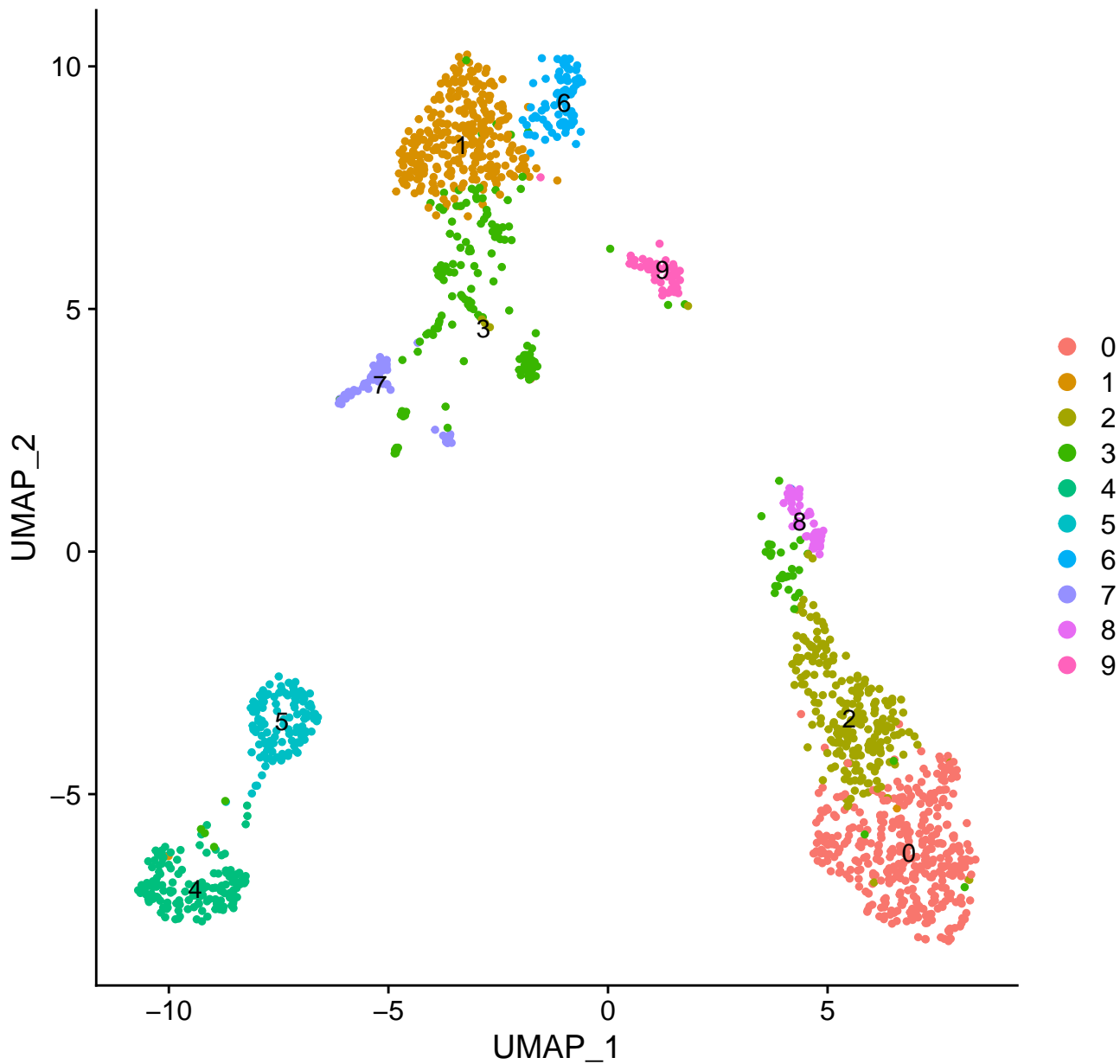
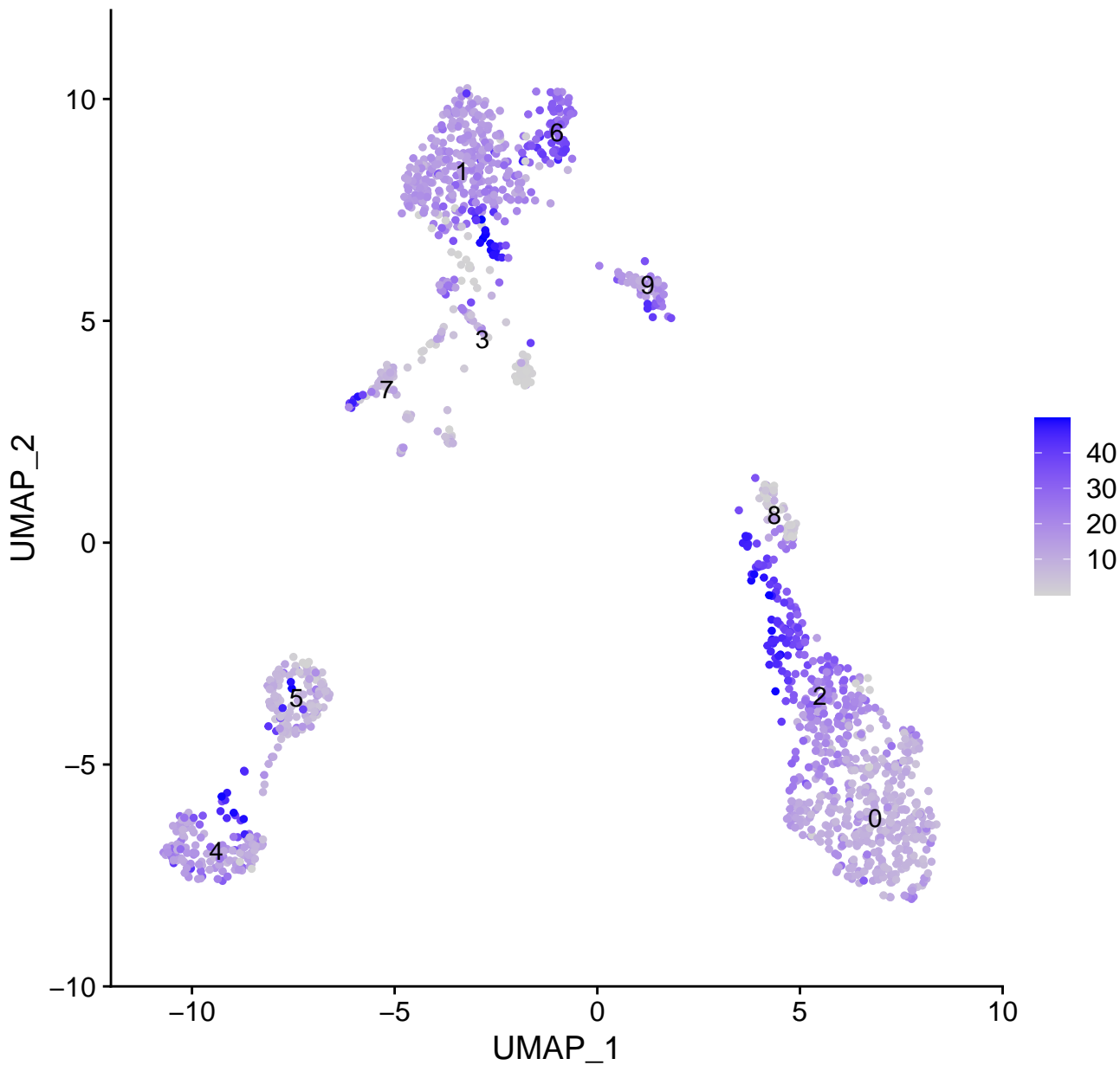


