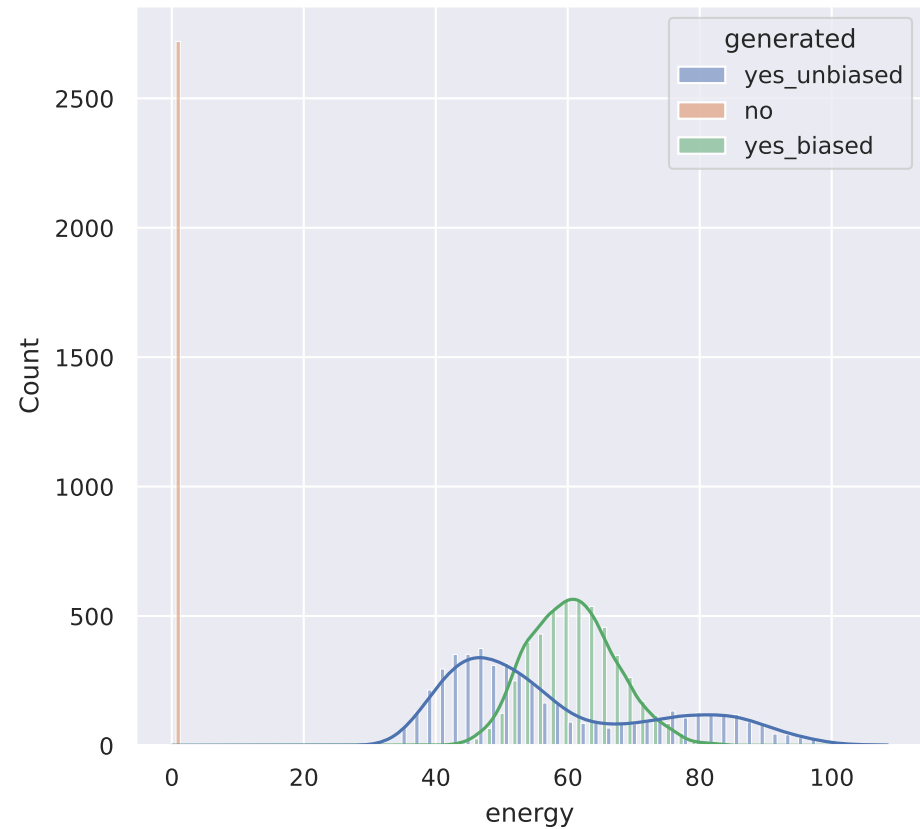
Numeric Feature : D_MFE



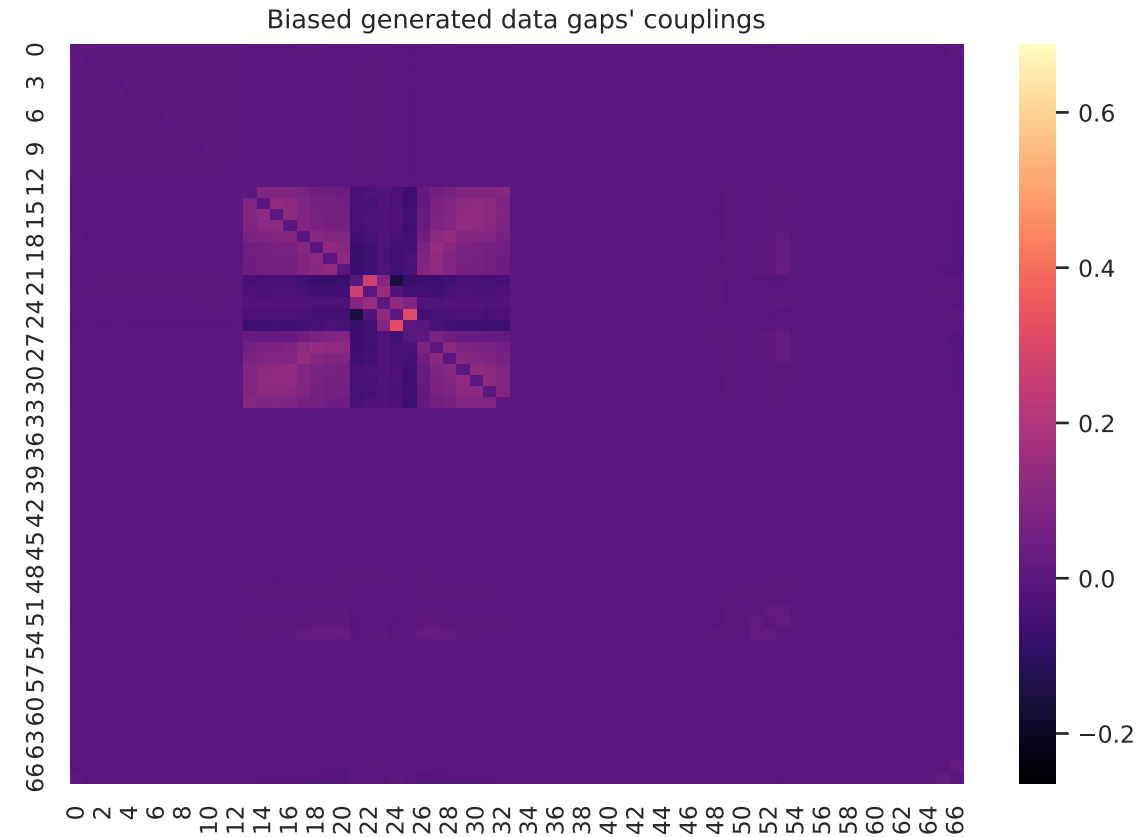
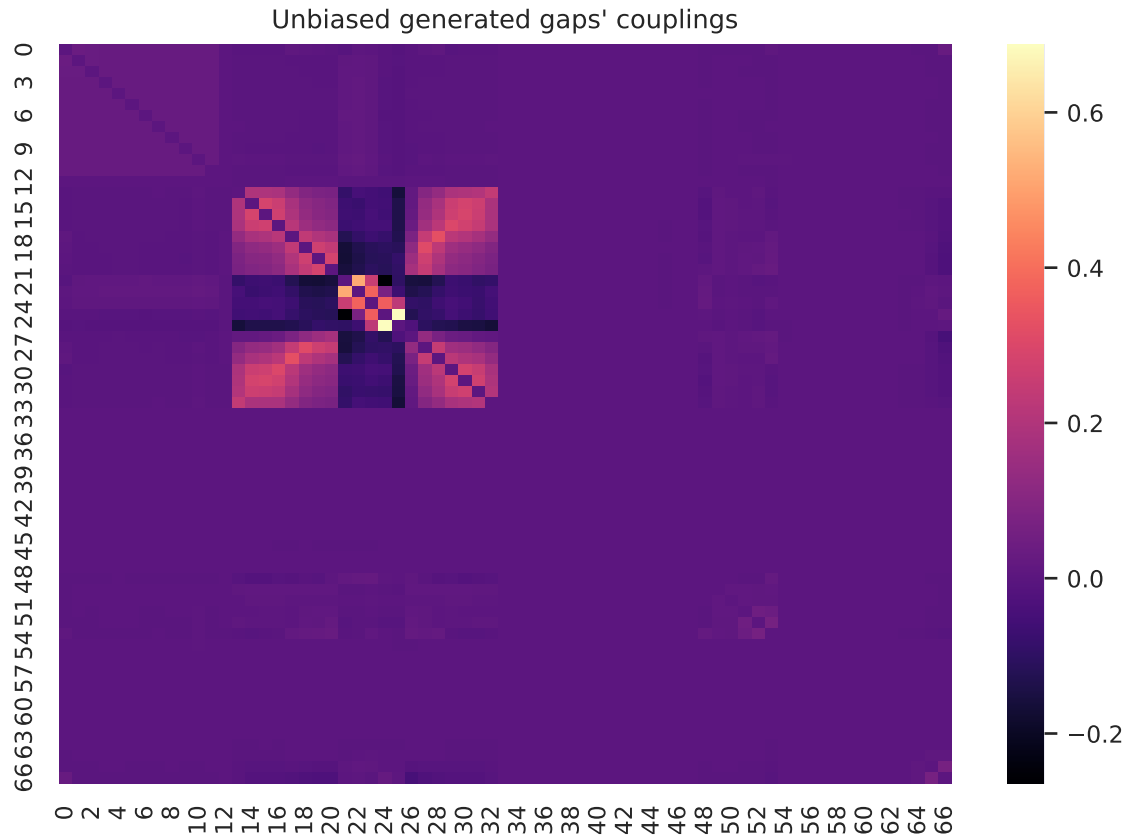
