

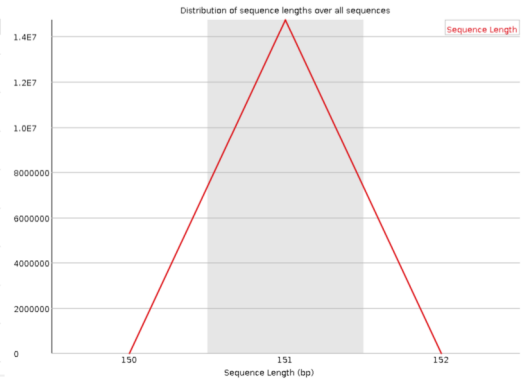
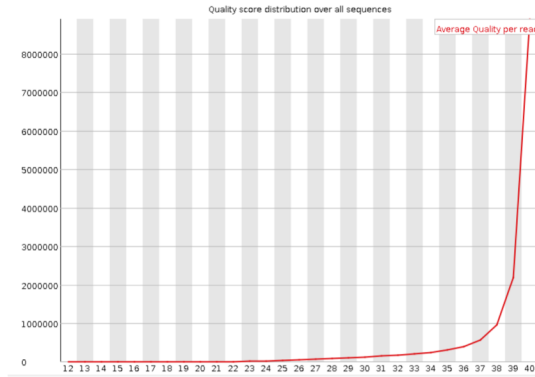
(A) *S. enterica*

SRR5866640 (150bp)

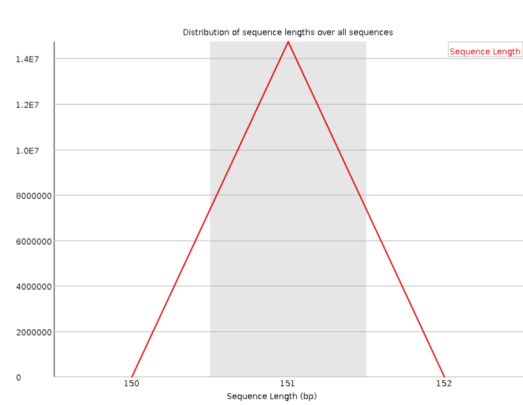
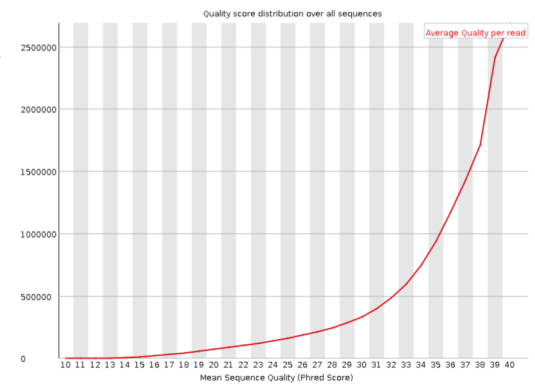
Per sequence quality scores

Sequence Length Distribution

Read 1

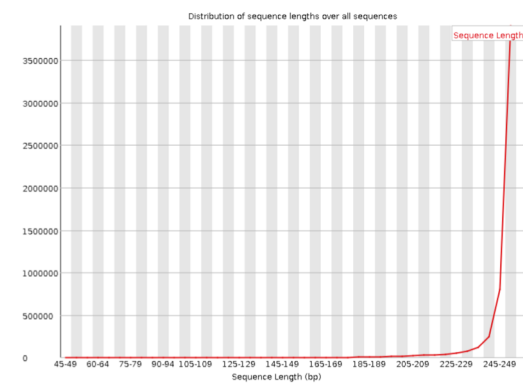
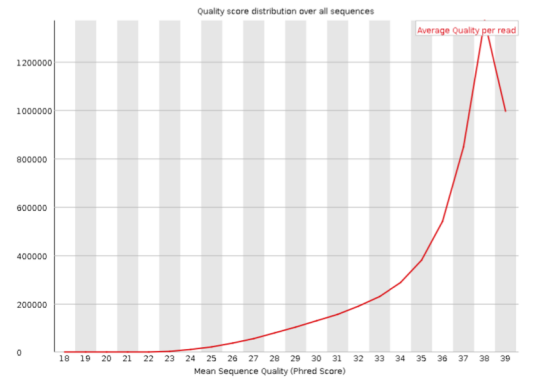