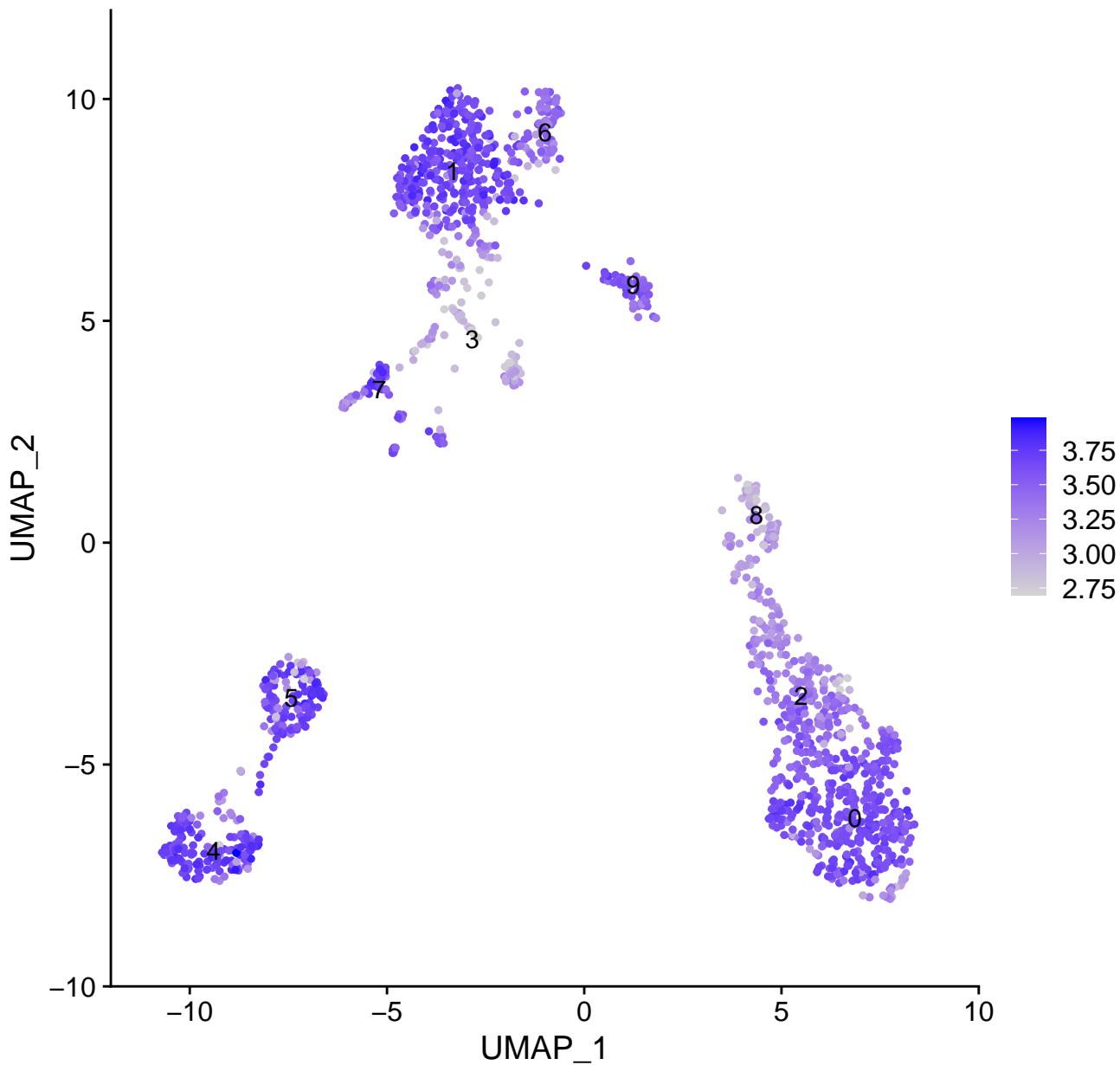
MS19035_Islet71 QC clusters



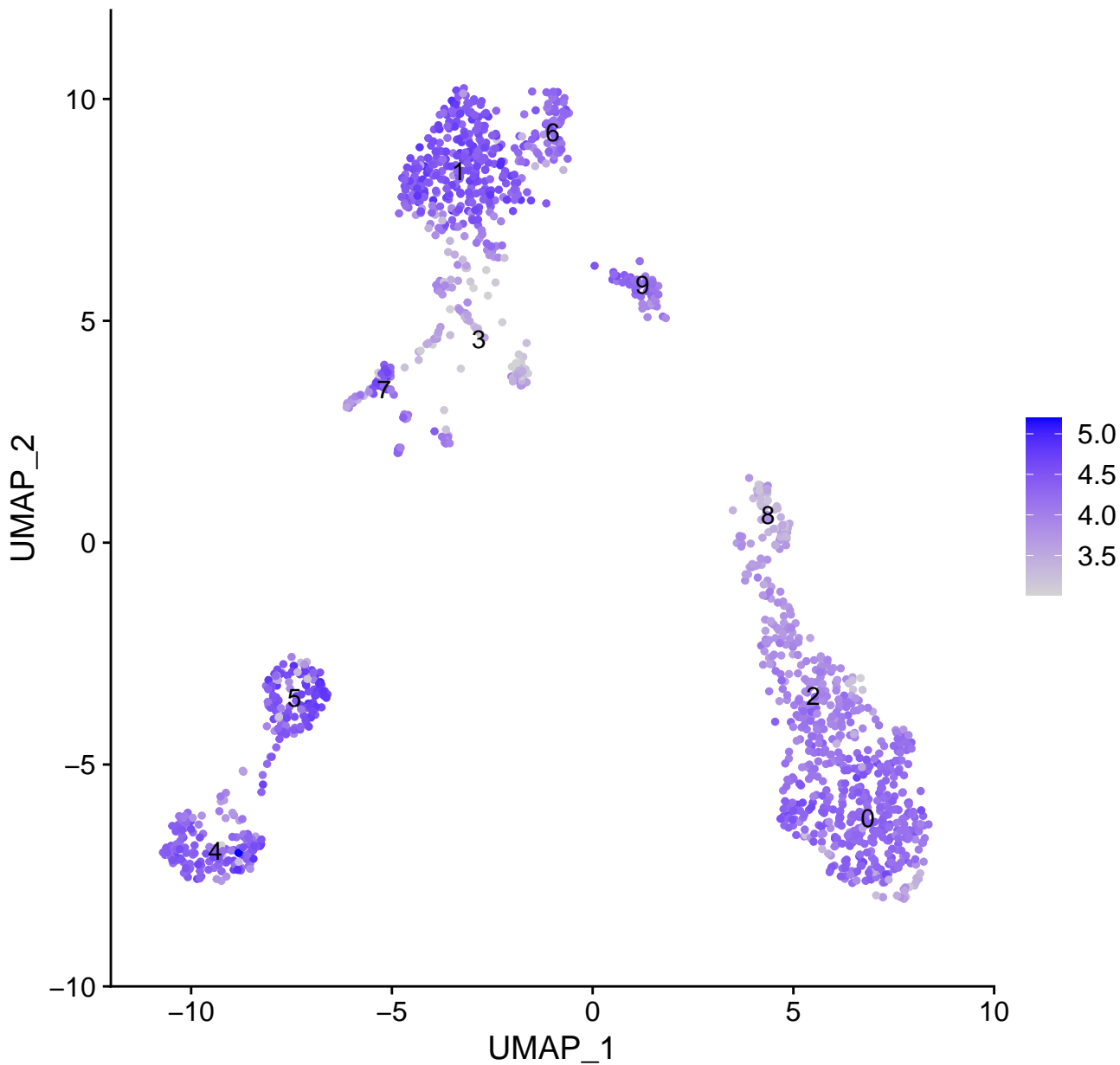
MS19035_Islet71 percent.mt



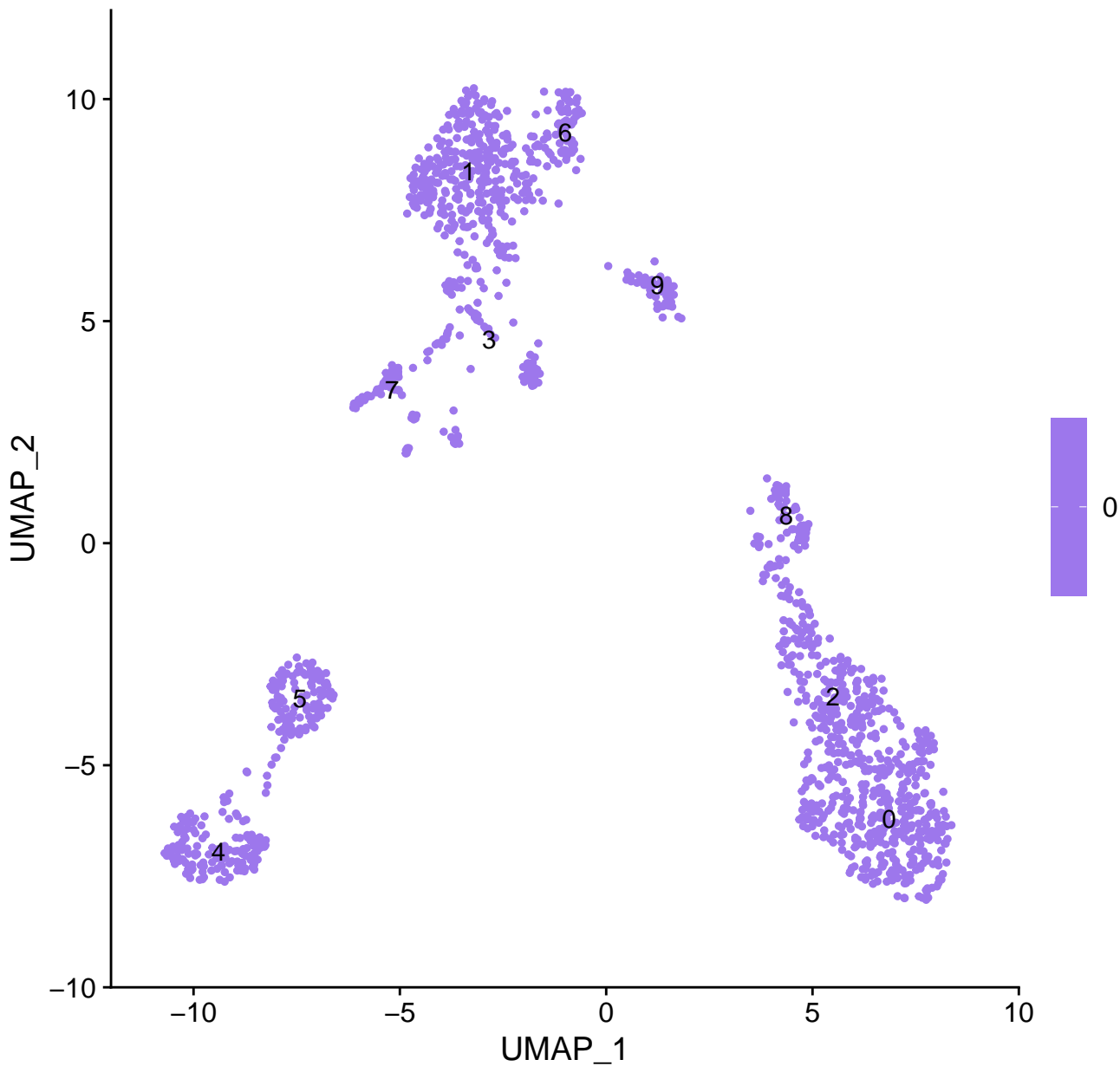
MS19035_Islet71 nFeature_RNA



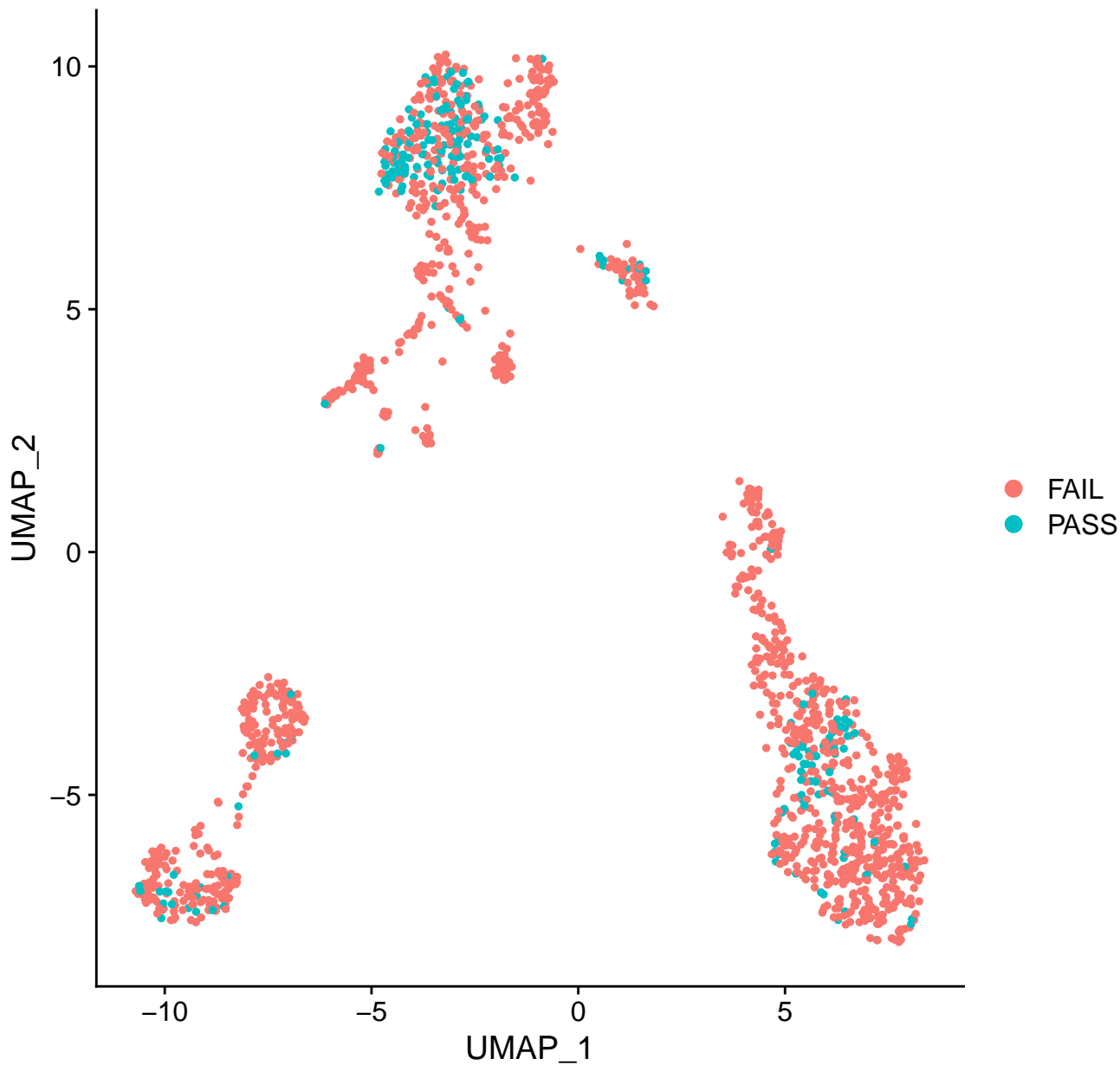
MS19035_Islet71 nCount_RNA



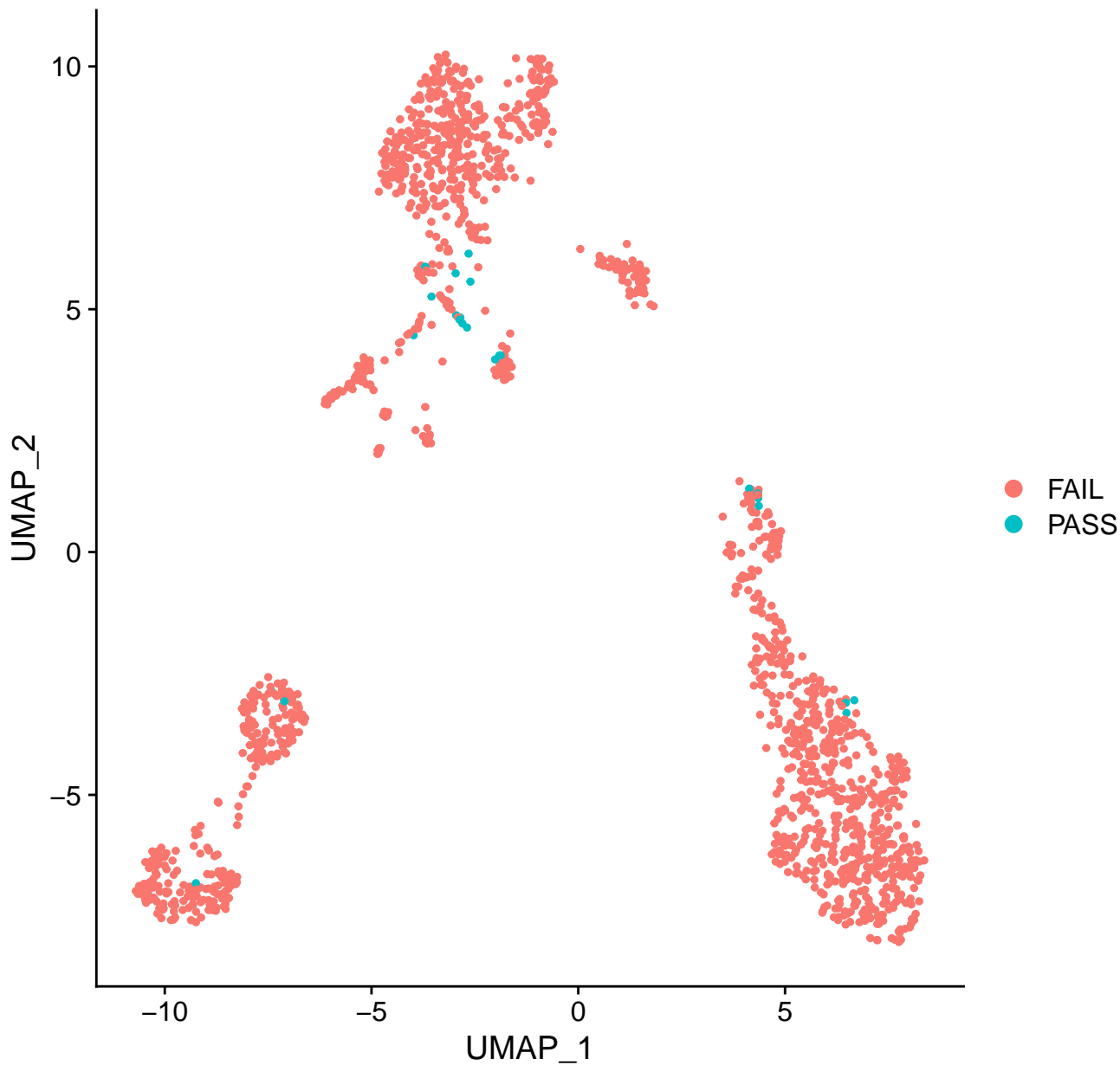
