
Sequencing analysis report

This document reports the sequencing results of the **EGF demo** project. There are **2** barcodes in the analysis. Each chapter details the results of one barcode, and within a chapter each section reports on one plasmid. Please see the Appendix on the last page for an explanation of the report.

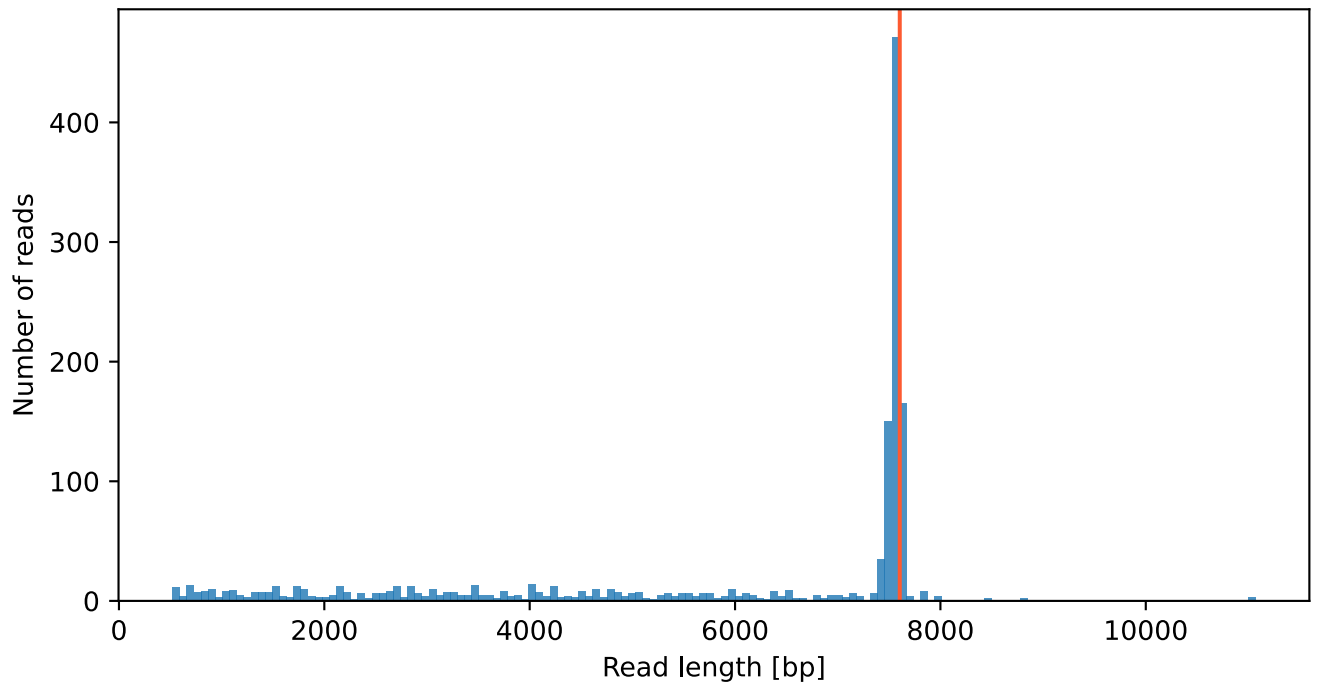
There are **1623** filtered reads in the analysed barcodes of this sequencing run.

Barcode: barcode01

Results for the **1** construct(s) in the pool: **1** / **0** / **0** / **0** (pass / warning / fail / low coverage).

Name	Result	Length [bp]	FASTQ reads	Coverage [x]
EGF2_2	PASS	7604	792	557

Histogram of the **792** FASTQ reads:



The vertical red lines show the expected construct lengths.