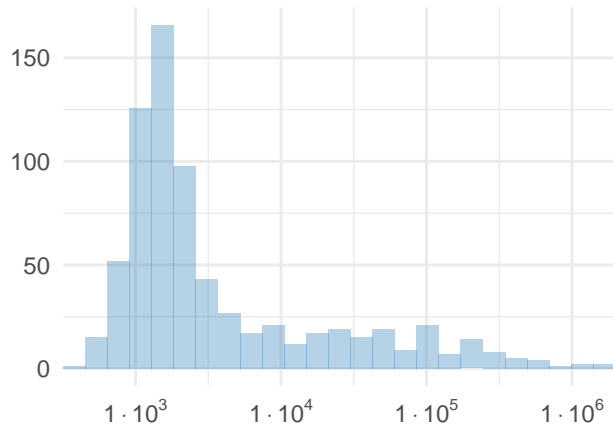
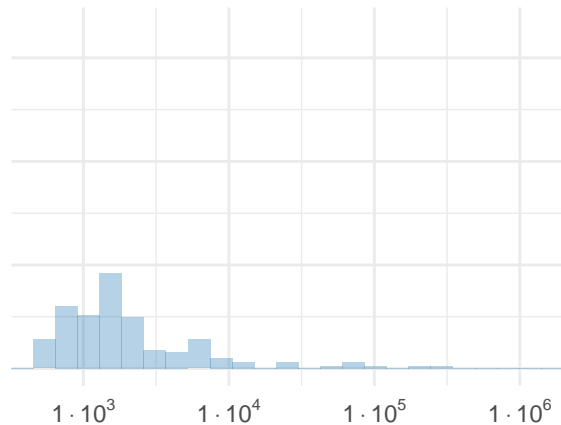


Number of Samples

HPV-16 +ve



HPV-16 -ve



Read Pairs Aligned to HPV-16 Contig