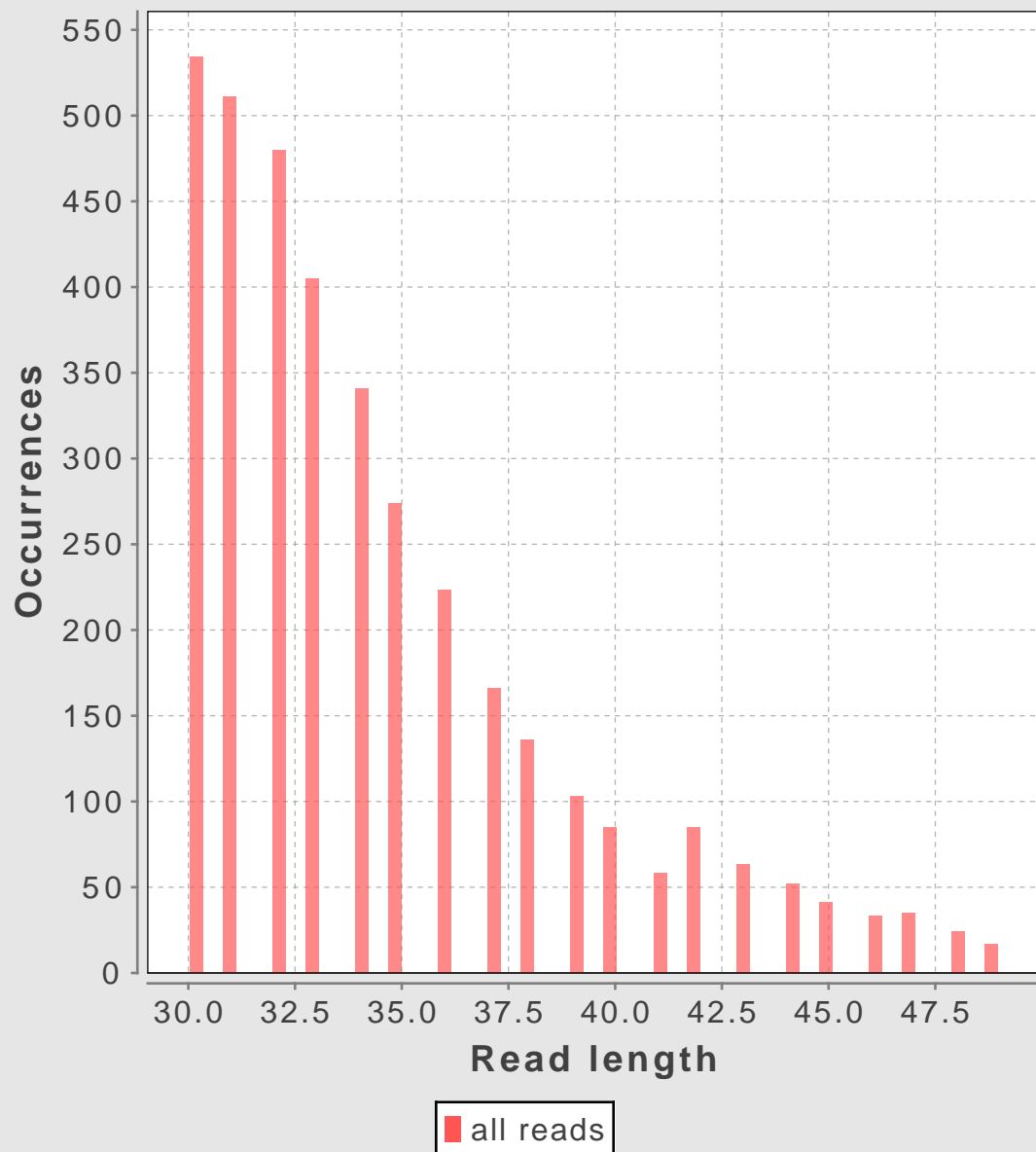


/data/analysis\_SimulatedAncientReads\_HQ439467/01-  
EAGERmapping/ANCIENT.SIM.CONT.COV-7.5.RUN-5/5-DeDup/ANCIENT.SIM.CONT.COV-  
7.5.RUN-5.truncated.MT\_realigned.mappedonly.sorted.qF.sorted.cleaned\_rmdup.sorted

Number of used reads: 3.666 (100.0% of all input reads) | Species: null

**Read length distribution**



**Read length distribution**

