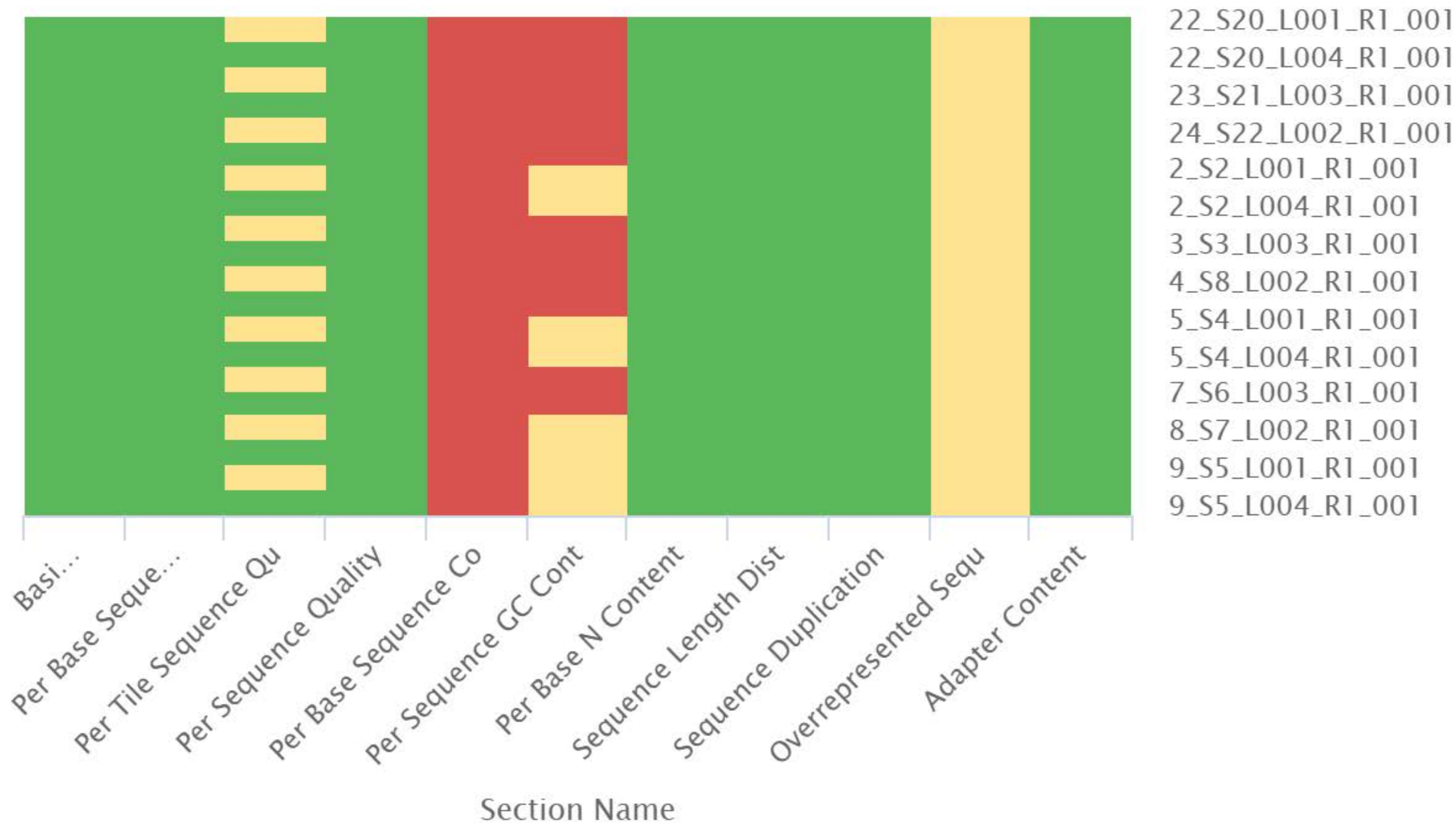
