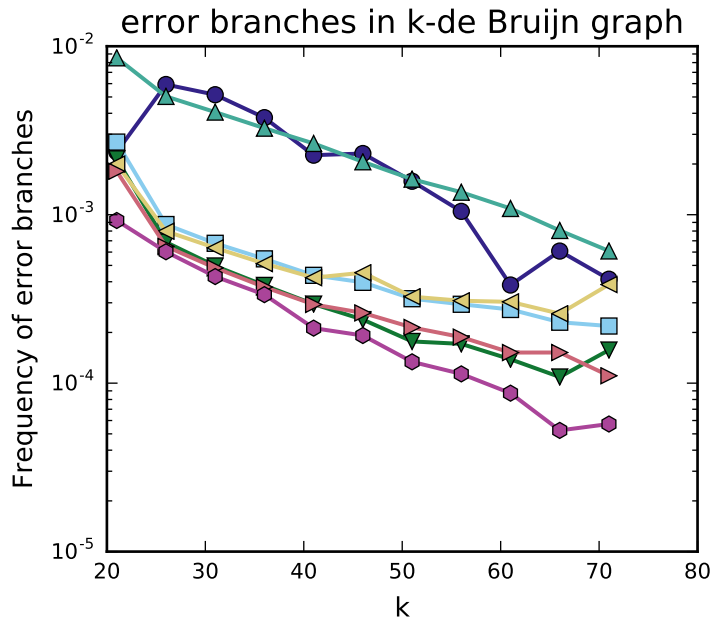
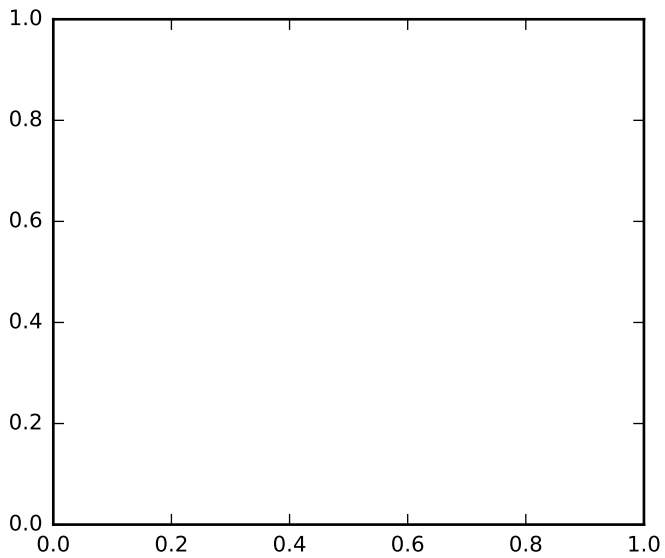
