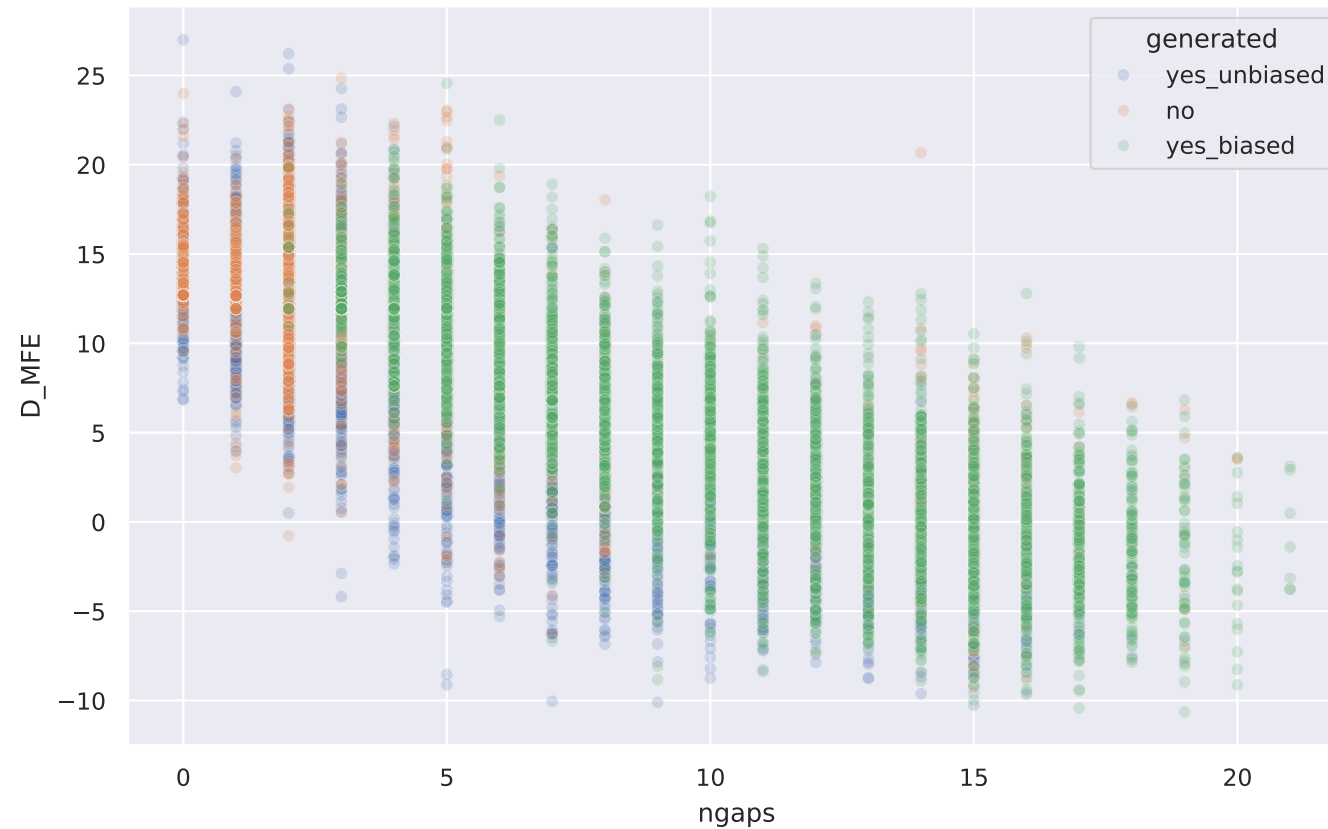
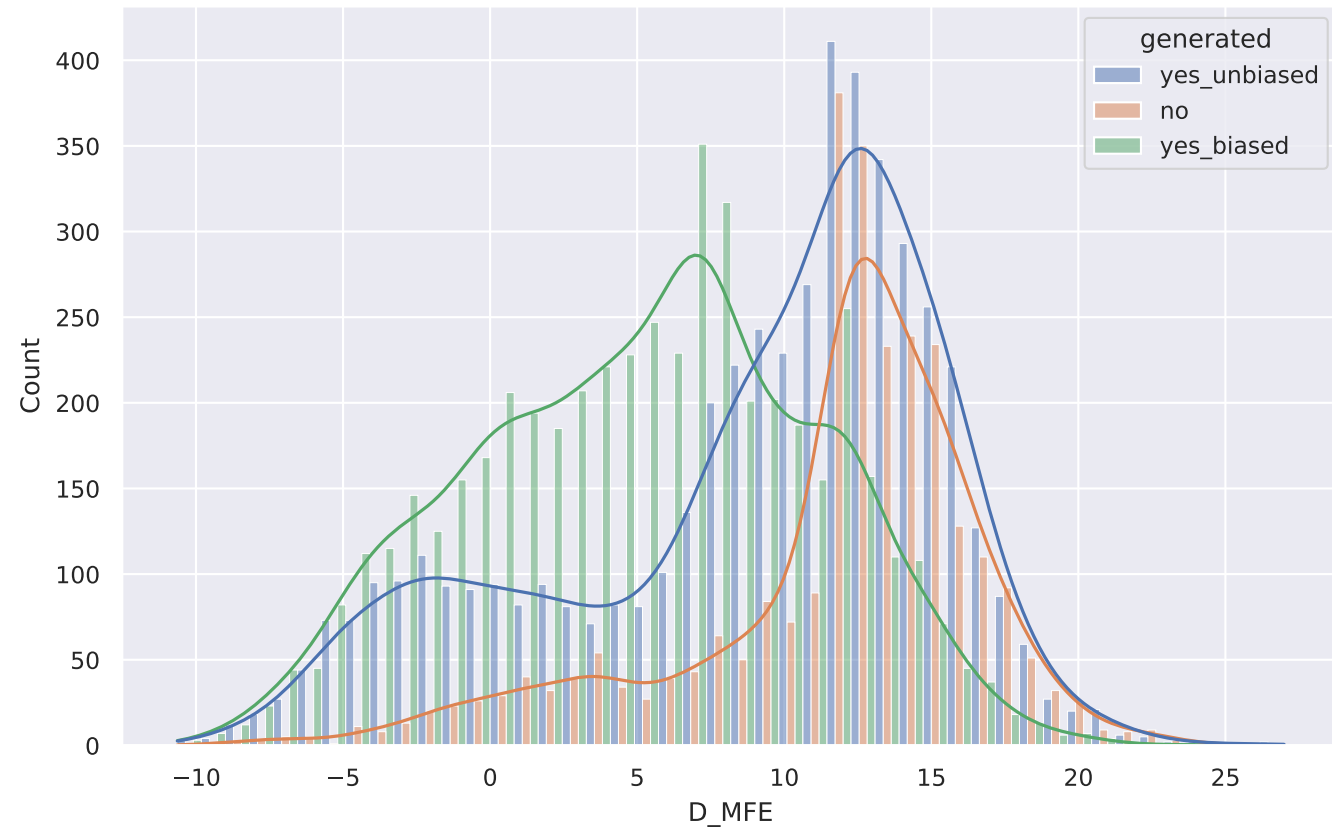
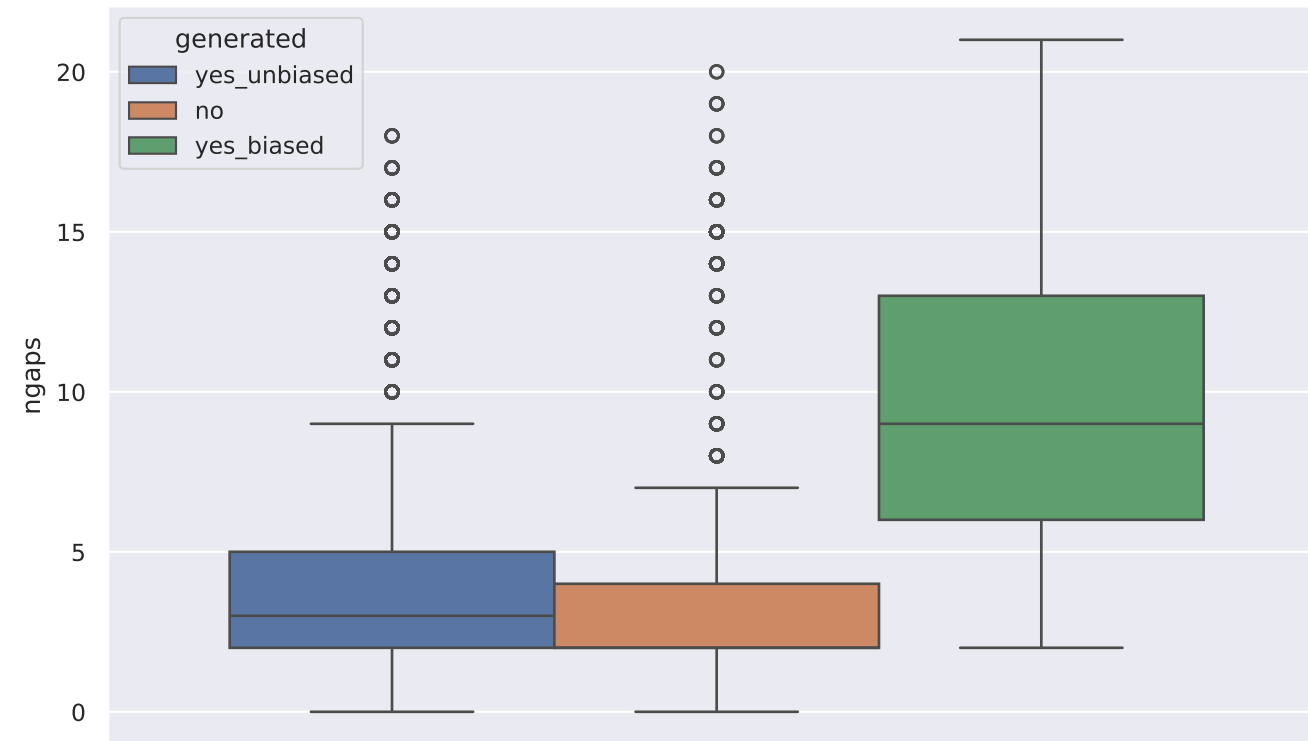
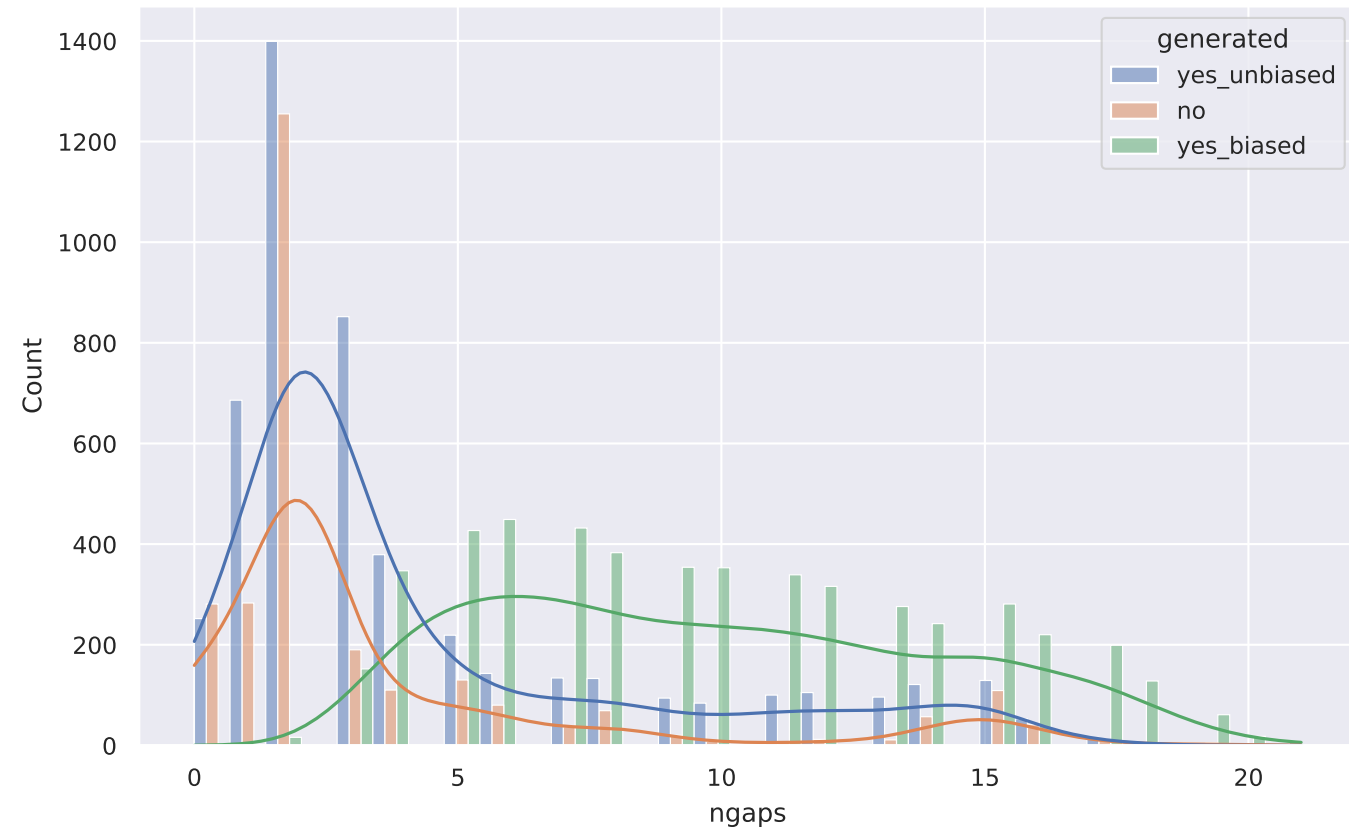


