

Resource Summary Report

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SYSB 3036 W02: Parsing FASTA files

DOI:10.17504/protocols.io.vfwe3pe

Type: Protocol

Proper Citation

Frank Aylward 2018. SYSB 3036 W02: Parsing FASTA files. protocols.io
dx.doi.org/10.17504/protocols.io.vfwe3pe

Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.vfwe3pe>

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Summary: Week 1 Introduction to parsing FASTA files. Commands to be entered into the command line are in bold. Here we will be using various base Unix commands such as head, tail, sort, wget, and others. We will also be using the seqkit tool to process FASTA files. The main page for seqkit is here: <https://github.com/shenwei356/seqkit>

Affiliations: Virginia Tech

Version: 1

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