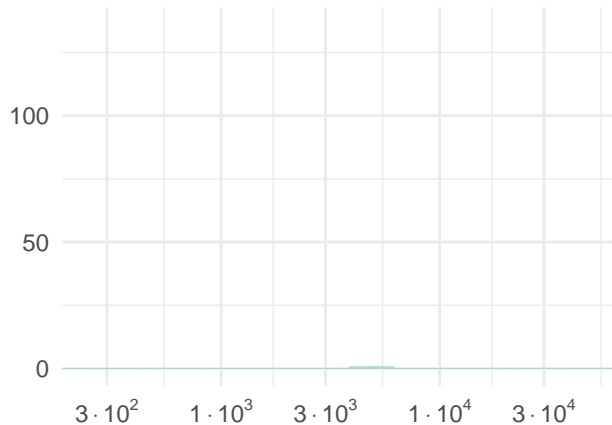
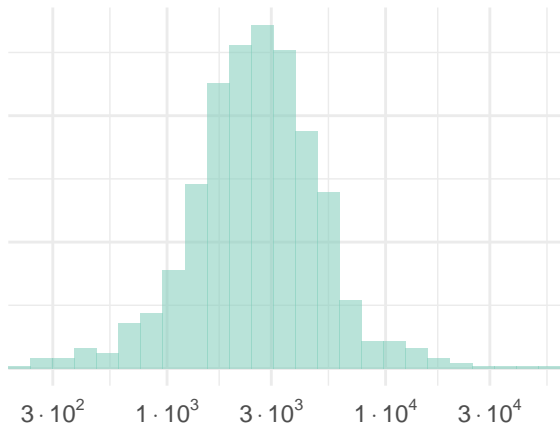


Number of Samples

HPV-18 +ve



HPV-18 -ve



Read Pairs Aligned to HPV-18 Contig