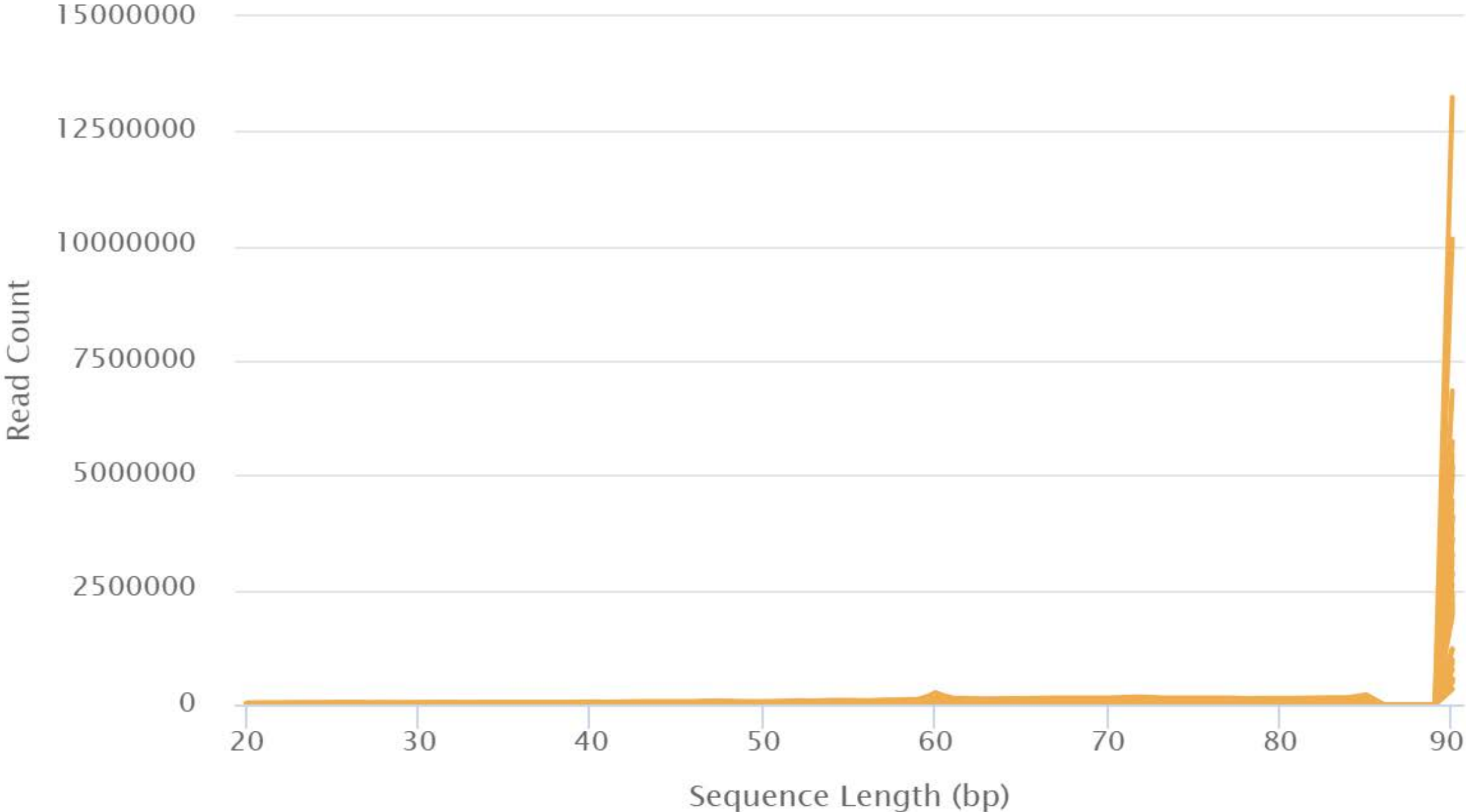
