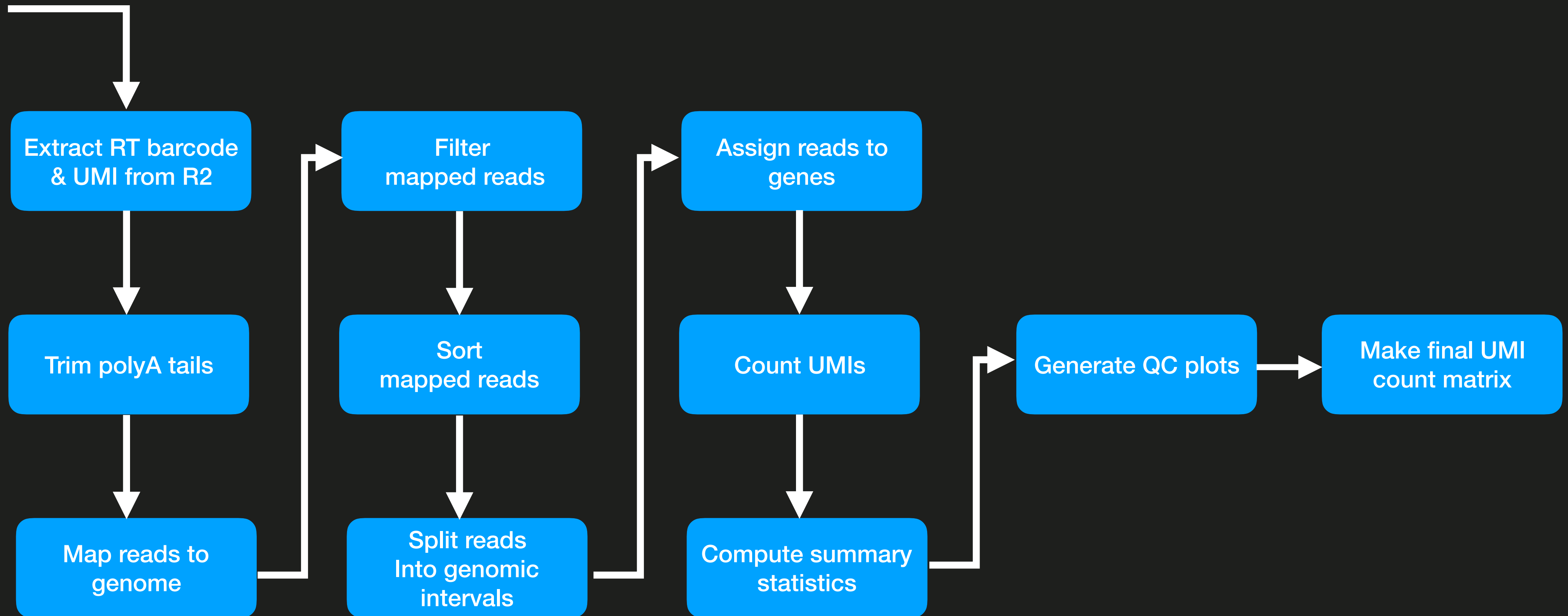


# The sci-RNA-seq pipeline

FASTQ files



**qlogin**