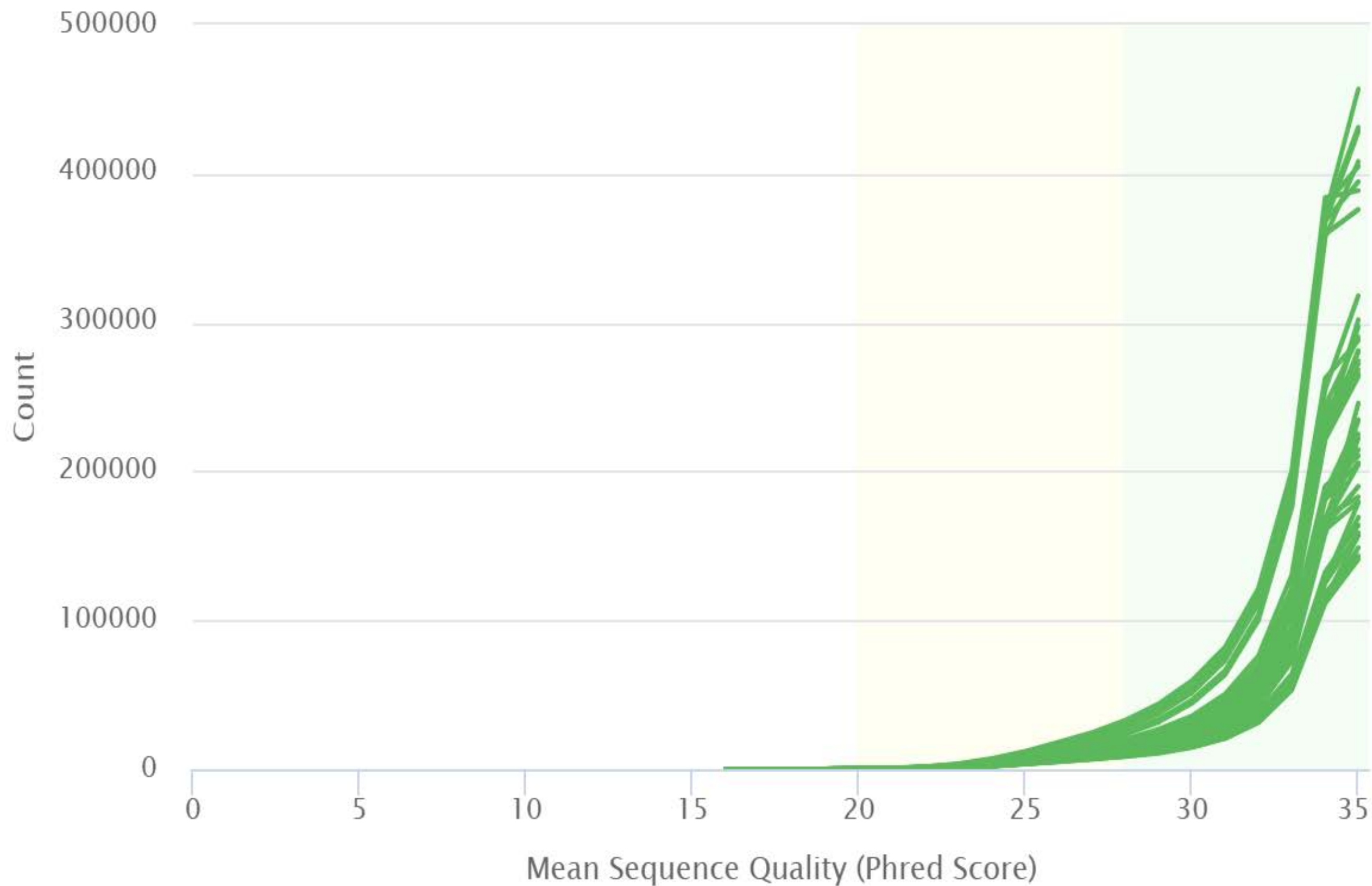


