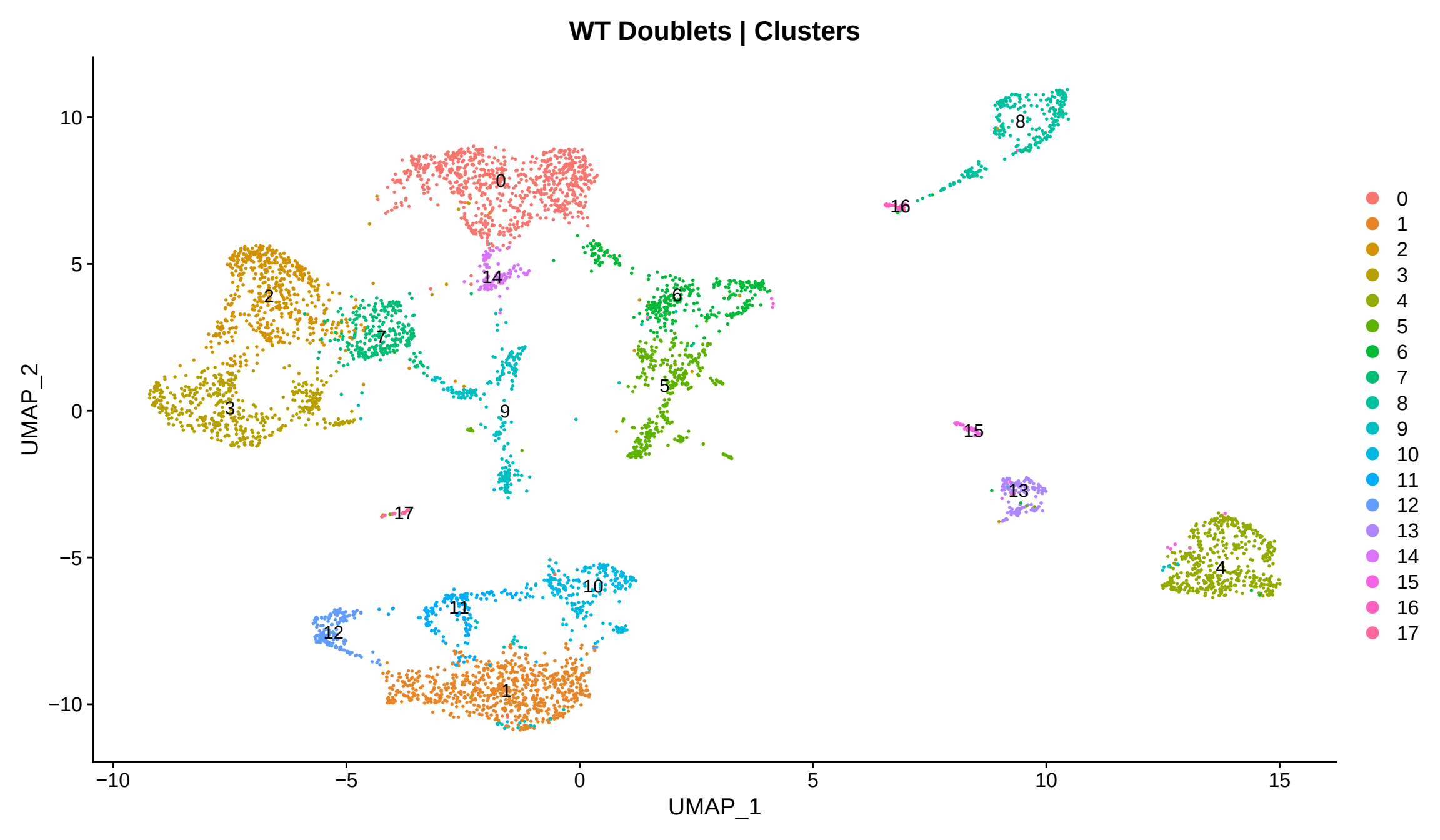
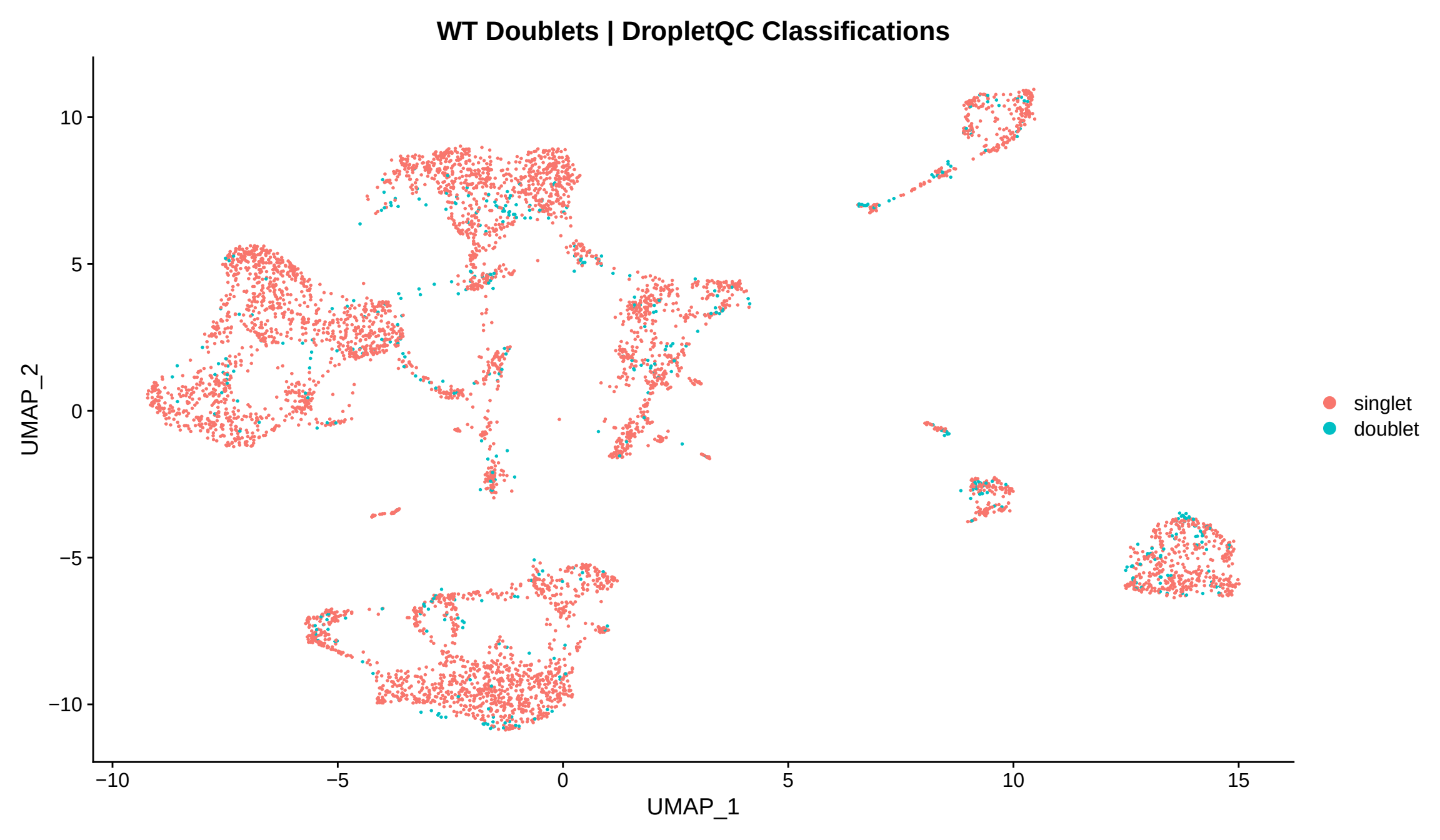
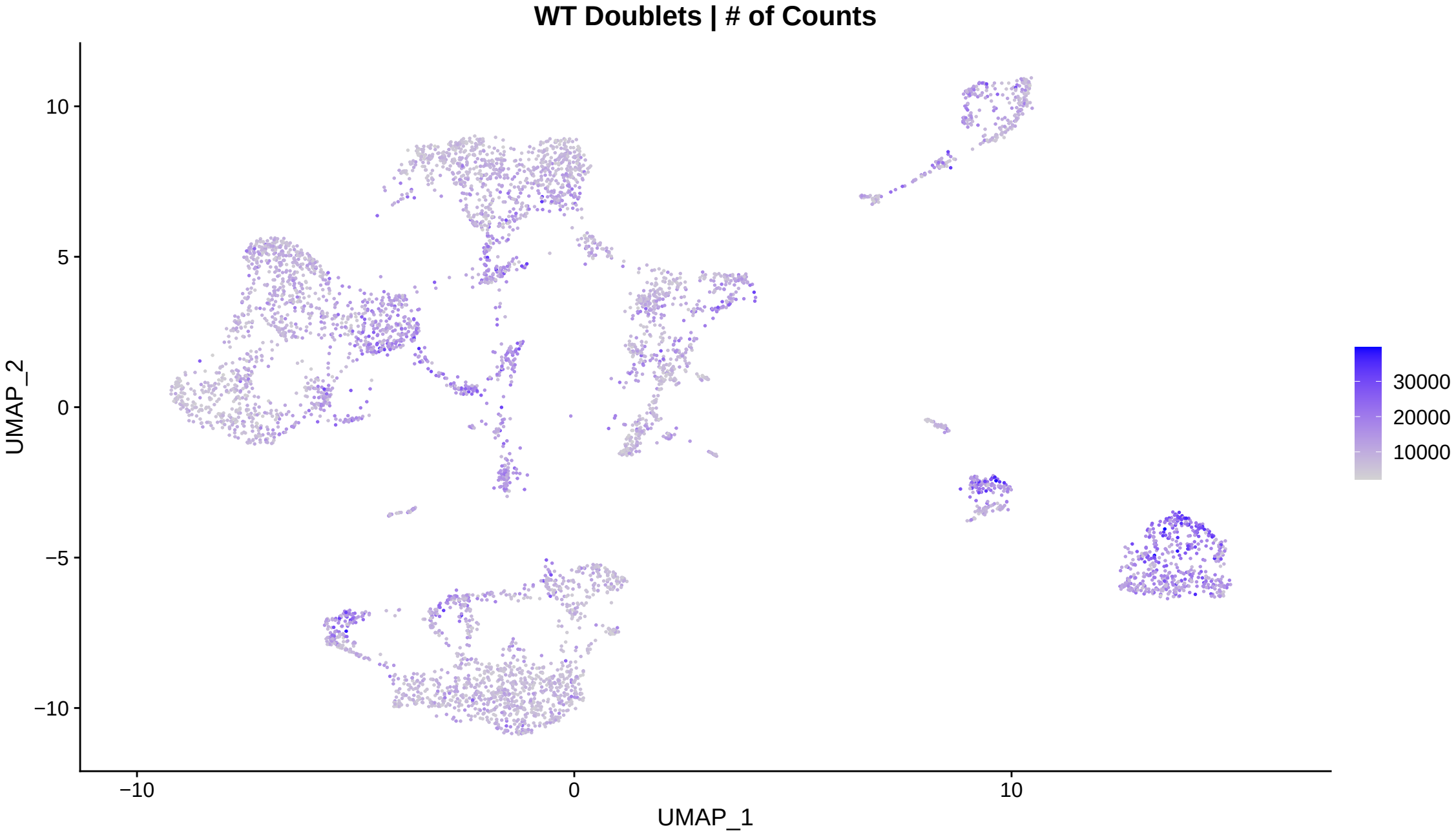