MS19035_Islet71 percent.hb



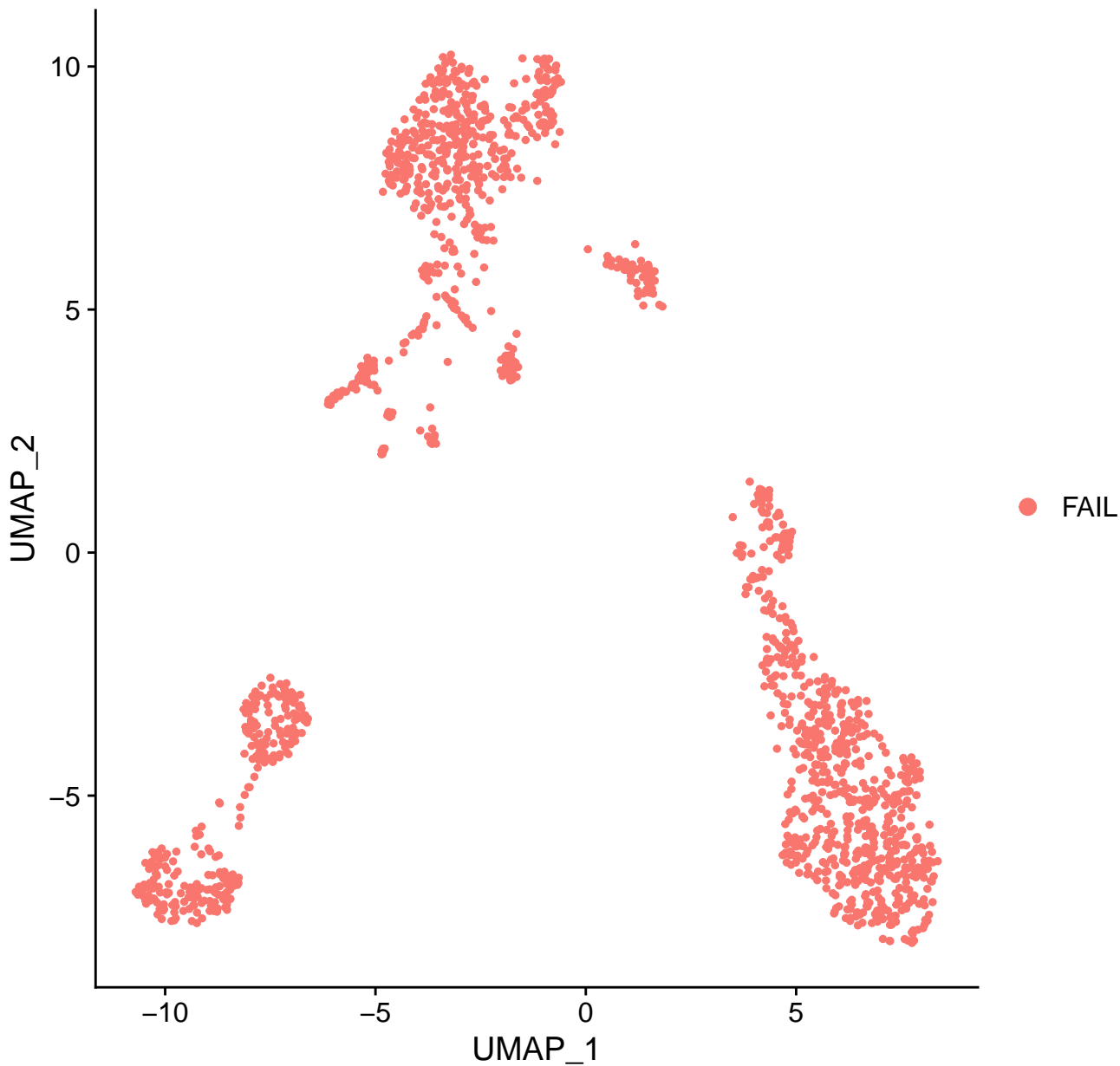
MS19035_Islet71 percent.mt (cutoffs)



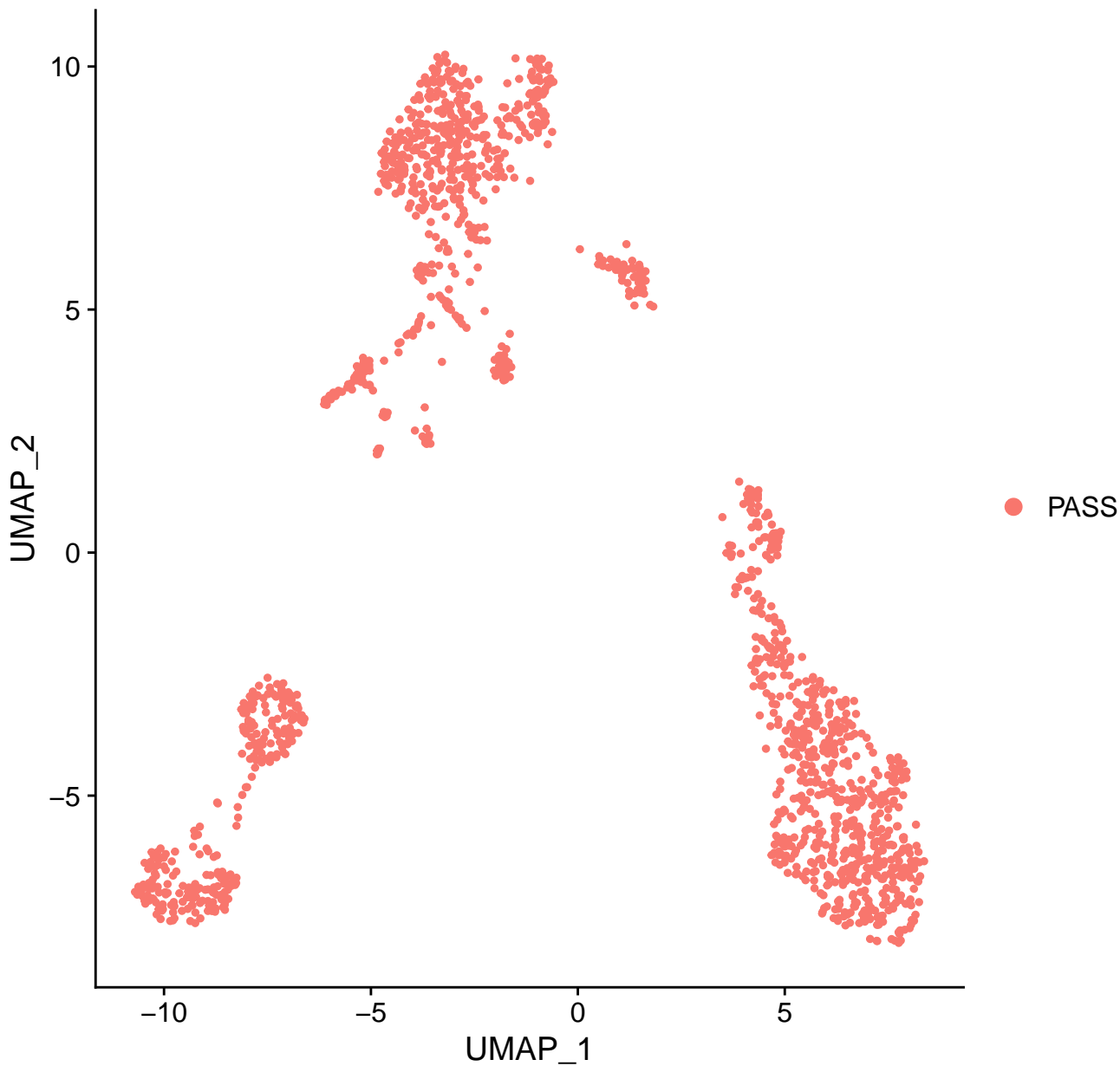
MS19035_Islet71 nFeature_RNA (cutoffs)



MS19035_Islet71 nCount_RNA (cutoffs)

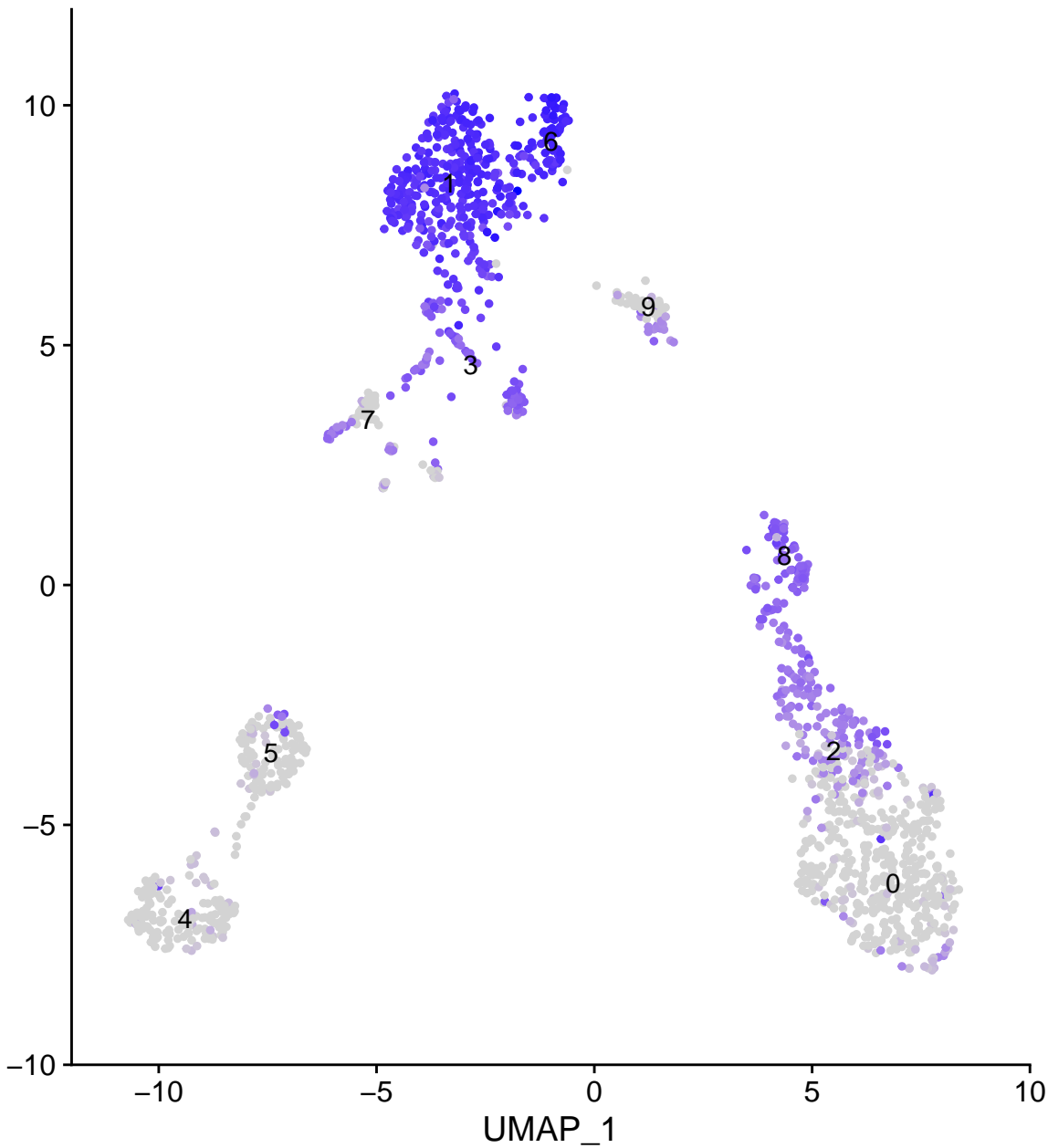


MS19035_Islet71 percent.hb (cutoffs)