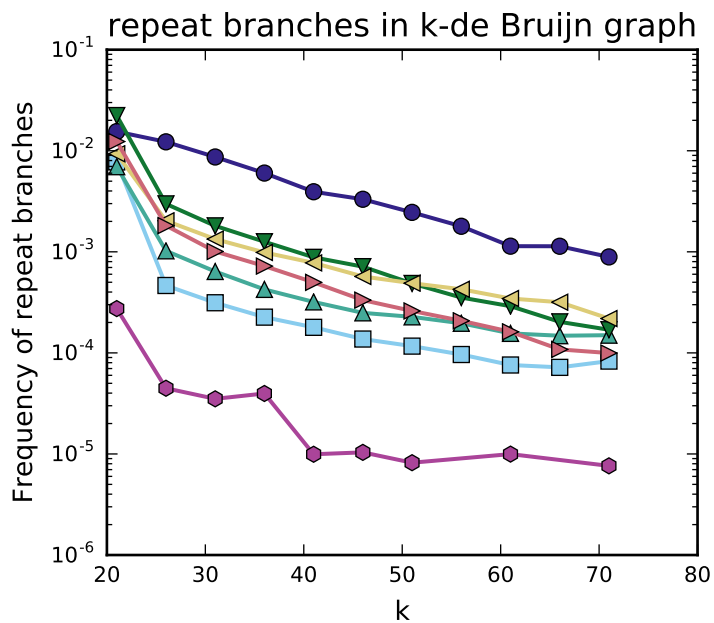
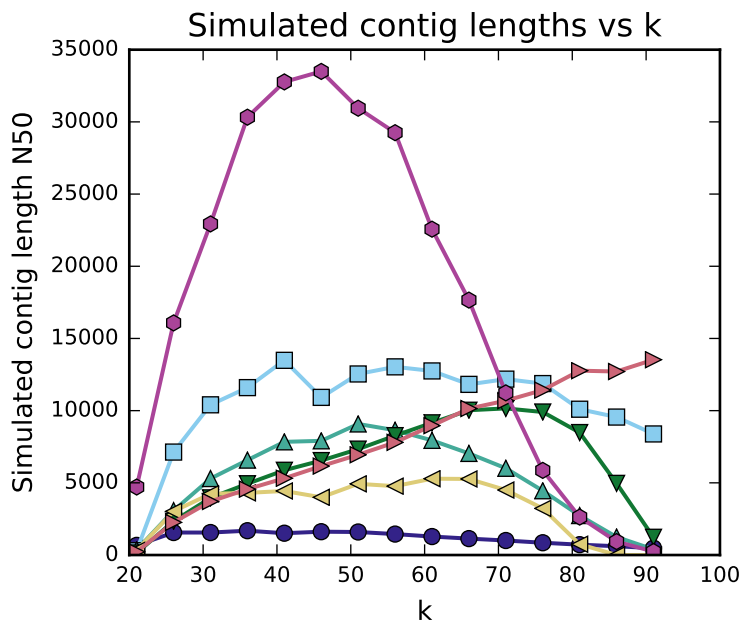
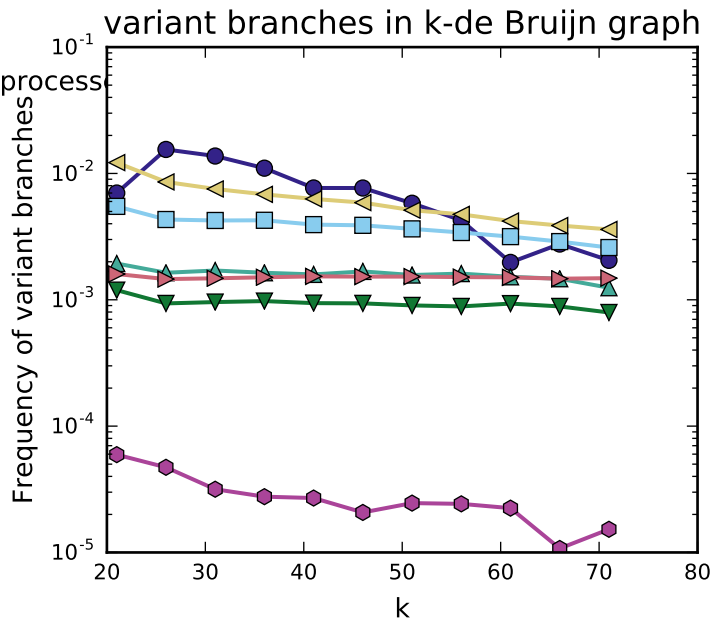
