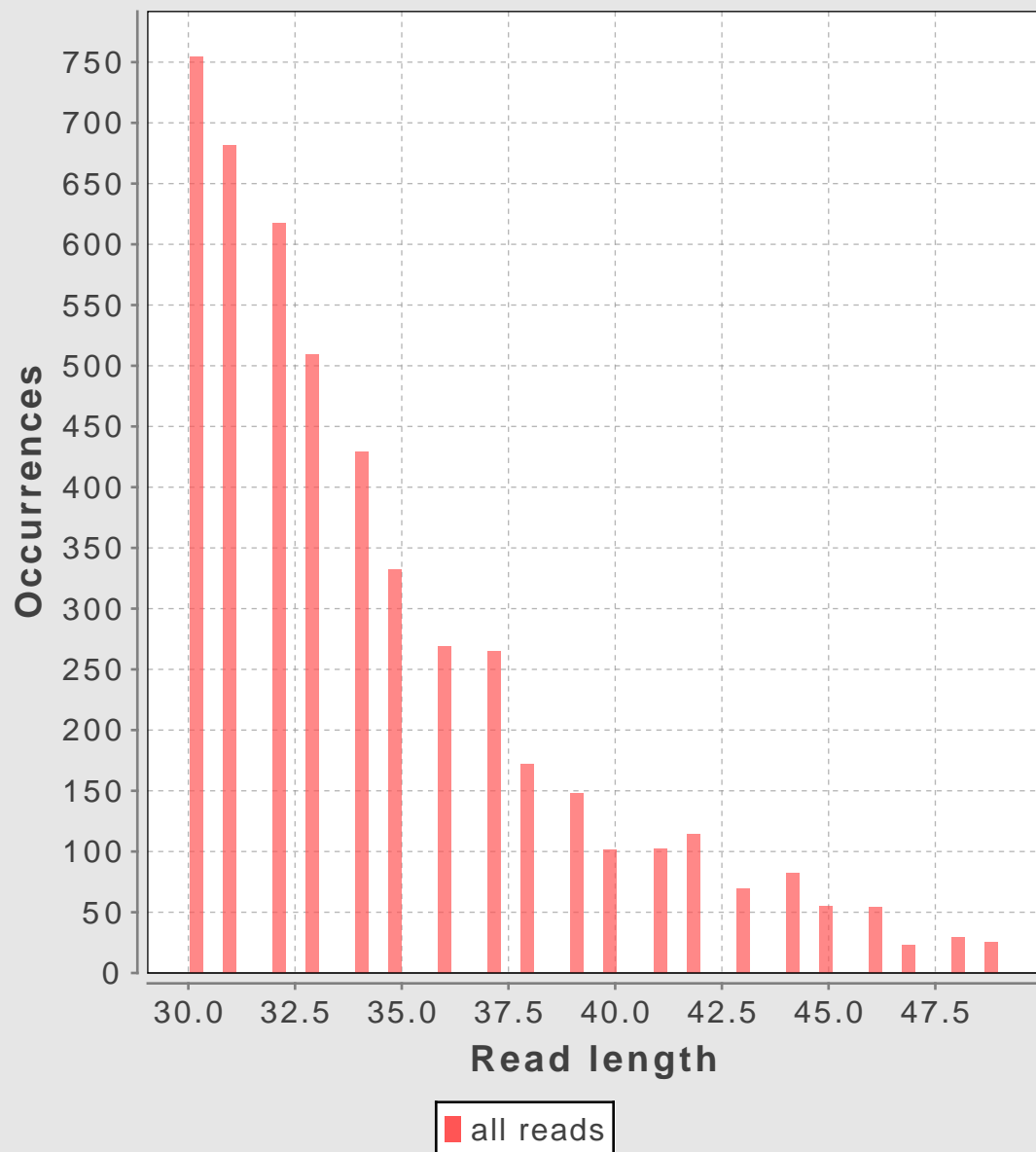


/data/analysis\_SimulatedAncientReads\_HQ439467/01-  
EAGERmapping/ANCIENT.SIM.CONT.COV-10.RUN-4/5-DeDup/ANCIENT.SIM.CONT.COV-  
10.RUN-4.truncated.MT\_realigned.mappedonly.sorted.qF.sorted.cleaned\_rmdup.sorted

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

## Read length distribution



## Read length distribution