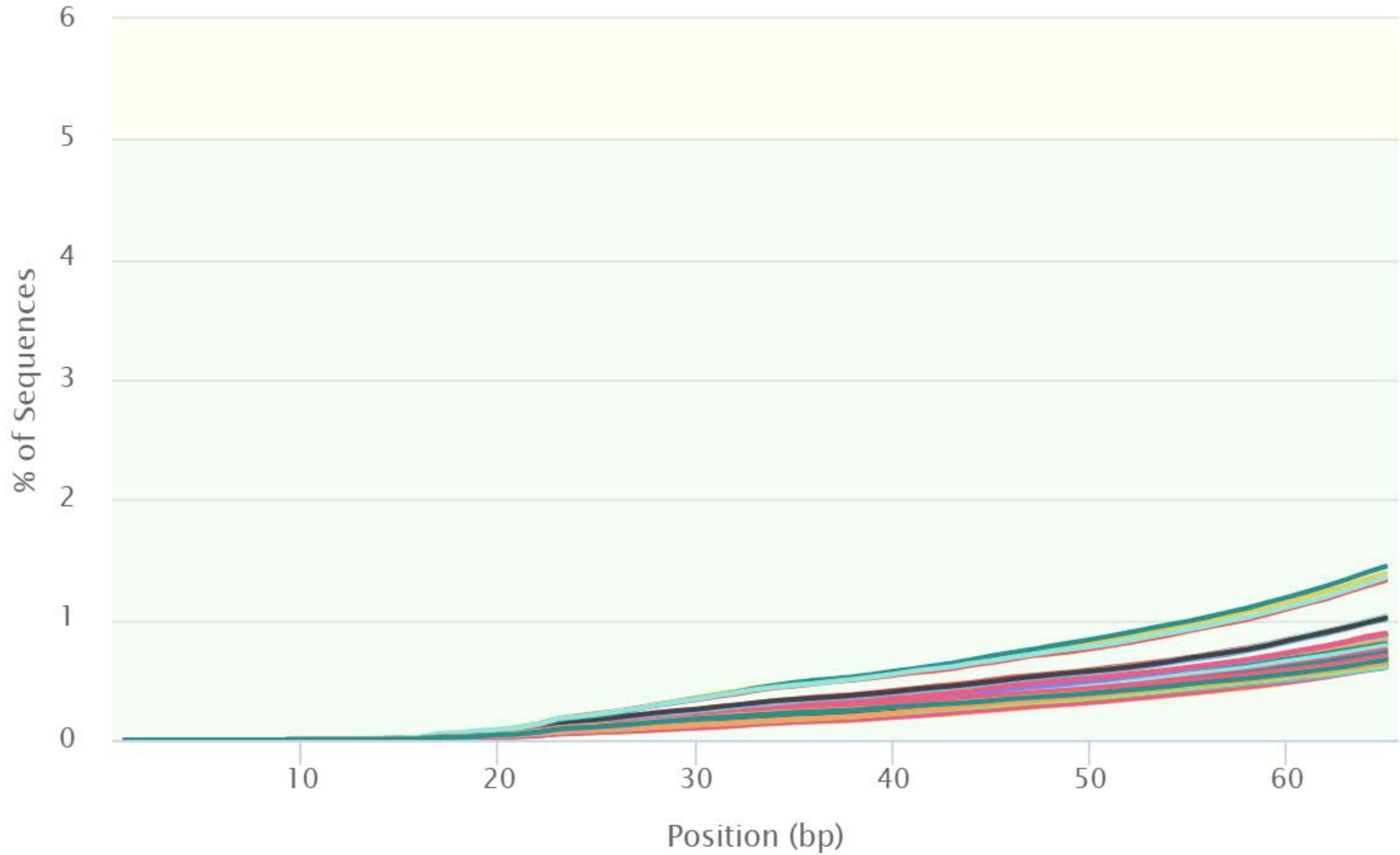


