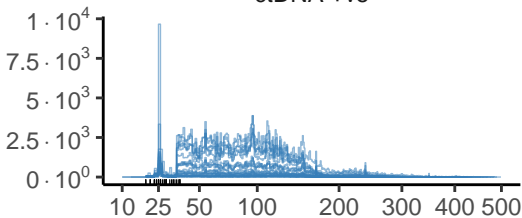
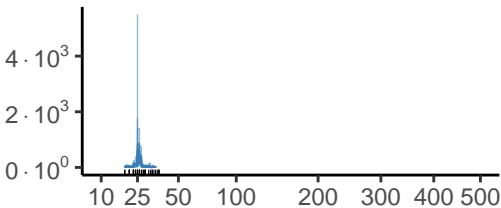


Number of Aligned Read Pairs

ctDNA +ve



ctDNA -ve



Insert Size (bp)