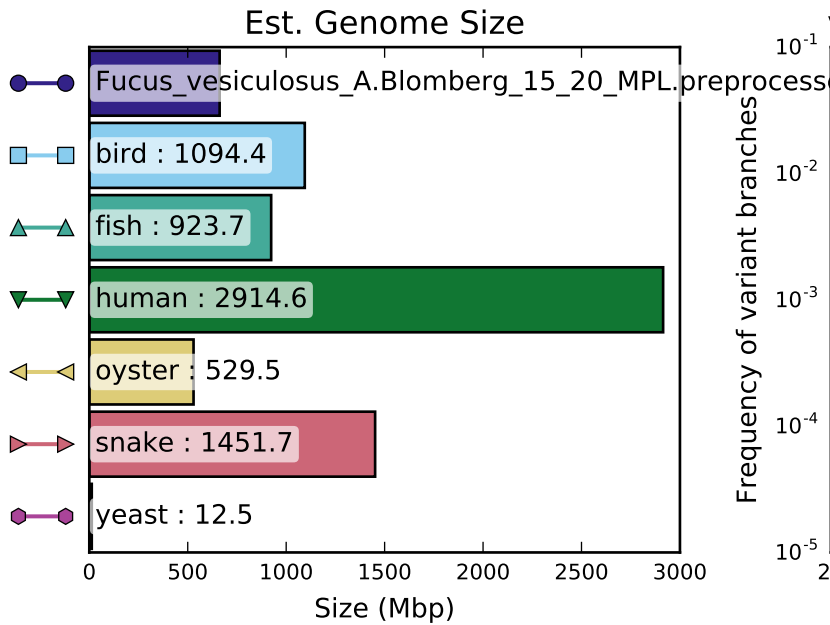
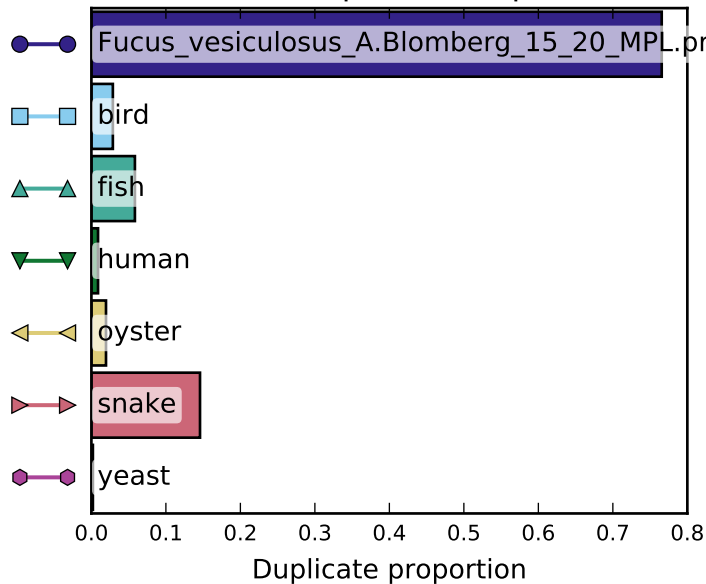


