/data/analysis_SimulatedAncientReads_HQ439467/01-EAGERmapping/ANCIENT.SIM.CONT.COV-7.5.RUN-3/5-DeDup/ANCIENT.SIM.CONT.COV-7.5.RUN-3.truncated.MT_realigned.mappedonly.sorted.qF.sorted.cleaned_rmdup.sorted

Number of used reads: 3.675 (100.0% of all input reads) | Specie: null

40.0

Read length

32.5

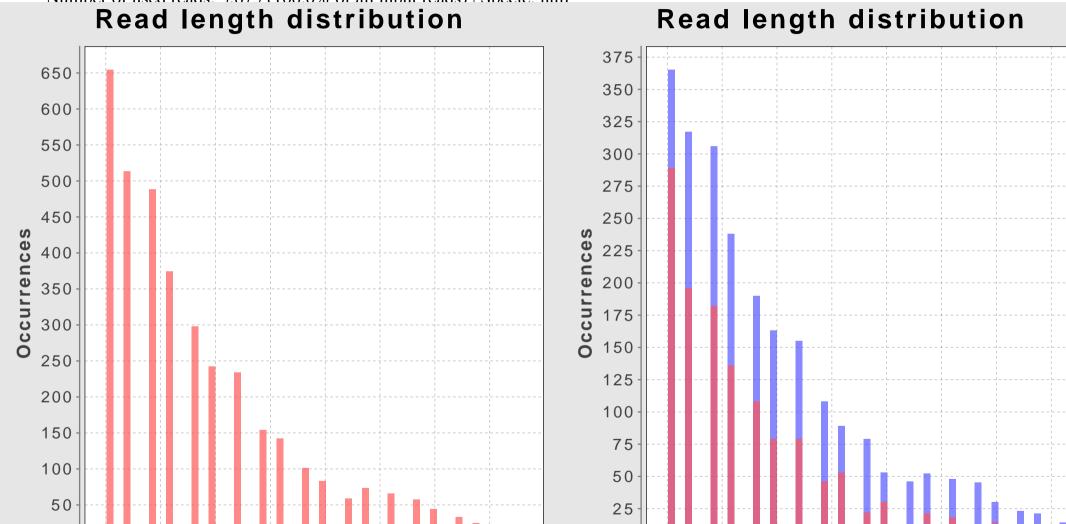
35.0

37.5

all reads

42.5

45.0 47.5



32.5

37.5

+ strand - strand

35.0

40.0

Read length

42.5

45.0 47.5