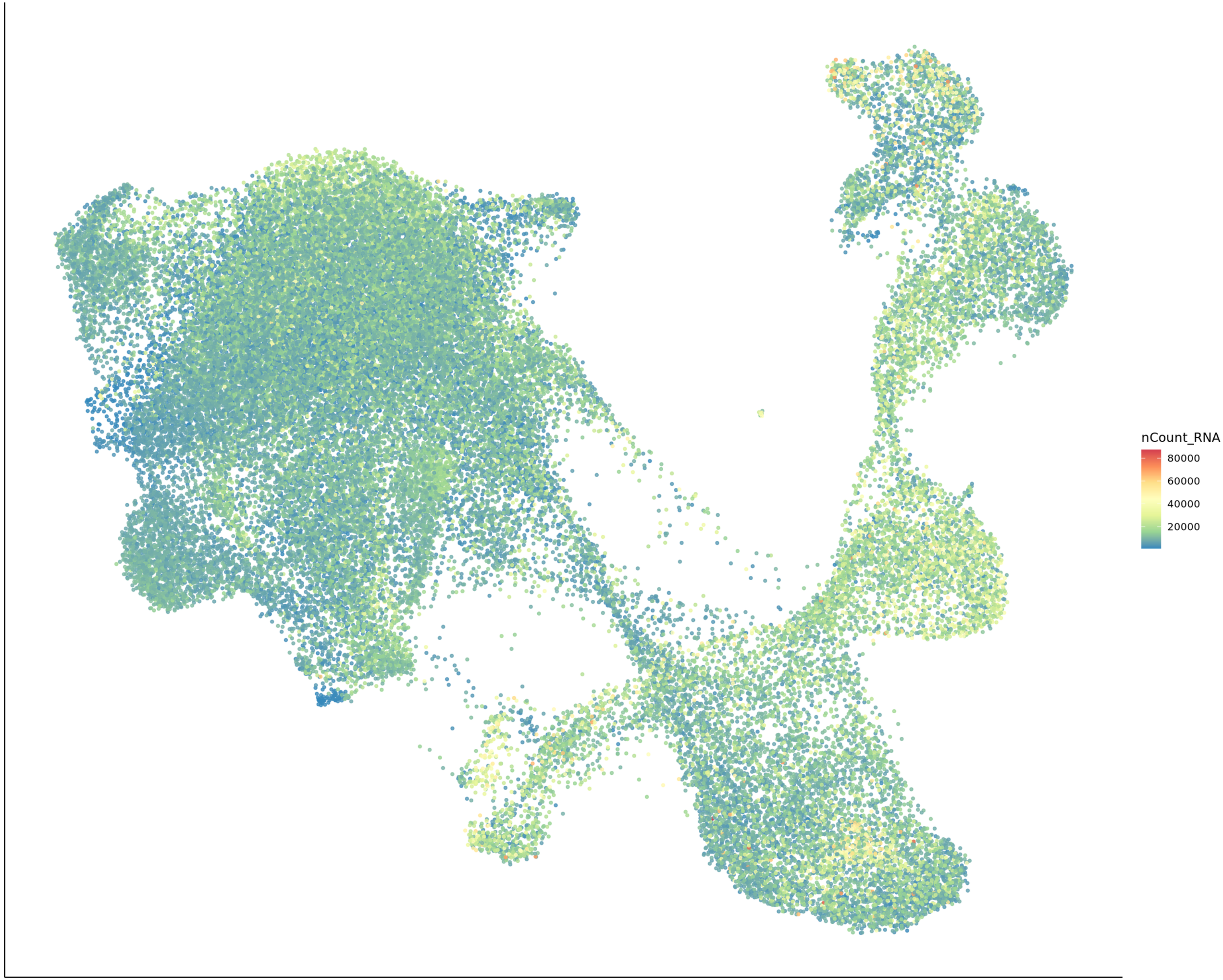