# Supplement VII

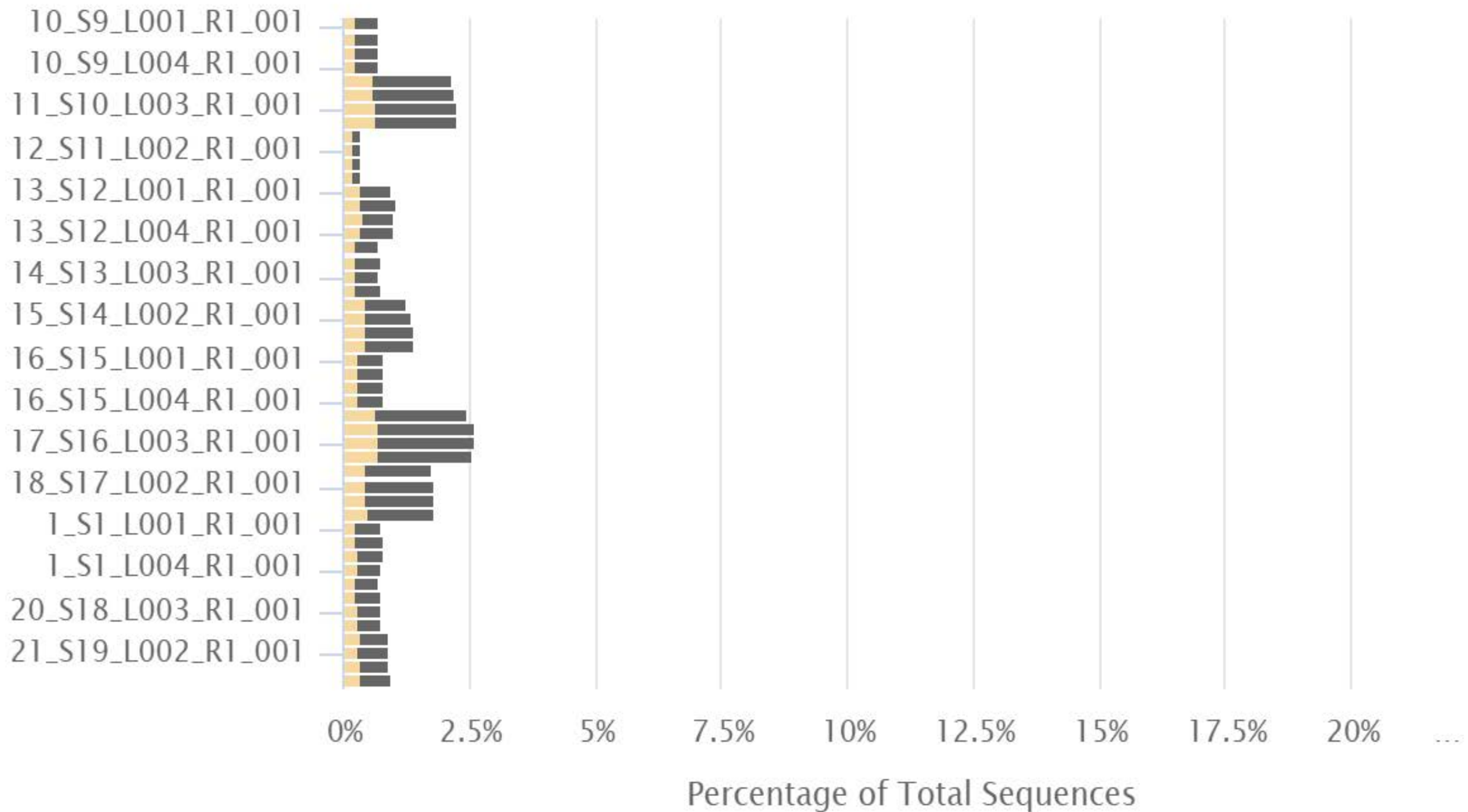
## FastQC: Sequence Length Distribution



# FastQC: Adapter Content



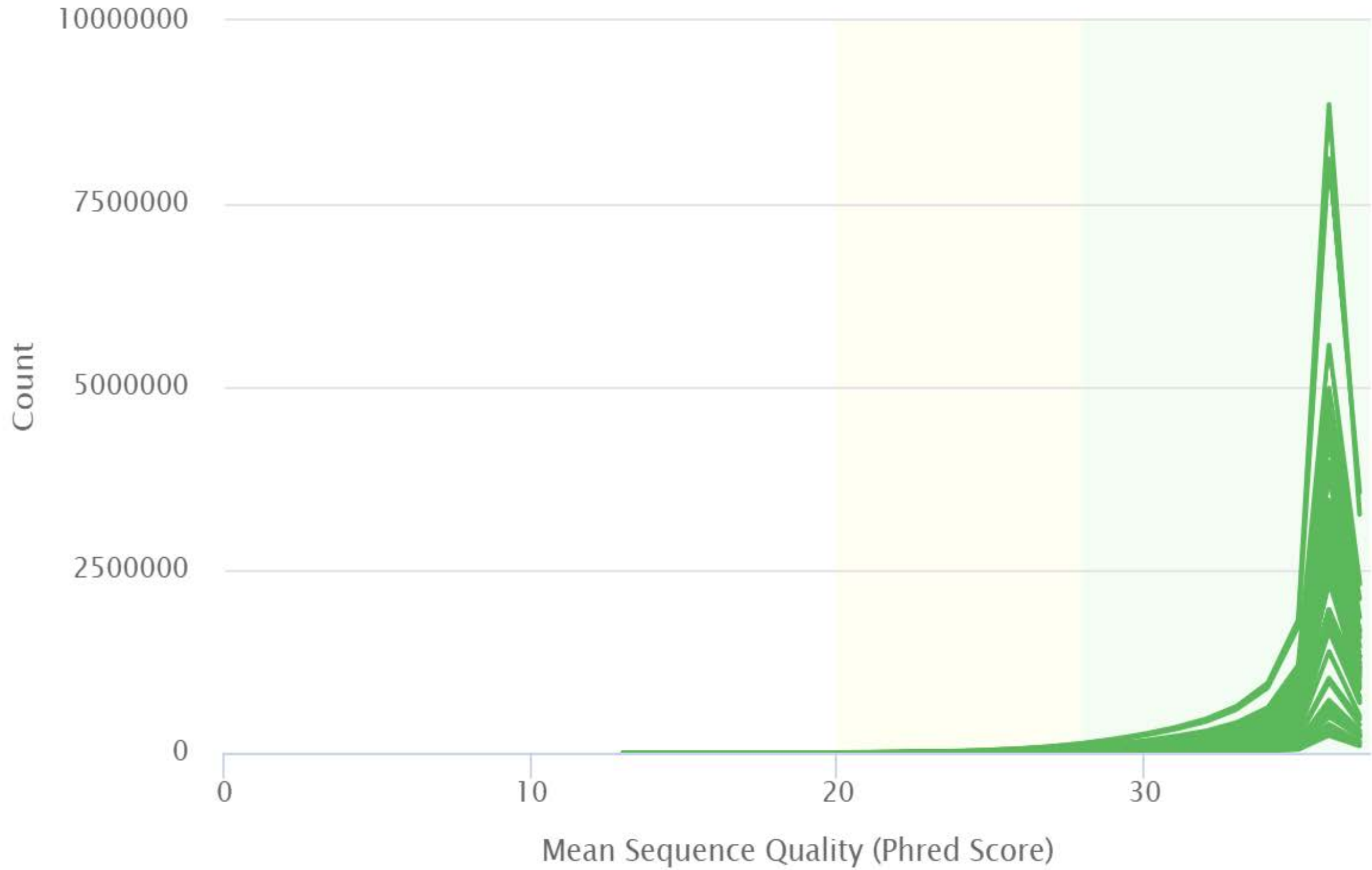
