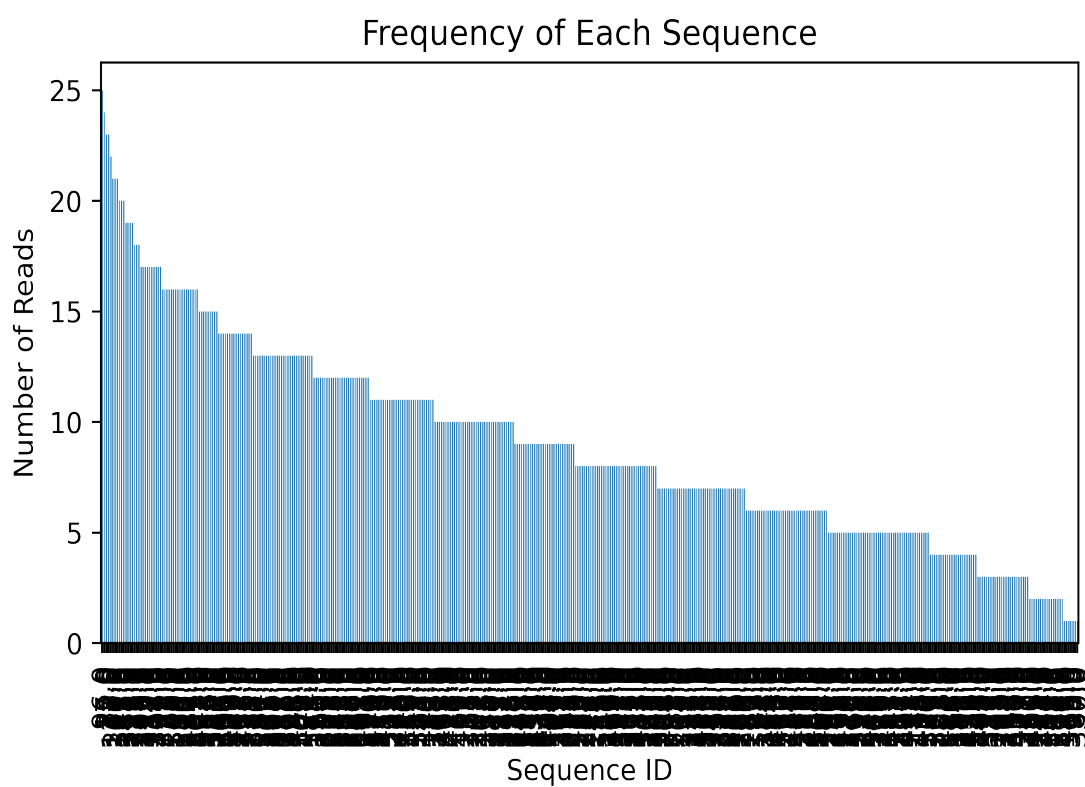


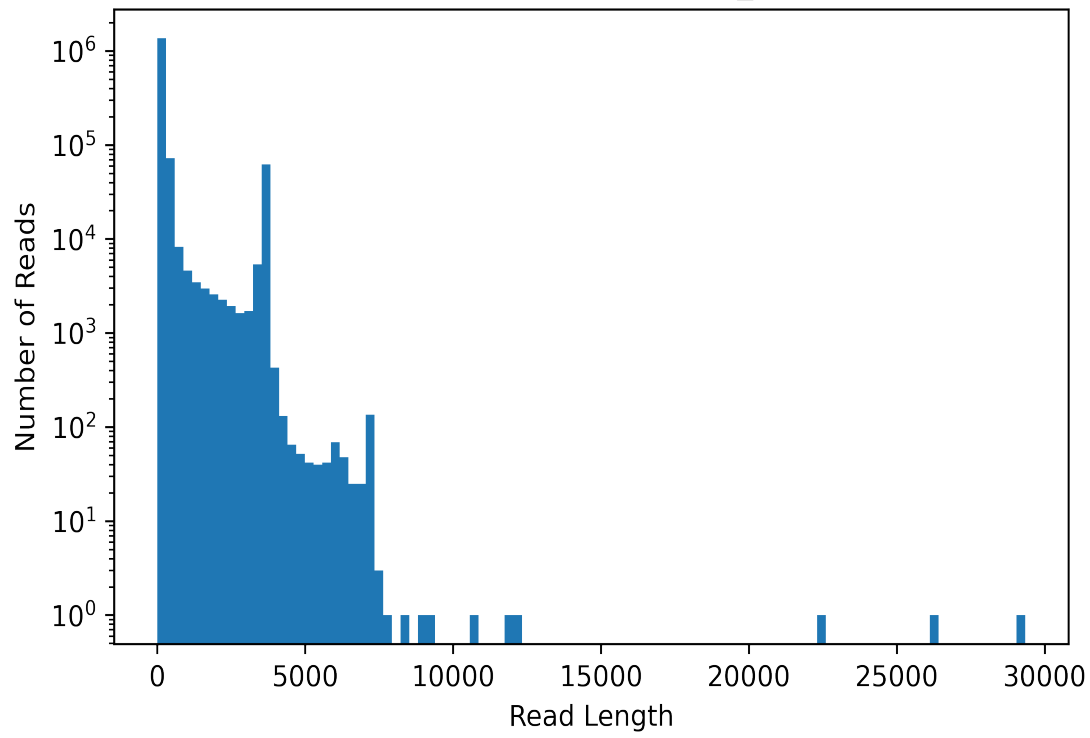
This document contains the analysis of the baseline CRISPR-Cas9 Similarity Search Project Experiment
performed on 20220907

Total Number of Reads in FASTQ_PASS directory: 1536905

Number of aligned reads (no duplicate reads):4052



Read Distribution of all FASTQ_PASS Data



Read Distribution of FASTQ_PASS Data Below Length 500