Read 2

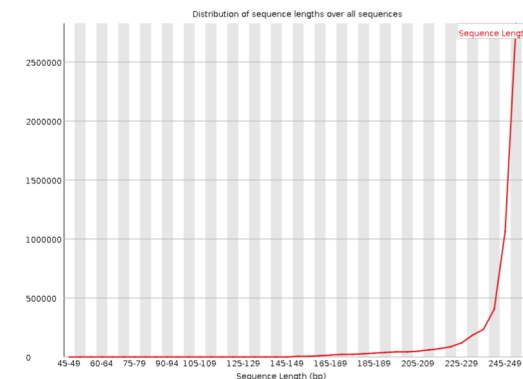
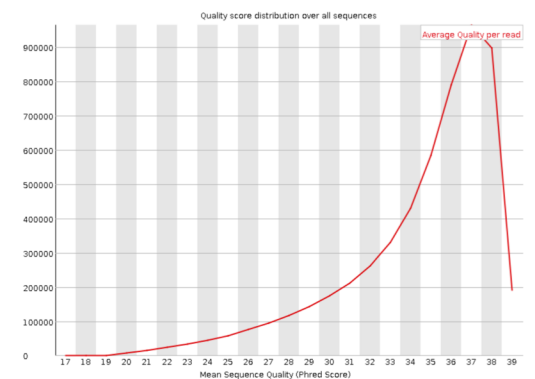


SRR6929558 (250bp)

Read 1



Read 2

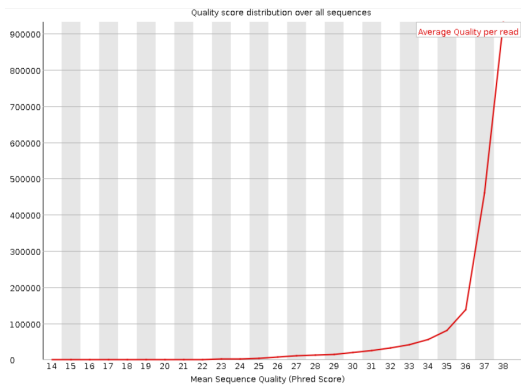


(B) *E. coli*

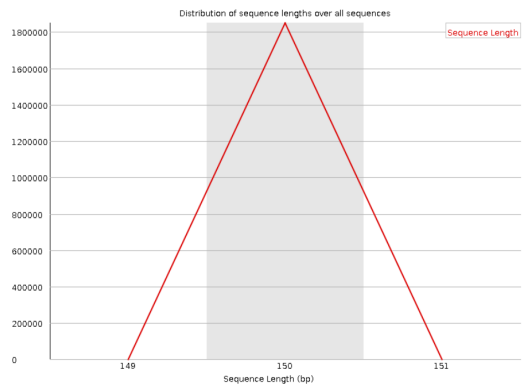
SRR6924239 (150bp)

Per sequence quality scores

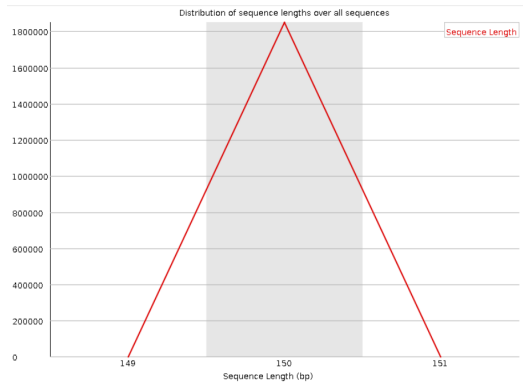
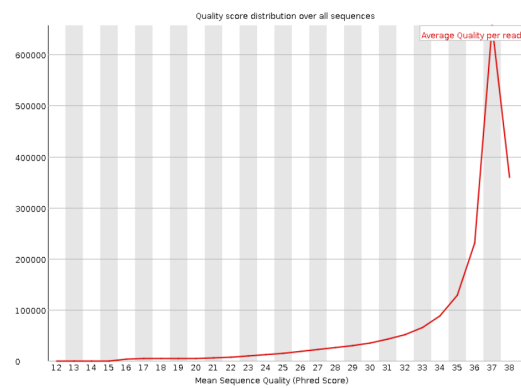
Read 1