SGA Preqc Results : fig1

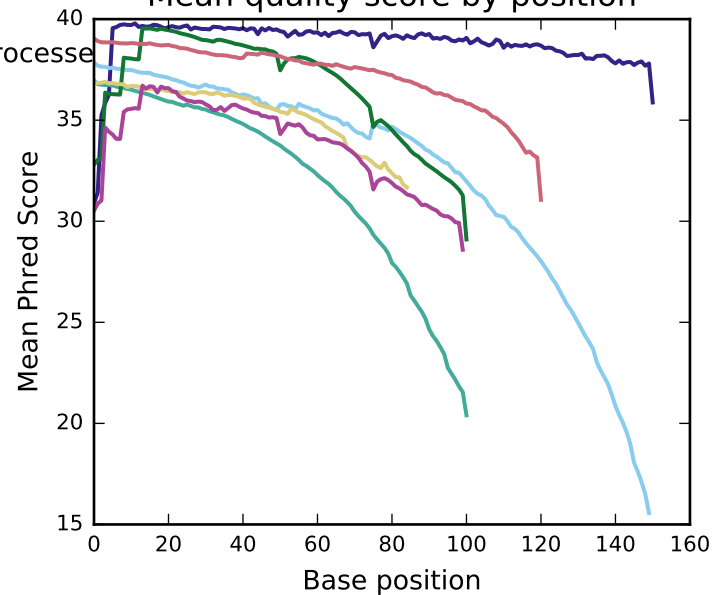


SGA Preqc Results : fig2

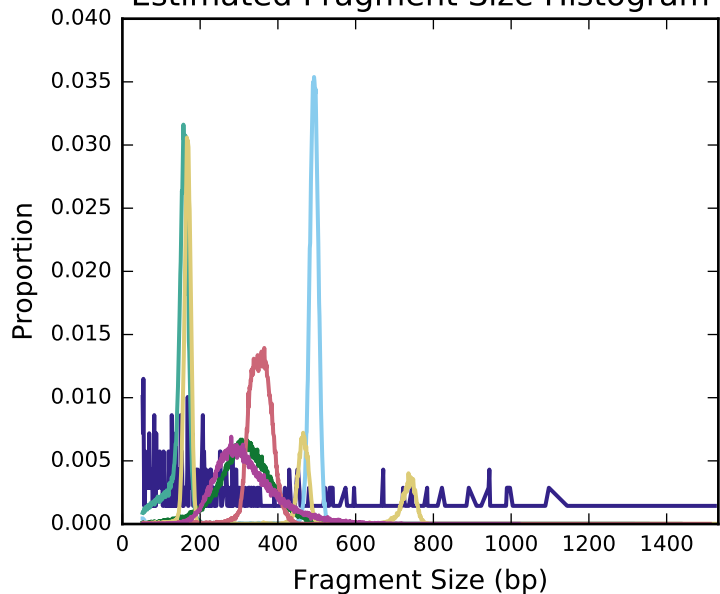
Est. PCR Duplicate Proportion



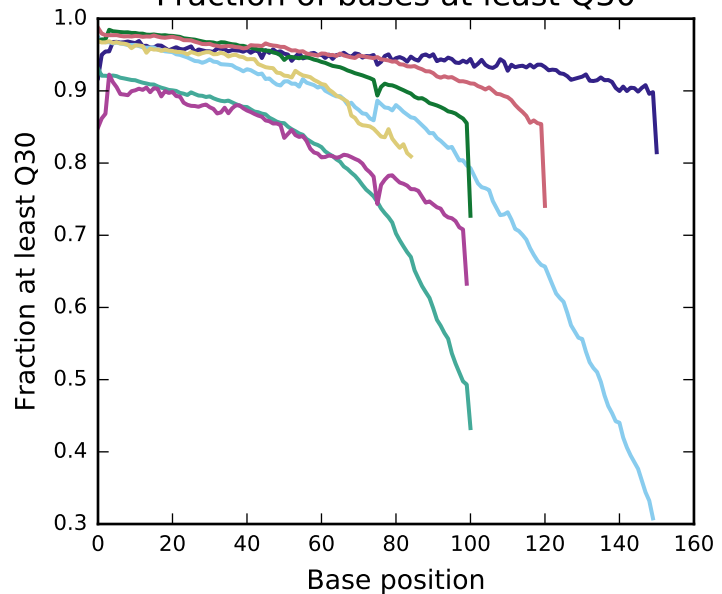
