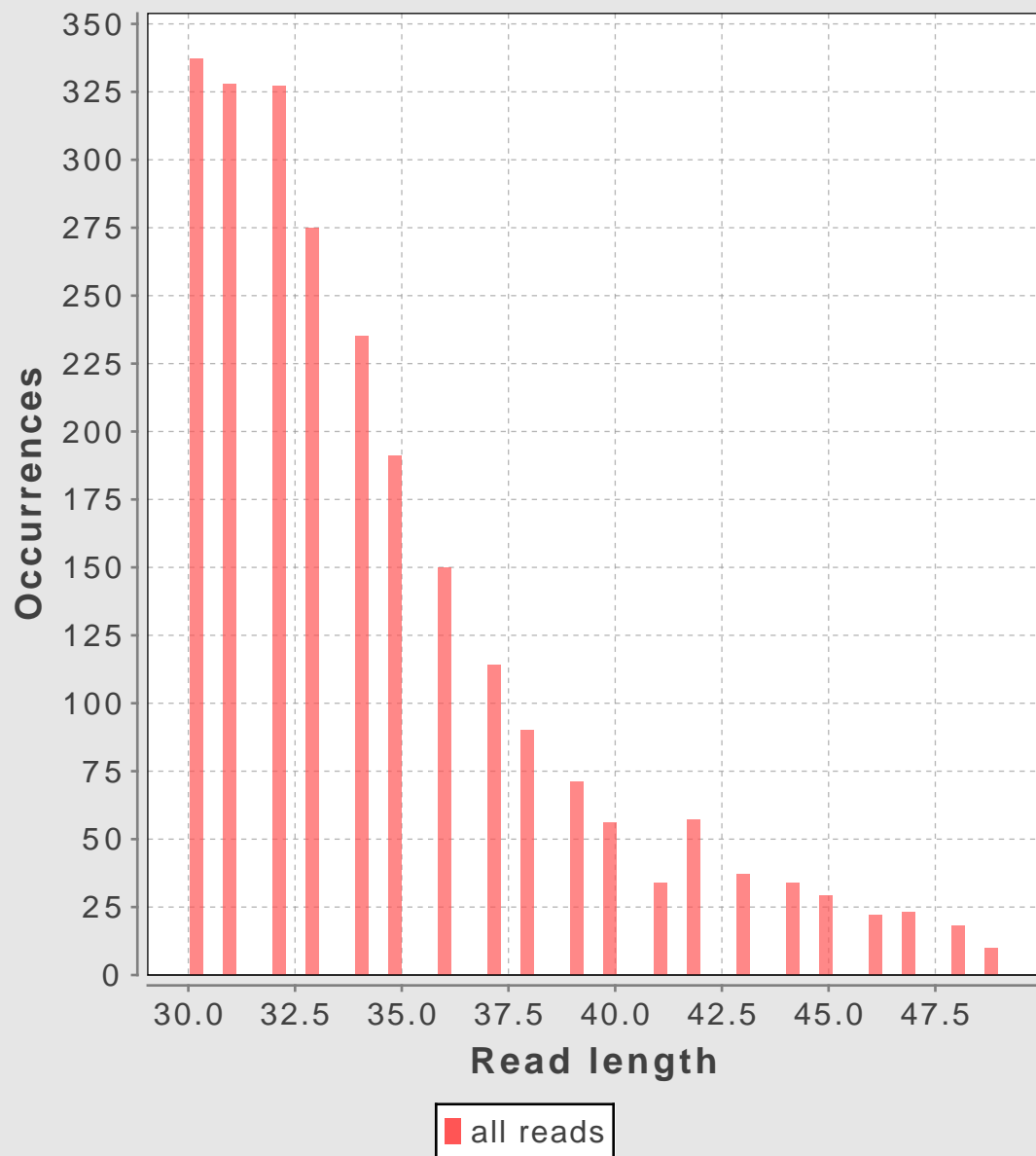


**/data/analysis_SimulatedAncientReads_HQ439467/01-
EAGERmapping/ANCIENT.SIM.CONT.COV-5.RUN-5/5-DeDup/ANCIENT.SIM.CONT.COV-
5.RUN-5.truncated.MT_realigned.mappedonly.sorted.qF.sorted.cleaned_rmdup.sorted**

Number of used reads: 2.438 (100.0% of all input reads) | Species: null

Read length distribution



Read length distribution

