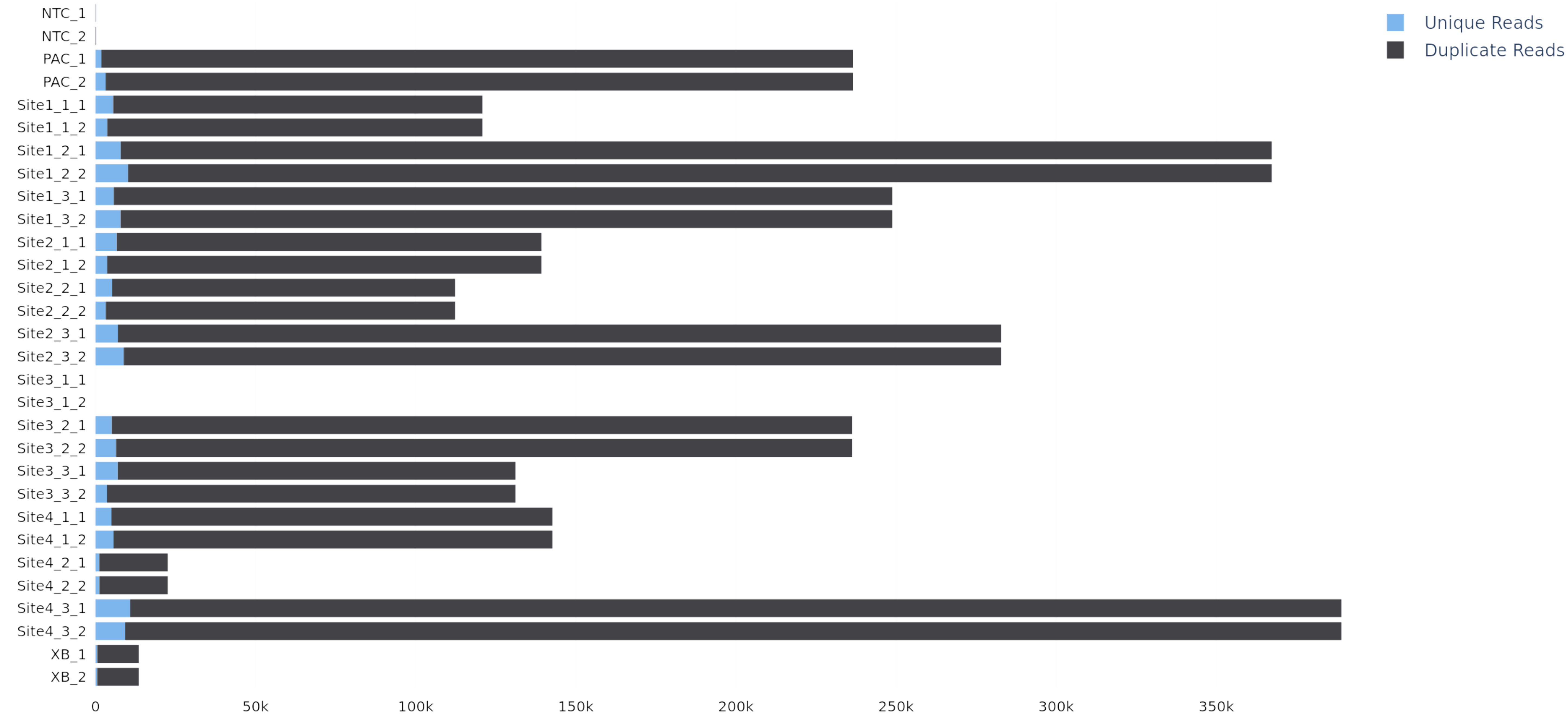


# FastQC: Sequence Counts



Number of reads