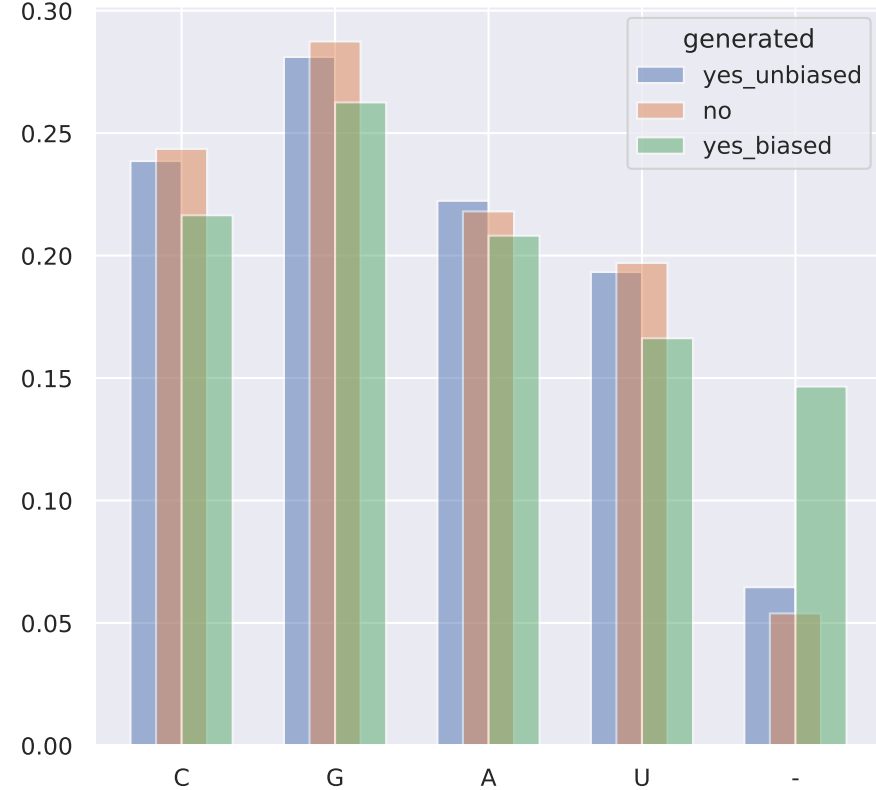
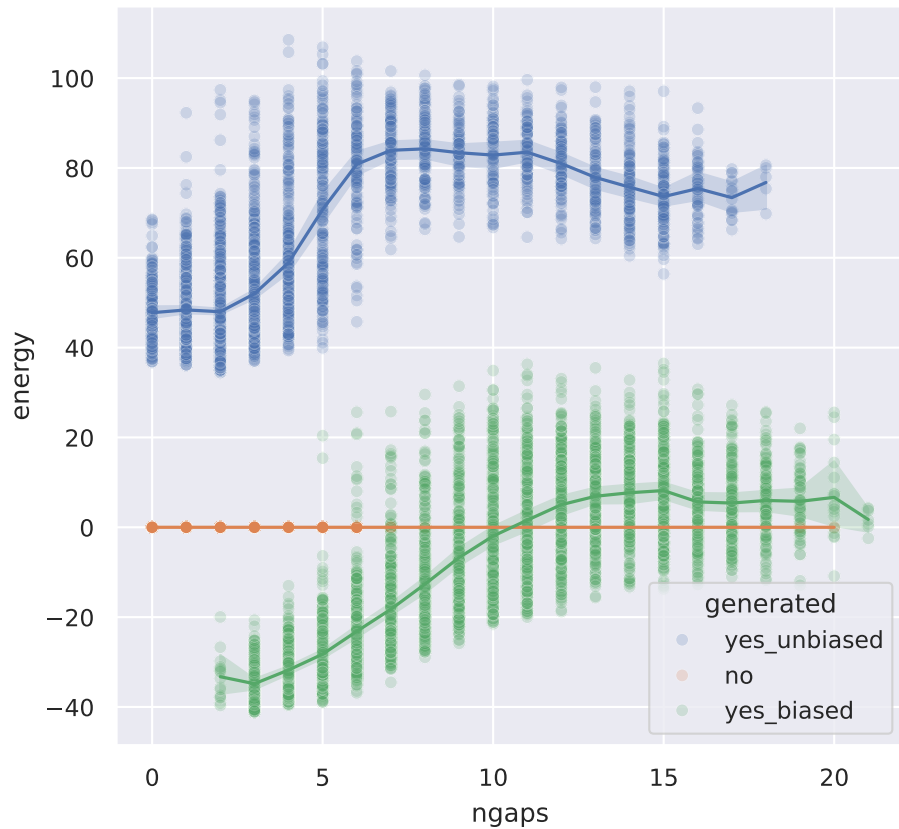
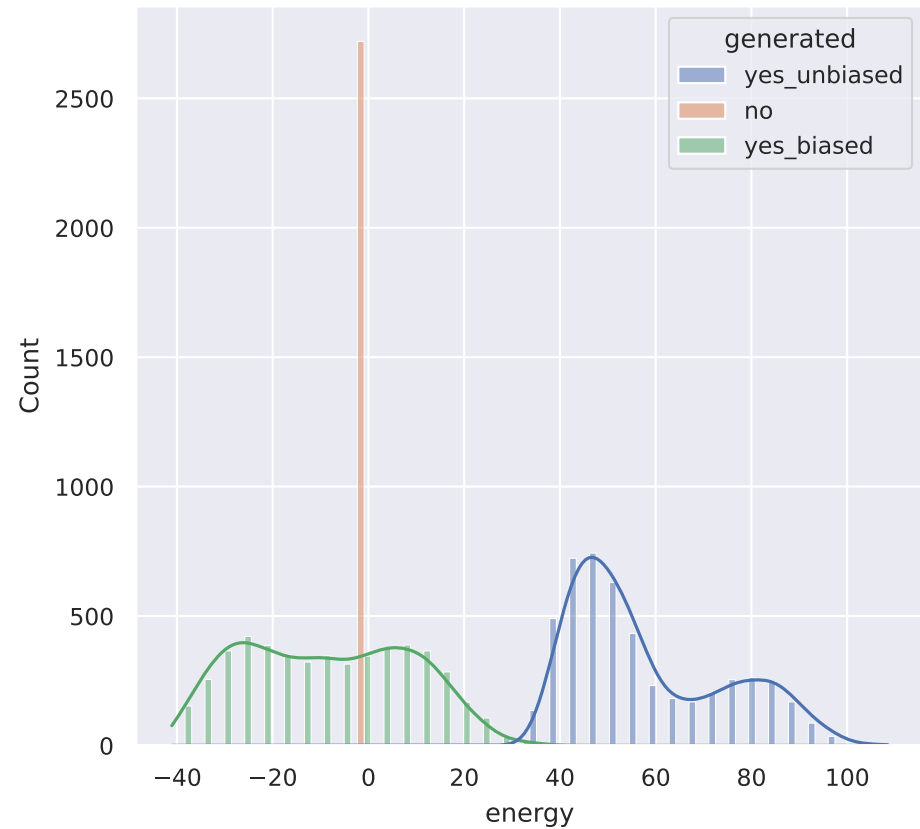
**Numeric Feature : D\_MFE**