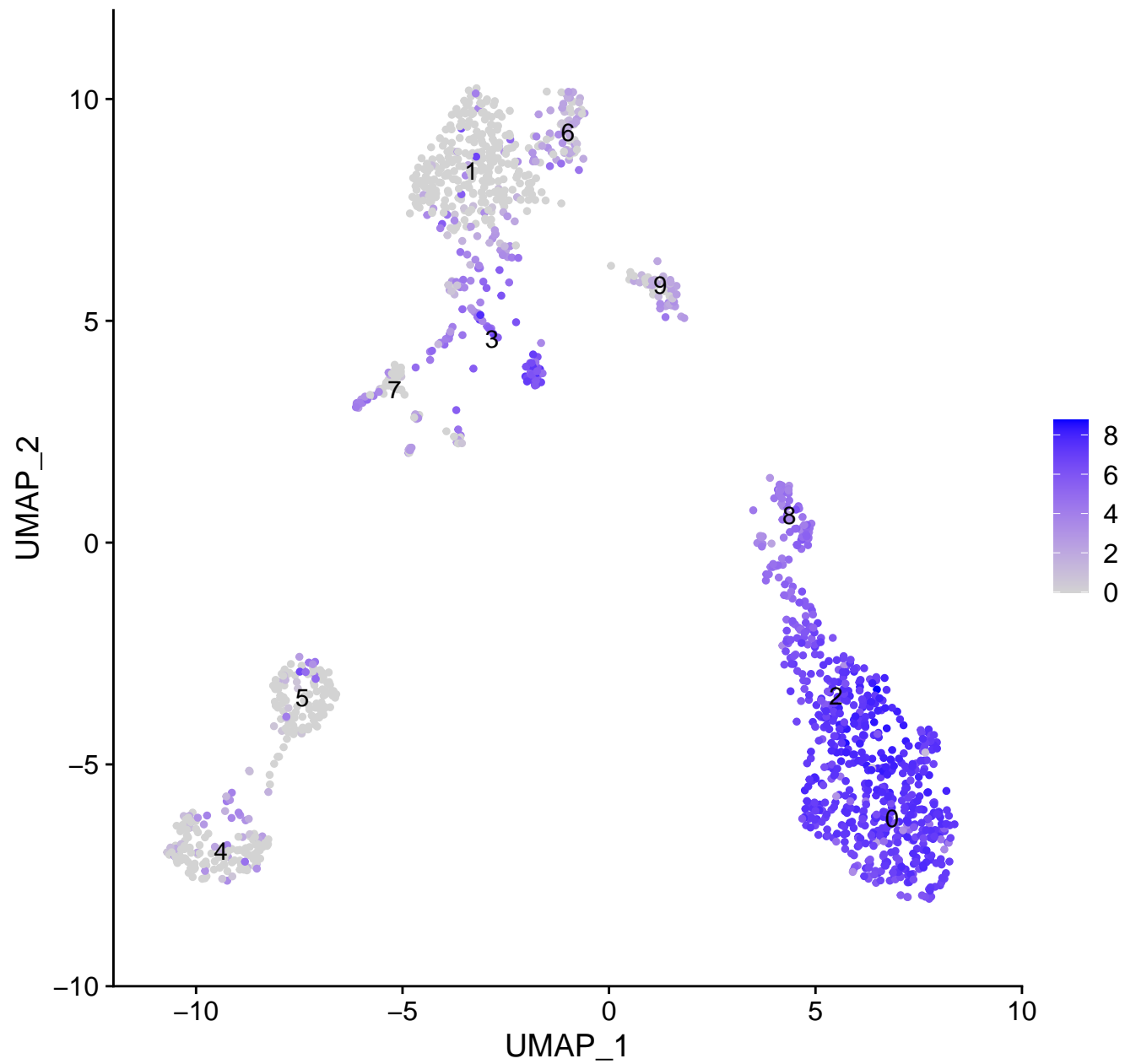


INS

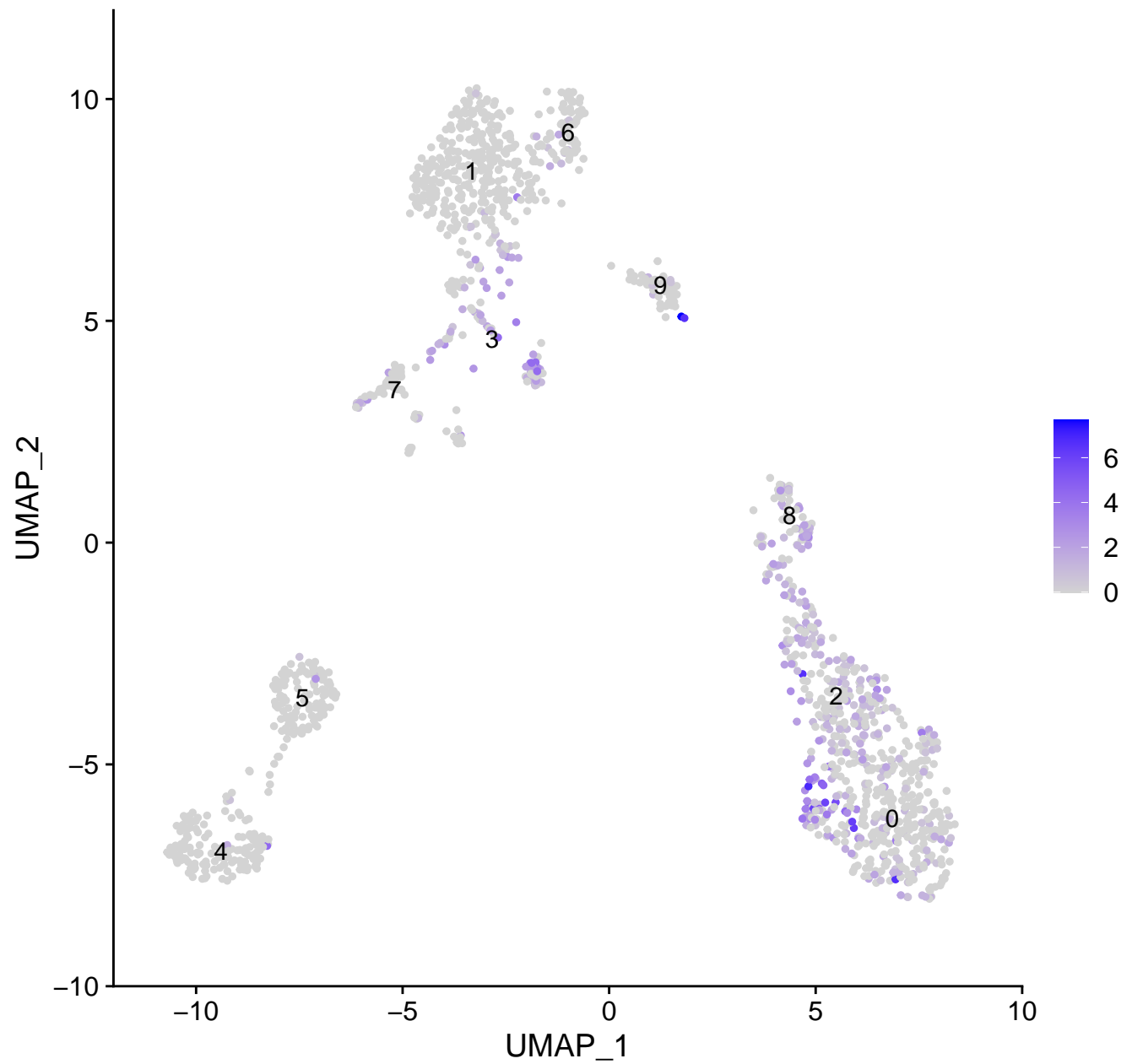
UMAP_2



GCG

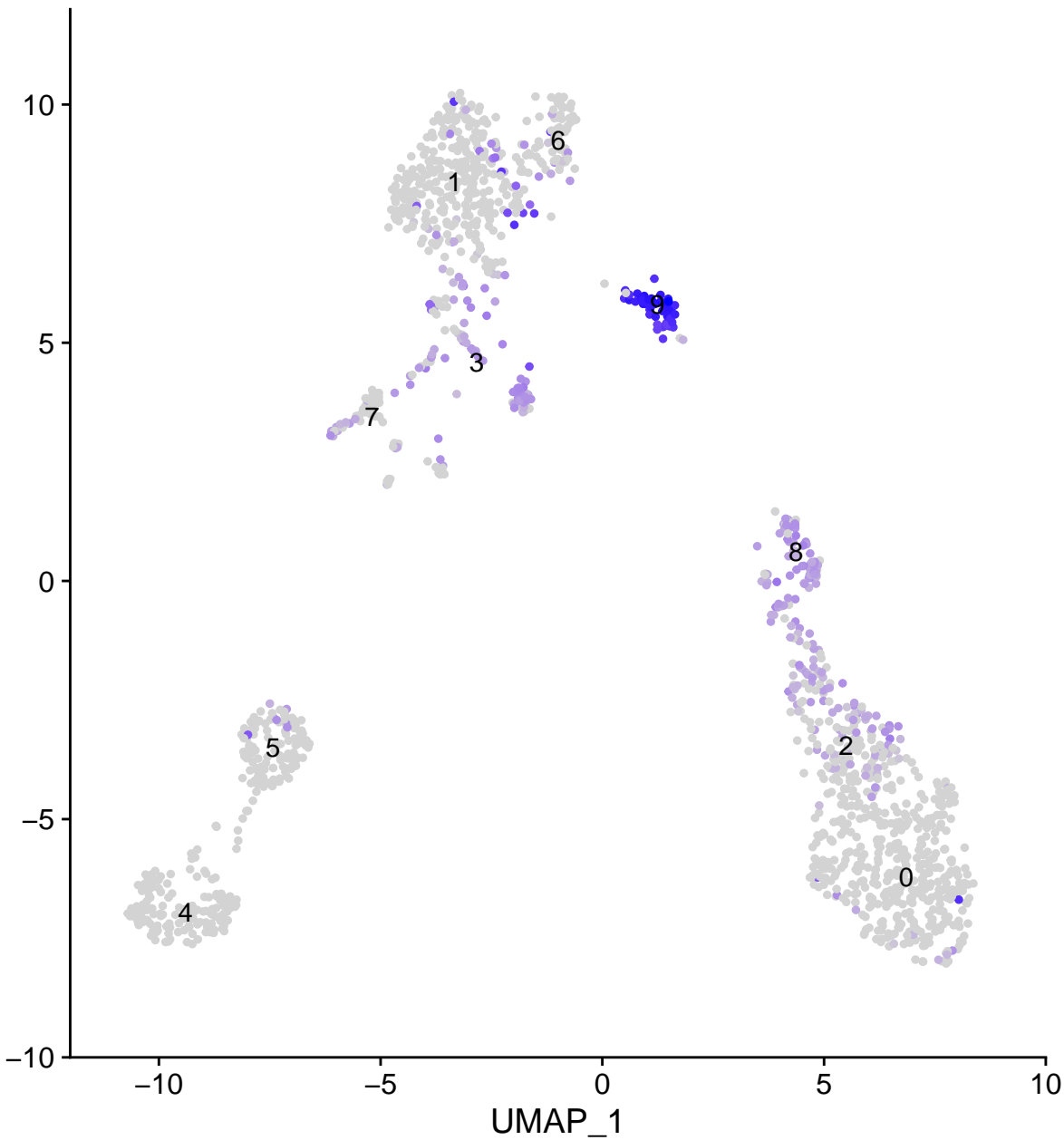


PPY