# Supplement X

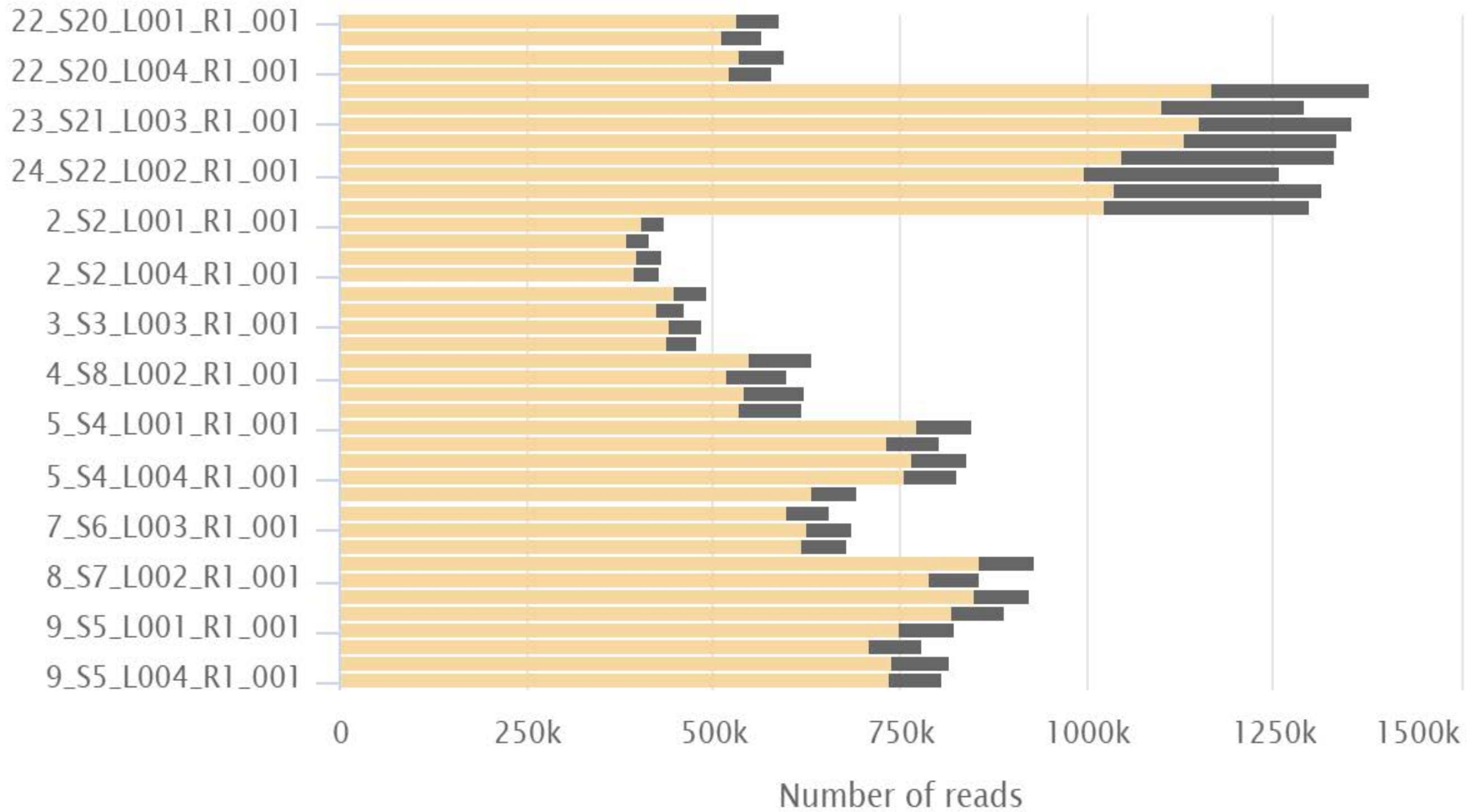
## FastQC: Status Checks



# FastQC: Per Sequence Quality