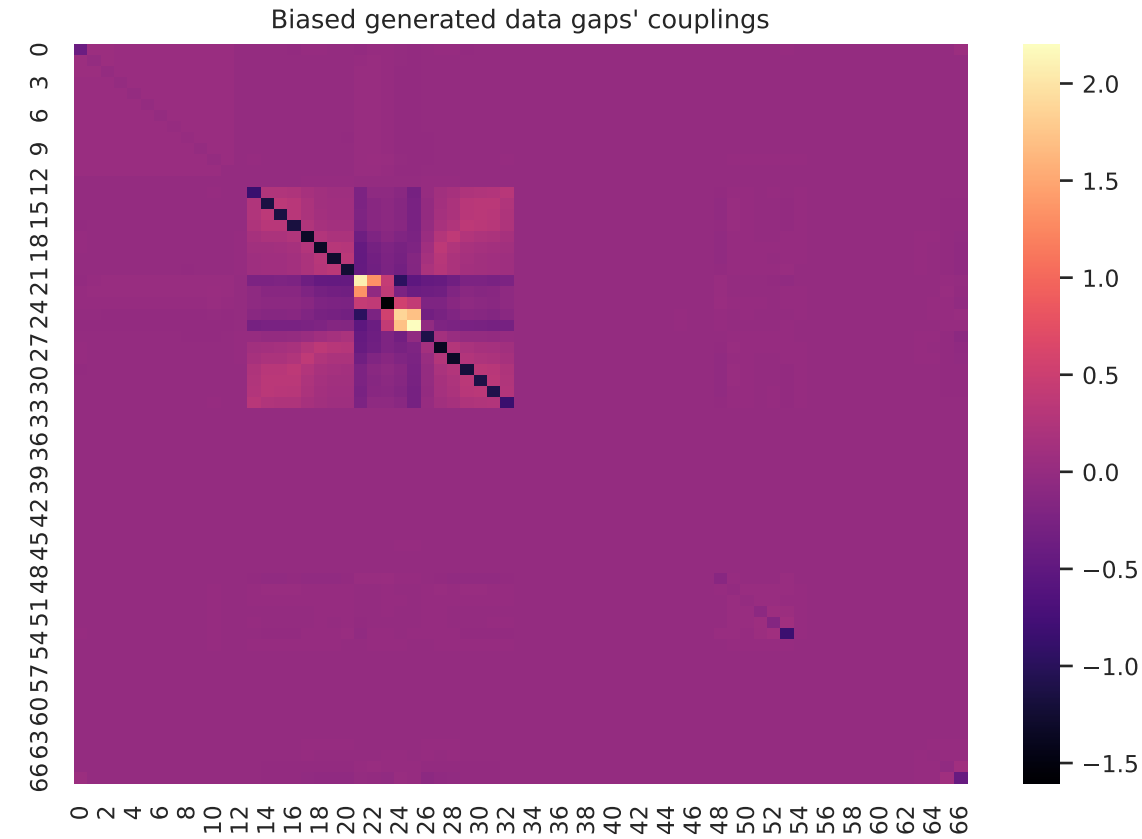
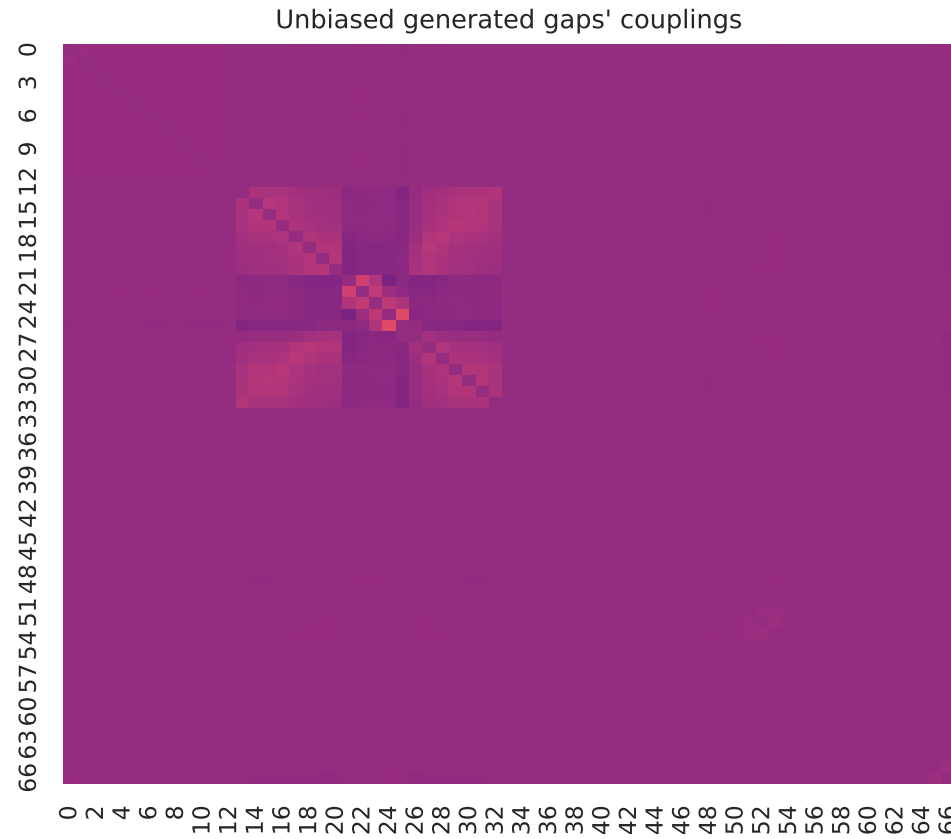
**Numeric Feature : ngaps**



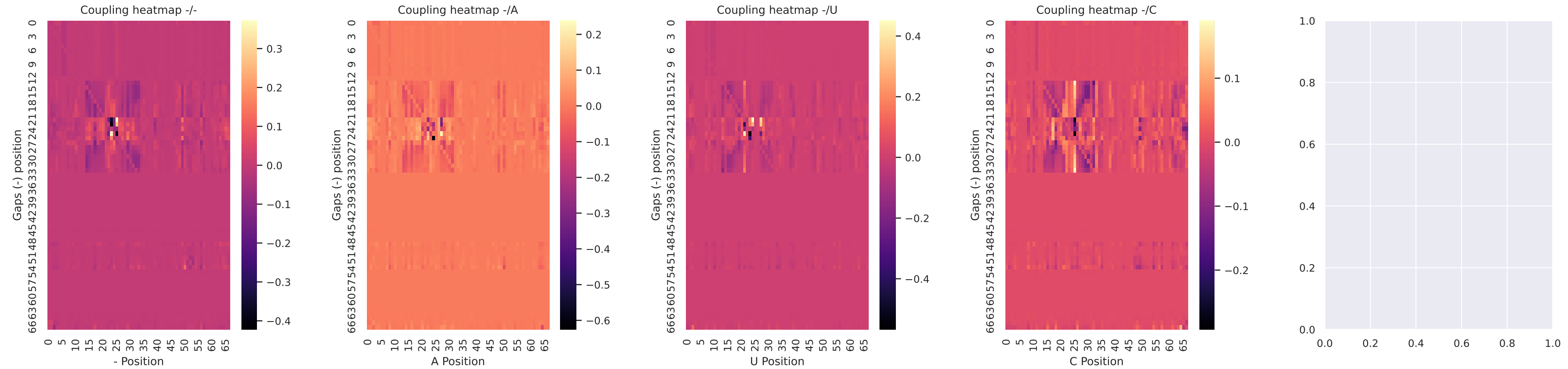
