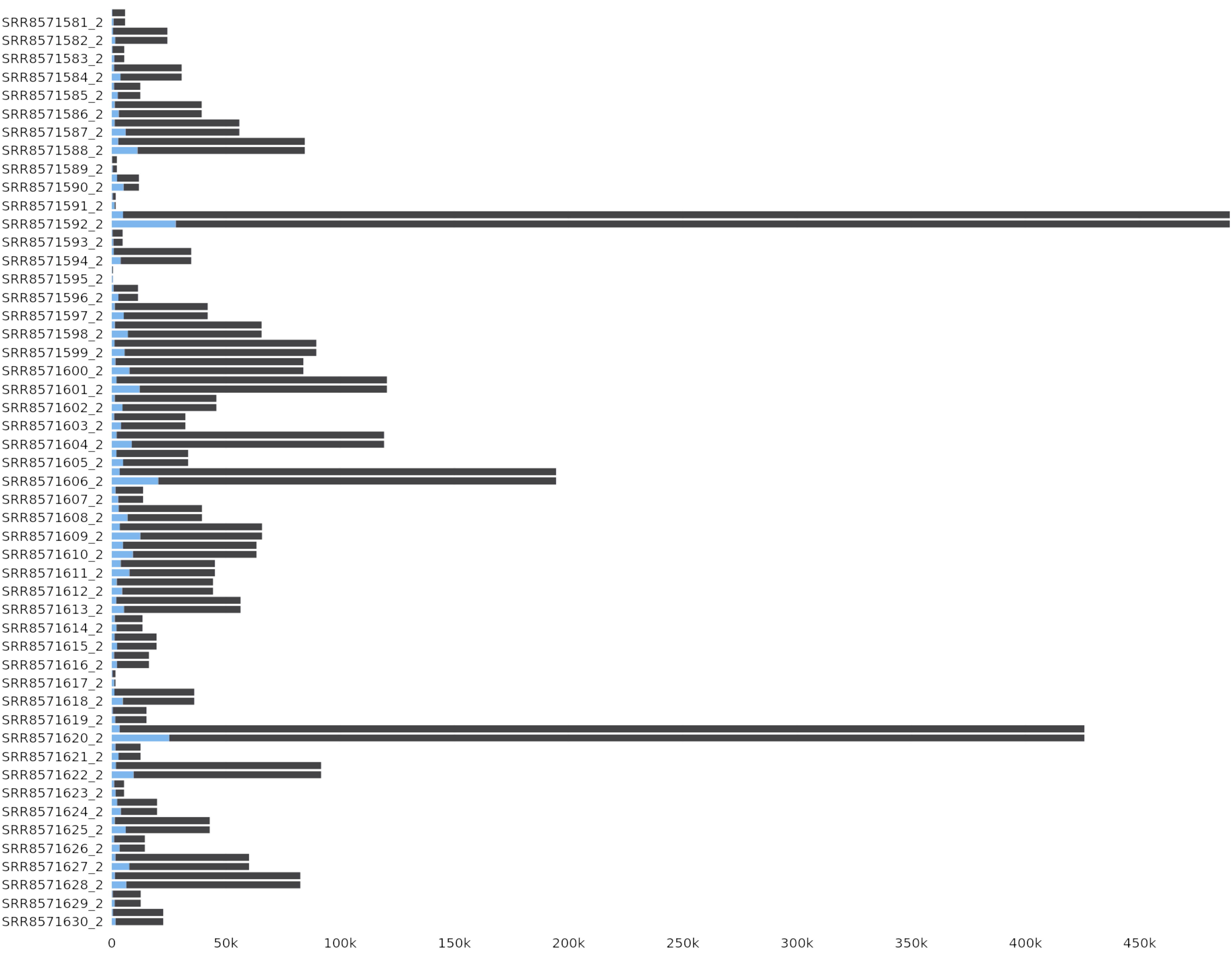


FastQC: Sequence Counts

Unique Reads
Duplicate Reads



Number of reads