Scores

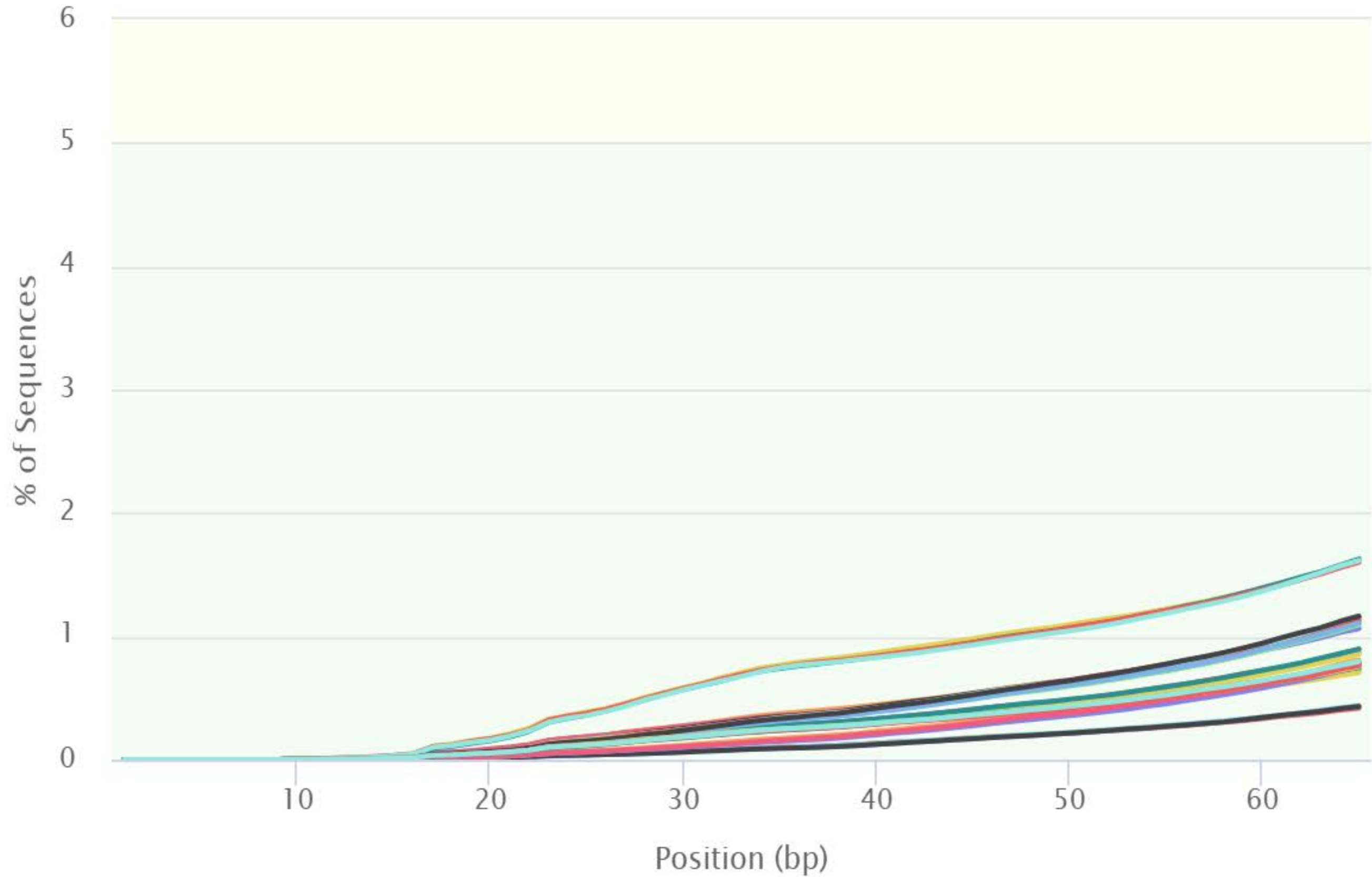


# FastQC: Sequence Counts

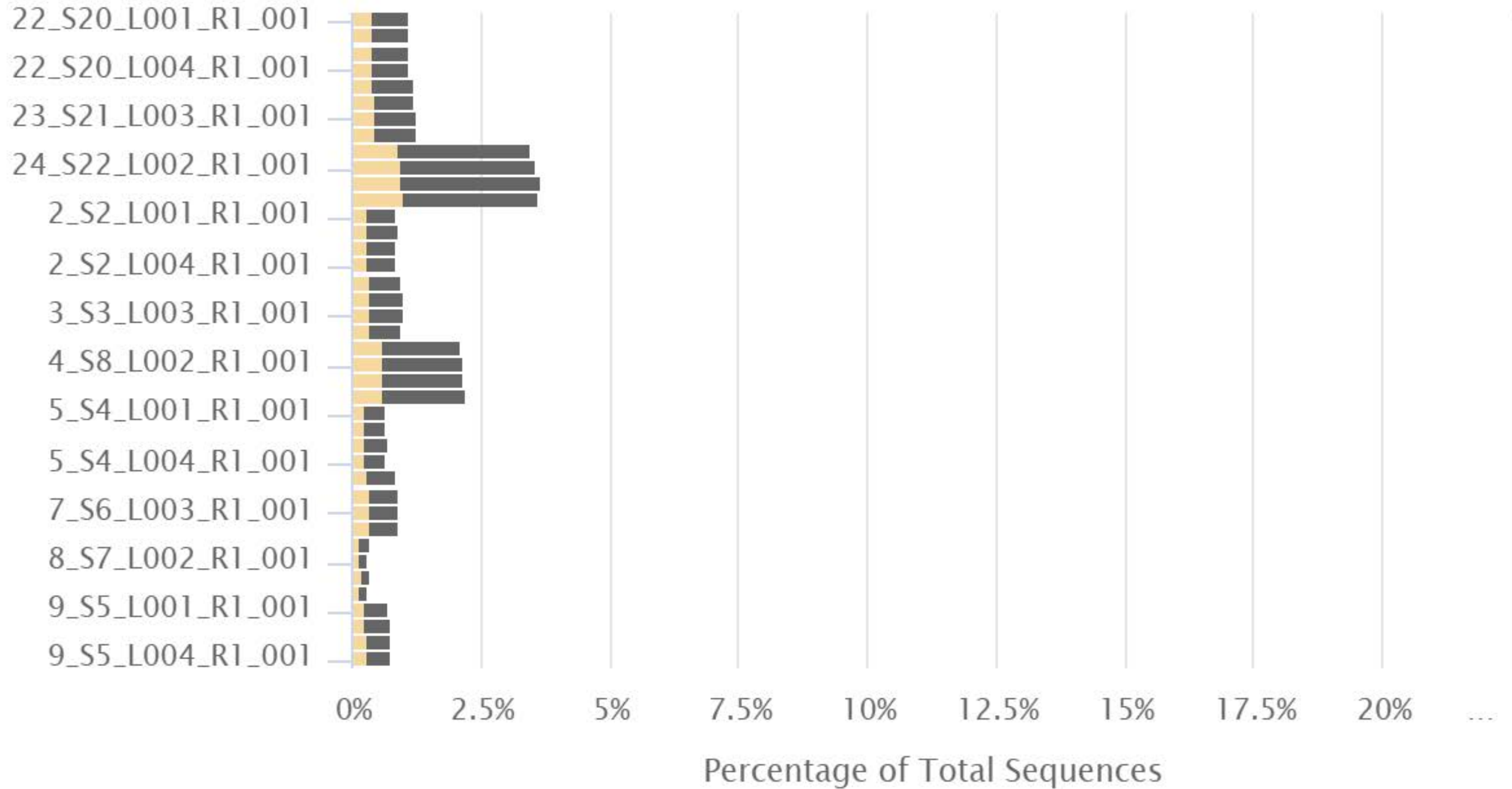