# Numeric Feature : Sequences Energy and Nucleic acid average frequency



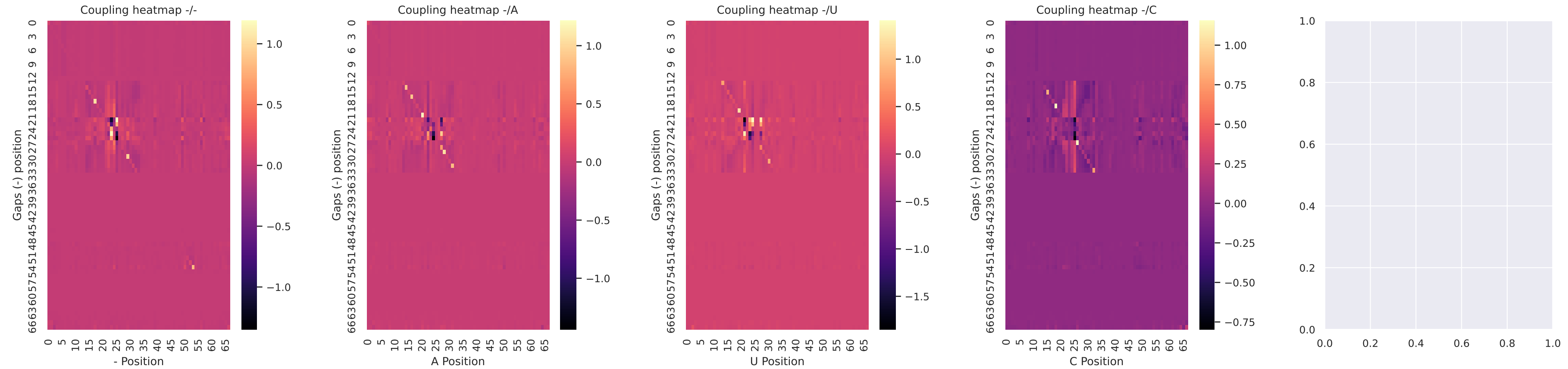
# Couplings heatmap for -/-