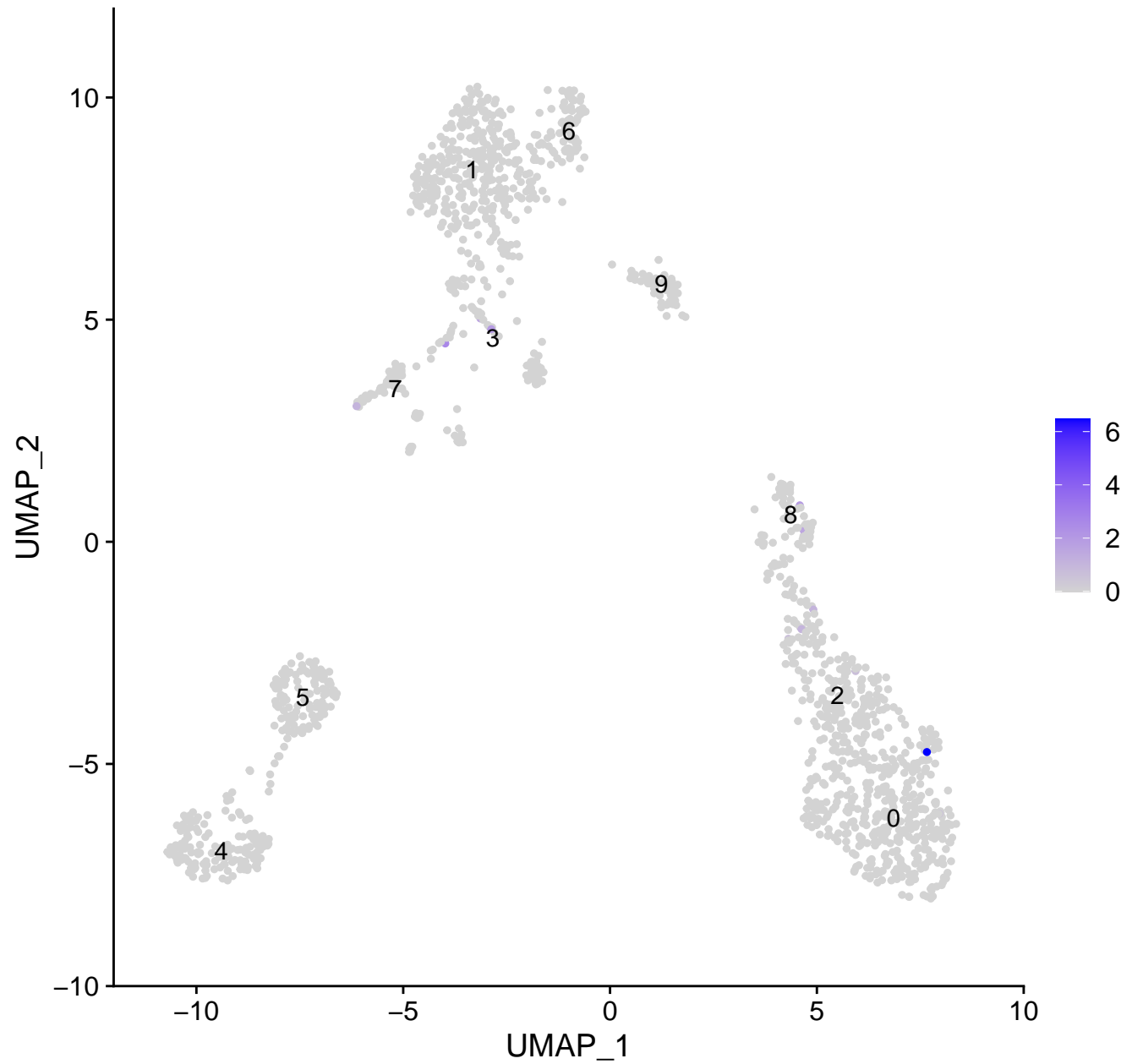


SST

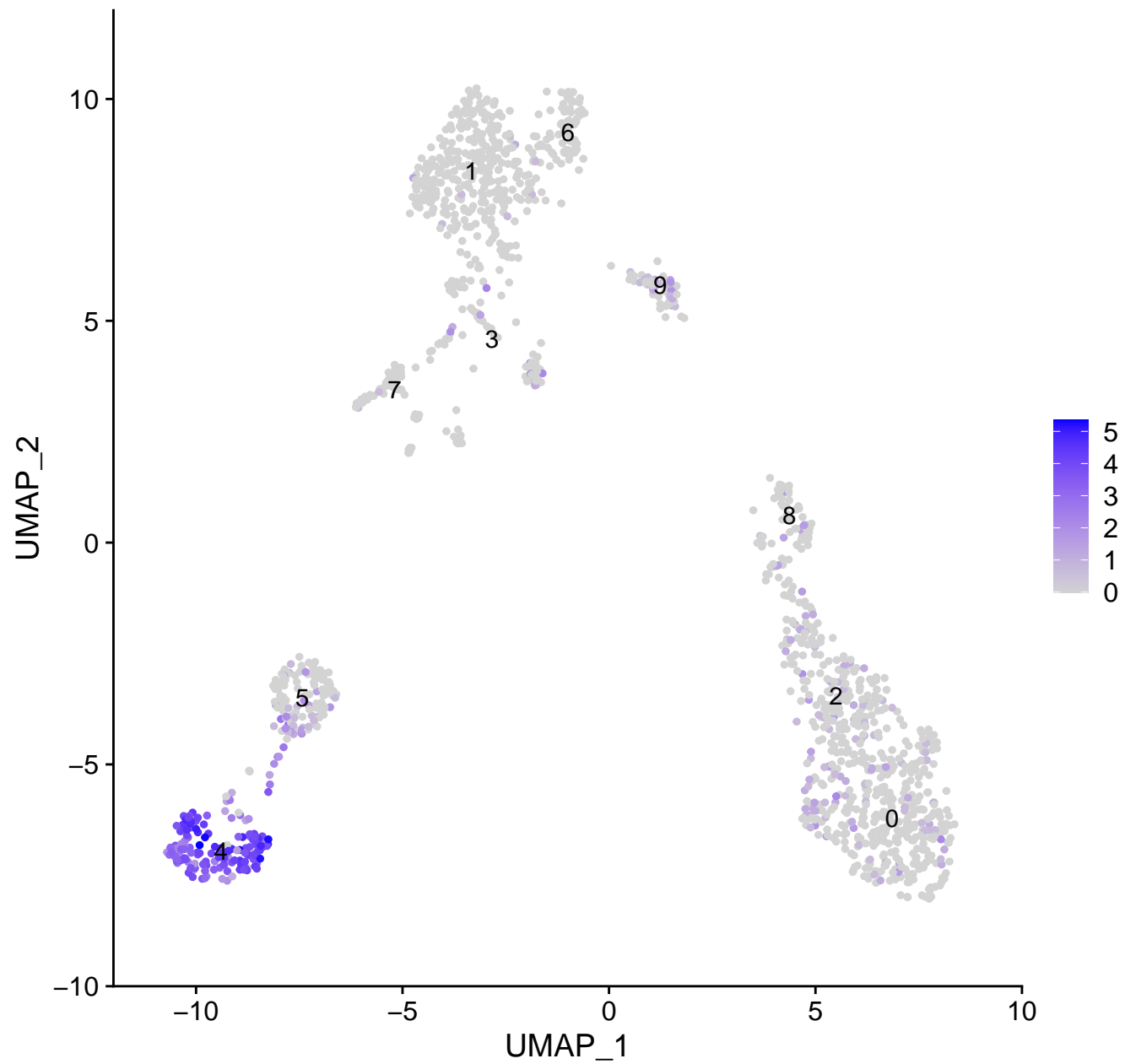
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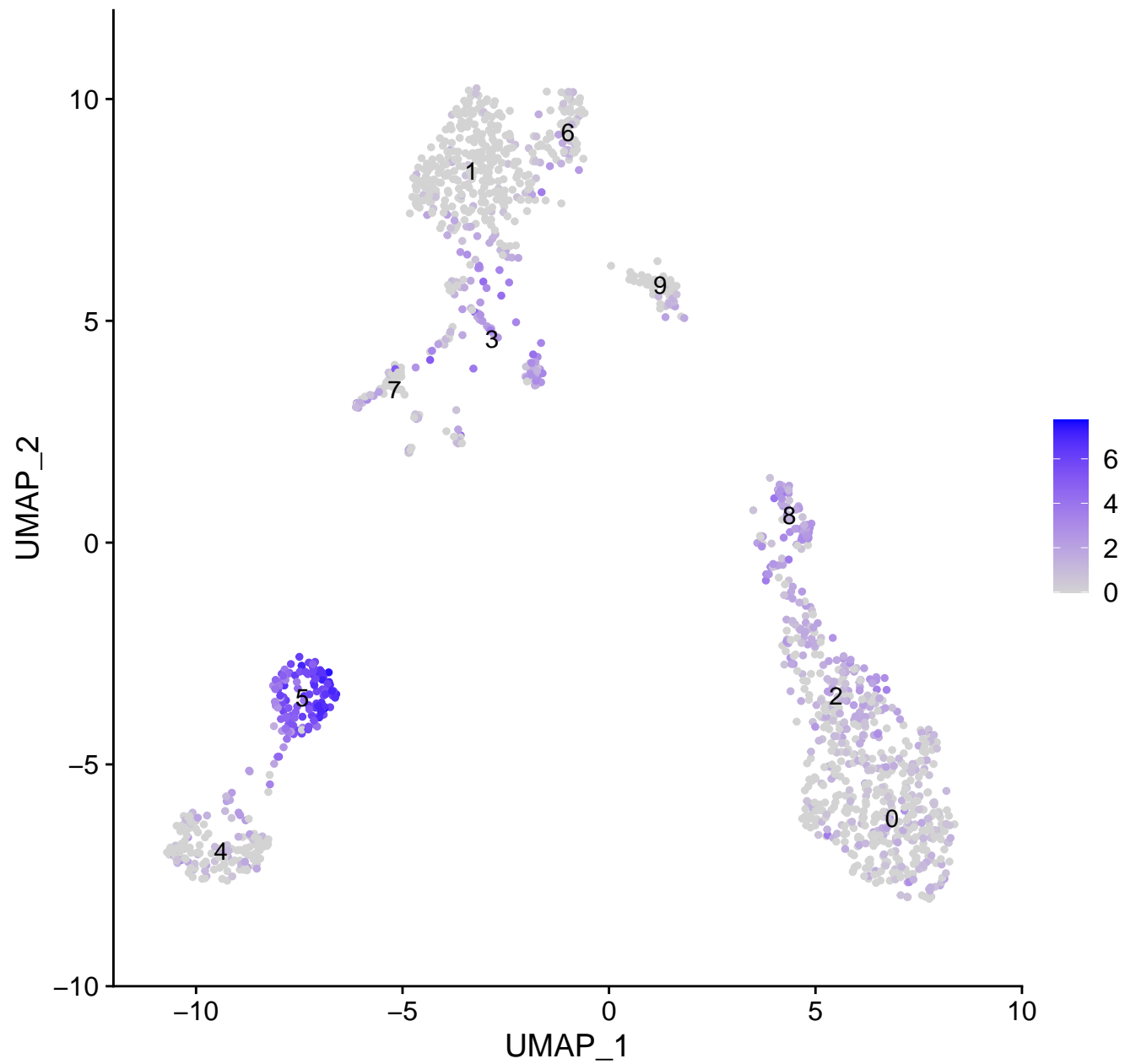
GHRL