Mean quality score by position



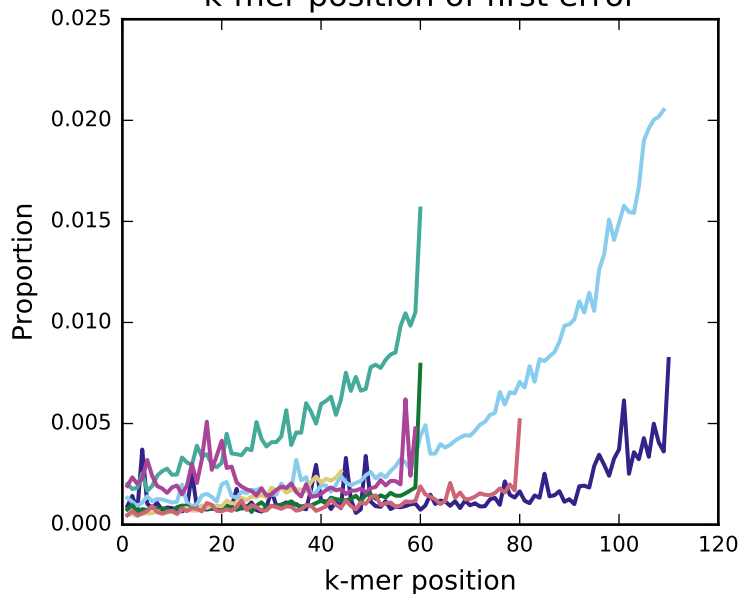
Estimated Fragment Size Histogram



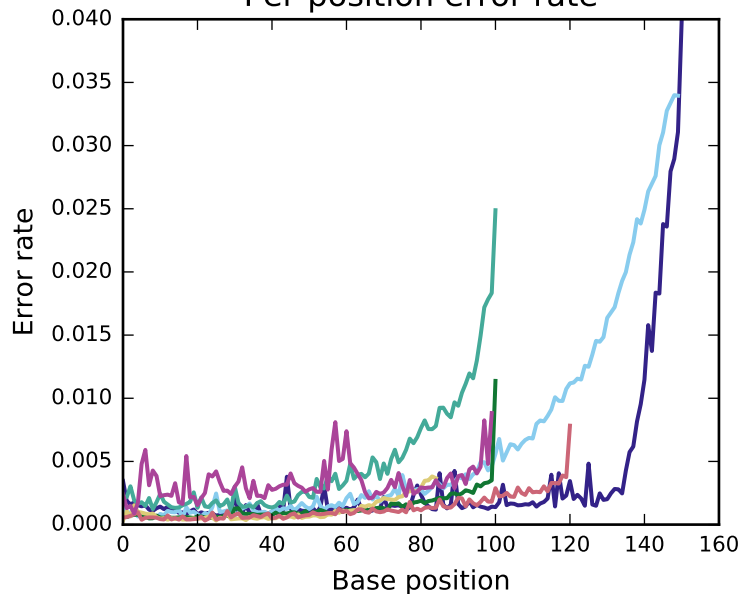
Fraction of bases at least Q30