Sequence Length Distribution

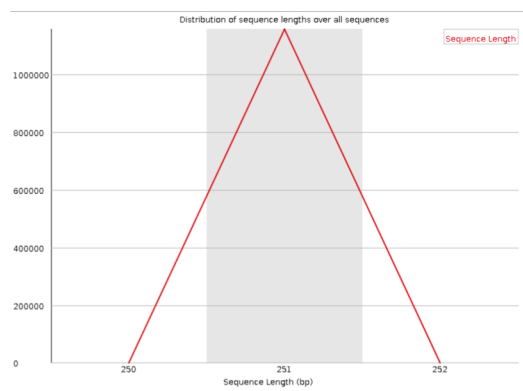
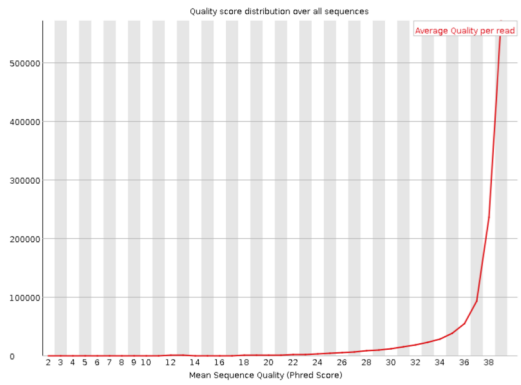


Read 2

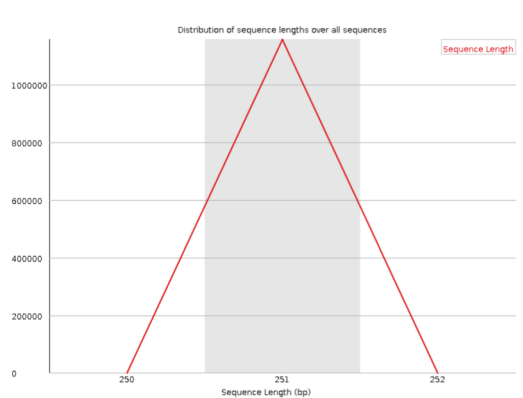
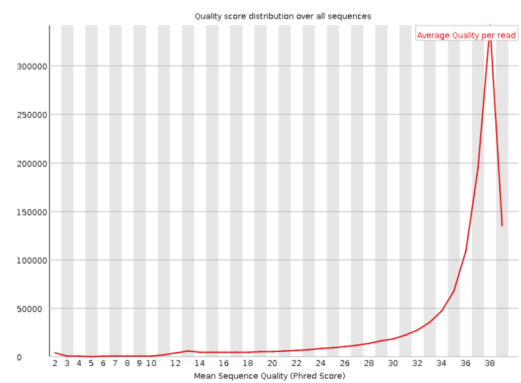


SRR3205757 (250bp)

Read 1



Read 2



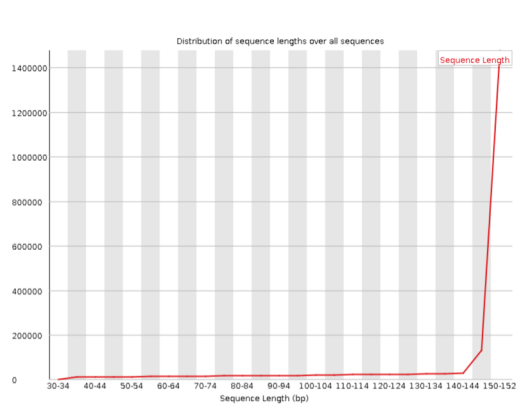
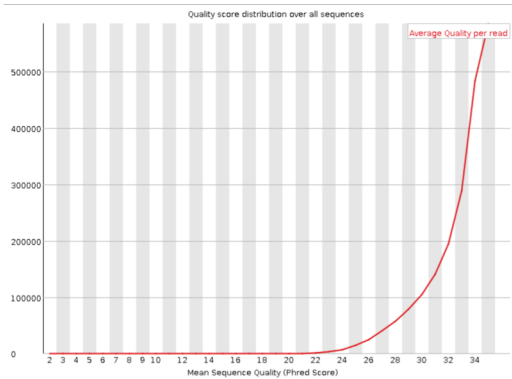
(C) *L. monocytogenes*

SRR3089759 (150bp)