Numeric Feature : ngaps



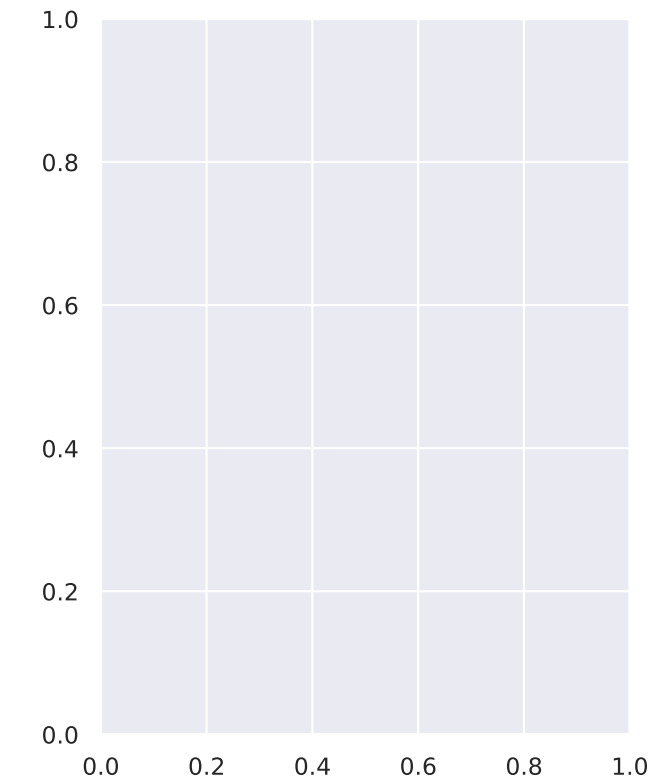
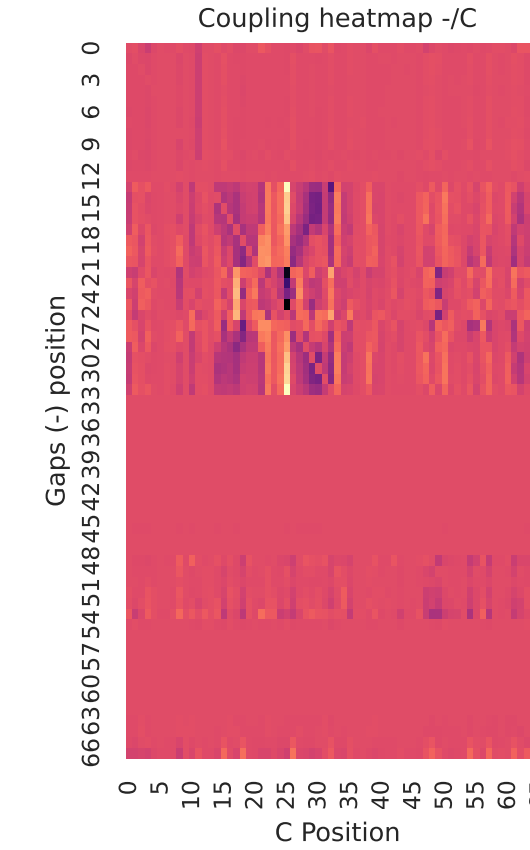
