

Samplesheet / Run Fastq File Cross Validator

Sequencing Run Directory: 20200825_A_0159_0208_0218_192233042_abridged

Date/Time Executed: Wed Sep 9 14:00:13 2020

Samplesheet found at:

20200825_A_0159_0208_0218_192233042_abridged/Samplesheet/20200825_A_0159_0208_0218_192233042SampleSheet.csv

Fastq directory at:

20200825_A_0159_0208_0218_192233042_abridged/Run

Num Sample IDs in samplesheet: 228

Num Sample IDs by paired fastq files: 226

Num Missing Fastq Samples IDs: 4

Num Excess Sample Sheet IDs: 2

Error: Sample ID count mismatch.

Sample IDs missing fastq.gz files:

0159.AZ03.32097.20200108.SP
0208.005012.20180608.BALL
0218.2210.20200309.MST.RNA
0218.2212A.20200309.MST.RNA

Sample IDs missing from sample sheet:

0208.003001.20170920.BALL
0218.2212.20200309.MST.RNA

Missing R1 Fastq Files:

0208.005012.20180608.BALL

Missing R2 Fastq Files:

0159.AZ03.32097.20200108.SP