# FastQC: Overrepresented sequences



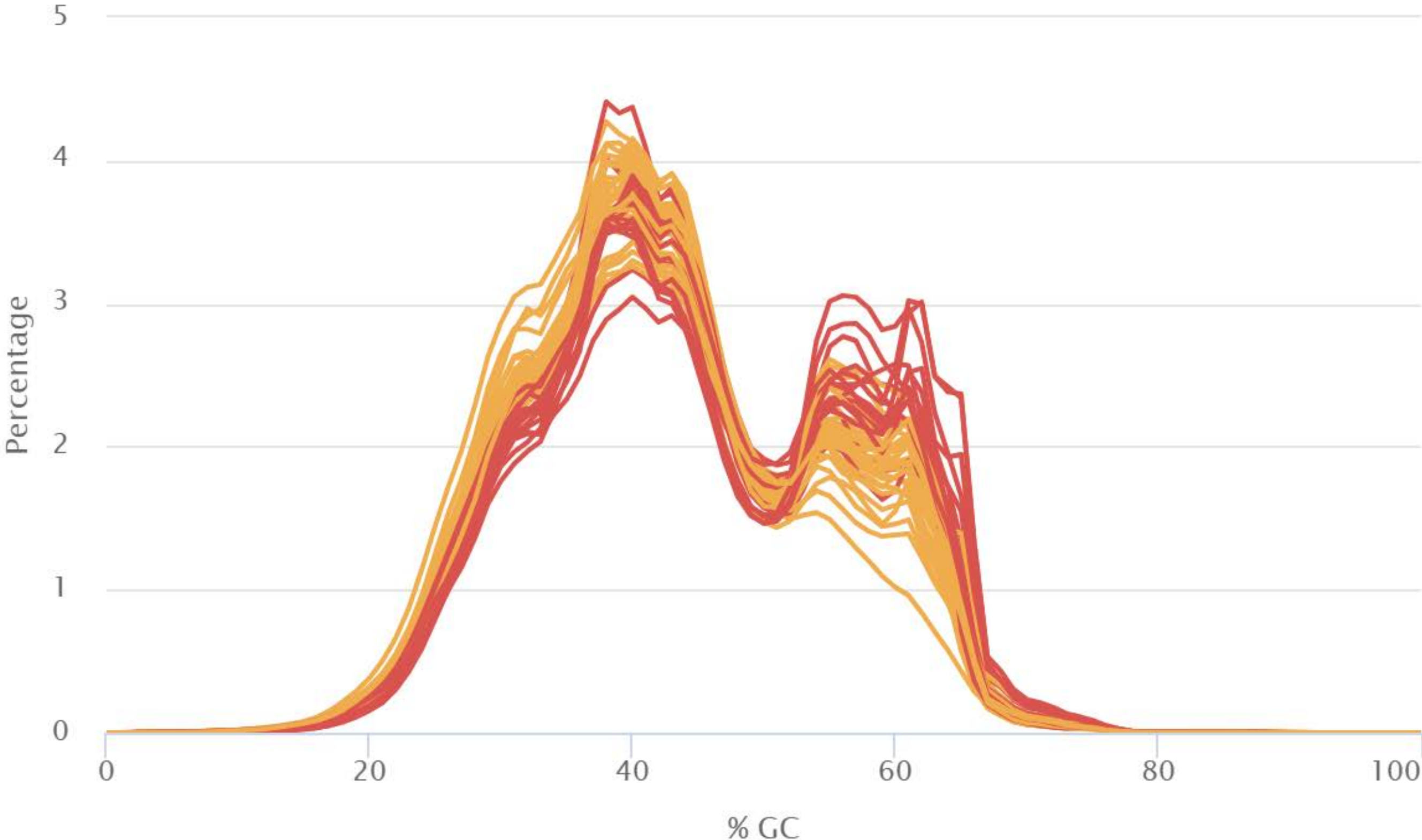
● Top over-represented sequence

● Sum of remaining over-represented sequences

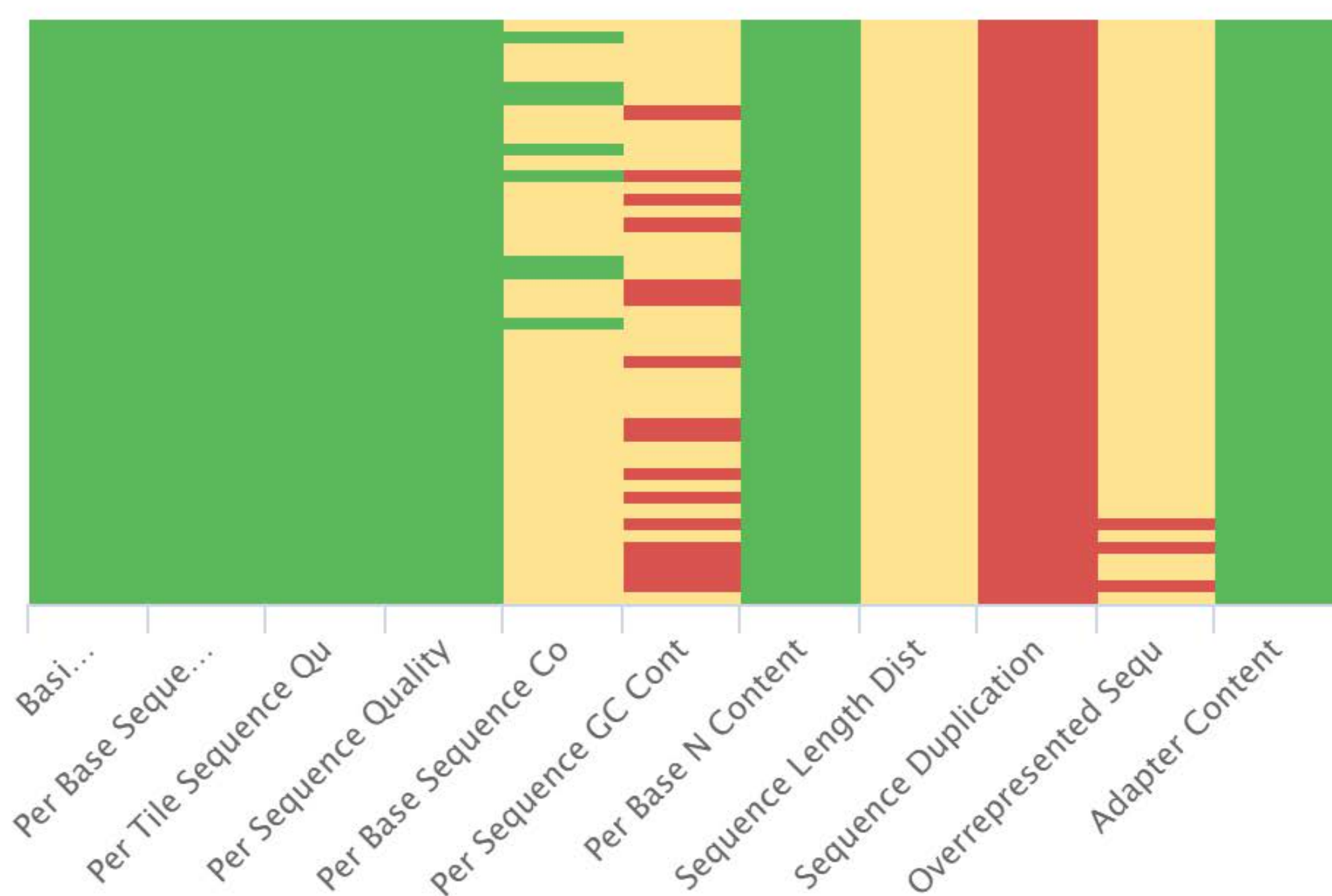
