1

Select a pipeline

TaxonomicProfiler

PathogenScreen

2

Run Name
wastewater

Pathogen Panel
Human Pathogenic Virus...

Cancel | 10 files selected | +

Blank_R1_001.fastq.gz
3.7 KB

Blank_R2_001.fastq.gz
4.4 KB

Kitchener_R1_001.fastq.gz
24 MB

Kitchener_R2_001.fastq.gz
28 MB

Waterloo_R1_001.fastq.gz
16 MB

Waterloo_R2_001.fastq.gz
18 MB

P2-1002-16S_S2_...001.fastq.gz
22 MB

SUBMIT

3

Run | wastewater | STATUS | SETUP

Uploads

Input Files

Uploading: 16%
2 of 10 files uploaded · 24 MB of 148 MB · 1m 44s left

5

Results

	Name	Pipeline	Created (Eastern Standard Time)	Finished (Eastern Standard Time)	Status	
☐	wastewater	pathogenscreen	1/25/2024, 2:11:45 PM	1/25/2024, 2:12:59 PM	completed	🔍 ⬇
☐	febrileDataset_pathScreen	pathogenscreen	1/25/2024, 11:45:30 AM	1/25/2024, 11:48:15 AM	completed	🔍 ⬇
☐	SRR7615344_taxProfiler	taxonomicprofiler	1/25/2024, 11:44:30 AM	1/25/2024, 11:46:34 AM	completed	🔍 ⬇
☐	ERR7912883_pathScreen	pathogenscreen	1/25/2024, 11:44:05 AM	1/25/2024, 11:44:56 AM	completed	🔍 ⬇

4

Pathogenscreen Report

Metadata

```
Run : wastewater
Status: completed
Created: 2024-01-25
```

Detections

Detections Table

Detections were generated using the selected Human Pathogenic Viruses pathogen panel with an abundance threshold of 0. A filtered abundance column is also provided for each sample, representing the estimated abundance after removal of Human reads.

Show 10 entries   Search: Severe acute respirat×

	Blank_T1			Waterloo_T1			Kitchener_T1		
	Reads	Abundance	Filtered Abundance	Reads	Abundance	Filtered Abundance	Reads	Abundance	Filtered Abundance
Organism									
Severe acute respiratory syndrome-related coronavirus	0.0	0.0	0.0	72.0	0.00102	0.00108	101.0	0.00082	0.00083

Showing 1 to 1 of 1 entries (filtered from 227 total entries)   Previous 1 Next