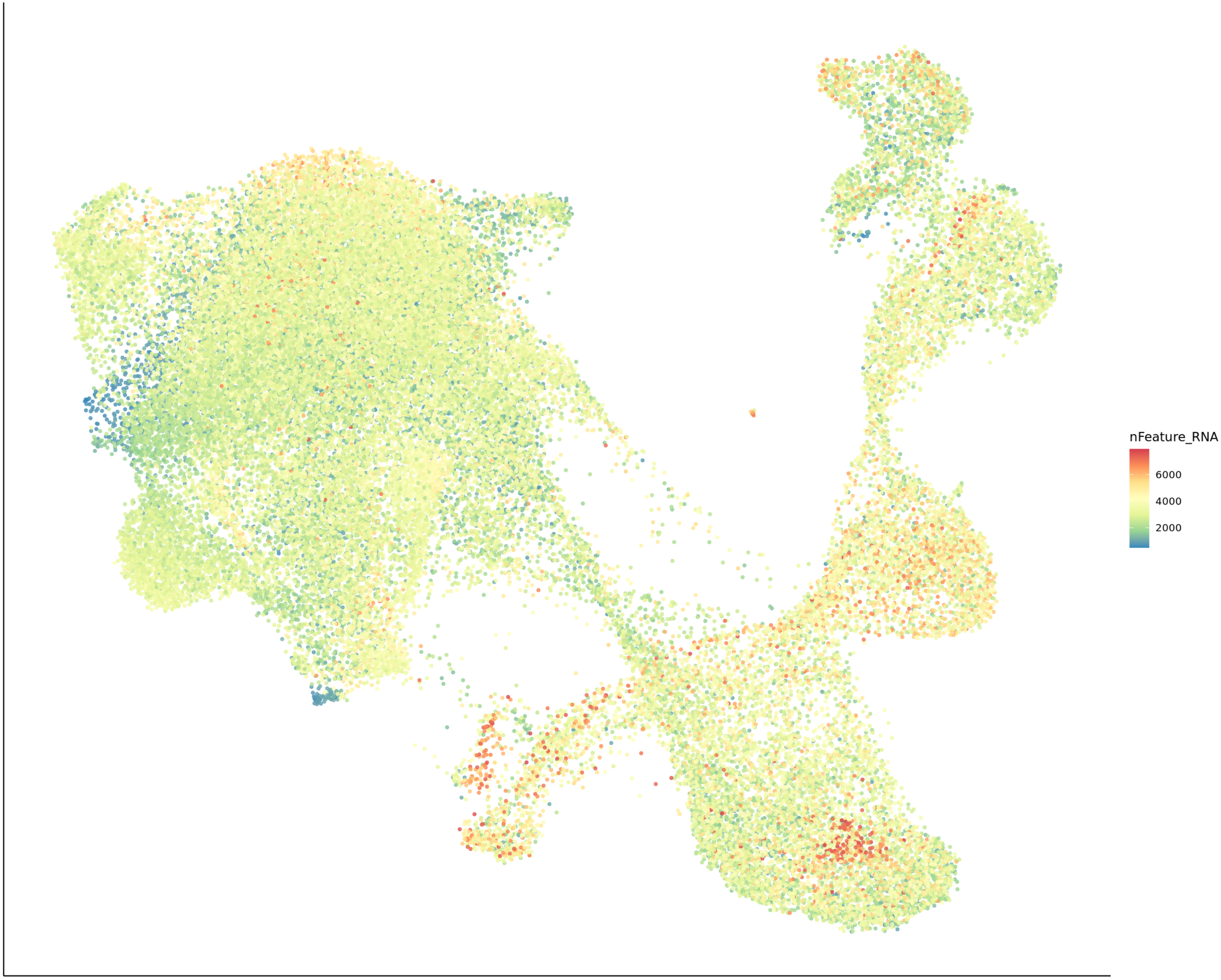