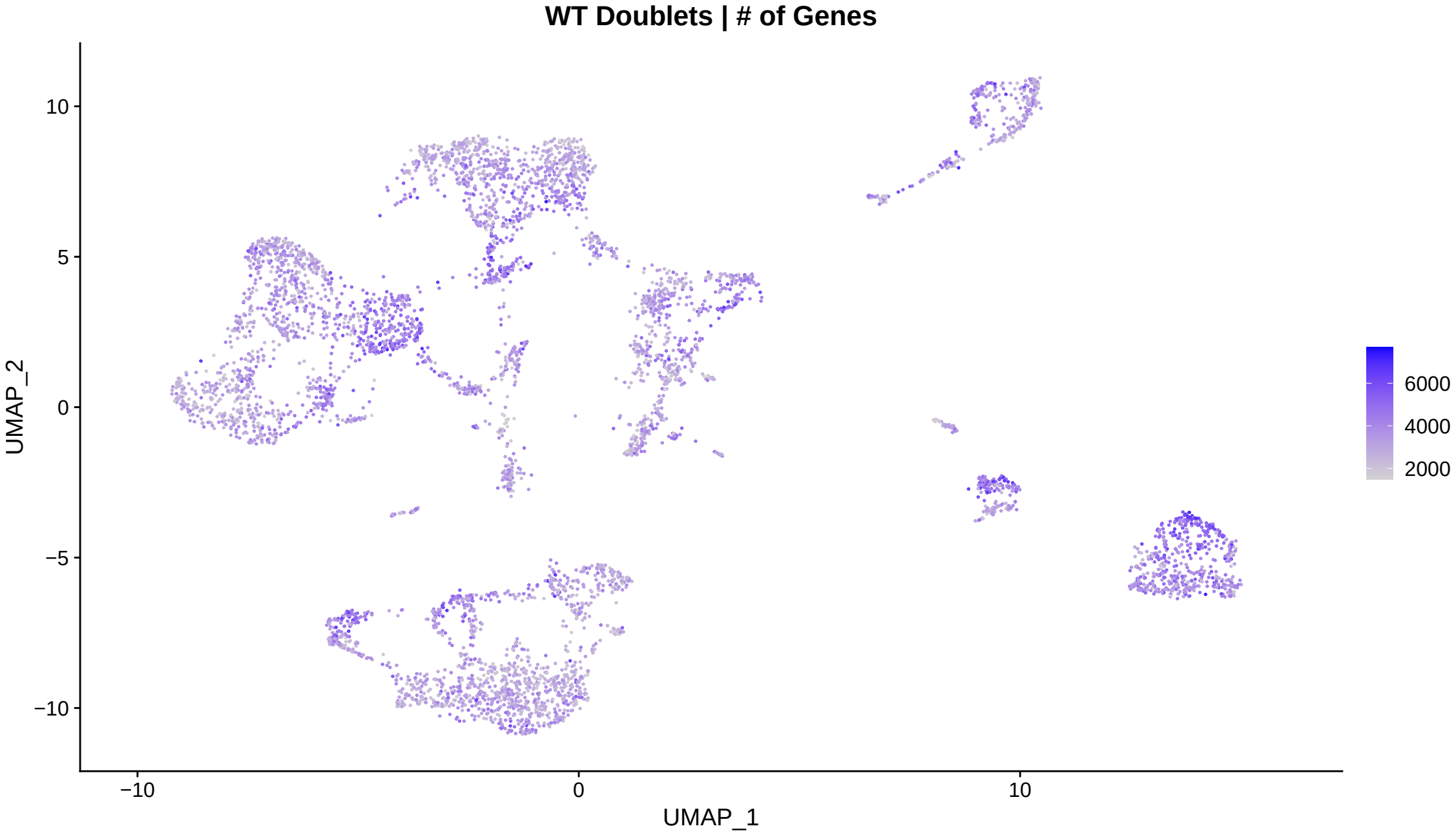
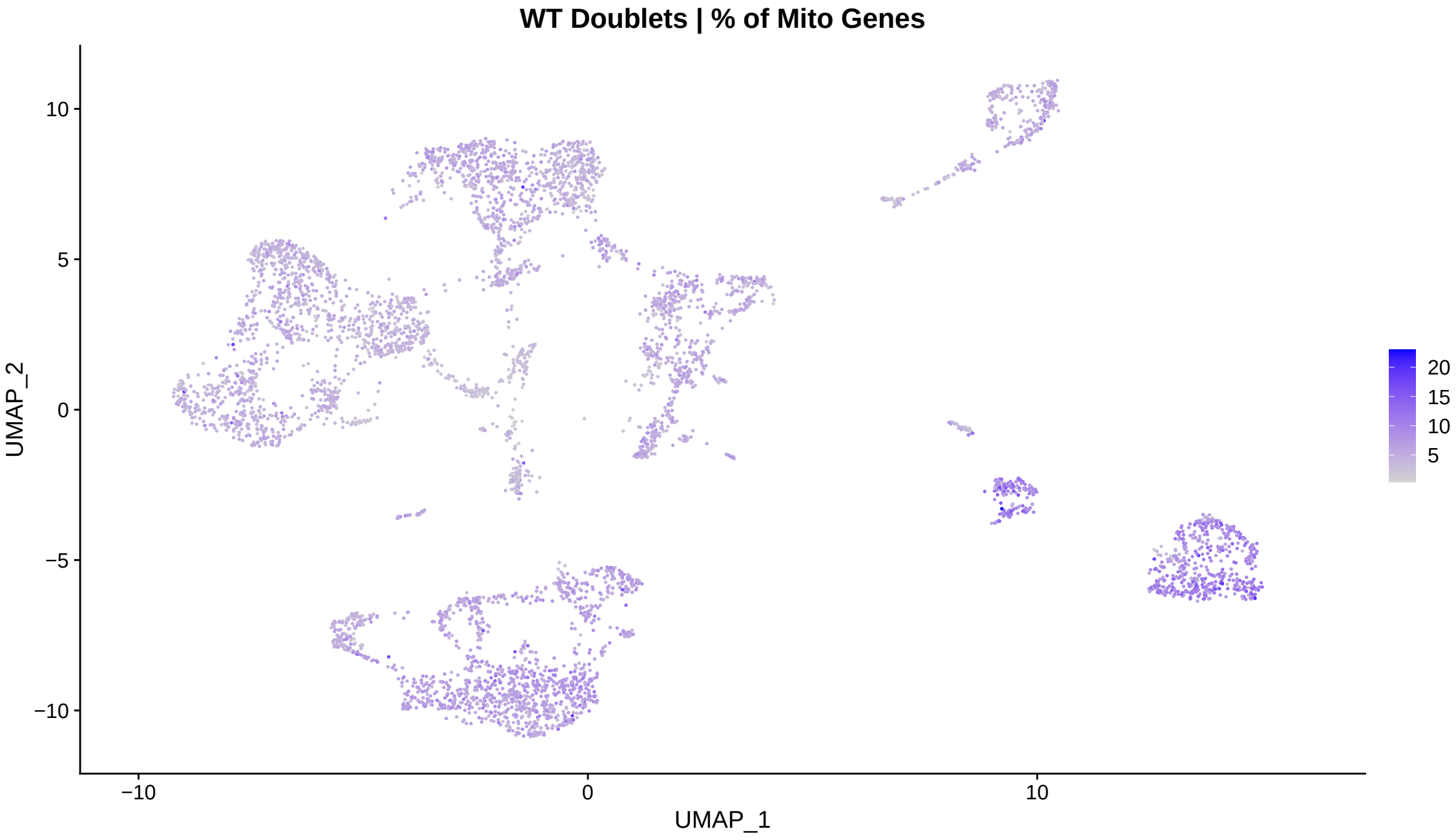
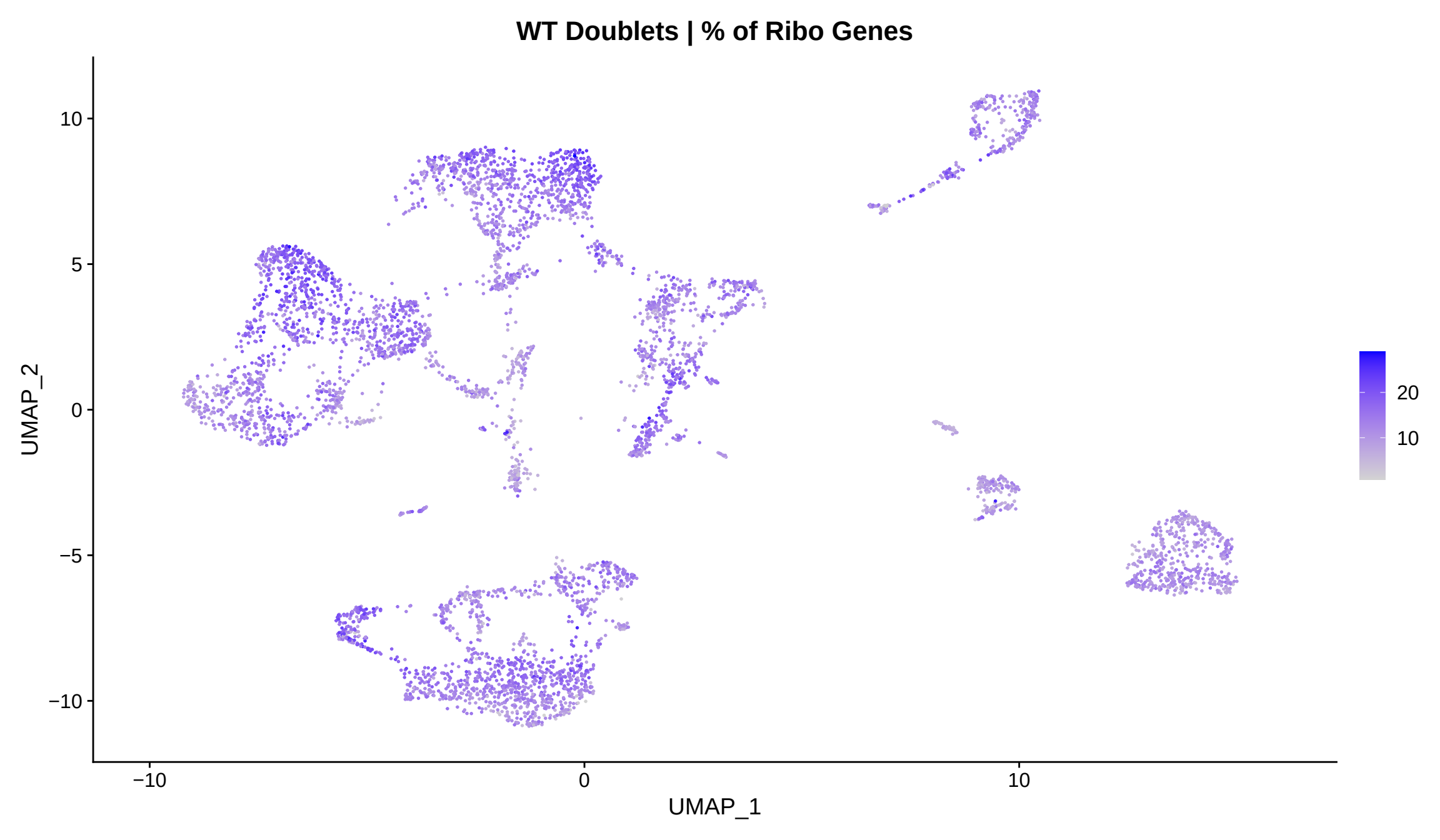

WT Doublets | Clusters