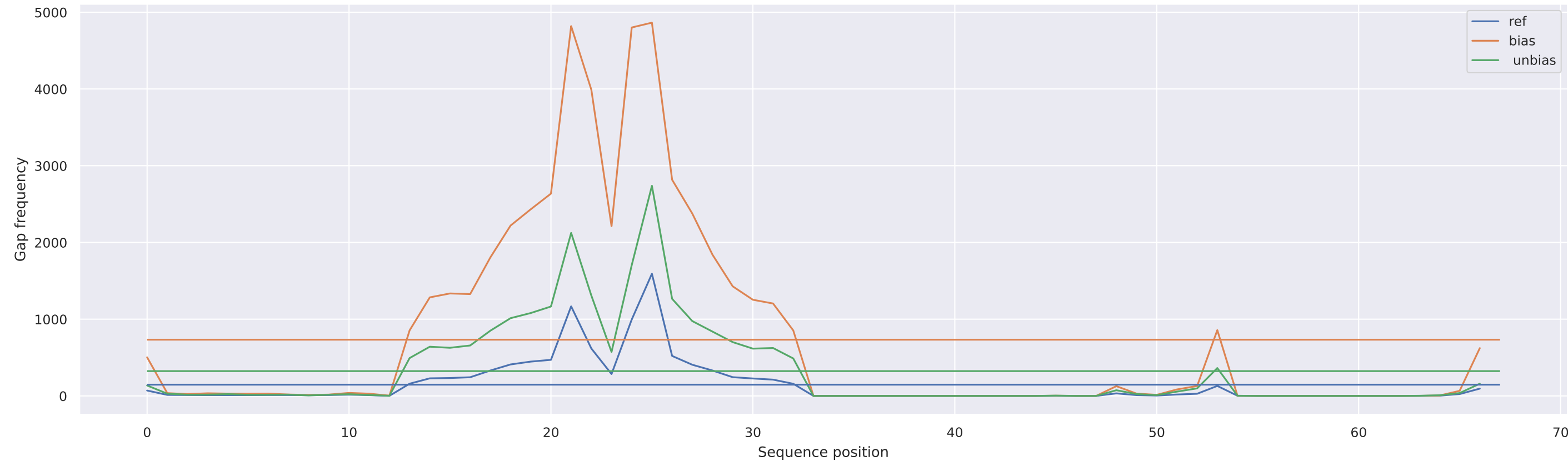
# Couplings heatmap for Unbiased

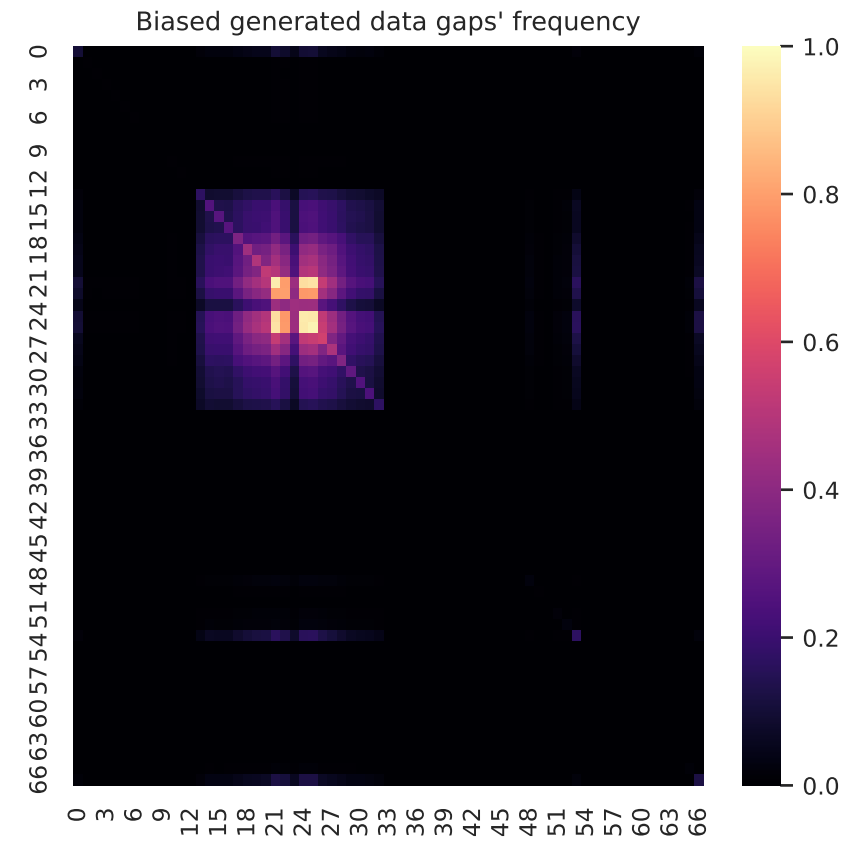
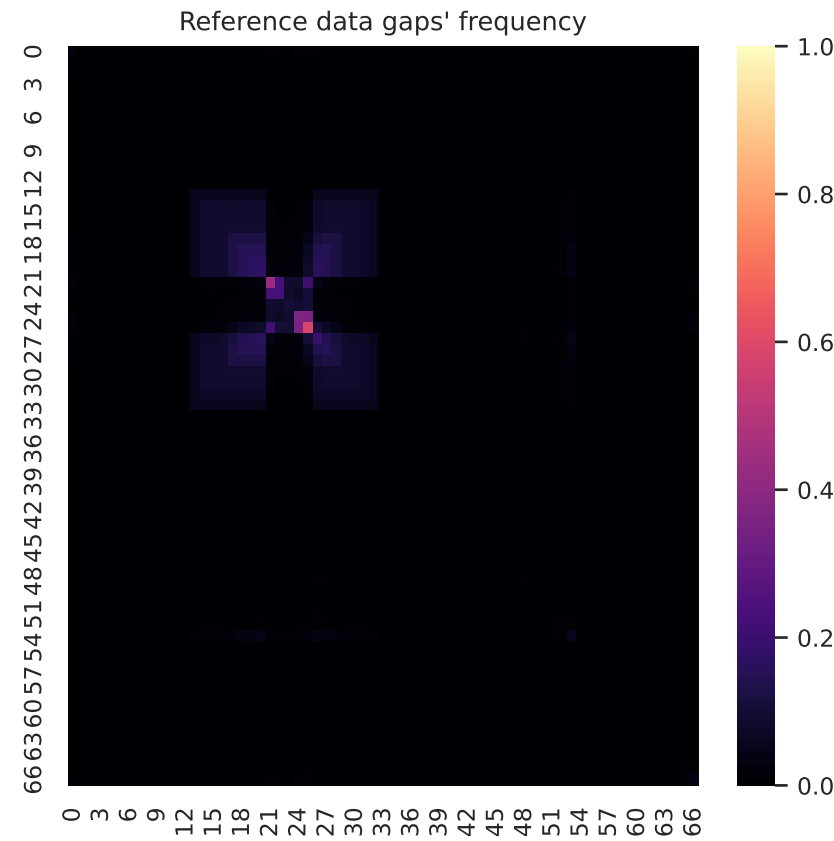