# FastQC: Per Sequence Quality Scores



# FastQC: Per Sequence GC Content



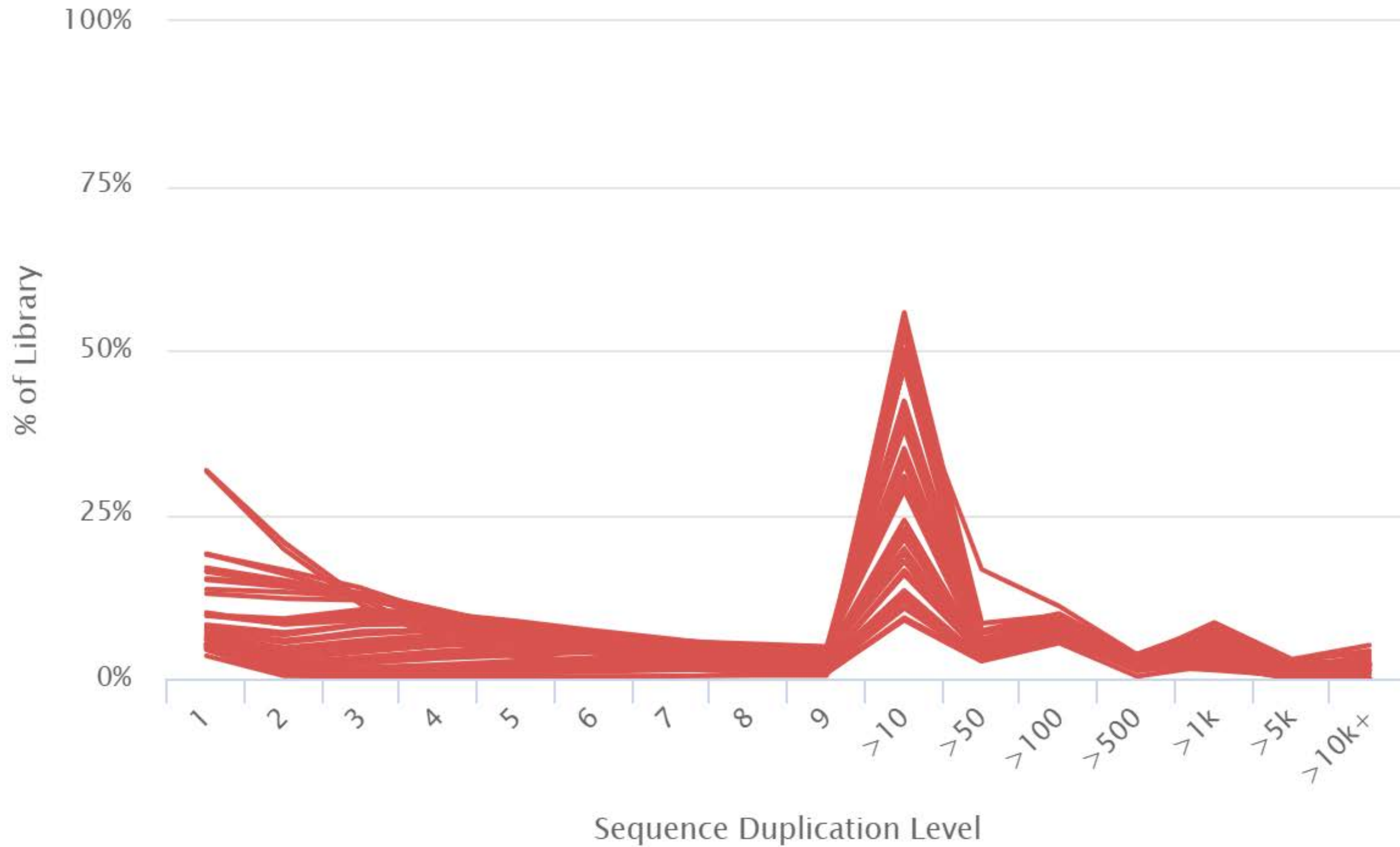
## FastQC: Status Checks



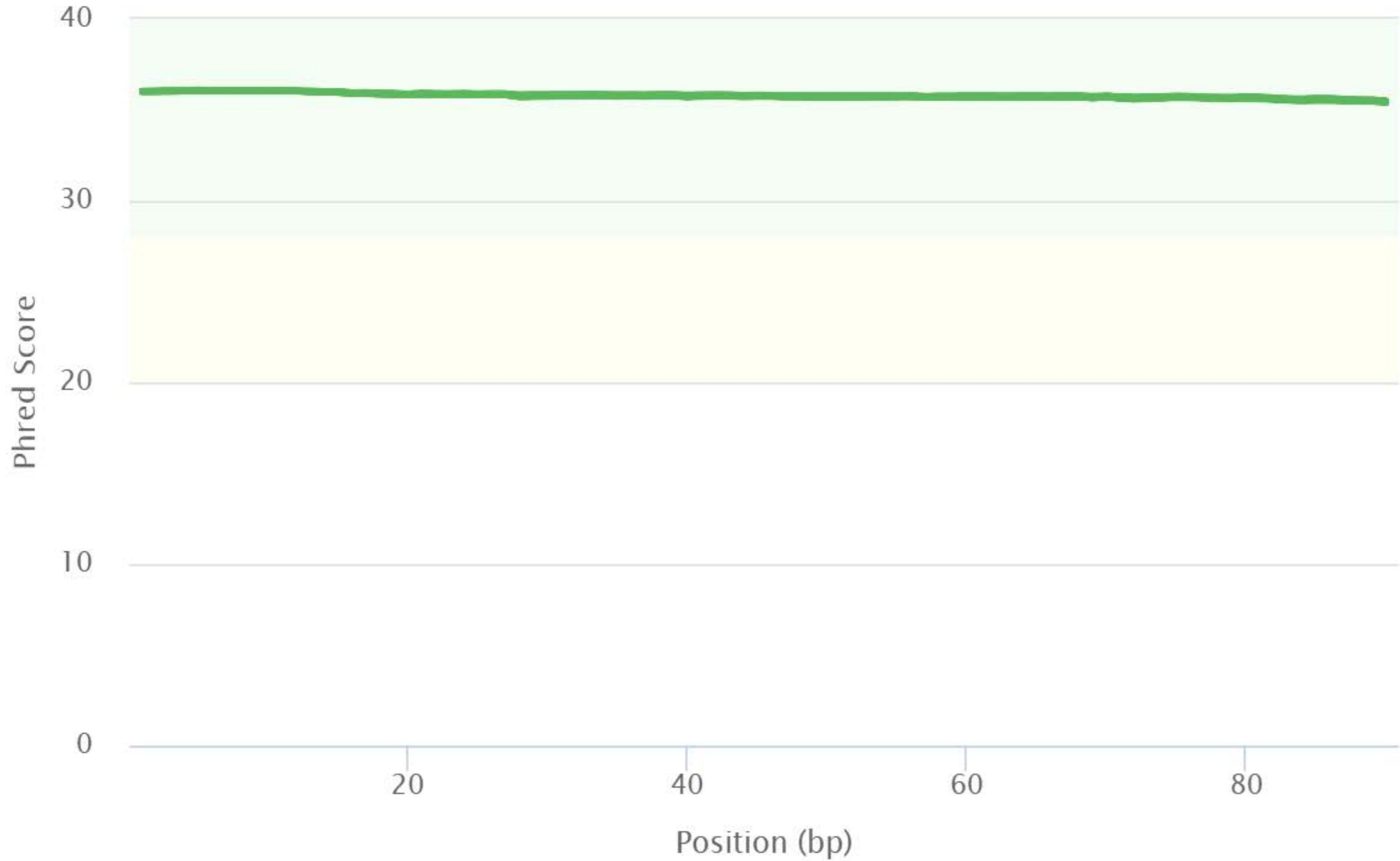
trim\_NA\_Eriksson\_53\_  
 trim\_NA\_Eriksson\_57\_  
 trim\_NA\_Eriksson\_60\_  
 trim\_NA\_Eriksson\_64\_  
 trim\_NA\_Eriksson\_69\_  
 trim\_NA\_Eriksson\_72\_  
 trim\_NA\_Eriksson\_76\_  
 trim\_NA\_Eriksson\_7\_S  
 trim\_NA\_Eriksson\_83\_  
 trim\_NA\_Eriksson\_87\_  
 trim\_NA\_Eriksson\_90\_  
 trim\_NA\_Eriksson\_94\_

