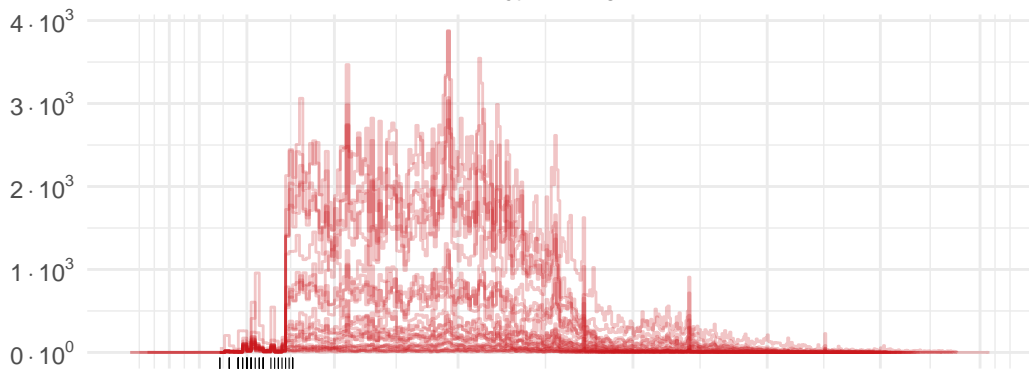


Number of Read Pairs

ctDNA +ve



ctDNA -ve



Insert size (bp)