

# Nextflow workflow report

# [condescending\_banach]

Workflow execution completed successfully!

#### Run times

26-Aug-2024 11:42:12 - 26-Aug-2024 12:00:43 (duration: 18m 31s)

21 succeeded

### **Nextflow command**

nextflow run main.nf -c nextflow.config -with-report
workflow\_report.html -with-trace trace.txt -with-timeline
timeline.html

CPU-Hours 1.1

 Launch directory
 /home/javeria/Downloads/dewgsimdocker

 Work directory
 /home/javeria/Downloads/dewgsimdocker/work

 Project directory
 /home/javeria/Downloads/dewgsimdocker

Script name main.nf

 Script ID
 16cbb10dcd80709c385bc0eea117d00a

 Workflow
 967c7e19-0938-4233-aa5d-b0e08570ad5d

session

Workflow profile standard

Workflow [annotate:staphb/prokka:latest,

container panaroo:staphb/panaroo:latest, dwgsim:dwgsimwith-ps:latest, snippy:staphb/snippy:4.6.0,

detectStopCodons:python-biopython-pandasps:latest, genedescription:python-biopython-

pandas-ps:latest]

Container engine docker

**Nextflow version** version 24.04.3, build 5916 (09-07-2024 19:35 UTC)

## Resource Usage

These plots give an overview of the distribution of resource usage for each process.

### **CPU**

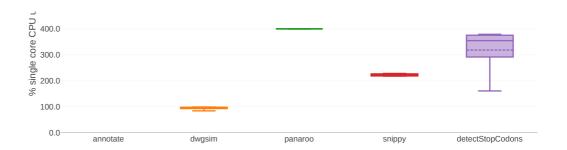
Raw Usage

% Allocated

CPU Usage



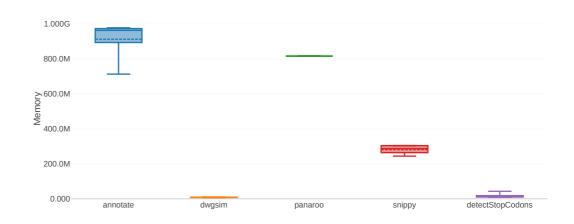




## Memory



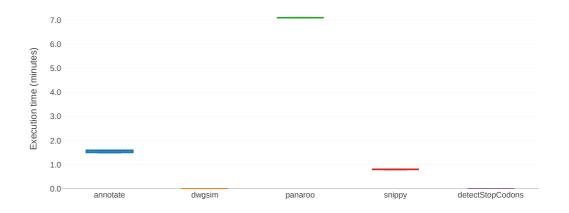
## Physical Memory Usage



## Job Duration



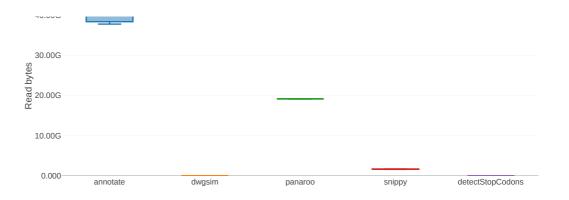
#### Task execution real-time





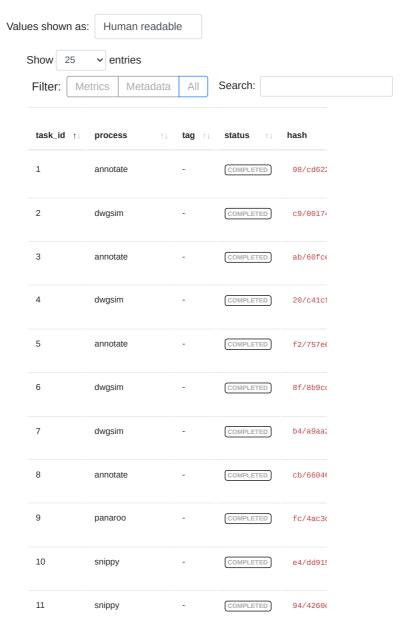
Number of bytes read



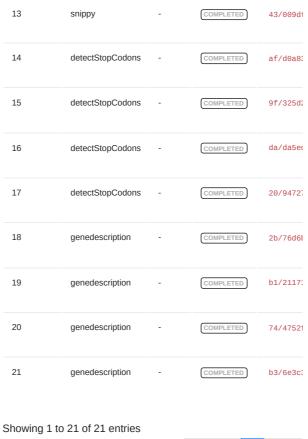


## **Tasks**

This table shows information about each task in the workflow. Use the search box on the right to filter rows for specific values. Clicking headers will sort the table by that value and scrolling side to side will reveal more columns.







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