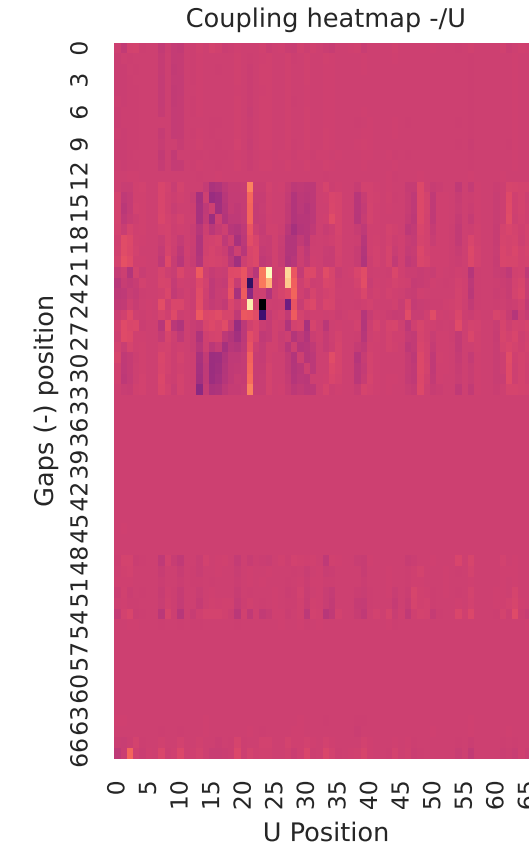
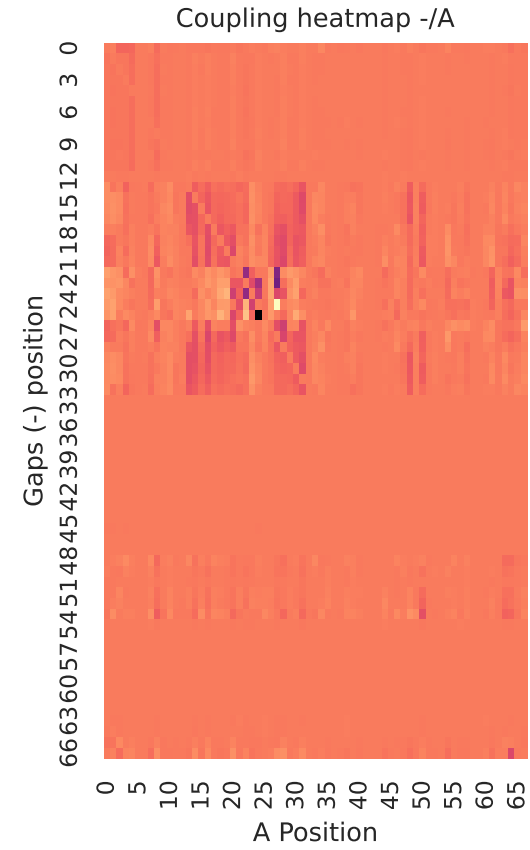
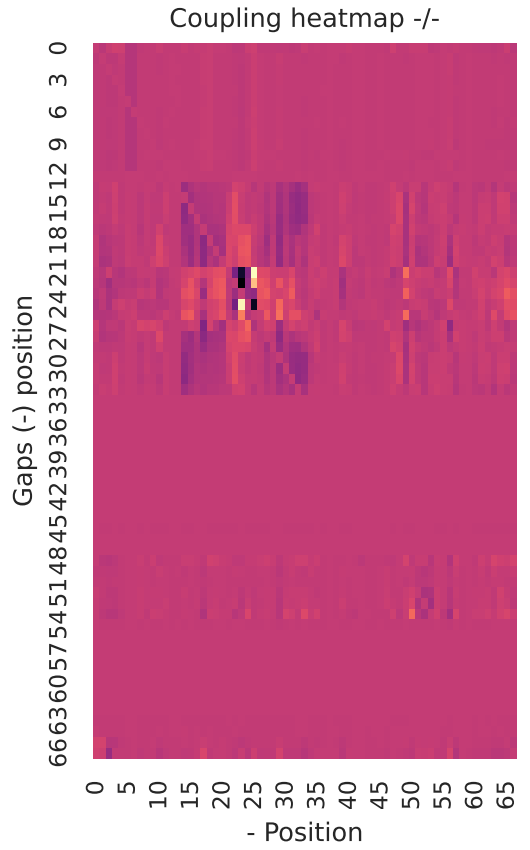
