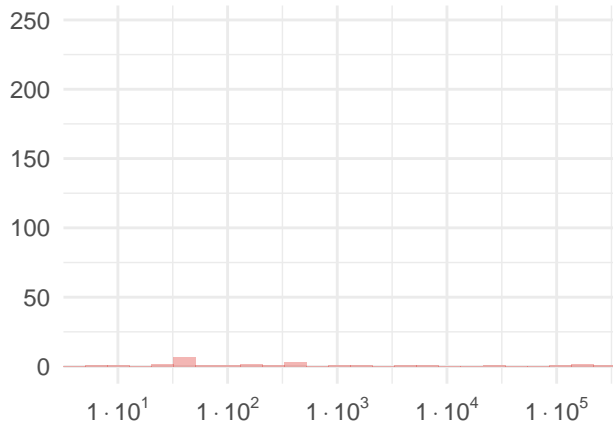
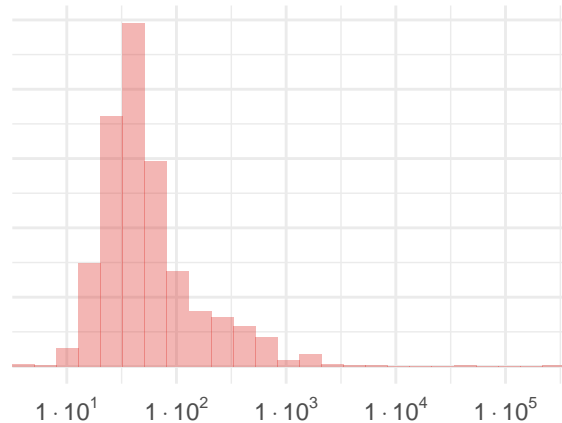


Number of Samples

HPV-35 +ve



HPV-35 -ve



Read Pairs Aligned to HPV-35 Contig