Section Name

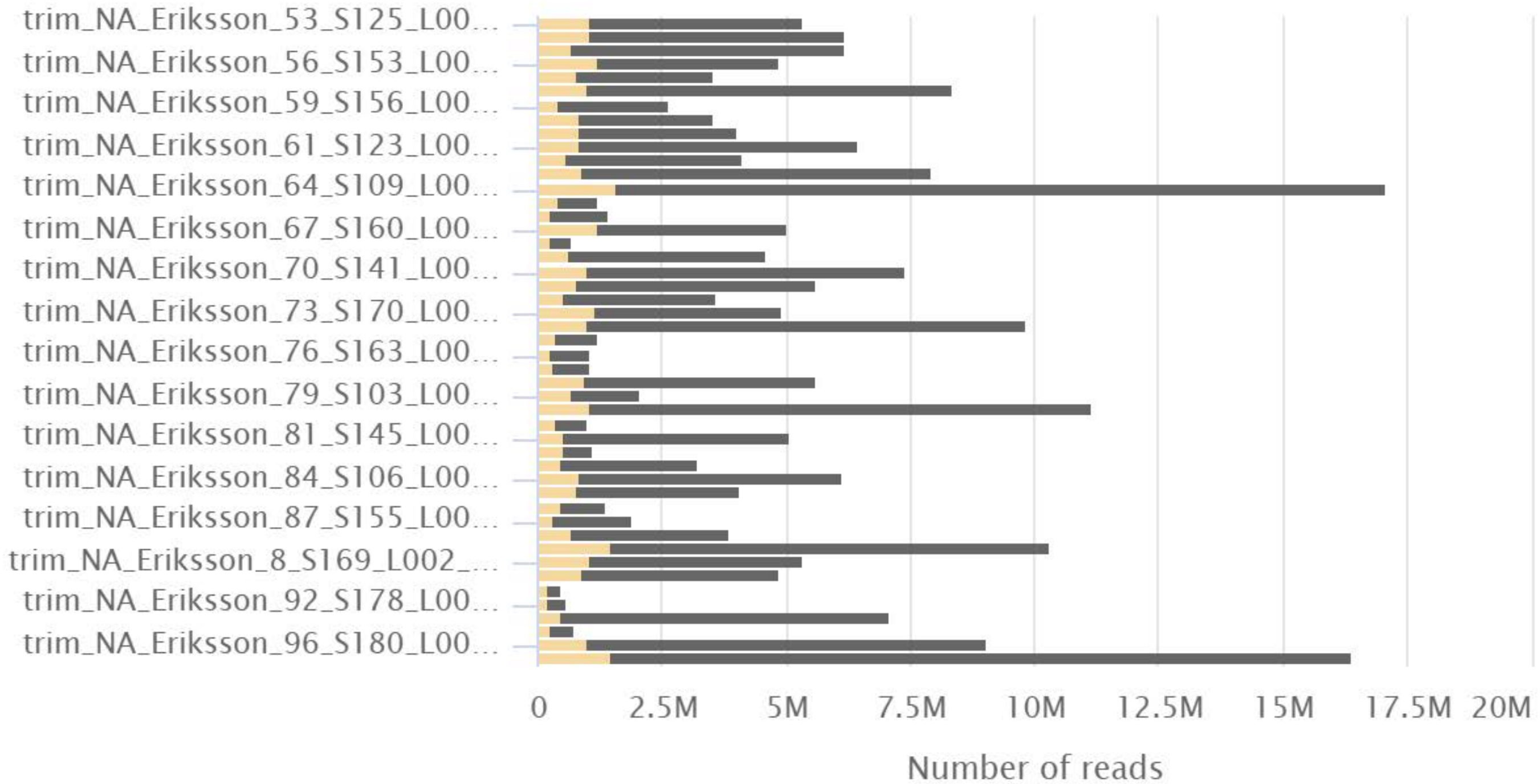
# FastQC: Sequence Duplication Levels



# FastQC: Mean Quality Scores



# FastQC: Sequence Counts



● Unique Reads ● Duplicate Reads