/data/analysis_SimulatedAncientReads_HQ439467/01-EAGERmapping/ANCIENT.SIM.CONT.COV-5.RUN-1/5-DeDup/ANCIENT.SIM.CONT.COV-5.RUN-1.truncated.MT_realigned.mappedonly.sorted.qF.sorted.cleaned_rmdup.sorted

Number of used reads: 2,449 (100.0% of all input reads) | Specie: null

Read identity distribution

Identitiy distribution