Numeric Feature : Sequences Energy and Nucleic acid average frequency



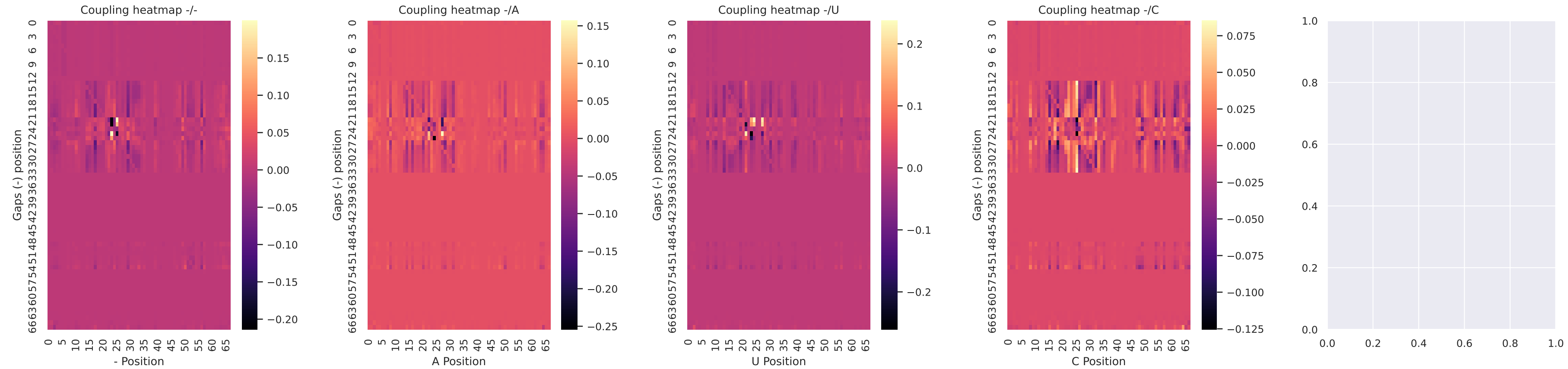
Couplings heatmap for -/-



