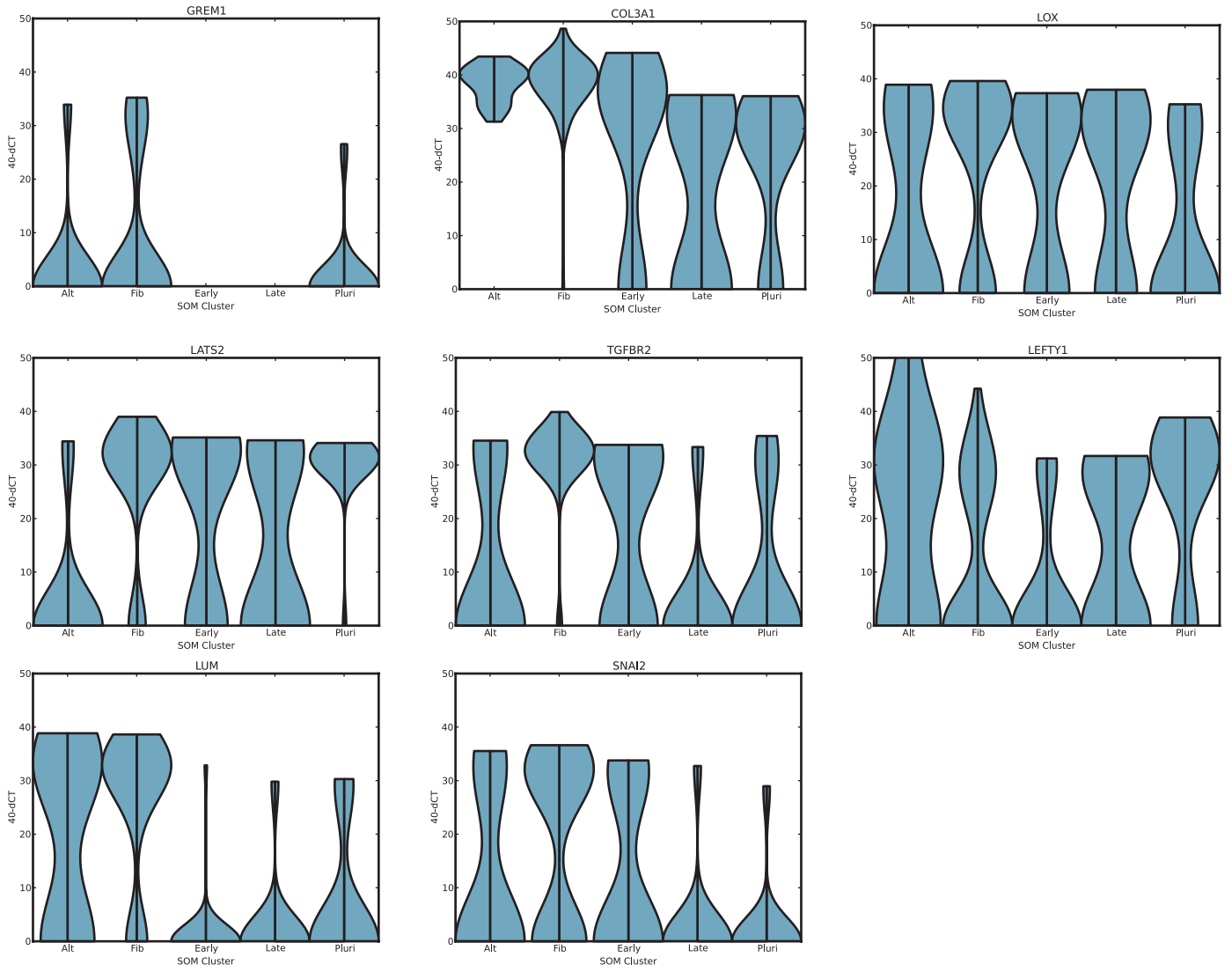
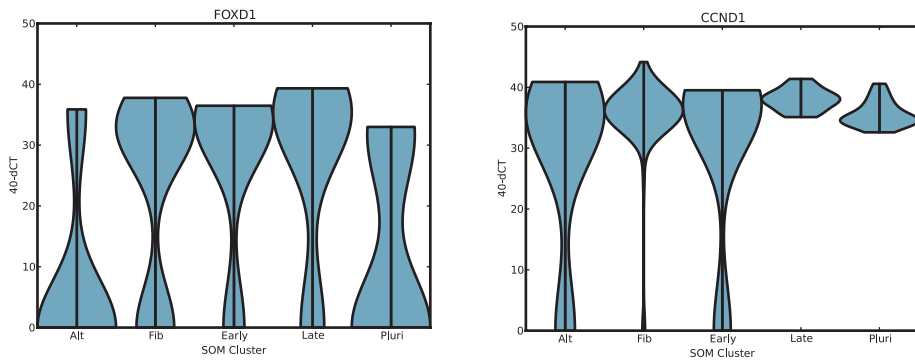


Figure S3

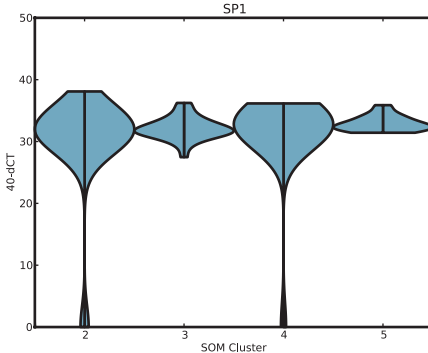
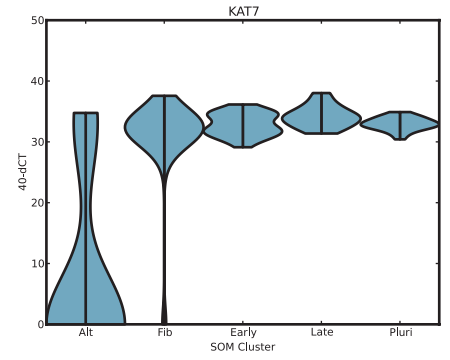
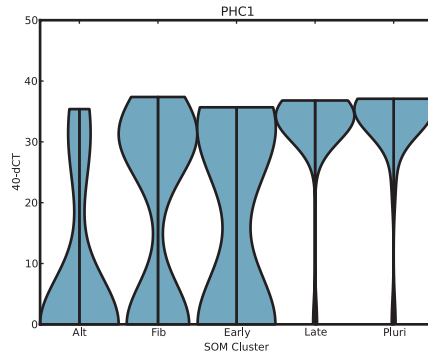
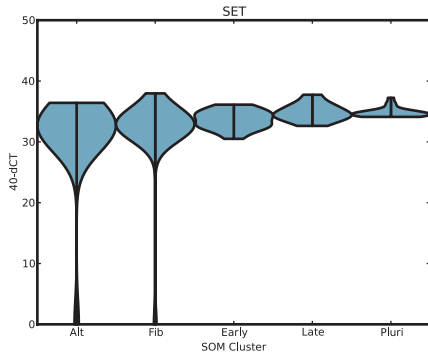
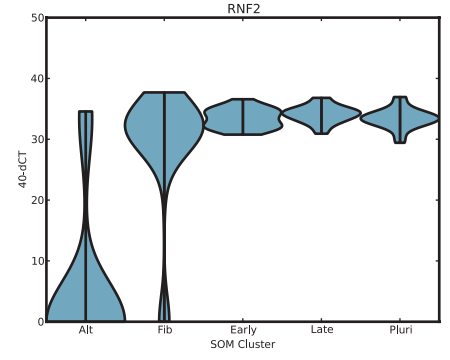