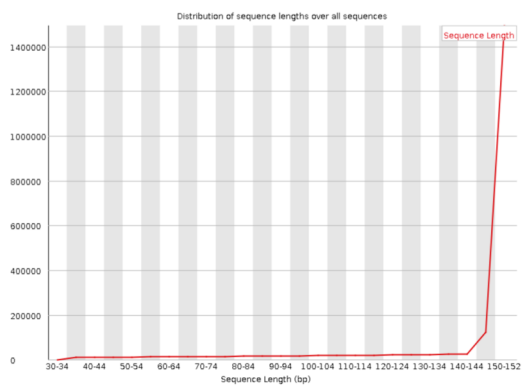
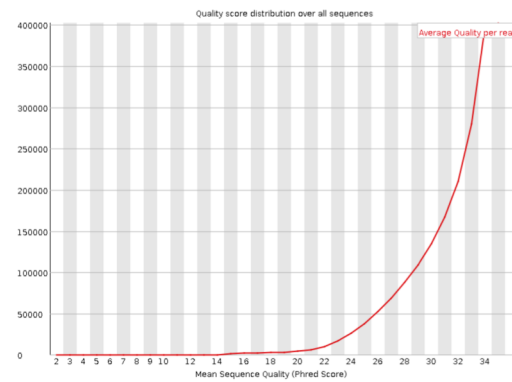
Per sequence quality scores

Sequence Length Distribution

Read 1

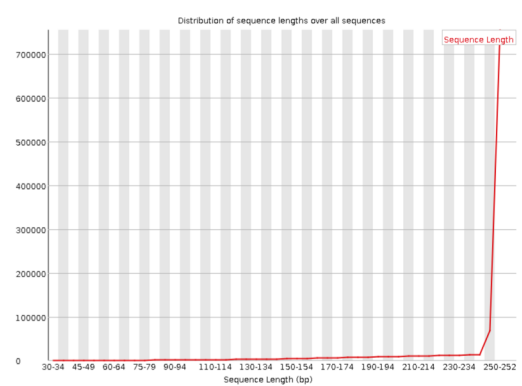
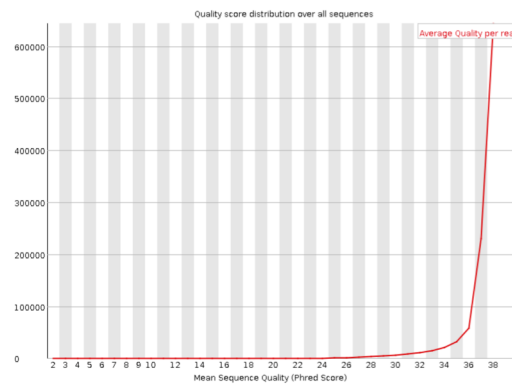


Read 2



SRR6347431 (250bp)

Read 1



Read 2

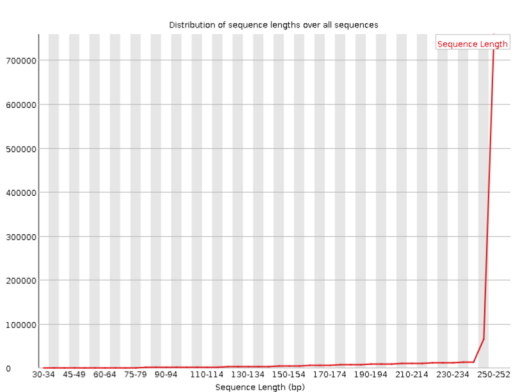
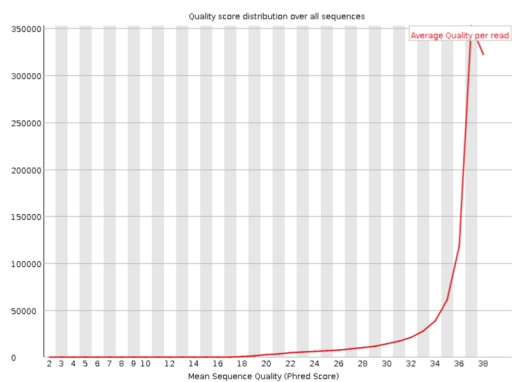


Figure S2. The real sequenced reads QC performed by using FastQC. The “Per sequence quality scores” and “Sequence Length Distribution” from QC reports of (A) *S. enterica* (SRR5866640 for 150 bp and SRR6929558 for 250 bp), (B) *E. coli* (SRR6924239 for 150 bp and SRR3205757 for 250 bp), and (C) *L. monocytogenes* (SRR3089759 for 150 bp and SRR6347431 for 250 bp) were shown.