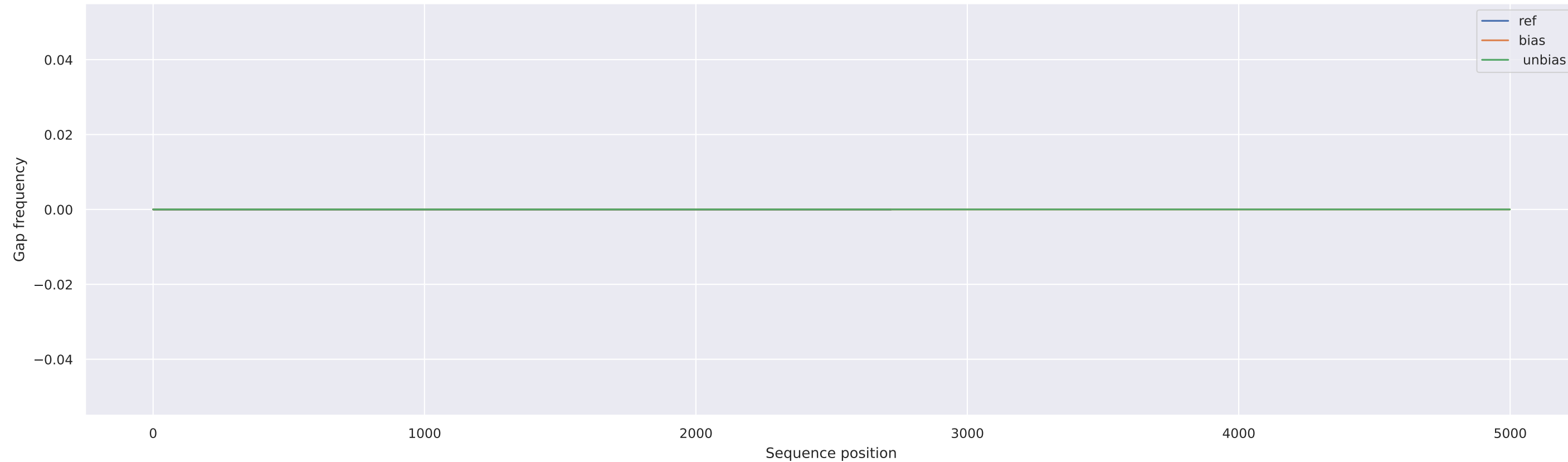
Couplings heatmap for Unbiased

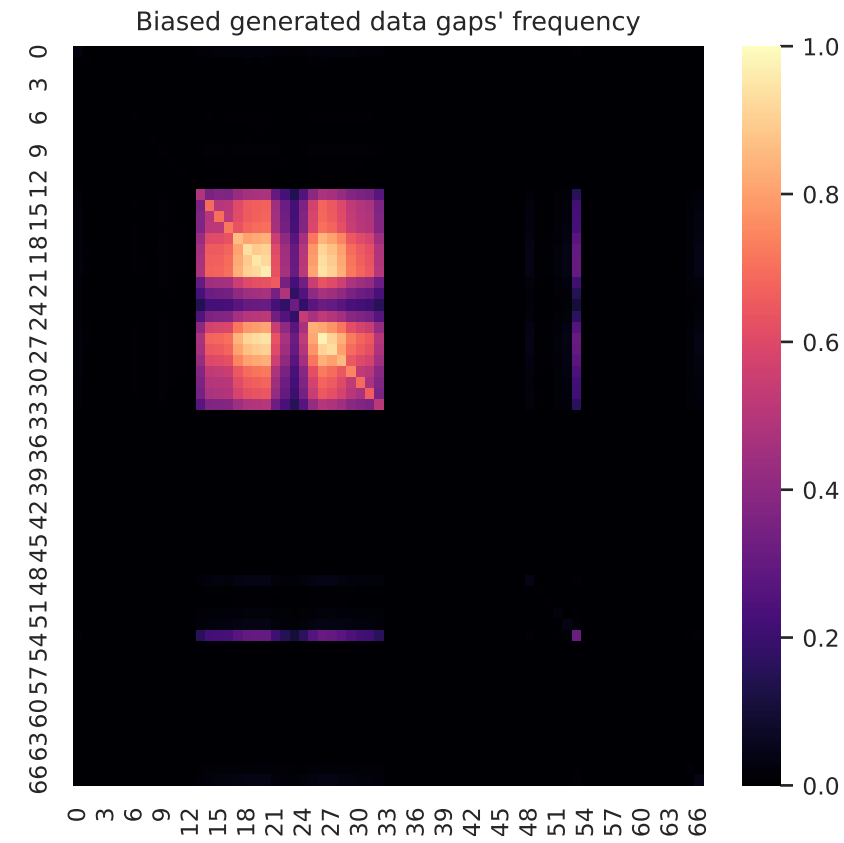