● Unique Reads ● Duplicate Reads

# FastQC: Adapter Content



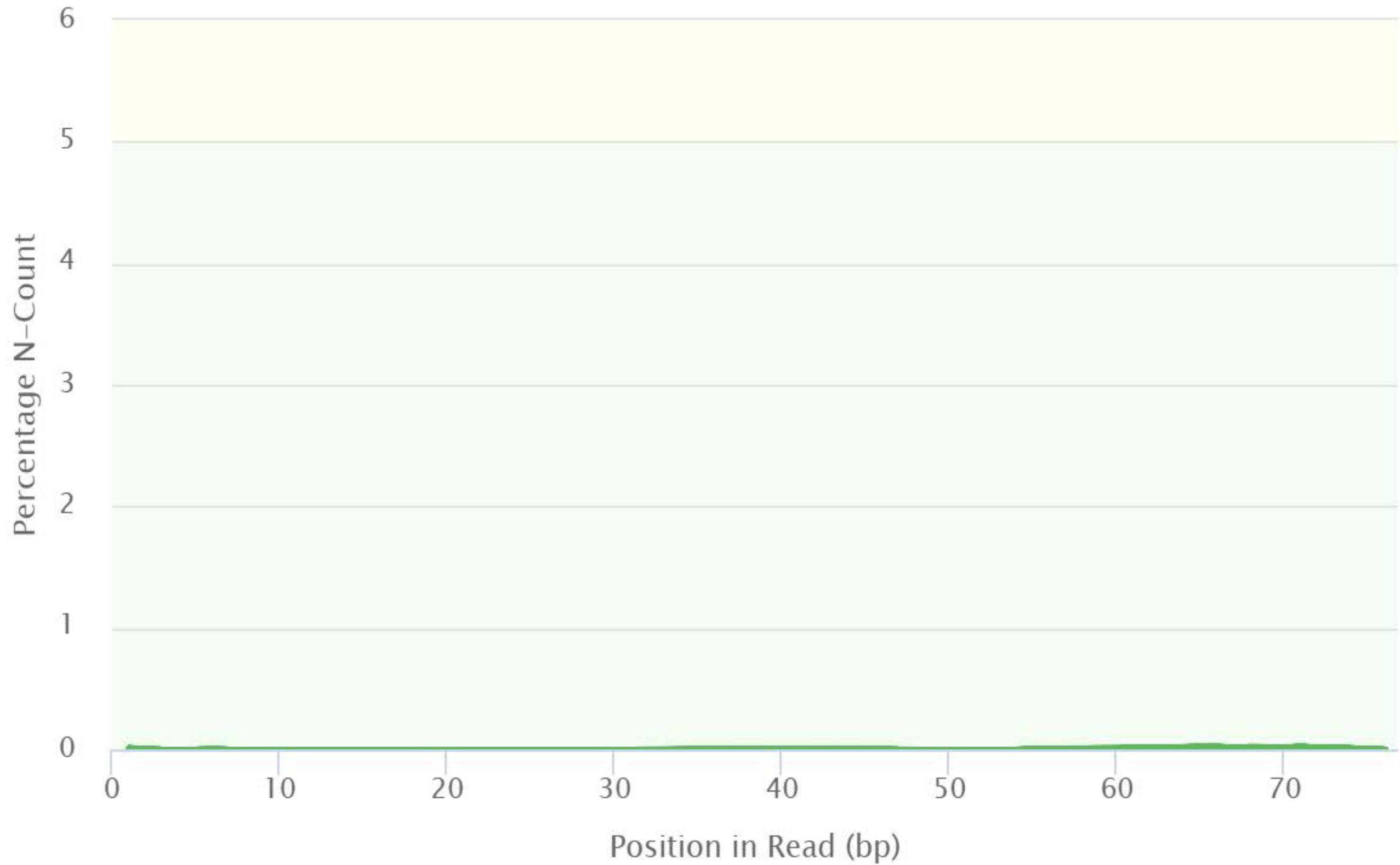
