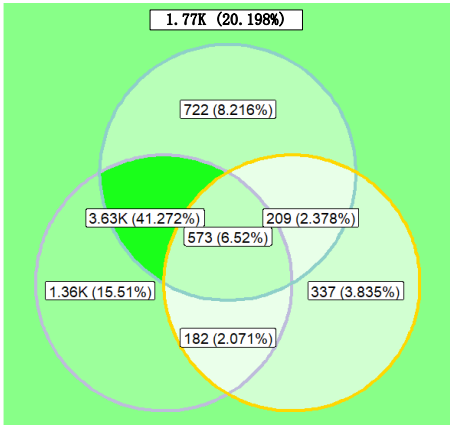
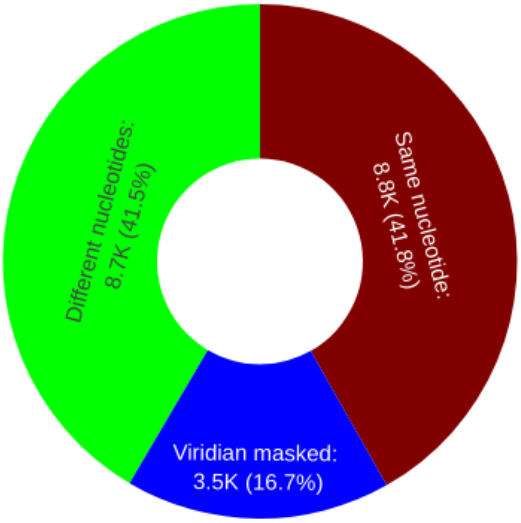
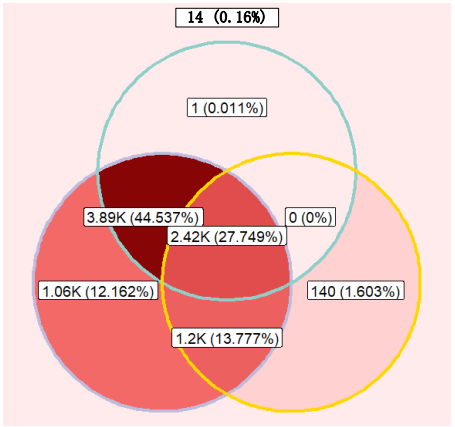
Sequence errors



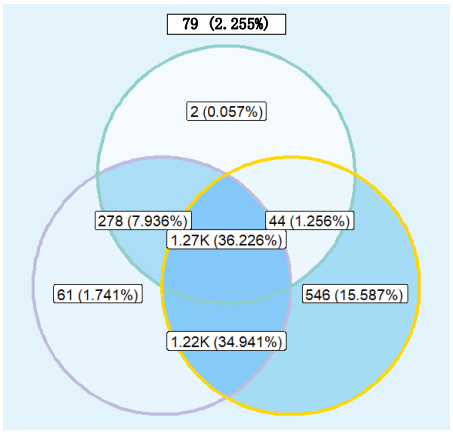


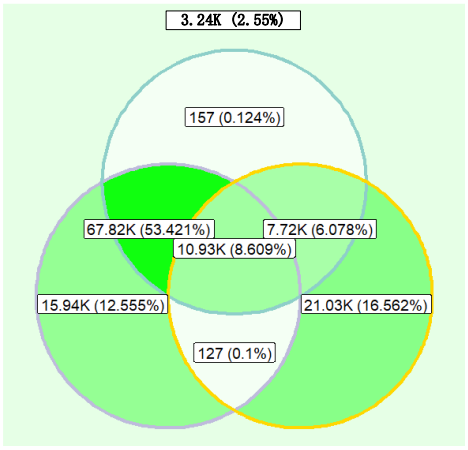


Coverage

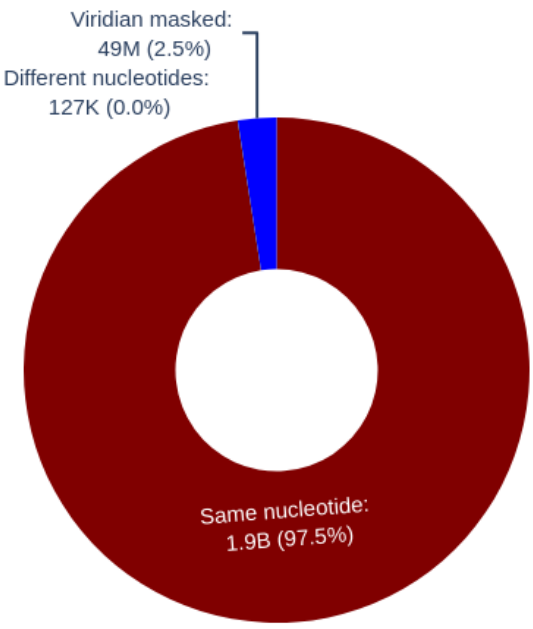
Strand Bias

Allele Frequency

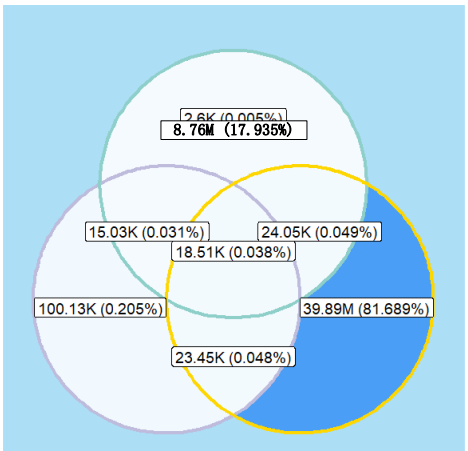
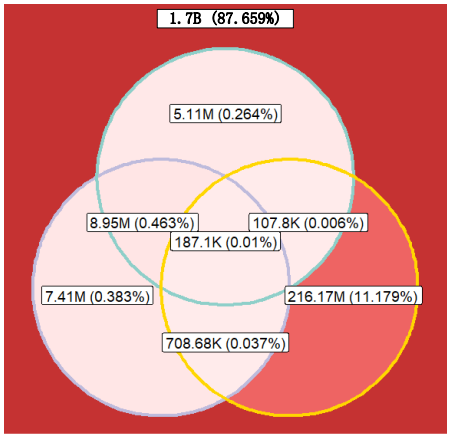




All Positions



Same nucleotide

Viridian masked

Different nucleotide