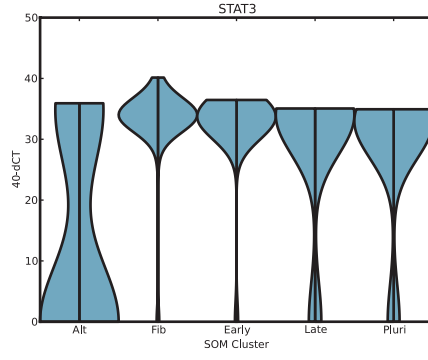
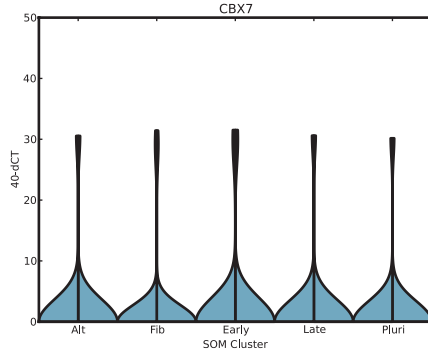
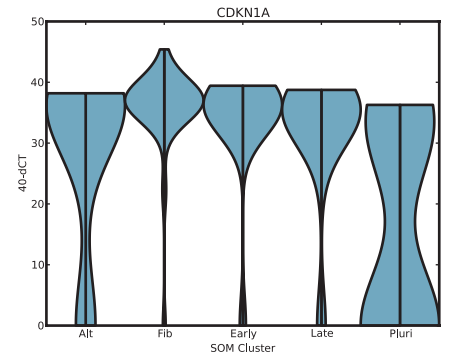
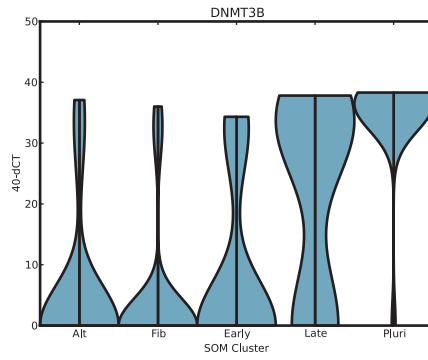
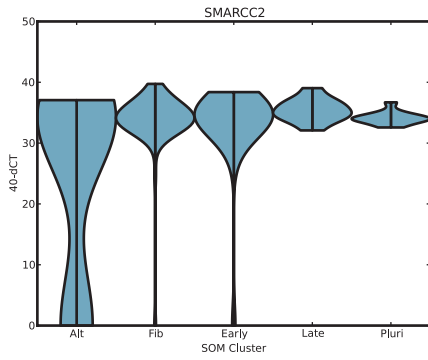
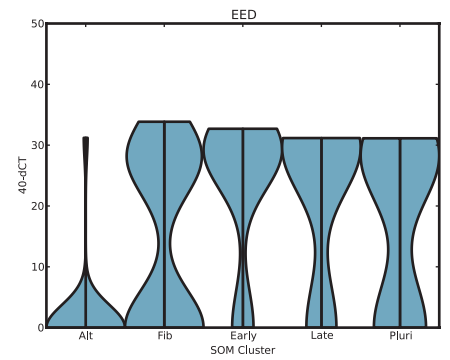
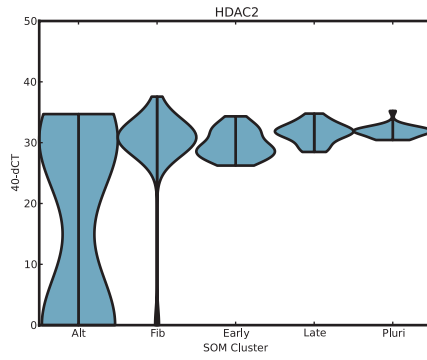