# FastQC: Overrepresented sequences



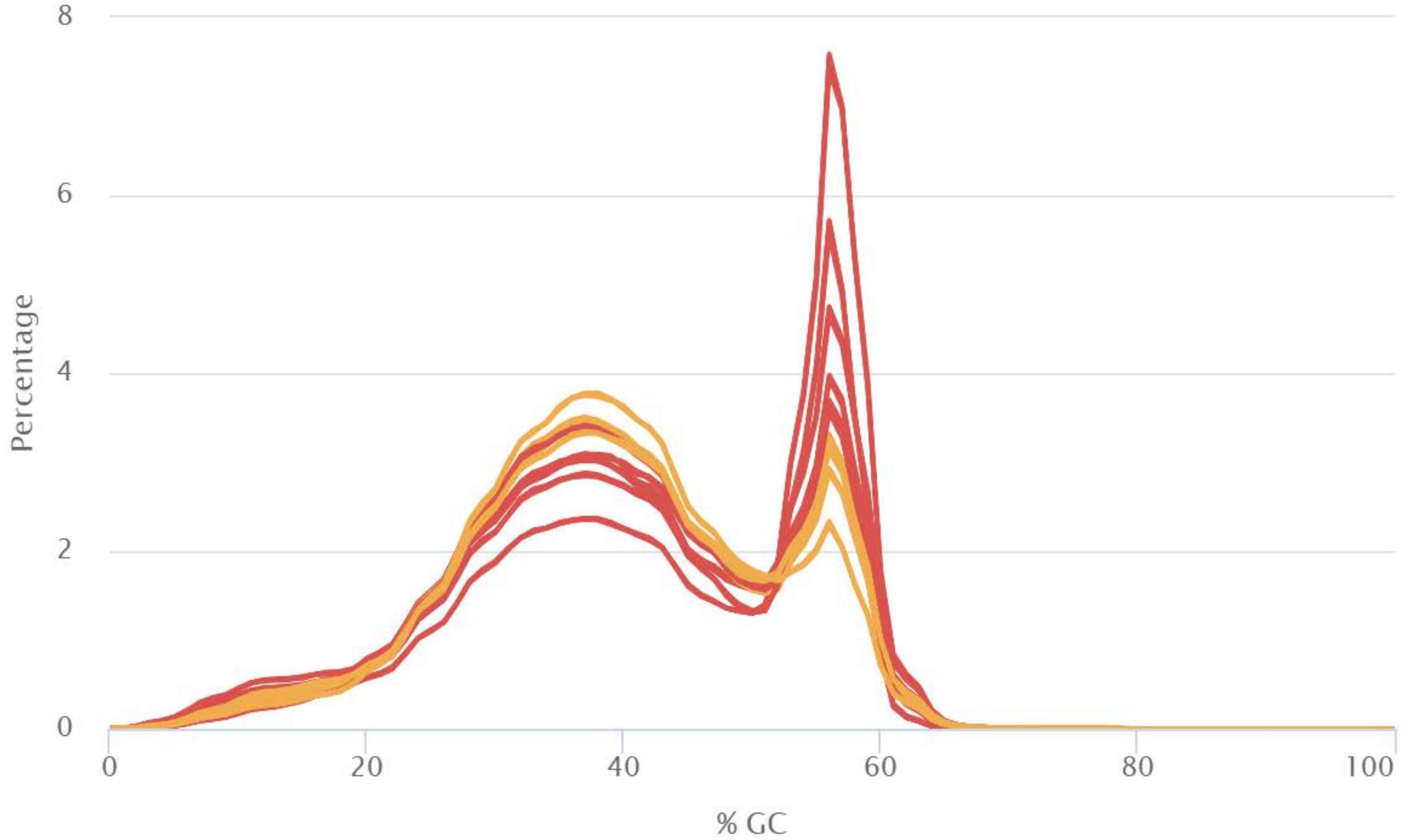
● Top over-represented sequence