k-mer position of first error

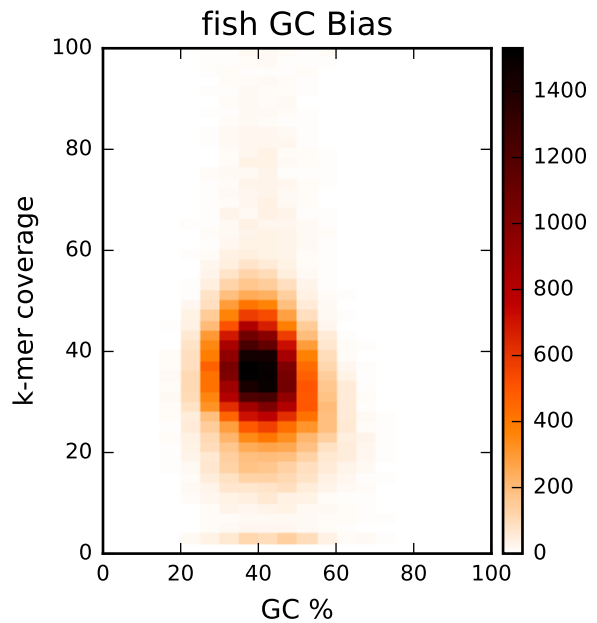
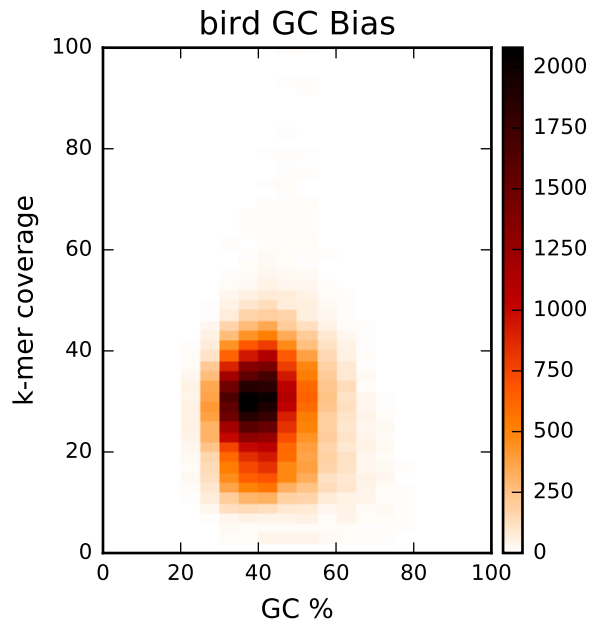
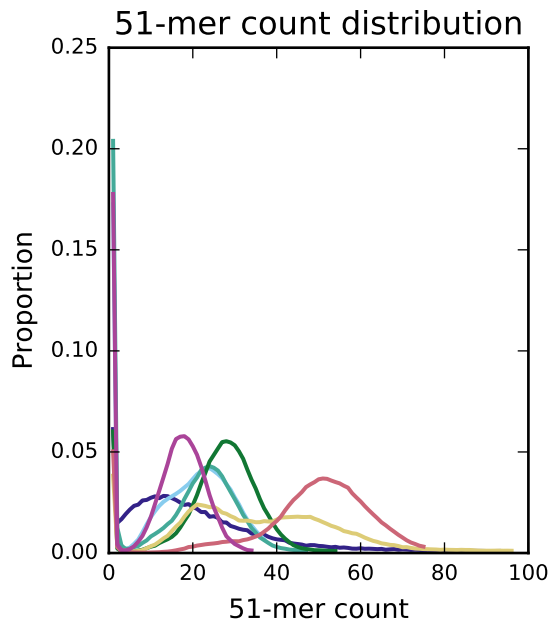
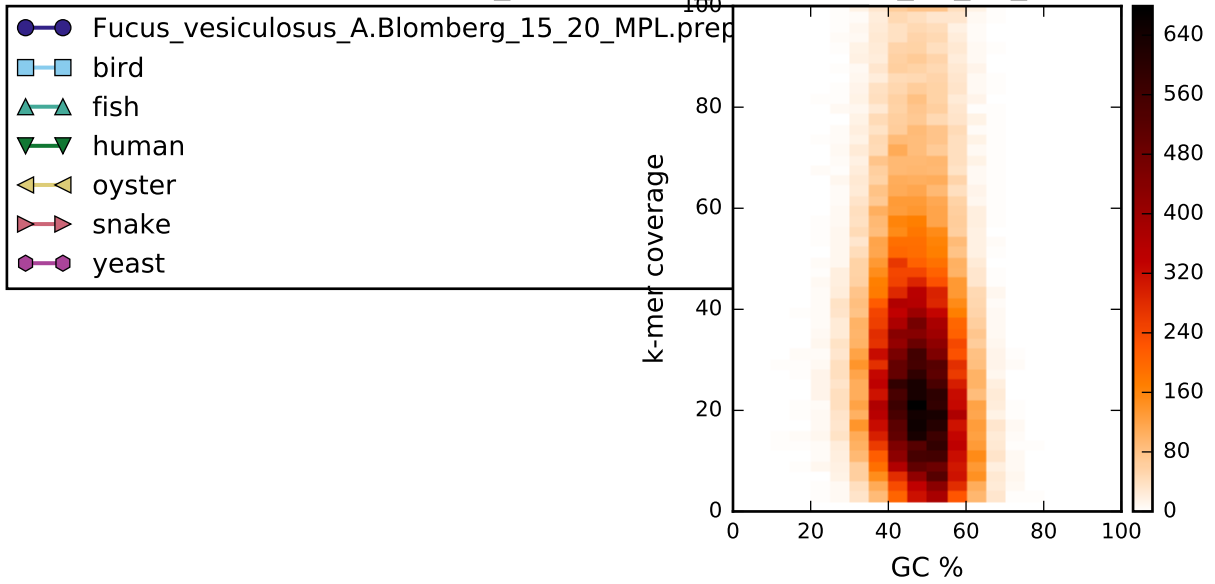


