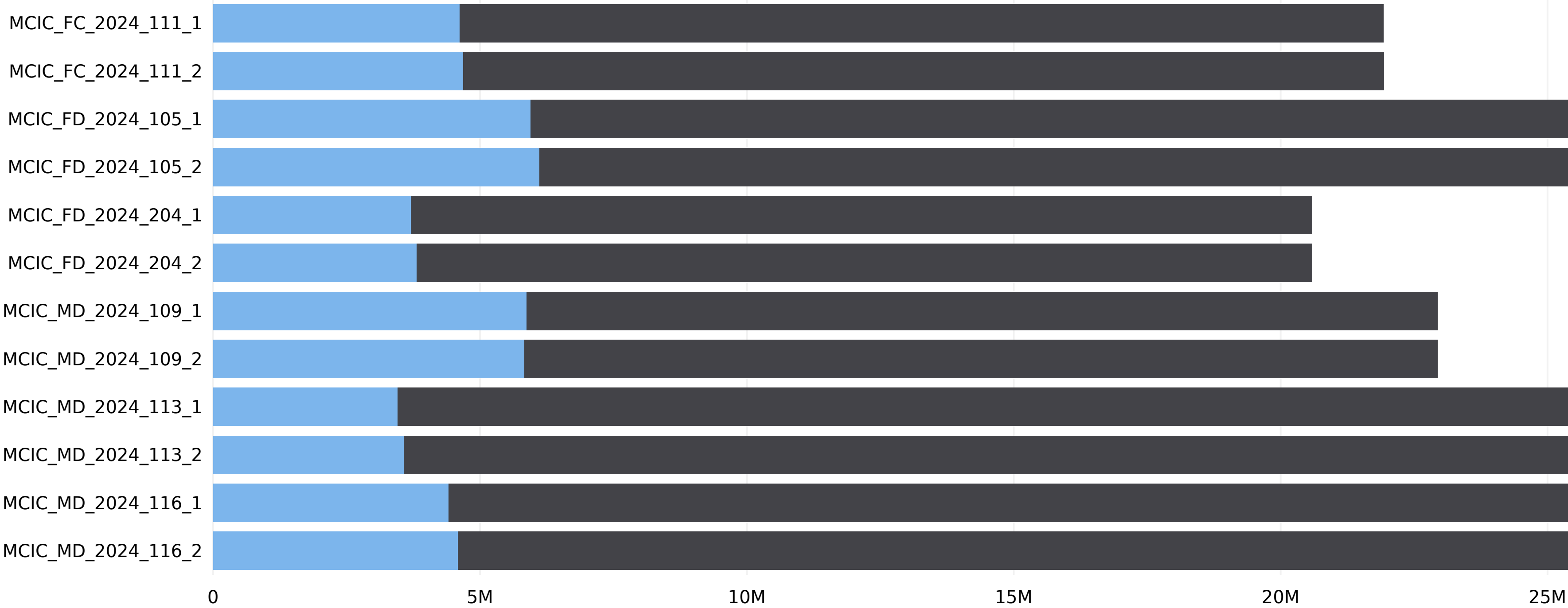


# FastQC: Sequence Counts

12 samples

Unique Reads  
Duplicate Reads



Number of reads