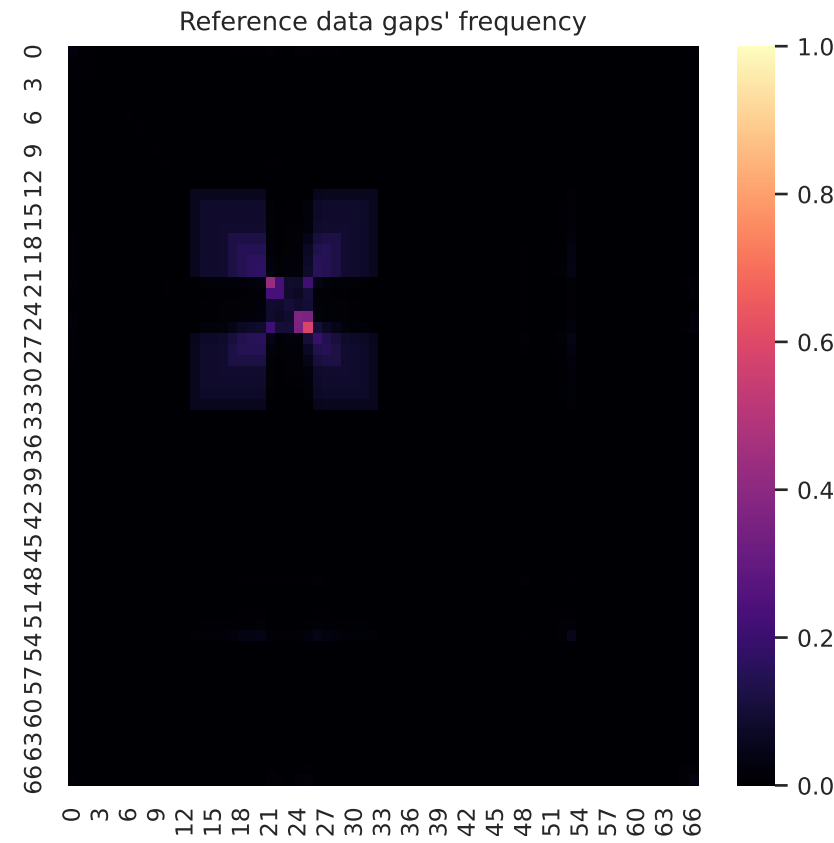
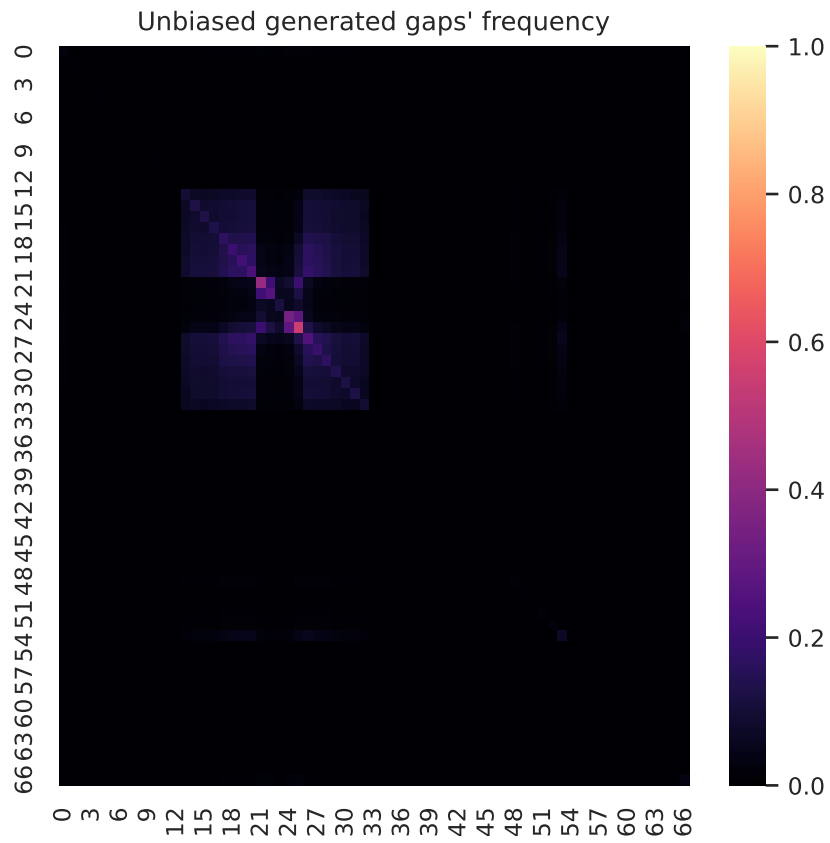


Couplings heatmap for biased

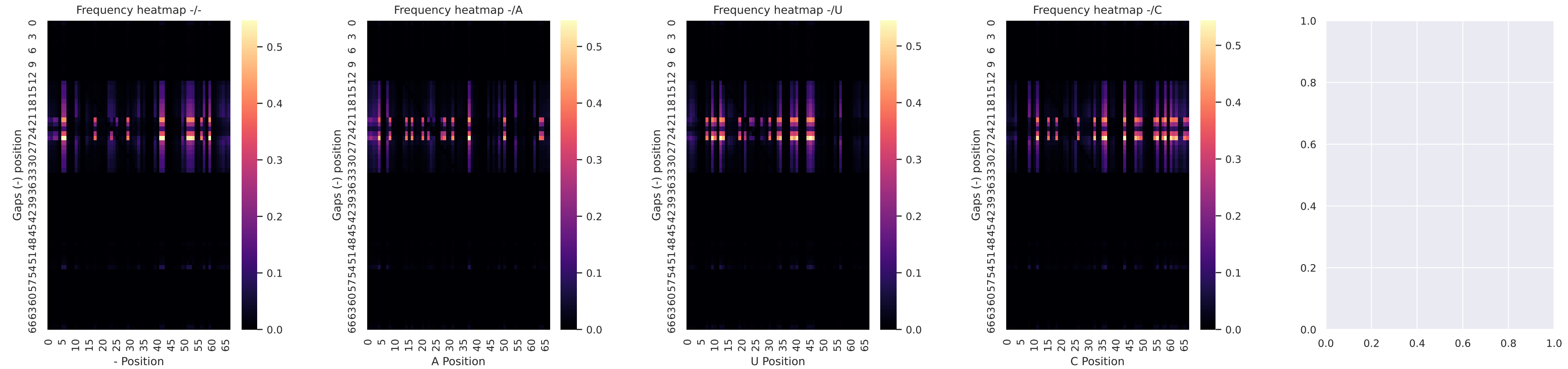


Gaps' frequency depending on position in the MSA's sequences





Frequency heatmap for Unbiased



Frequency heatmap for ref

