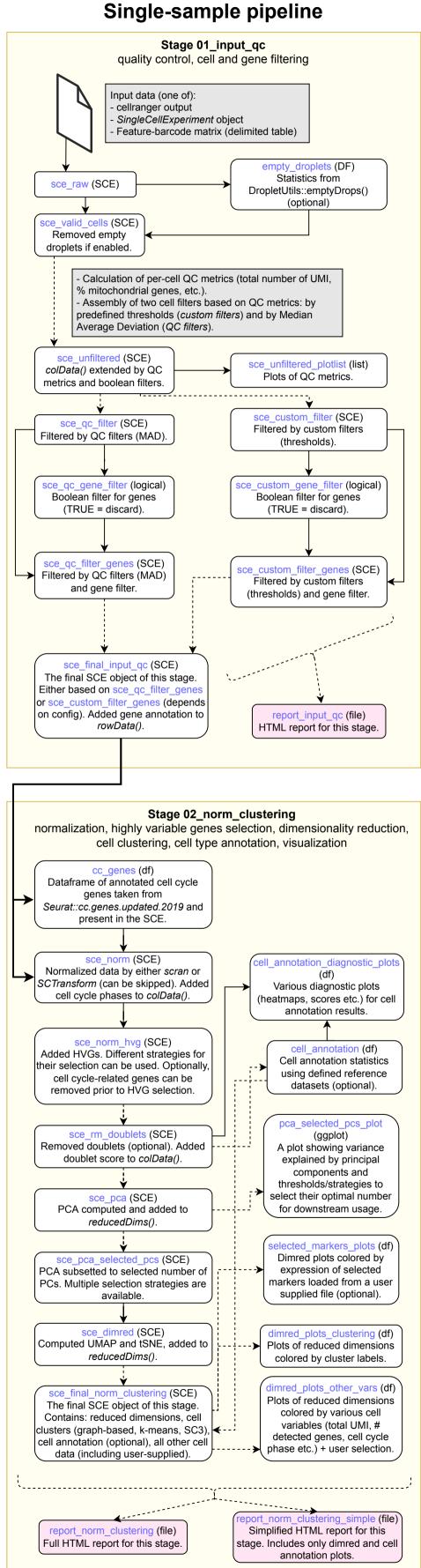
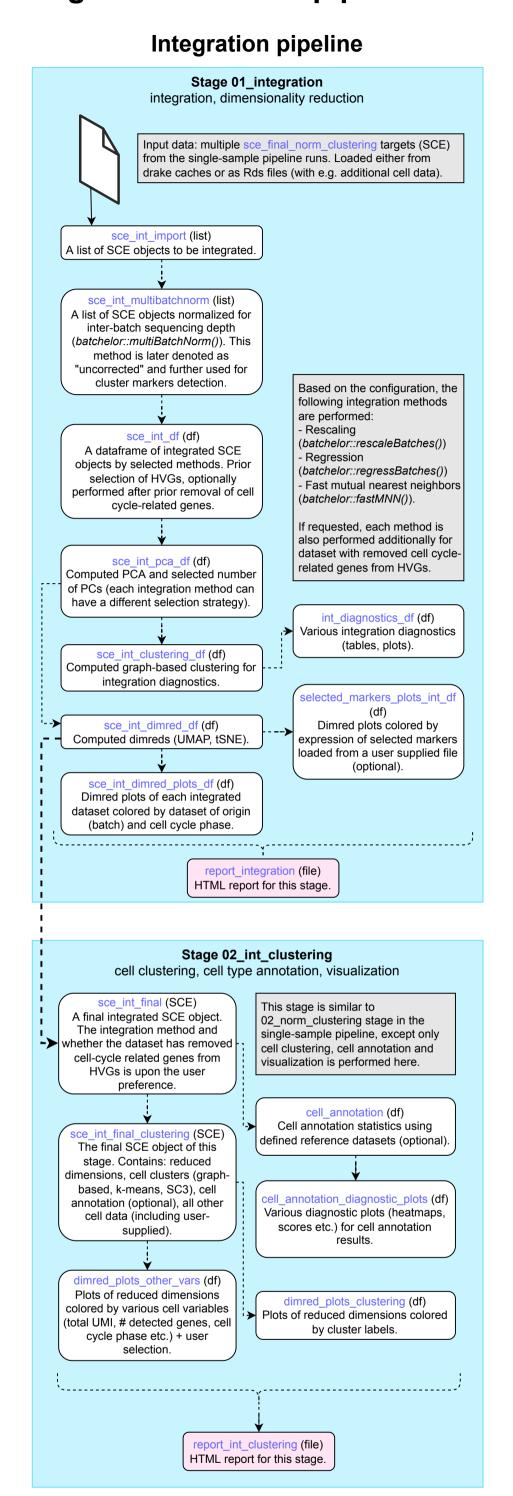
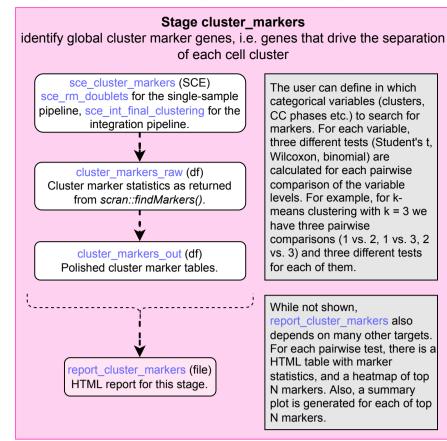
Supplementary Figure 2: Low-level diagram of scdrake pipelines





Common pipelines

stages common to both single-sample and integration pipelines



Stage contrasts

differential gene expression between cell clusters

The contrasts stage is very

stage, except all outputs

similar to the cluster_markers

(heatmaps, tables) are related to

individual pairwise comparisons

report_contrasts (file)

HTML report for this stage.

sce contrasts (SCE)

sce_rm_doublets for the single-sample

pipeline, sce int final clustering for the

integration pipeline.

Differential expression (DE) statistics as

returned from scran::findMarkers().

Polished DE tables.