Per-position error rate



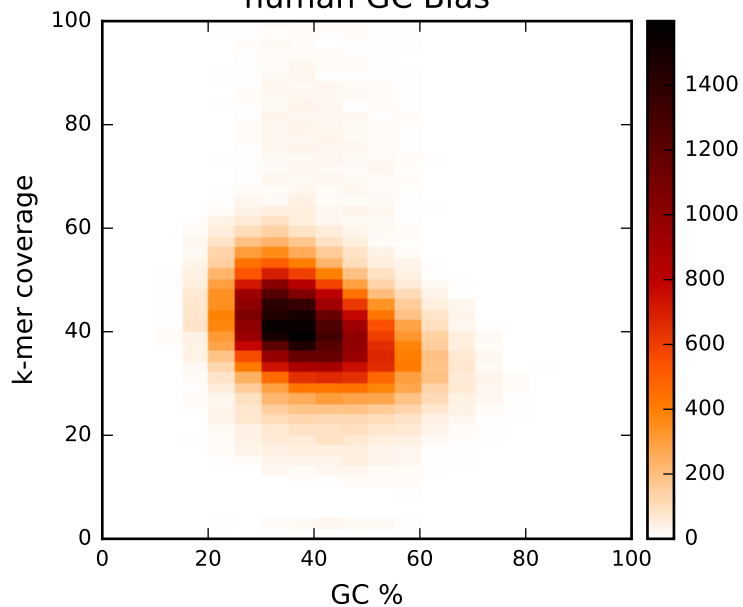
SGA Preqc Results : fig3

Fucus_vesiculosus_A.Blomberg_15_20_MPL.preprocessed GC Bi

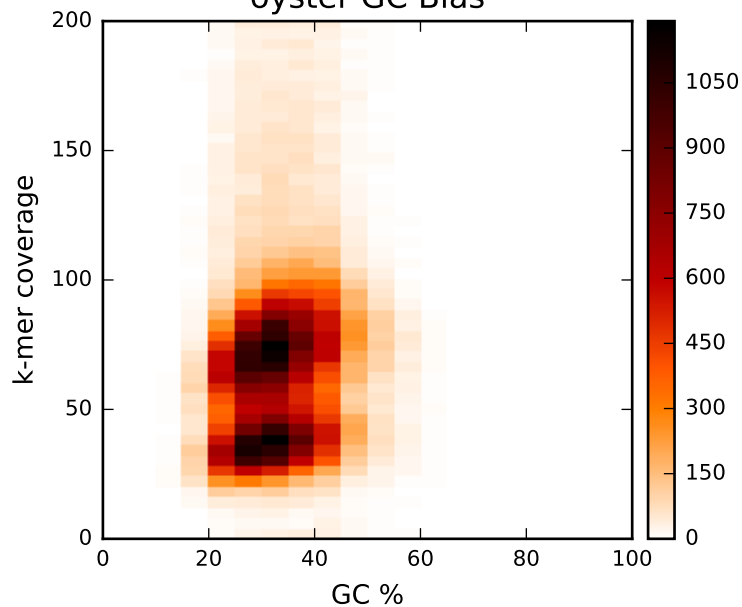


SGA Preqc Results : fig4

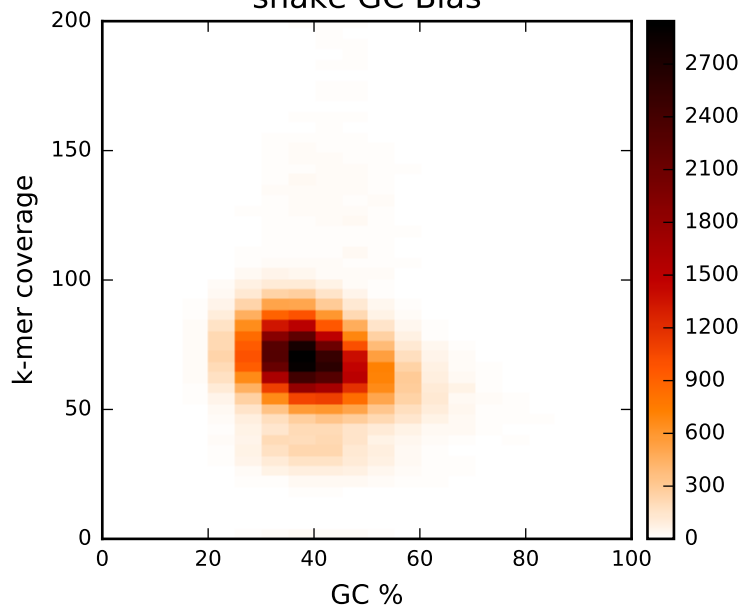
human GC Bias



oyster GC Bias



snake GC Bias



yeast GC Bias

