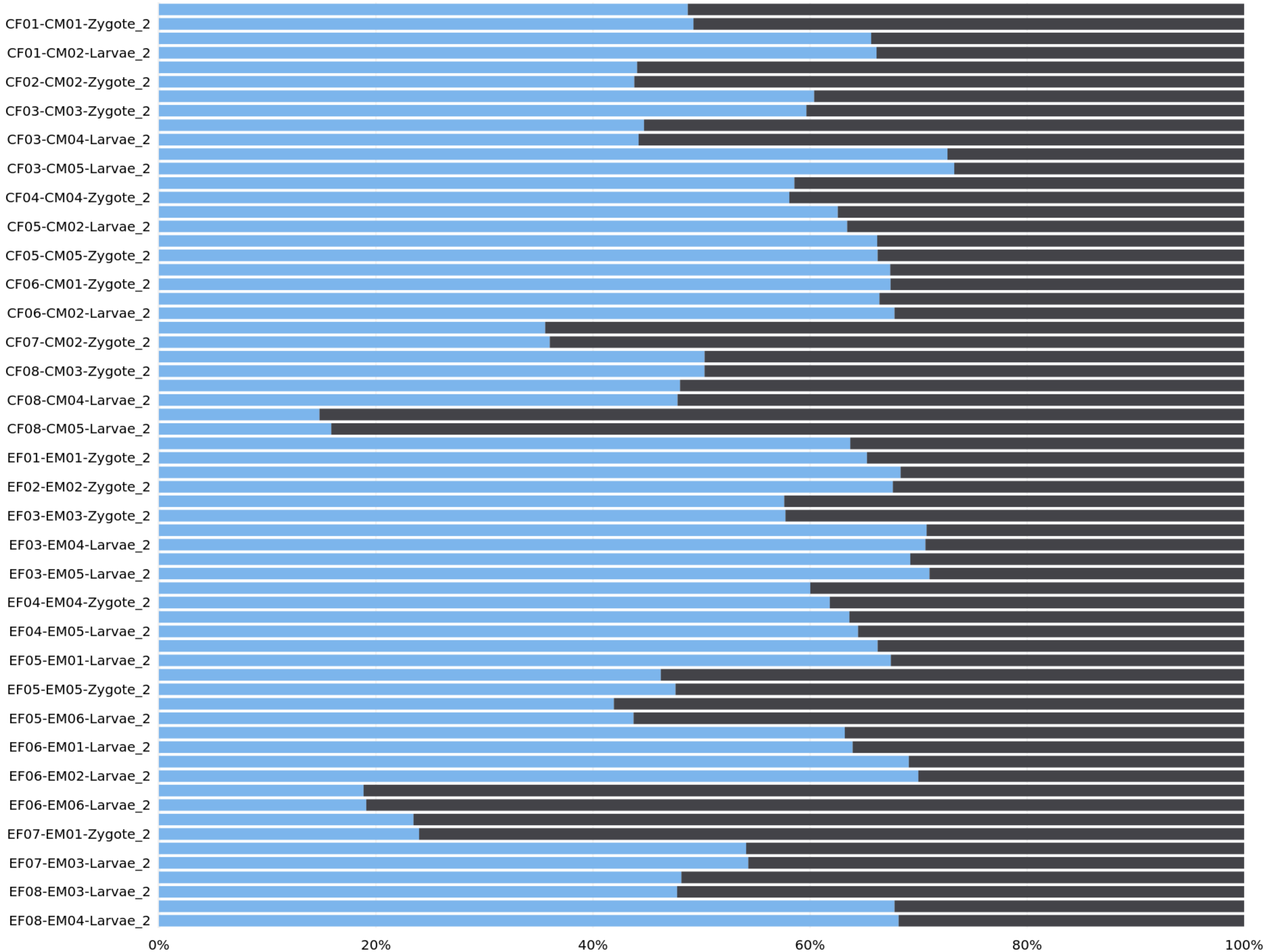


FastQC: Sequence Counts

64 samples

Unique Reads
Duplicate Reads



0%

20%

40%

60%

80%

100%

Number of reads