

# ABC - Report

nf-codon-align

## Pipeline Information

Git Commit: *hash*

Version: x.y.z

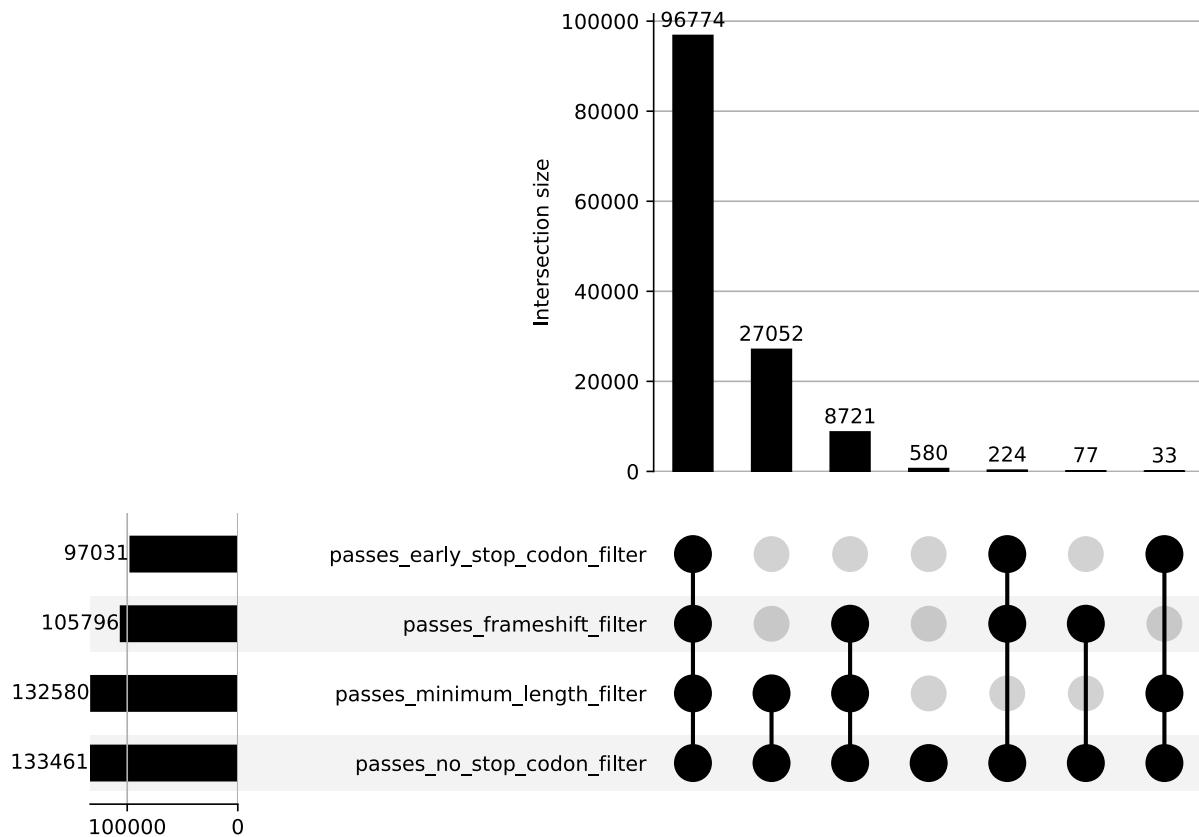
## Overview

This pipeline run contained **100** files. **10** files made it through the pipeline. We lost **10** sequences, representing **10%** of all sequences.

