**Meaning of the naming of the files**

The name of the file has a unique identifier part, a project name, and an internal label. Below is an example:

C19-MB-11\_V1\_R3C19OR\_DVB\_S7\_R2\_001.fastq.gz.aspera-env

*Break down:*

C19-MB-11\_V1\_R3: unique identifier of the sample

C19OR: Project name (Covid19 organoids)

DVB\_S7\_R2\_001: Internal label

*Break down of the unique identifier:*

C19-MB-11\_V1\_R3

C19: Covid 19 (the same for all the samples)

MB: The type of organoid (MB, for midbrain)

11: The individual from where the cells are derived (it can be 11 or 15)

V1: The variant of the Sars-Cov-2 used (it can be Un, for untreated, or V1, for variant 1)

R3: The replicate of that condition (it can be R1, R2, R3, or R4)