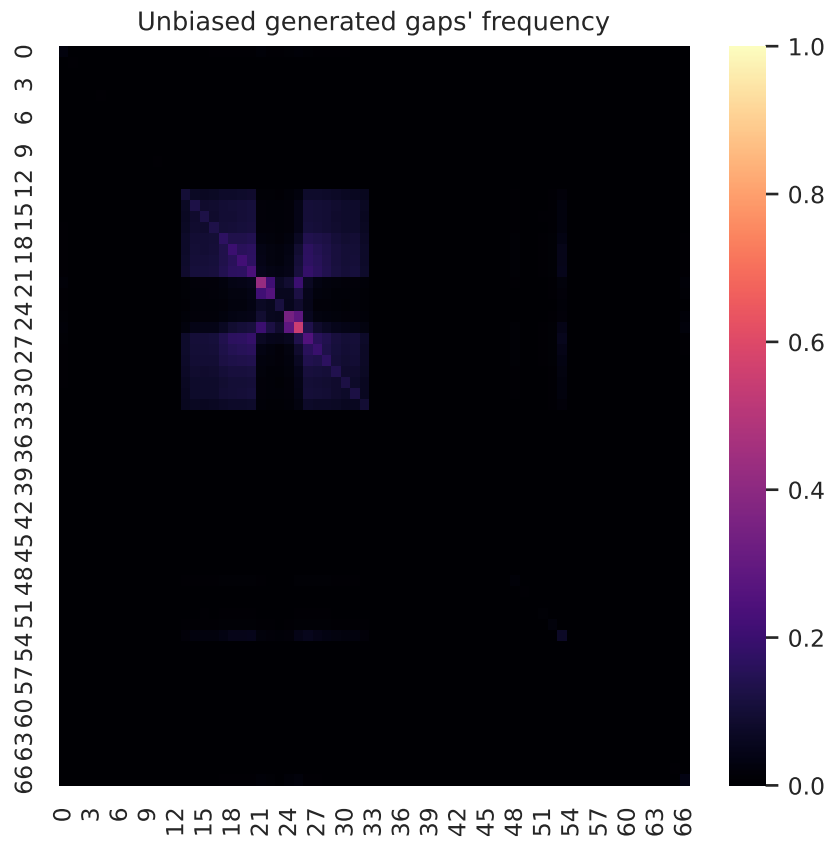


# Couplings heatmap for biased



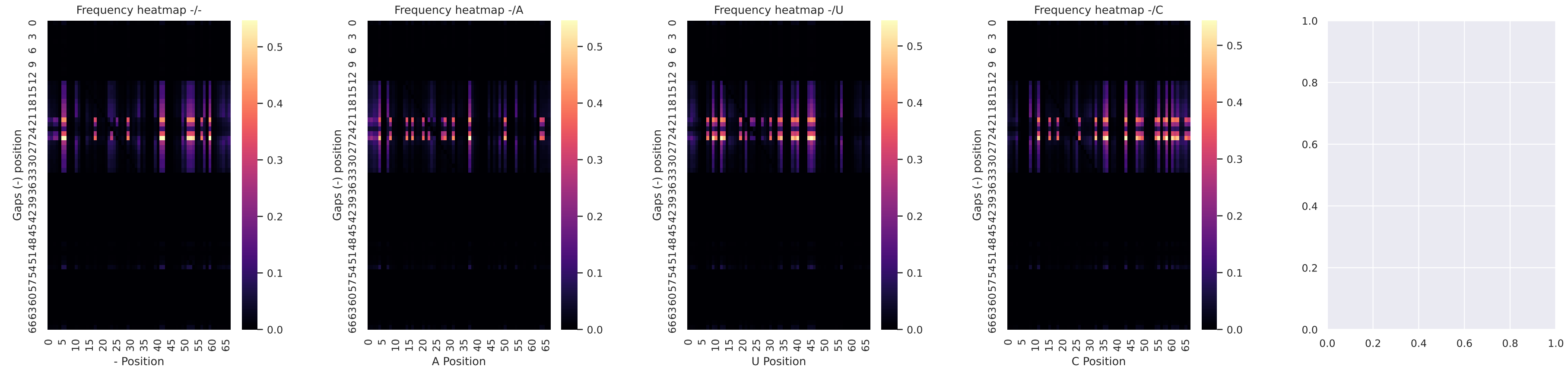
Gaps' frequency depending on position in the MSA's sequences