Pipeline Point	Sequence Count
Pre	100
Post	19

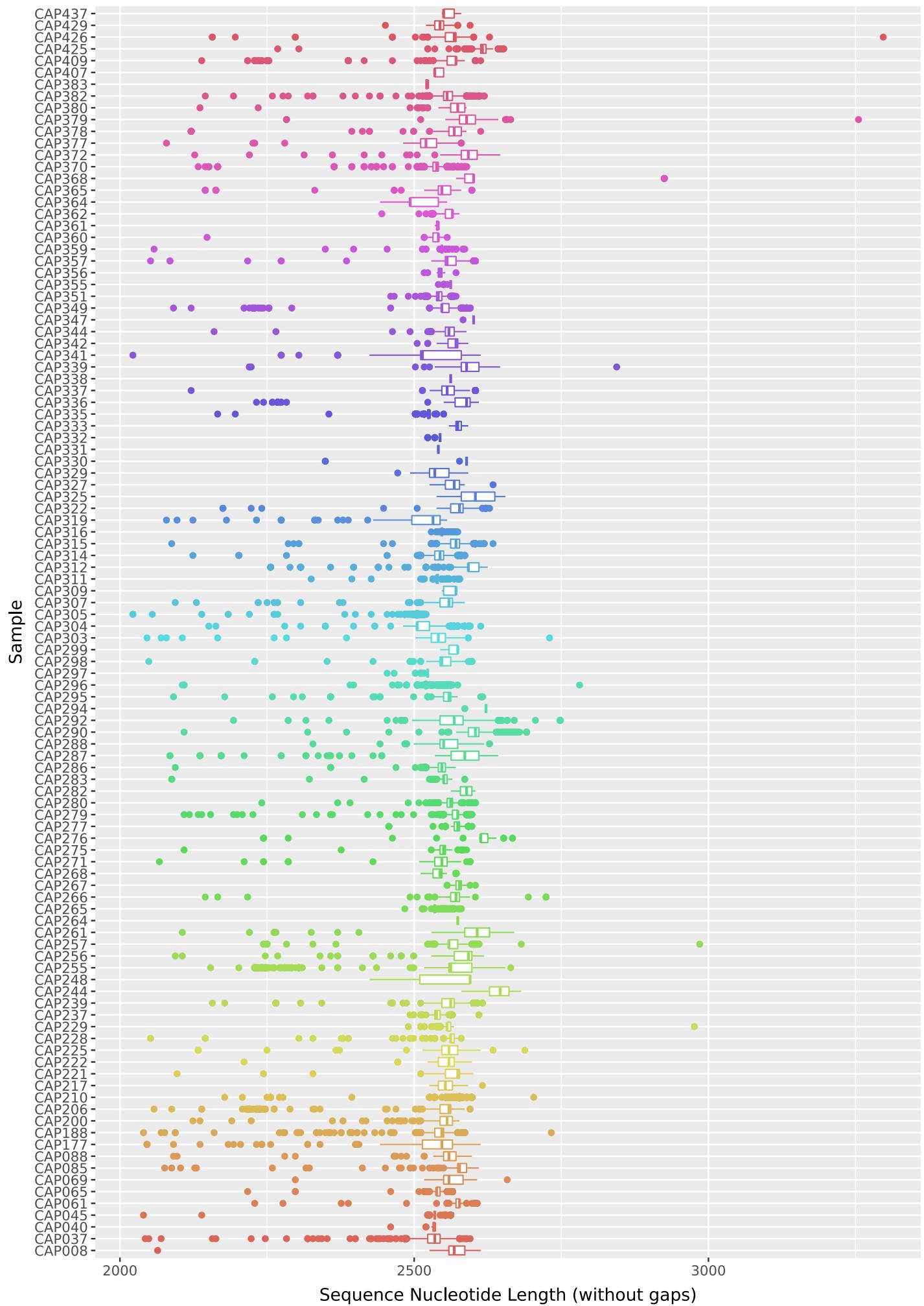
## Sequence Loss Overview

This plot is an UpSet plot, which indicates the various reasons why sequences were lost from the pipeline. A circle is **filled in** if that test was passed.