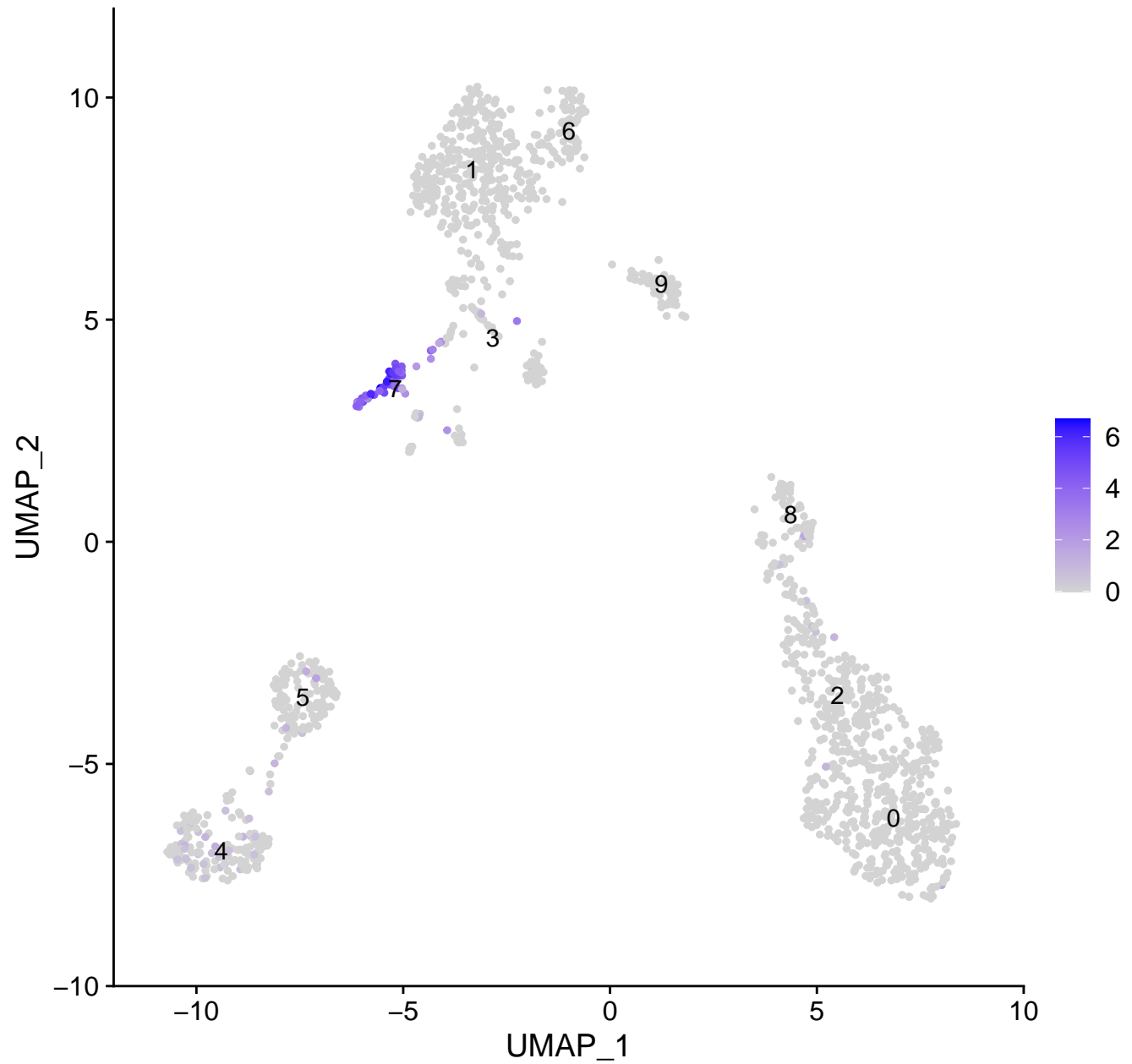
KRT19



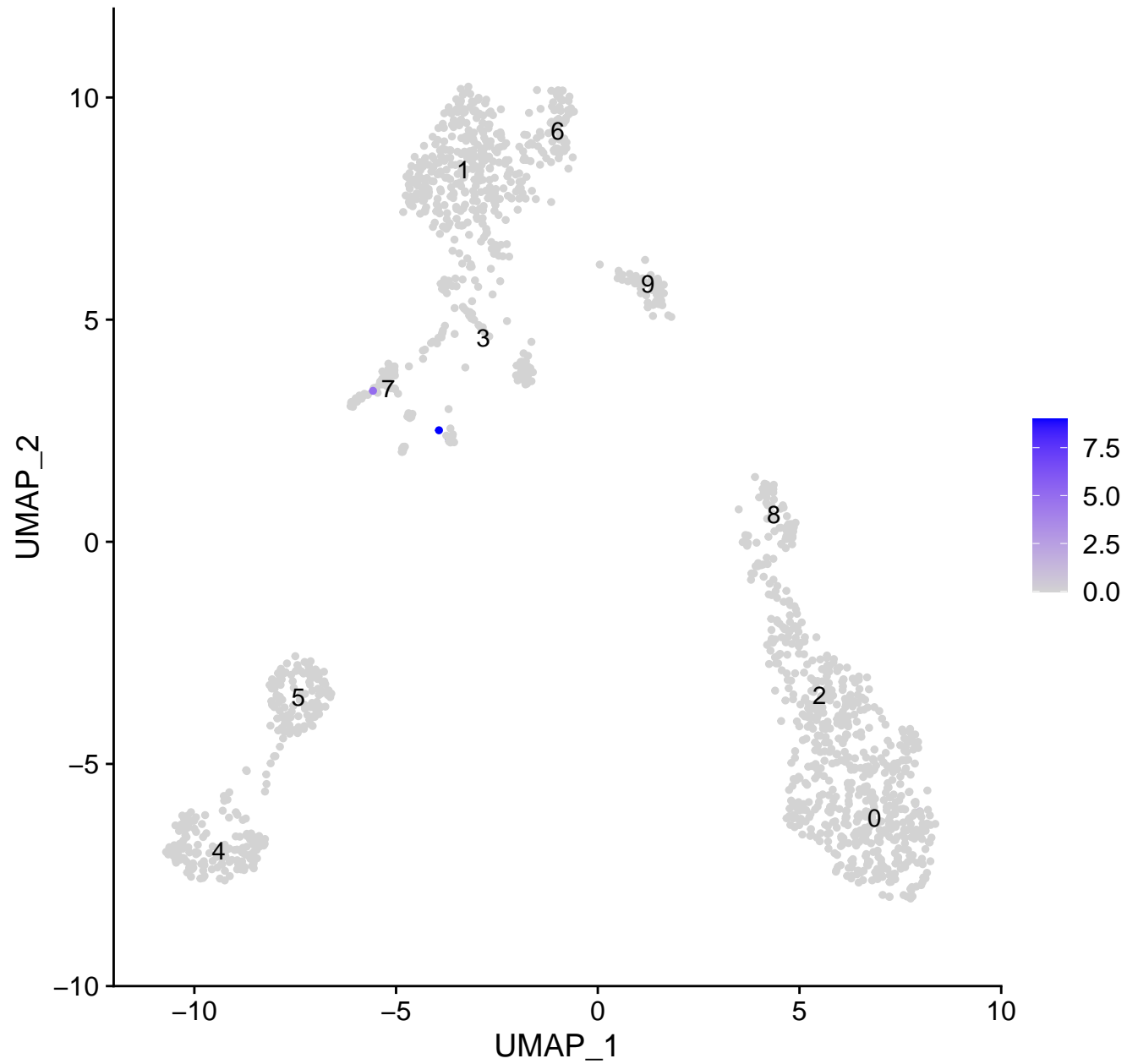
REG1B



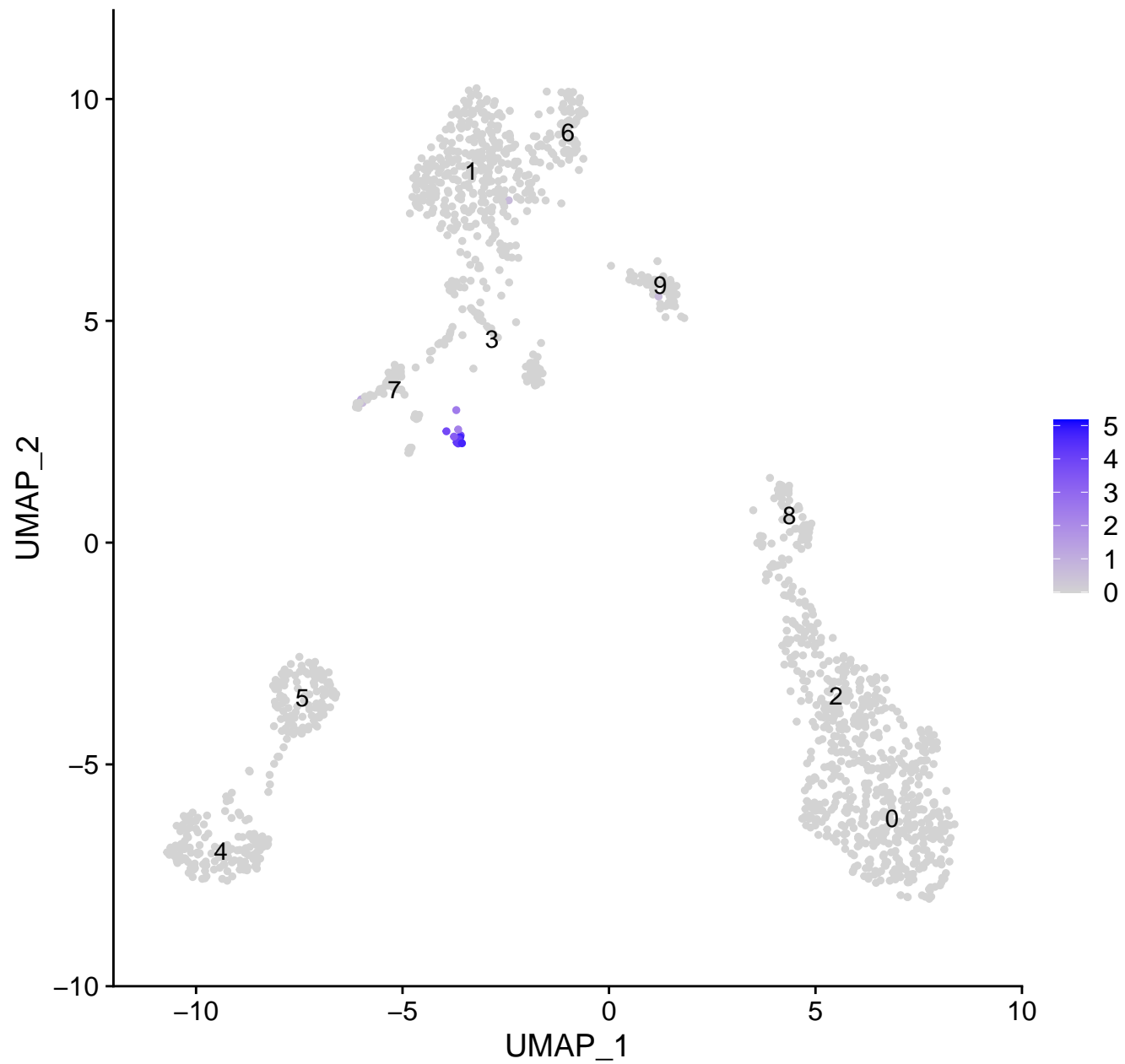
COL1A1



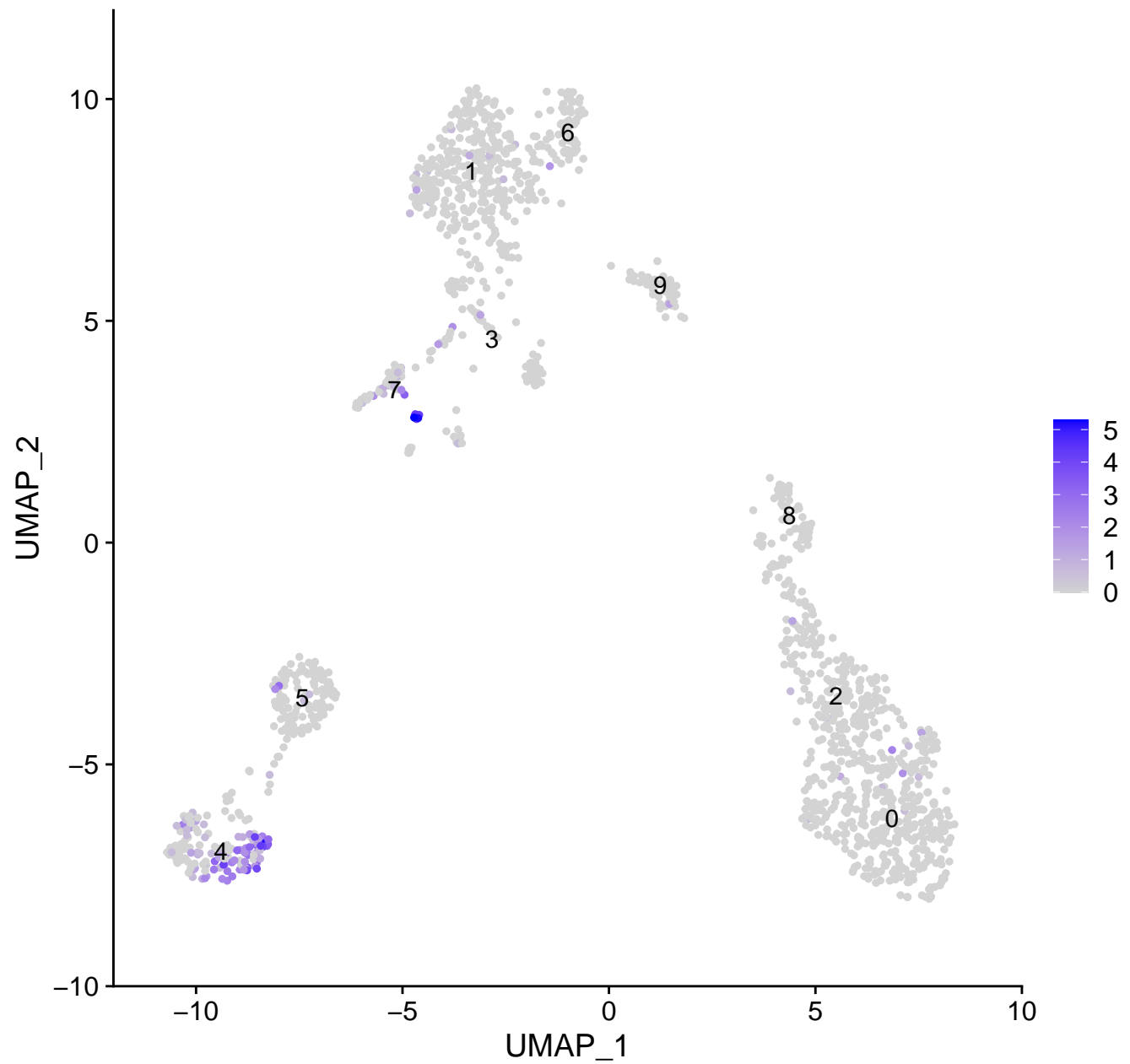
rna_FABP4



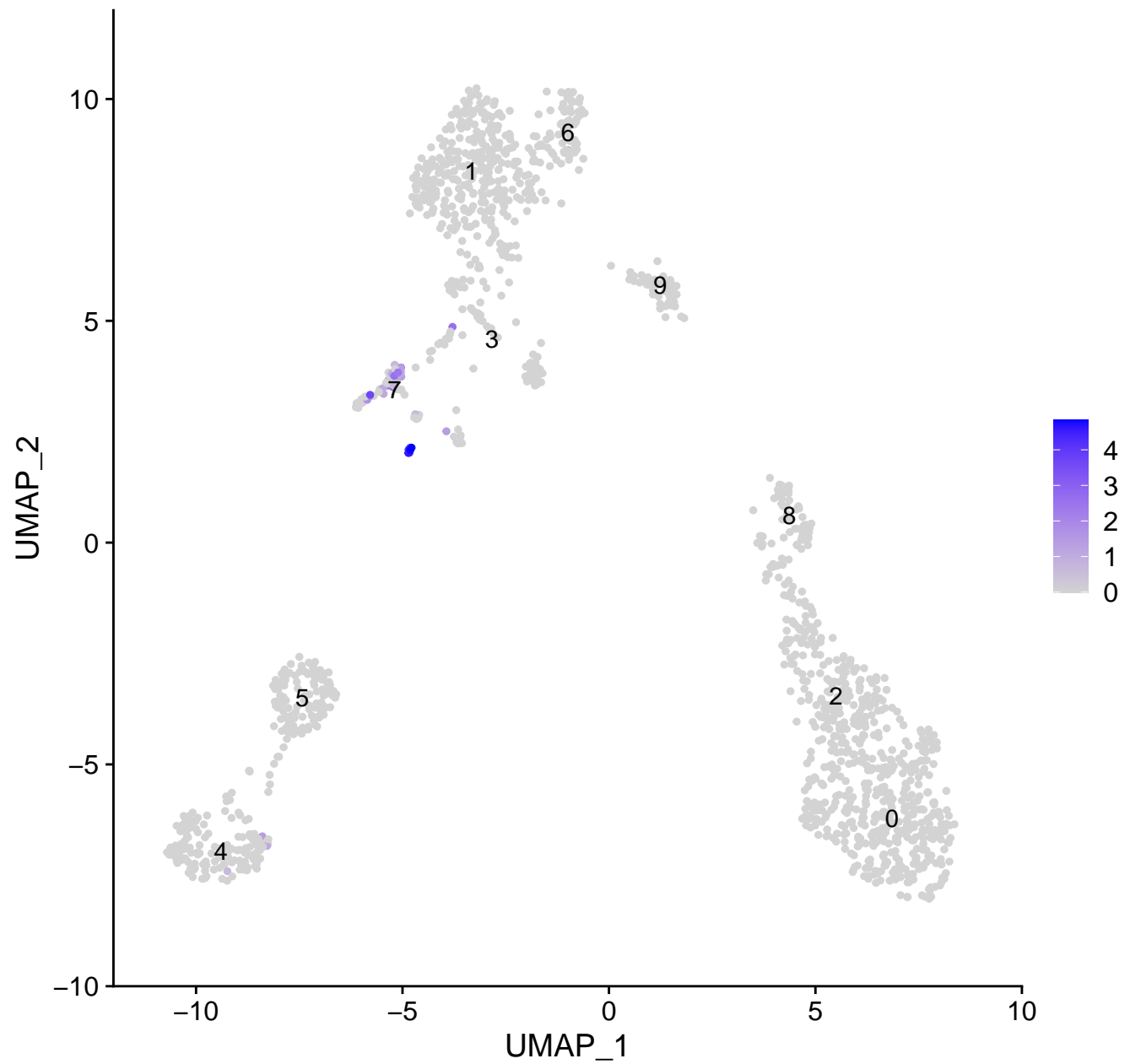