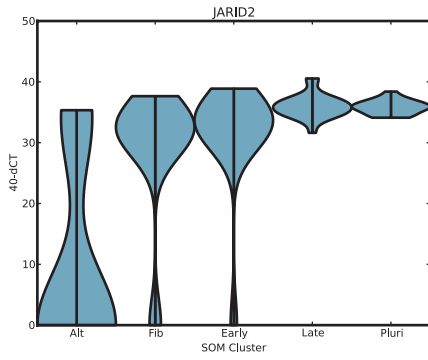
## Fibroblast



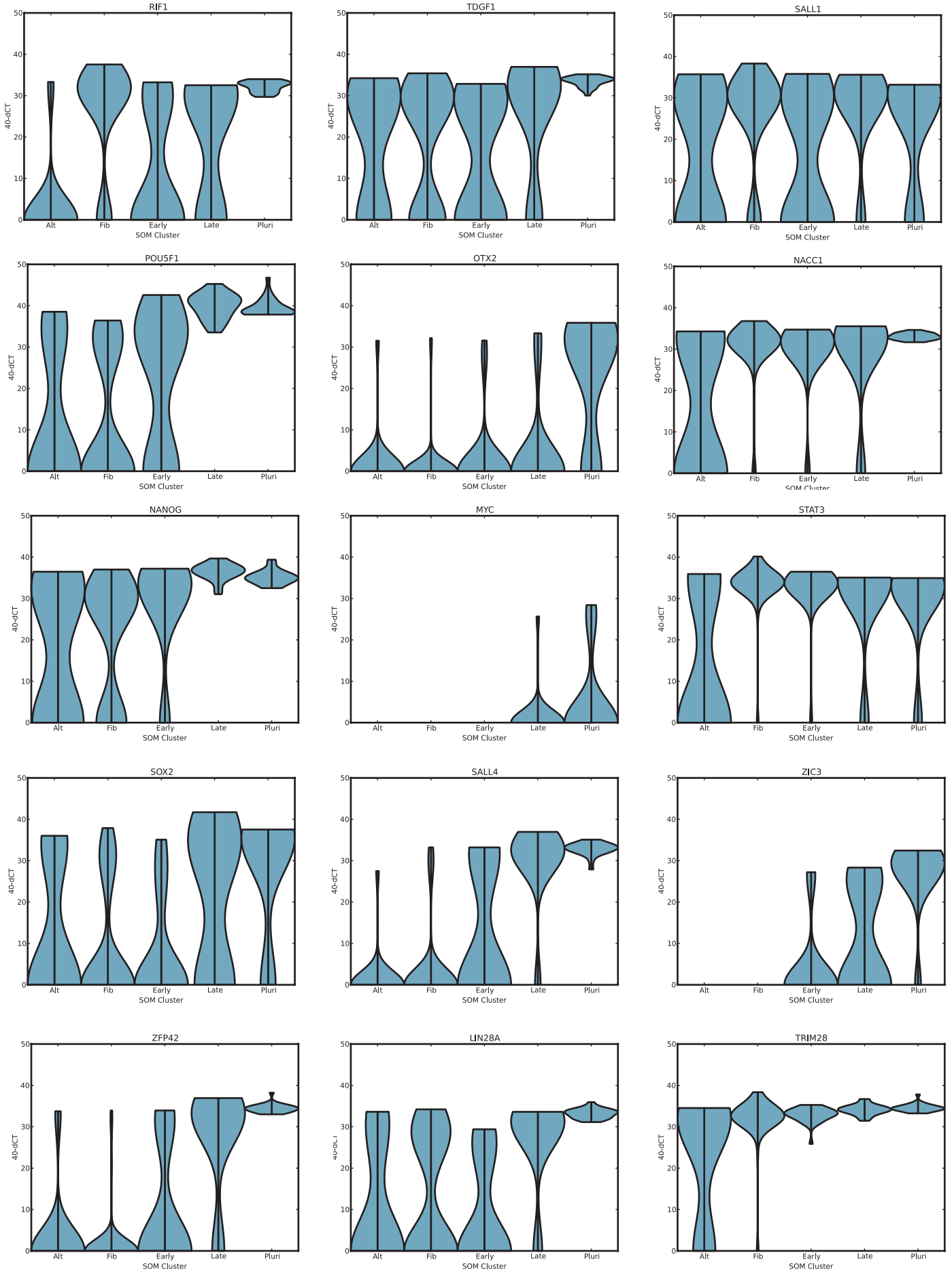
## Intermediate



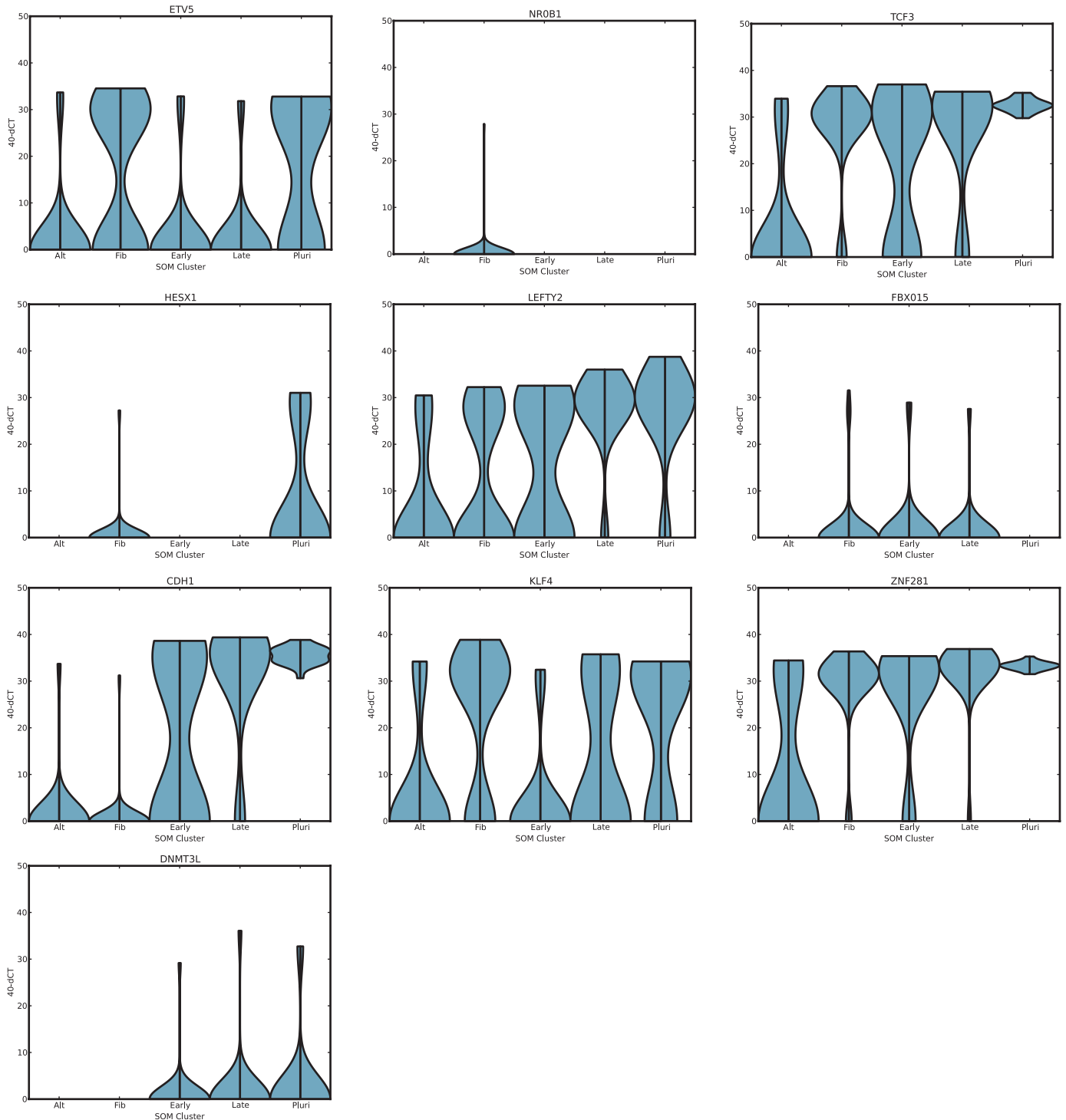
# Chromatin Modifiers / Cell Cycle



# Pluripotency



## Pluripotency (cont.)



**Figure S3:** Violin Plots Depicting Quantitative Changes in Gene Expression During Reprogramming

Cells were separated by SOM grouping (outliers excluded) and gene expression levels were plotted for each cell within the group. The width of the violin represents the distribution of cells across expression levels. For the majority of genes, there is a clear inflection point in these graphs where a large number of cells up or down regulate a given transcript when transitioning from one SOM group to another, eventually coalescing around high or low expression levels as cells approach pluripotency and cellular phenotypes stabilize.