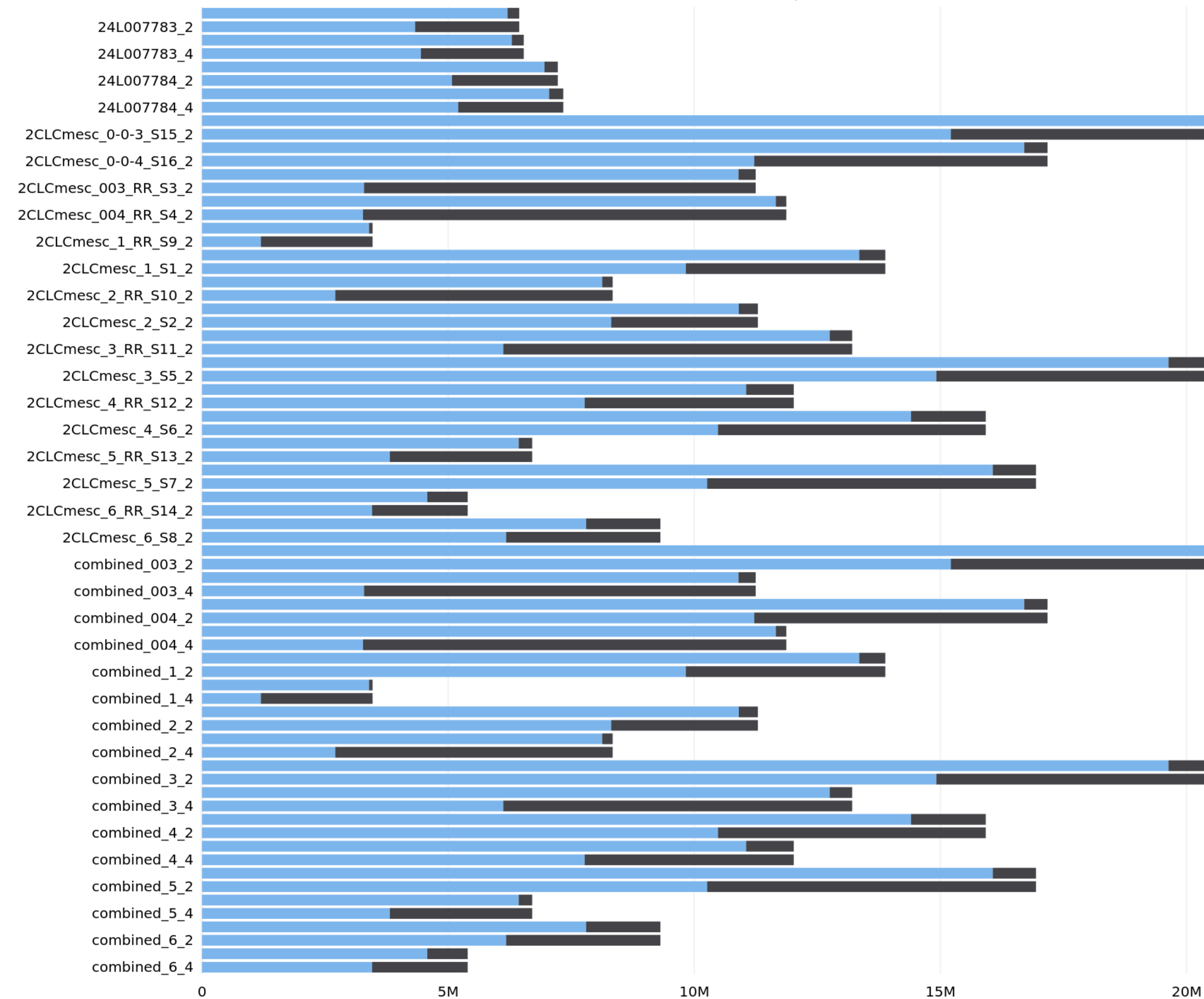


# FastQC: Sequence Counts

72 samples

Unique Reads  
Duplicate Reads



Number of reads