PLVAP



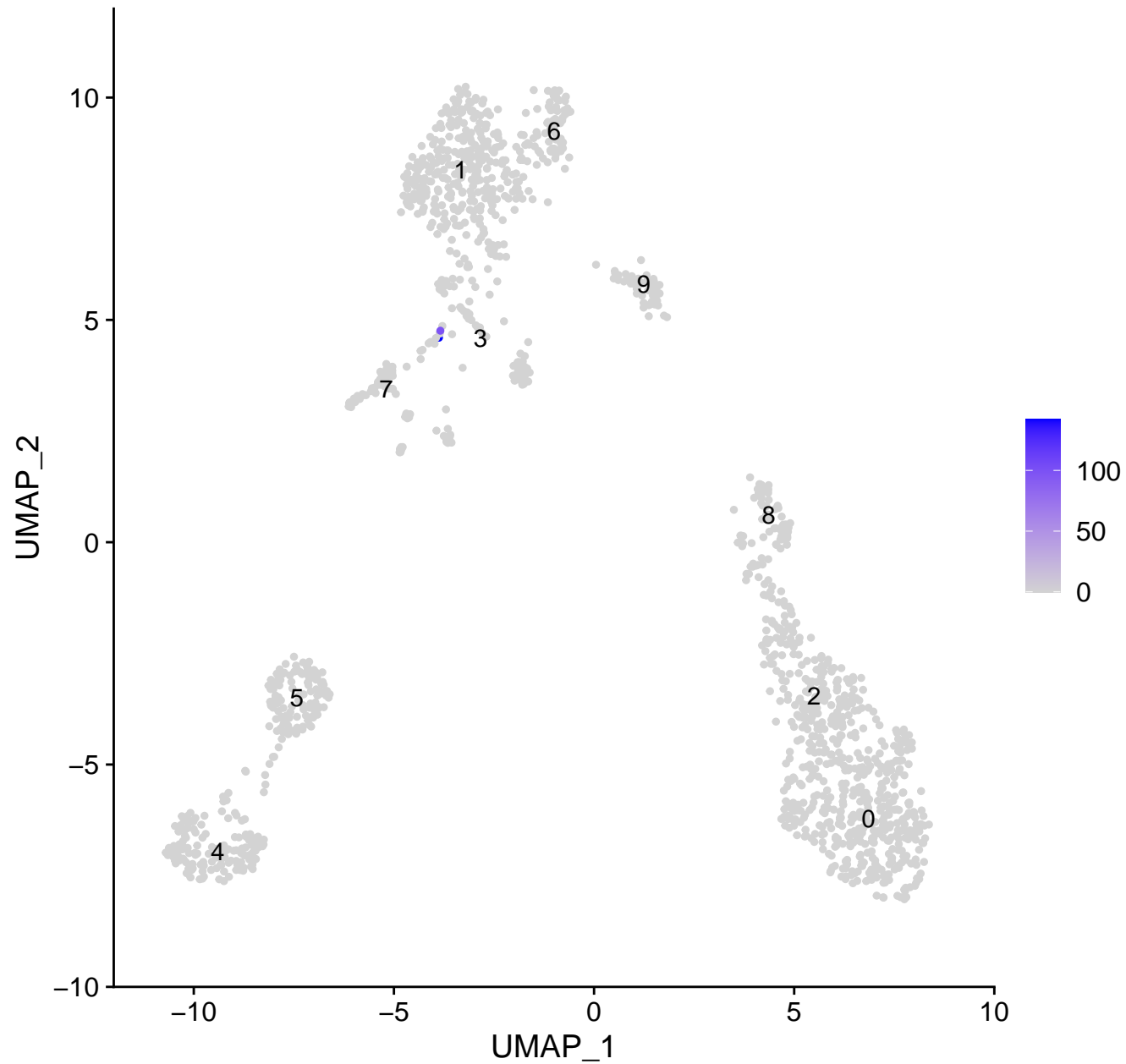
CRYAB



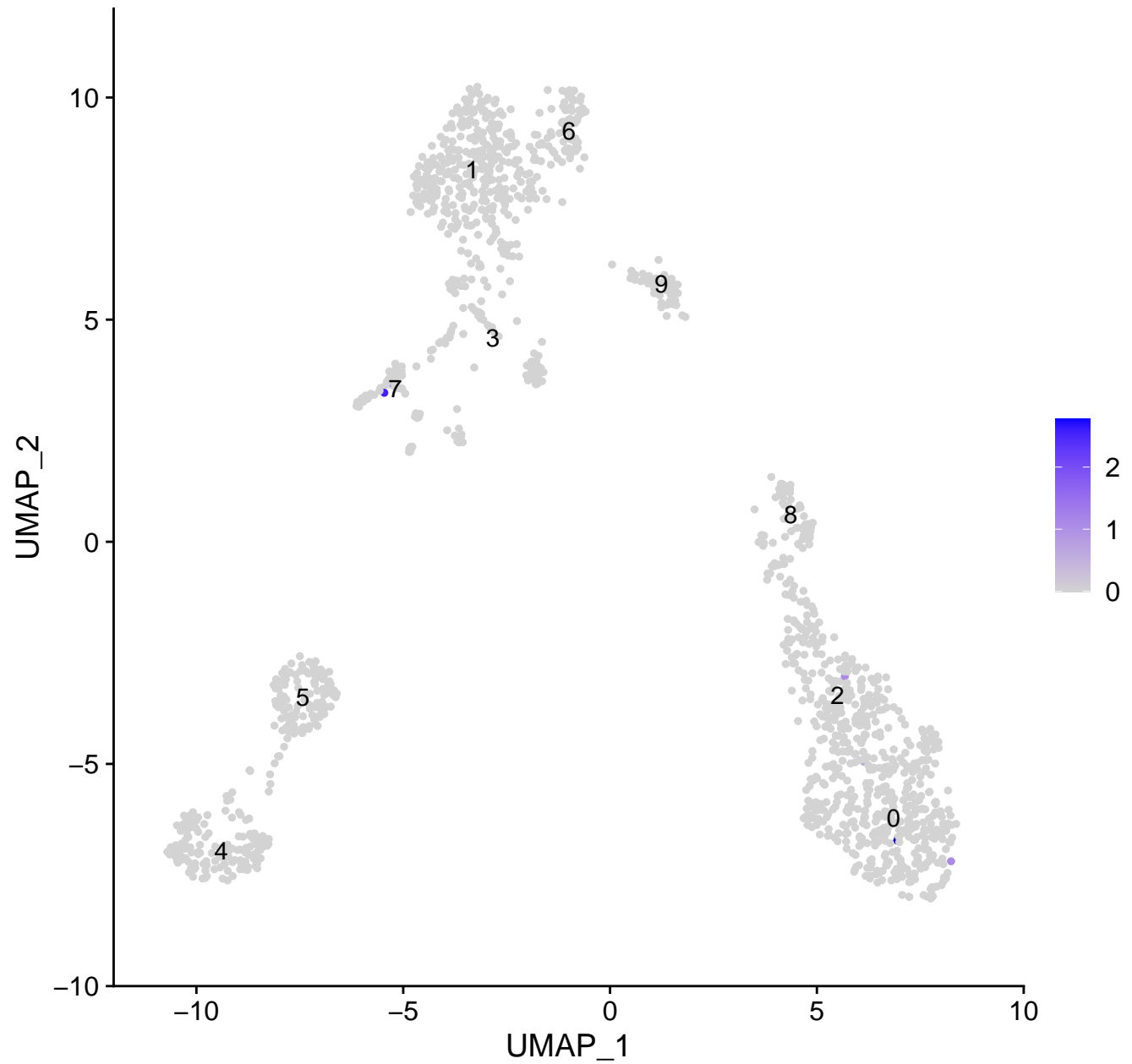
APOE



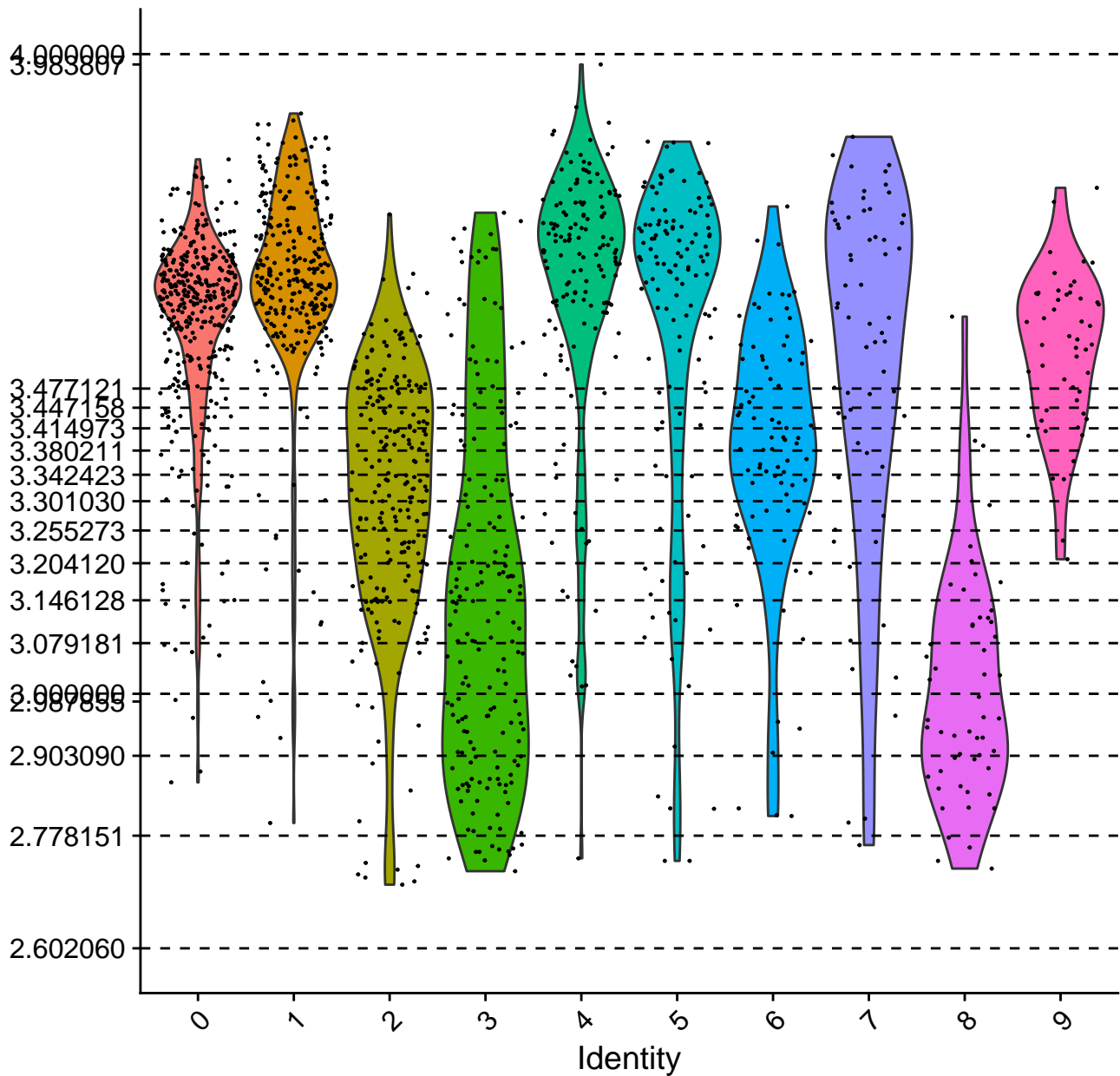
rna_TPSB2



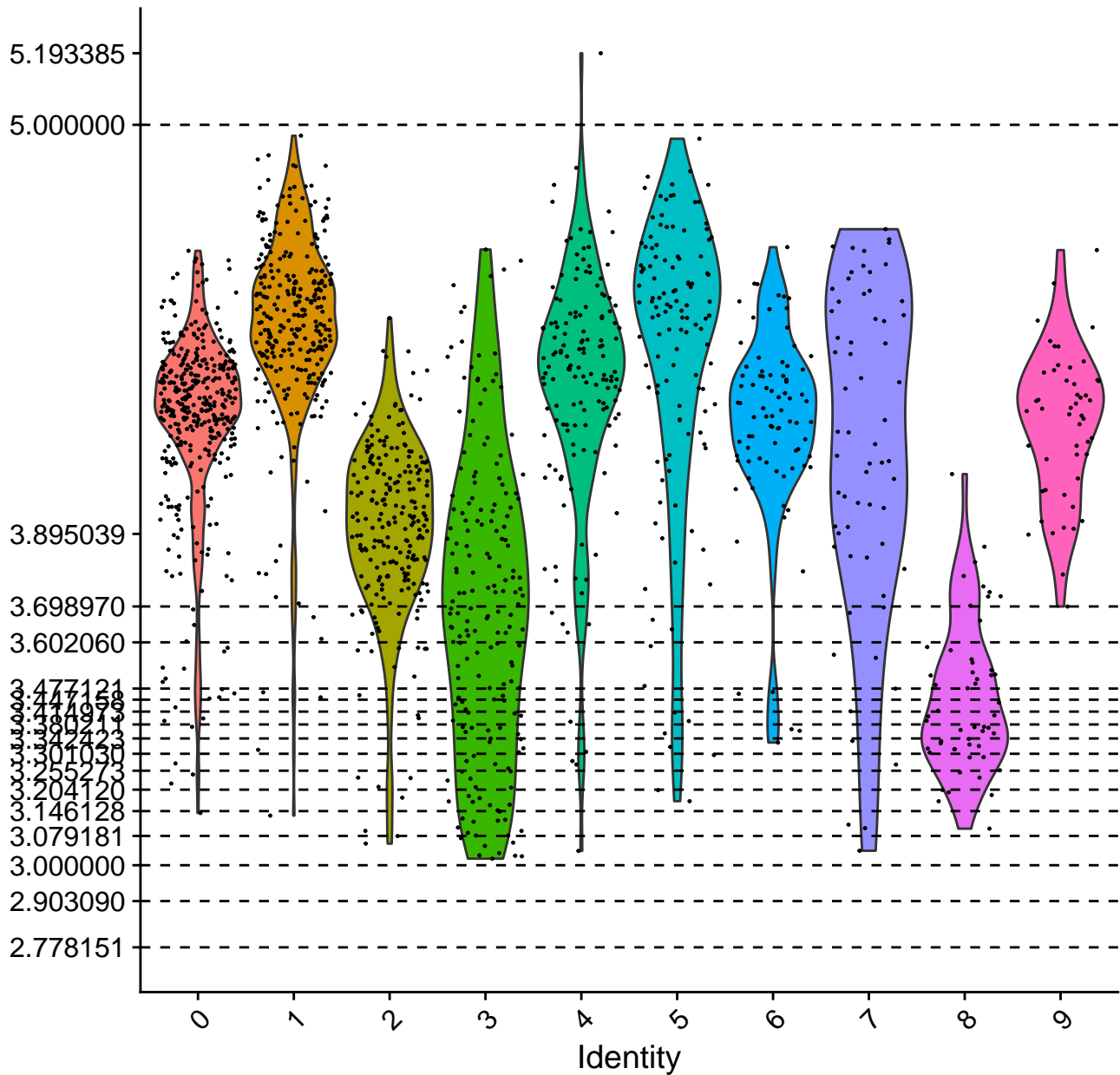
TOP2A