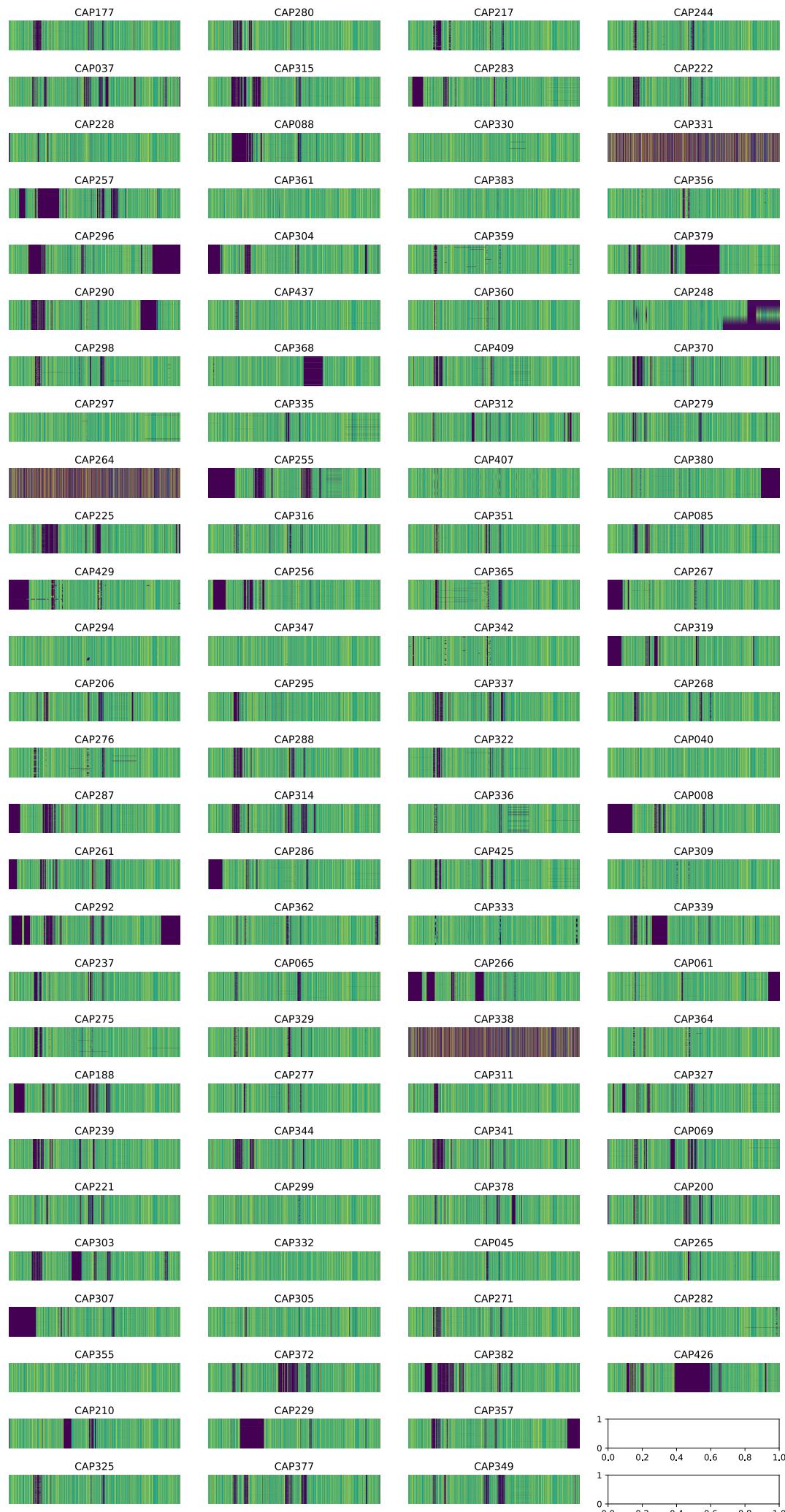


## Sequence Length

This plot shows the sequence length distribution for the sequences **that pass filter**.



# Alignment Quality Overview



## Pipeline Run Info

The following table shows the parameters that were used to run this pipeline version.

(a: "abc", b: "bce")