

Start Aggregation
(run_mapseq_aggr.sh)



Source project_config.txt



Run Cell Ranger RNA Aggr (background)
(run_cellranger.aggr_rna.sh)

Run Cell Ranger ATAC Aggr (background)
(run_cellranger.aggr_atac.sh)



Wait for Aggr jobs



Demultiplex RNA with Seurat
(run_seurat.demux_rna.R)

Demultiplex ATAC/ASAP with Seurat
(run_seurat.demux_atac.R)



End Aggregation
(RDS Objects Created)