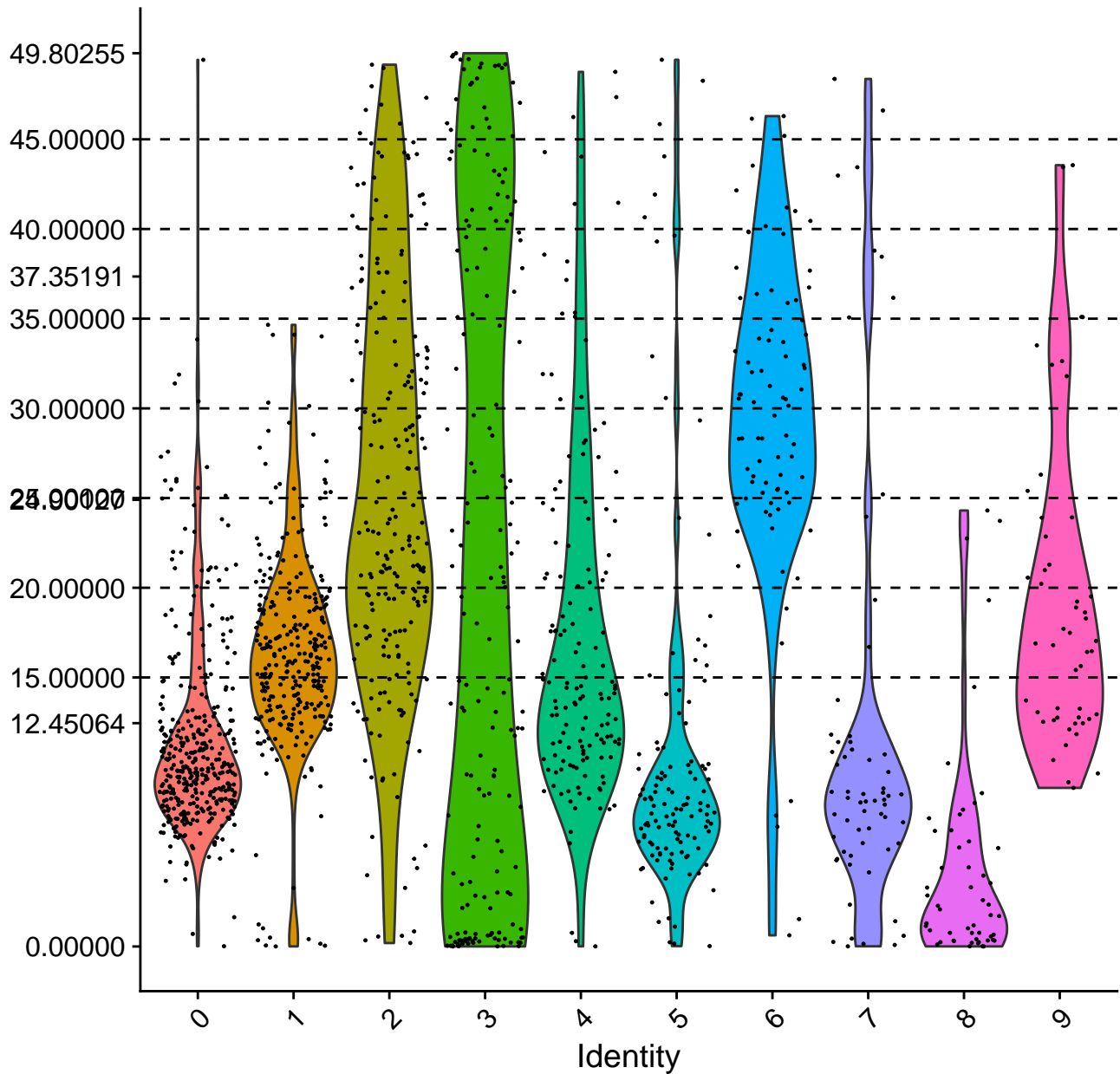
Number of features for MS19035_Islet71 per cluster



Number of counts for MS19035_Islet71 per cluster



% of MT reads for MS19035_Islet71 per cluster



% of HB reads for MS19035_Islet71 per cluster