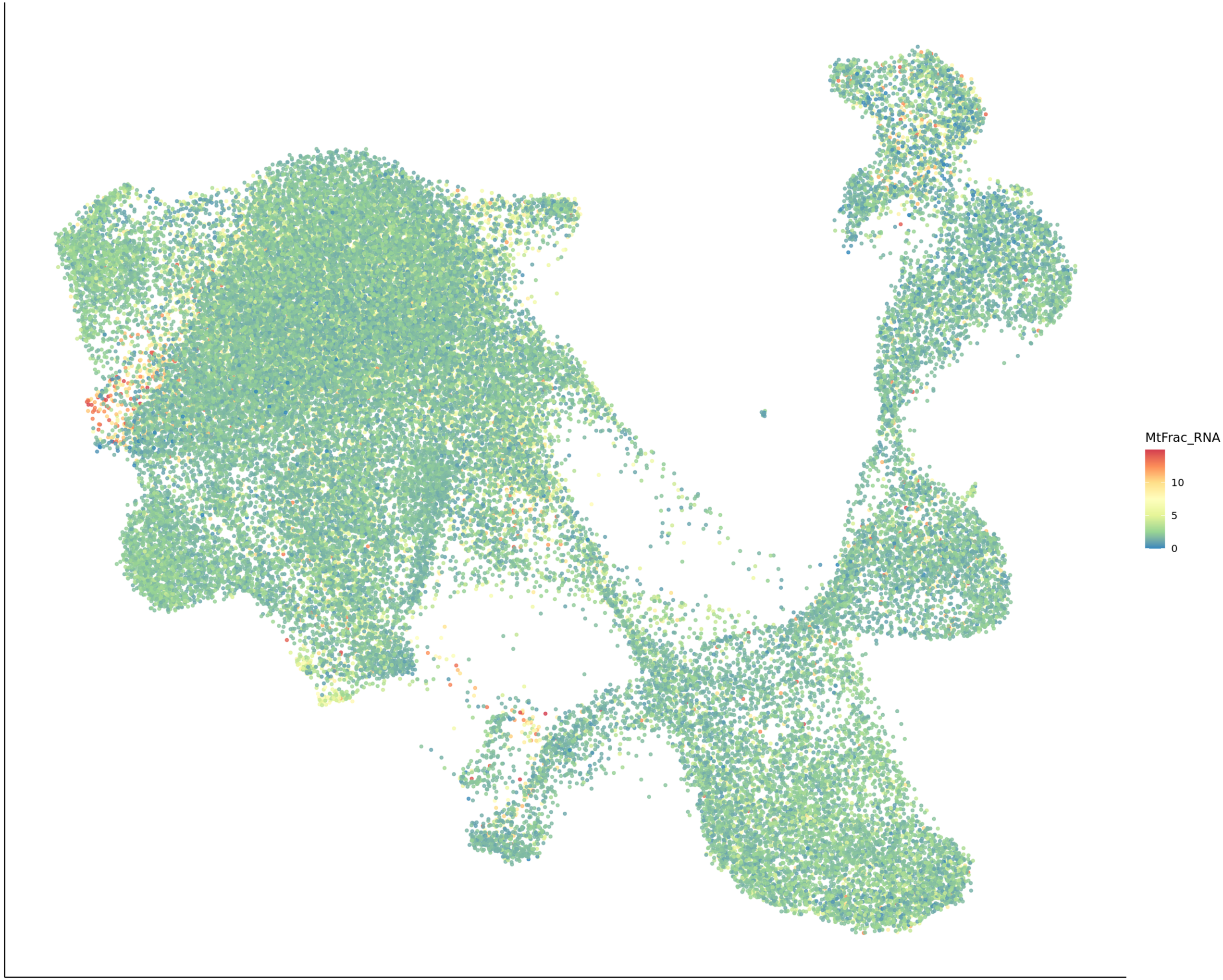