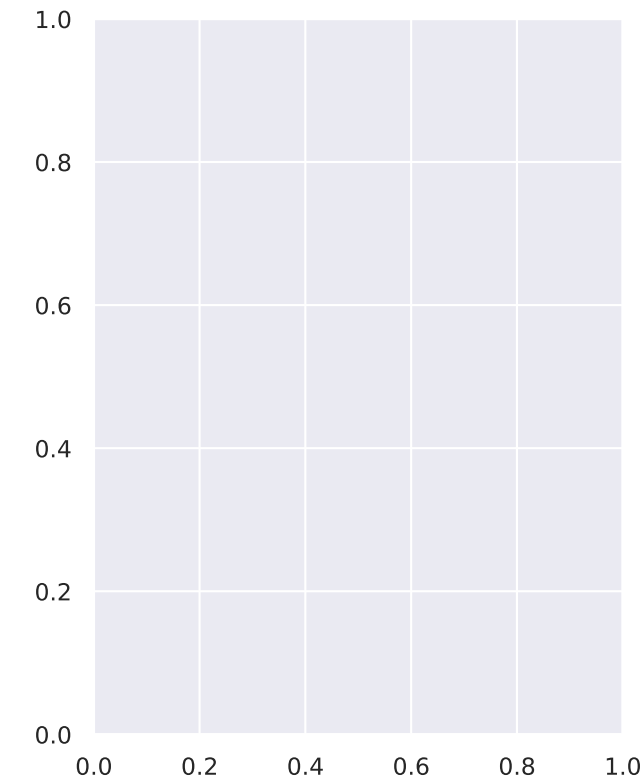
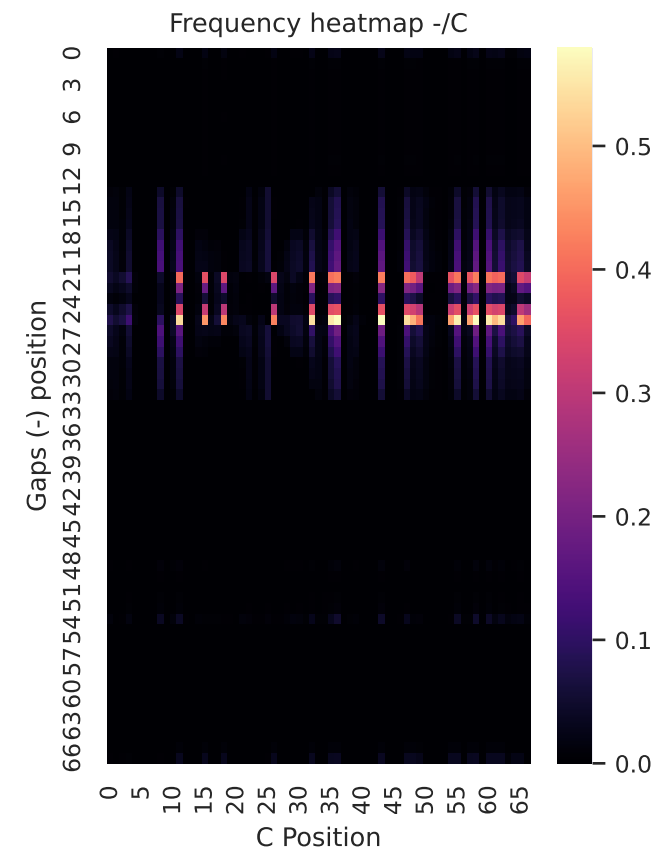
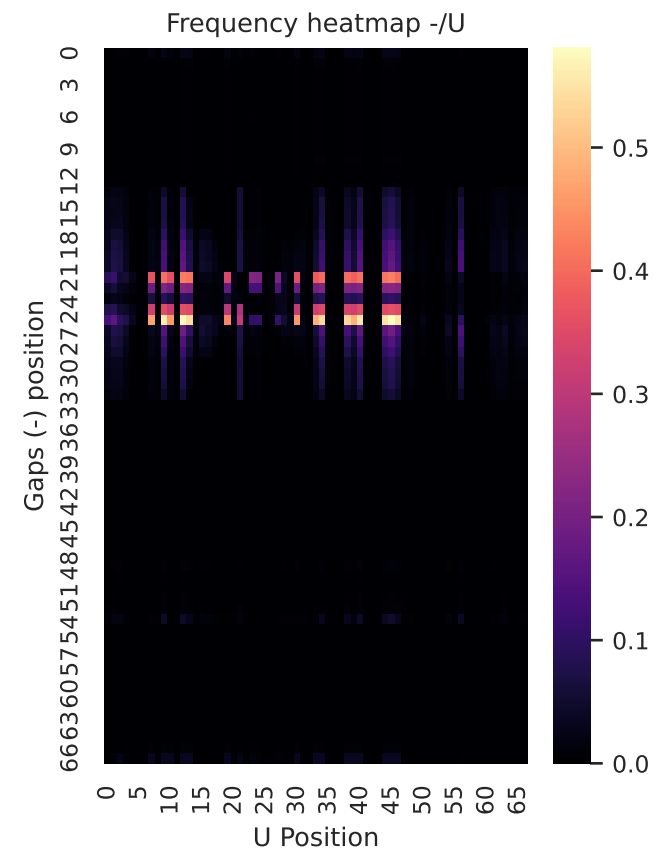
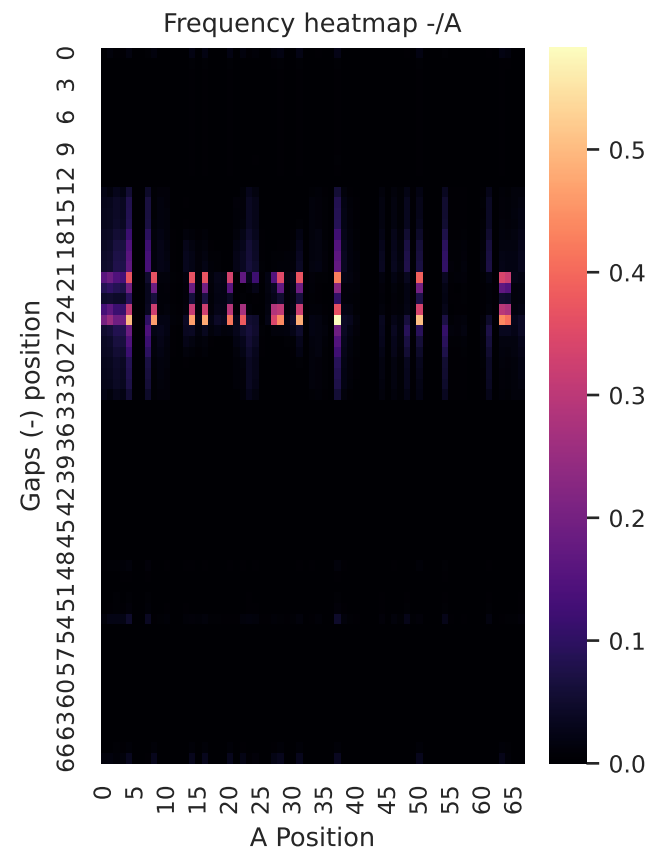
