

FASTQ files

Server

Sample 1

Sample 2

Sample 3

Sample 4

Sample 5

DAC_RNAseq_process()

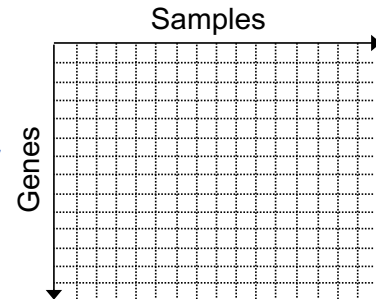
Job 1

Job 2

Job 3

Job 4

Job 5



Parallel processing