

/data/analysis_SimulatedAncientReads_HQ439467/01-EAGERmapping/ANCIENT.SIM.CONT.COV-10.RUN-4/5-DeDup/ANCIENT.SIM.CONT.COV-10.RUN-4.truncated.MT_realigned.mappedonly.sorted.qF.sorted.cleaned_rmdup.sorted

Number of used reads: 4.830 (100.0% of all input reads) | Species: null

Read identity distribution

