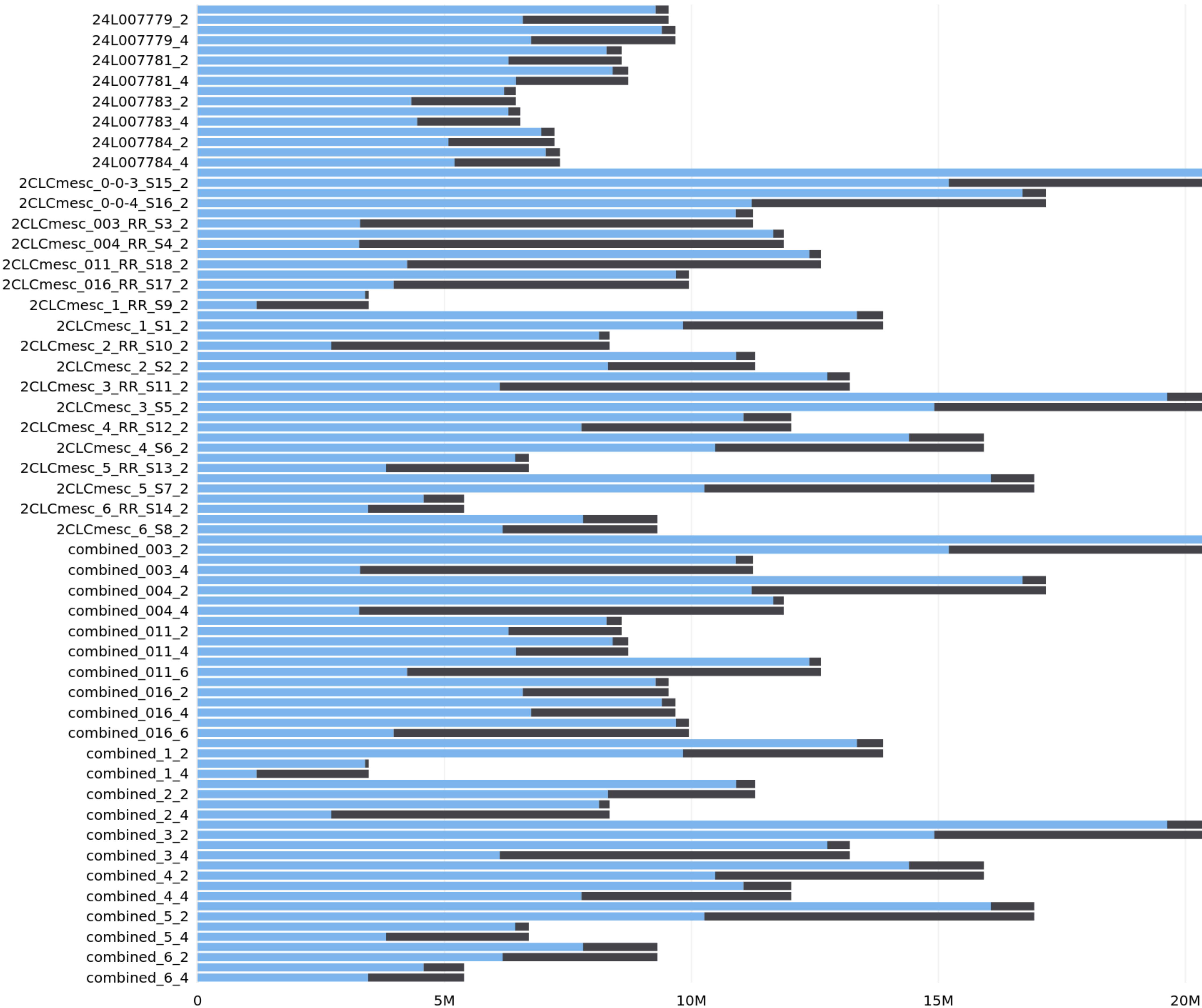


# FastQC: Sequence Counts

96 samples

Unique Reads  
Duplicate Reads



0

5M

10M

15M

20M

Number of reads