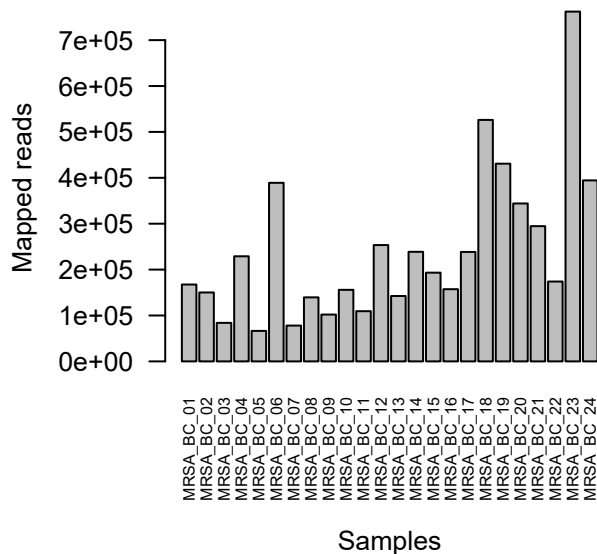
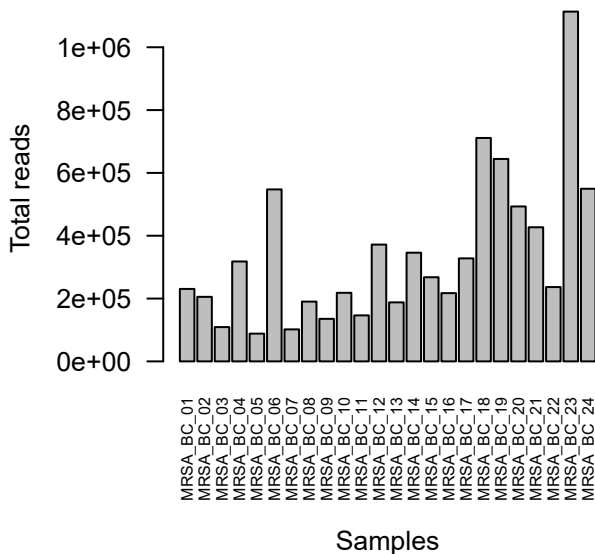
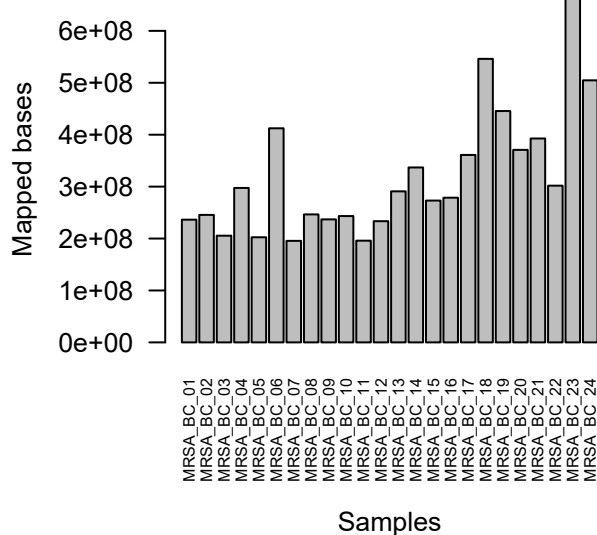
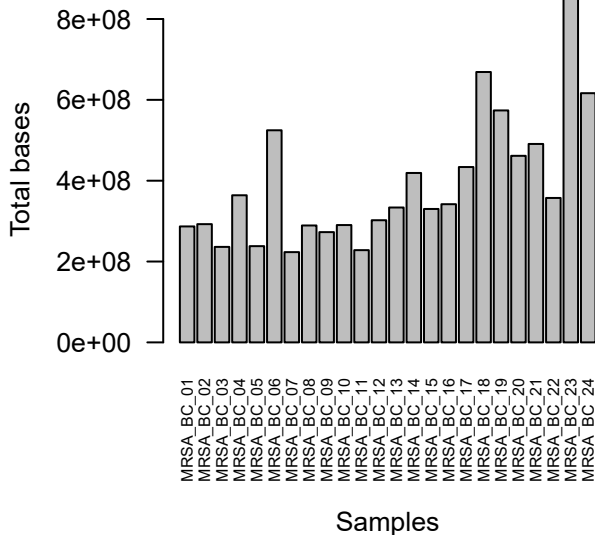
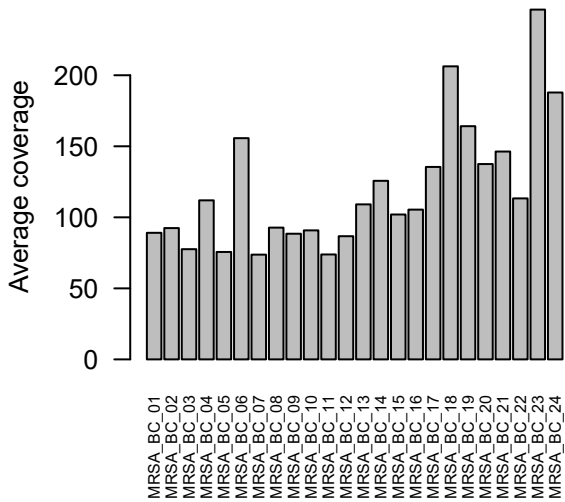
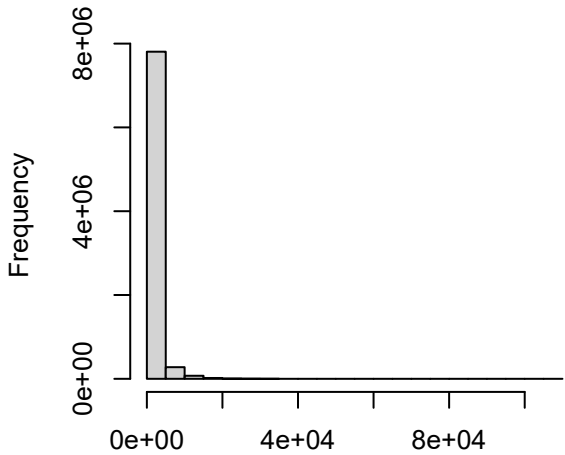
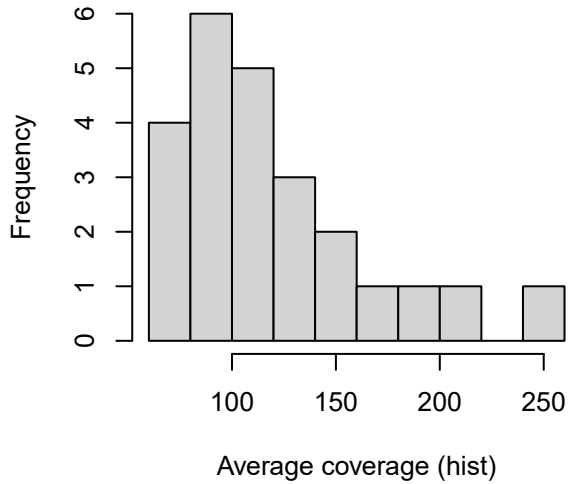


Basic statistics (MRSA, Nanopore-only)

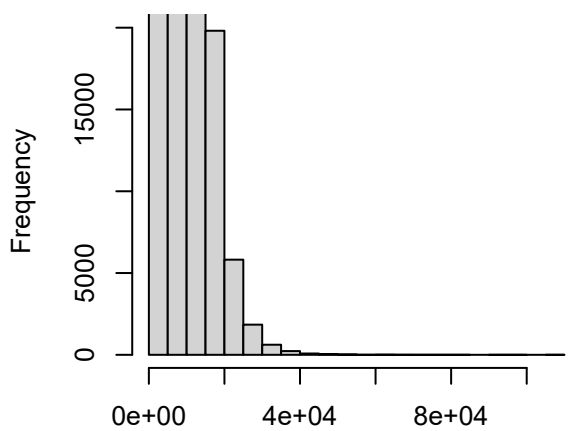




Samples



Read length (hist)



Read length (hist, cutoff)

Supplementary Figure 1: Basic statistics of the Nanopore-only validation experiment 1 on MRSA data. Shown are the number of total bases, the number of mapped bases, the number of total reads, the number of mapped reads and the average coverage, each per sample, as well as frequency histograms of the average coverage and the read length.