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MY FASTQS

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CW

basepawsD100

SUCCEEDED

CLIENT ID

basepawsD100

GENCOVE ID

2fe95c47-725e-46c4-b7e3-1acbbf37f558

Has data: 2020-03-07T06:28:38.086434Z

Running: 2020-03-07T06:28:38.718236Z

Succeeded: 2020-03-07T07:20:30.276377Z

QC Overview

VALID FASTQ FORMAT

StatusSucceeded

BASES SEQUENCED, MINIMUM

502.8M

Observed value502,801,914

Threshold270,000,000

StatusSucceeded

BASES SEQUENCED, MAXIMUM

502.8M

Observed value502,801,914

Threshold16,200,000,000

StatusSucceeded

BASES DEDEDUPICATED AND ALIGNED, MINIMUM

506.7M

Observed value506,706,048

Threshold270,000,000

StatusSucceeded

KNOWN SITES COVERED BY A READ, MINIMUM

5.3M

Observed value5,349,418

Threshold4,500,000

StatusSucceeded

RAW DATA AVAILABILITY

Alignment, BAI file

Alignment, BAM file

Input, FASTQ file (R1)

Input, FASTQ file (R2)

HELP

LOG OUT

https://web.gencove.com/projects/dbec3e55-a0c8-4d74-a5bd-815ded7137e7/samples/2fe95c47-725e-46c4-b7e3-1acbbf37f558

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Imputation, CSI file	
Imputation, TBI file	
Imputation, VCF file	