UMAP_2
UMAP_1
10
5
0
−5
−10
−10
−5
0
5
10
15
0
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17

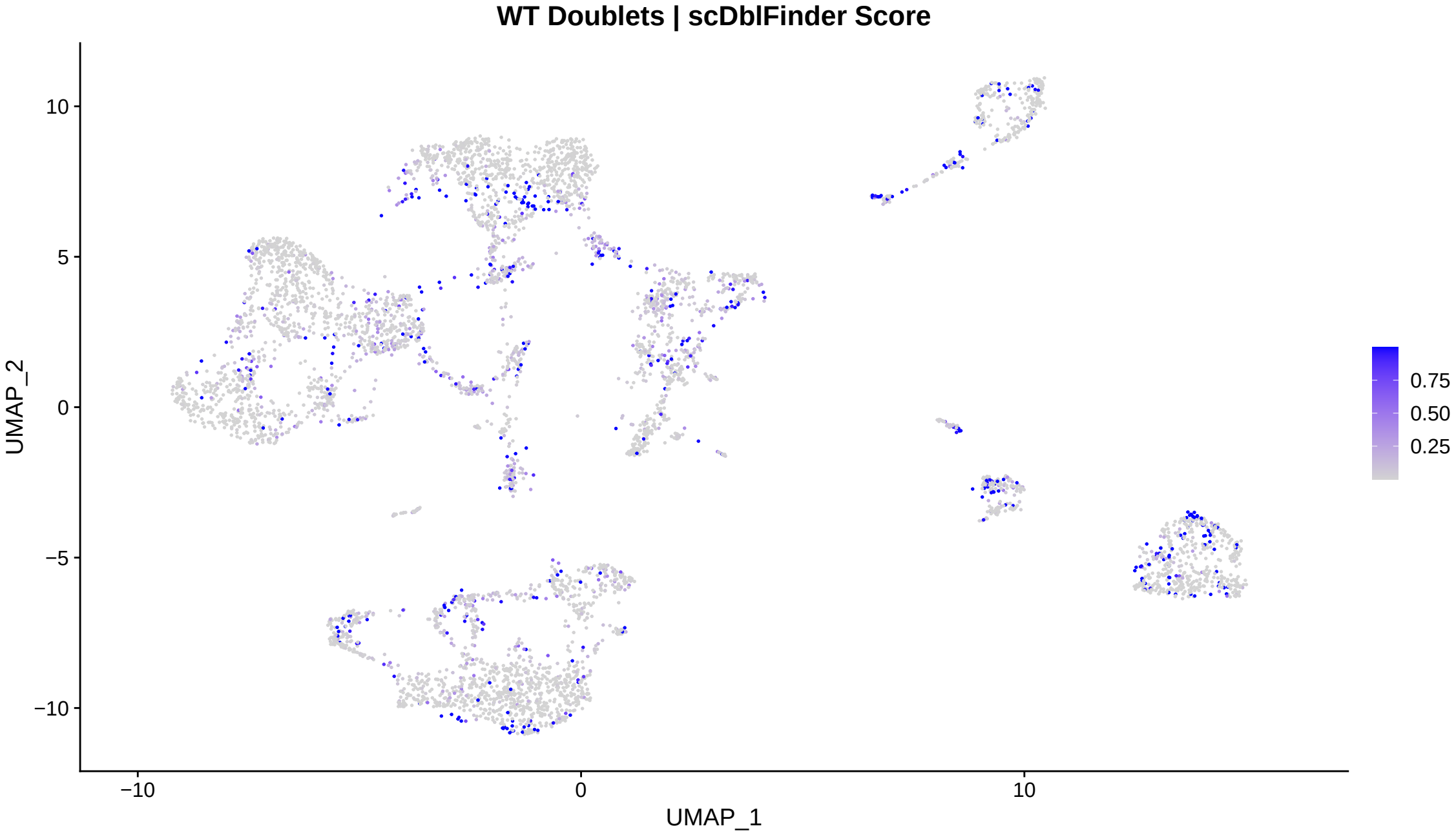

WT Doublets | DropletQC Classifications
UMAP_2
UMAP_1
10
5
0
−5
−10
−10
−5
0
5
10
15
singlet
doublet


WT Doublets | # of Counts
UMAP_2
UMAP_1
10
5
0
−5
−10
−10
0
10
30000
20000
10000


WT Doublets | # of Genes
10
5
0
−5
−10
UMAP_2
−10
0
10
UMAP_1
6000
4000
2000


WT Doublets | % of Mito Genes
10
5
0
−5
−10
UMAP_2
−10
0
10
UMAP_1
20
15
10
5


WT Doublets | % of Ribo Genes
10
5
0
−5
−10
UMAP_2
−10
0
10
UMAP_1
20
10


WT Doublets | scDblFinder Score
UMAP_2
UMAP_1
10
5
0
−5
−10
−10
0
10
0.75
0.50
0.25