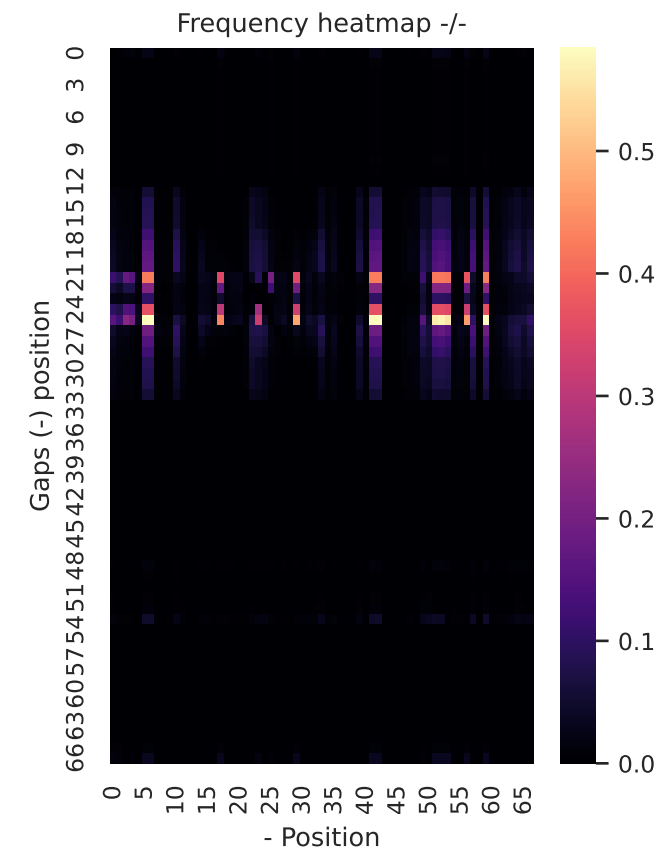




# Frequency heatmap for Unbiased



# Frequency heatmap for ref



# Frequency heatmap for biased

