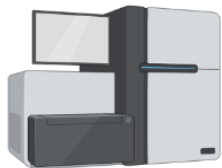


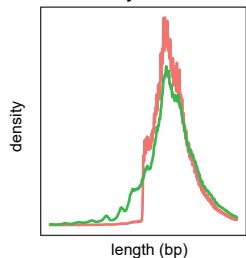


cfDNA WGS

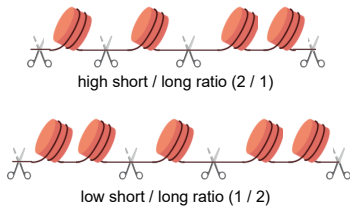


illumina HiSeq

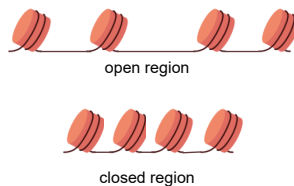
Quality control



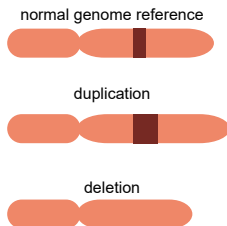
Fragment size ratio



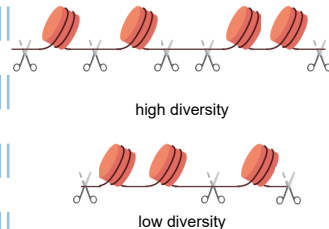
TSS relative coverage



Copy number variance



Promoter fragmentation entropy

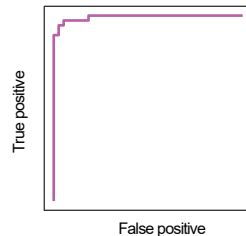


Promoter region

Machine learning



Performance of the model



Endonuclease site



Feature pattern



Model performance