● Sum of remaining over-represented sequences

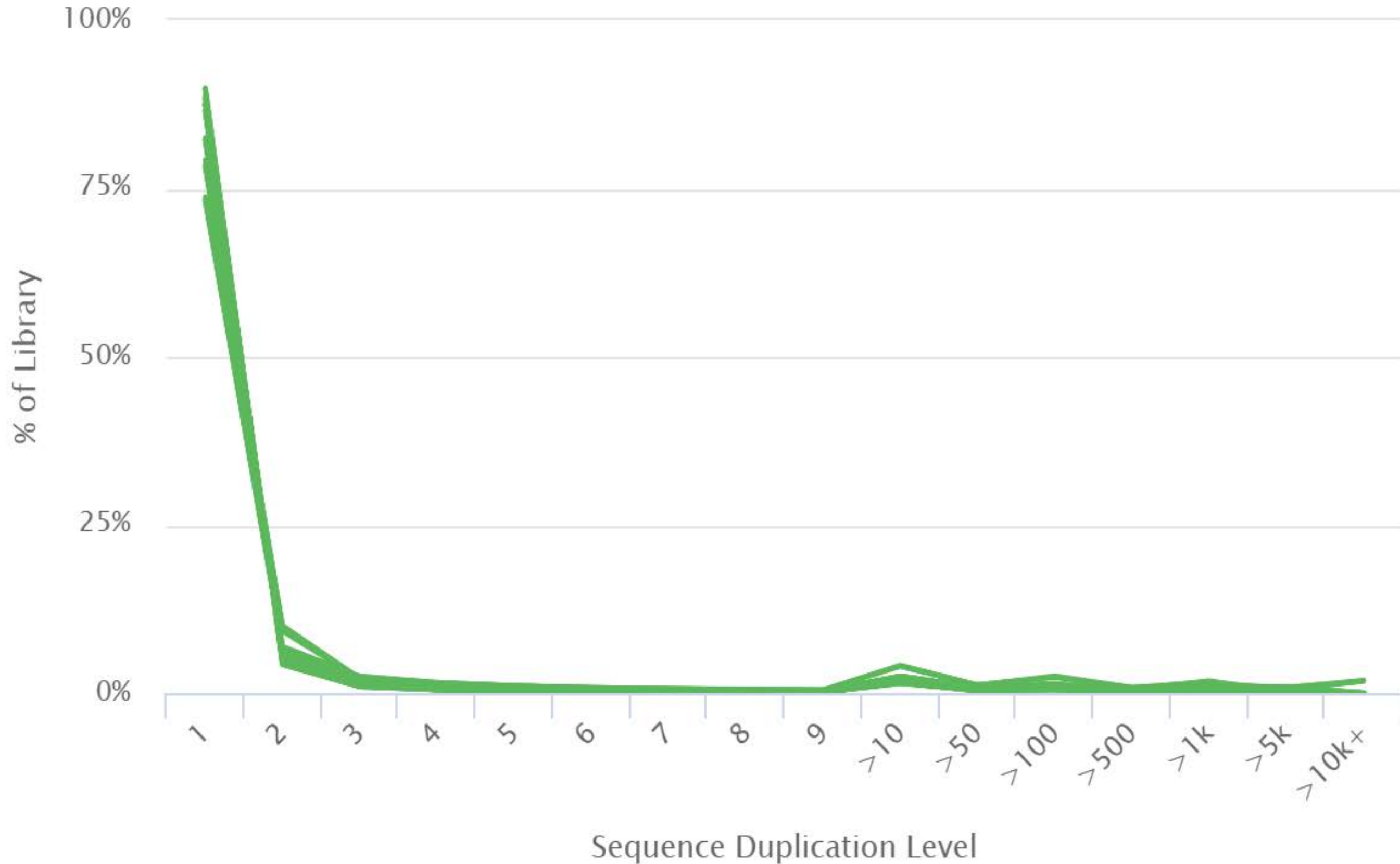
# FastQC: Per Base N Content



# FastQC: Per Sequence GC Content



# FastQC: Sequence Duplication Levels



# FastQC: Mean Quality Scores

