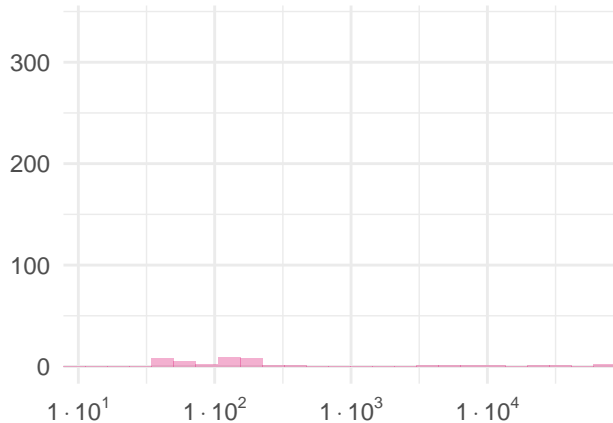
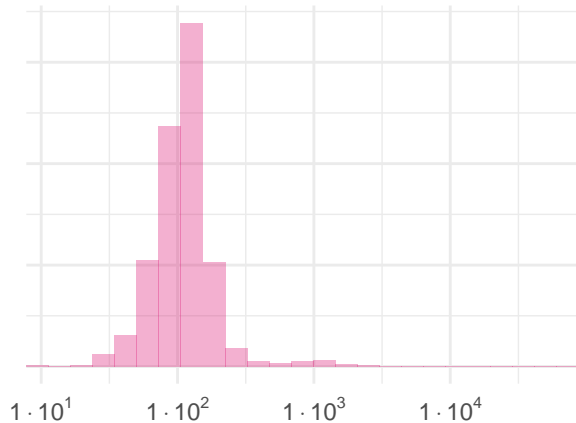


Number of Samples

HPV-33 +ve



HPV-33 -ve



Read Pairs Aligned to HPV-33 Contig