



## Command Line Variables Reference

In this document, we have provided a reference for you to fill in as you progress through the instructions for [Genome Assembly on the Command Line](#) and [Genome Analysis on the Command Line](#).

Absolute path of Bioinformatics pipeline: \_\_\_\_\_  
<sequencing-path>

Name of project directory: \_\_\_\_\_  
<project-directory-name>

Absolute path of project directory: \_\_\_\_\_  
<project-path>

Number of processors: \_\_\_\_\_  
<number-of-processors>

Absolute path of background dataset: \_\_\_\_\_  
<background-dataset-path>