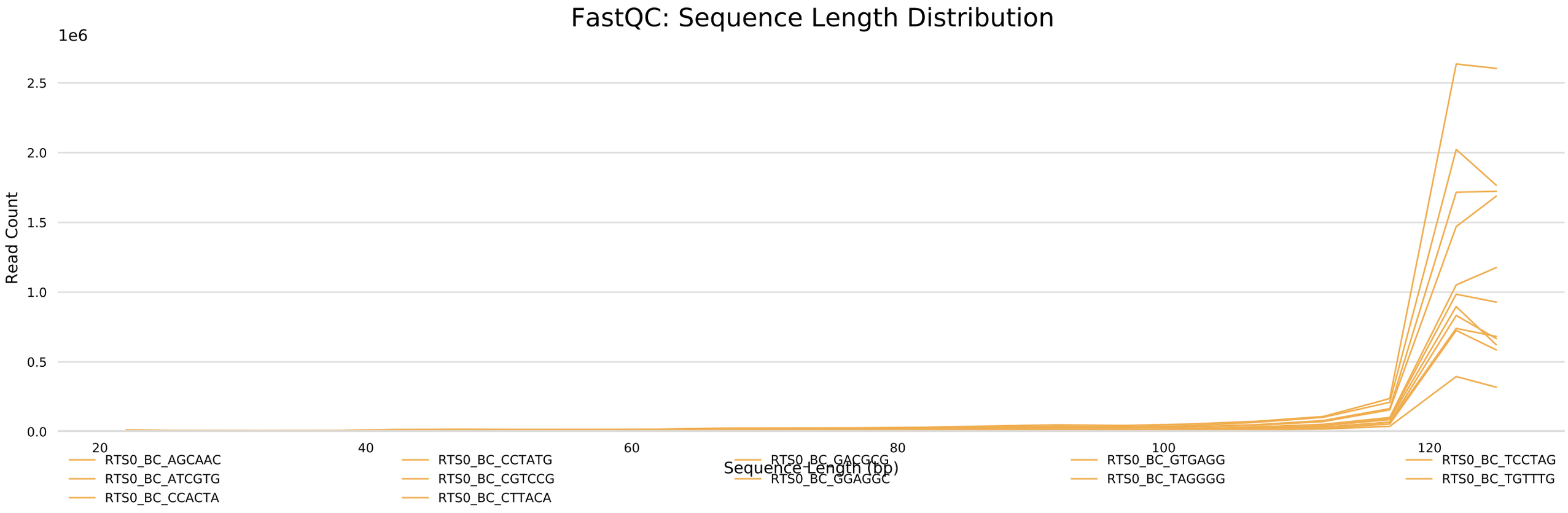
FastQC: Sequence Length Distribution
1e6
2.5
2.0
1.5
1.0
0.5
0.0
Read Count
20
40
60
80
100
120
Sequence Length (bp)
RTS0_BC_AGCAAC
RTS0_BC_ATCGTG
RTS0_BC_CCACTA
RTS0_BC_CCTATG
RTS0_BC_CGTCCG
RTS0_BC_CTTACA
RTS0_BC_GACGCG
RTS0_BC_GGAGGC
RTS0_BC_GTGAGG
RTS0_BC_TAGGGG
RTS0_BC_TCCTAG
RTS0_BC_TGTTTG