

**/data/analysis\_SimulatedAncientReads\_HQ439467/01-EAGERmapping/ANCIENT.SIM.CONT.COV-10.RUN-2/5-DeDup/ANCIENT.SIM.CONT.COV-10.RUN-2.truncated.MT\_realigned.mappedonly.sorted.qF.sorted.cleaned\_rmdup.sorted**

Number of used reads: 4.849 (100.0% of all input reads) | Species: null

## Read identity distribution