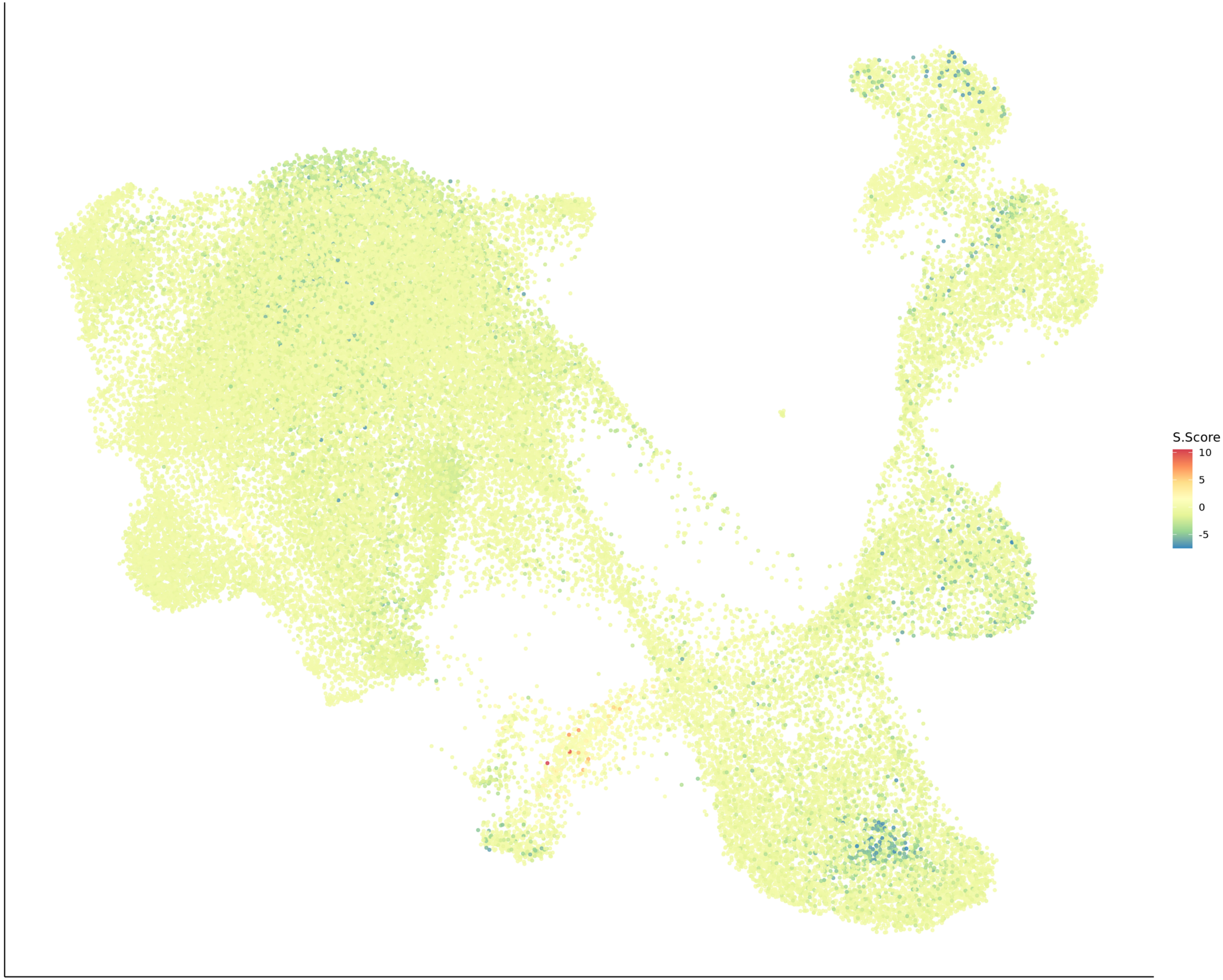
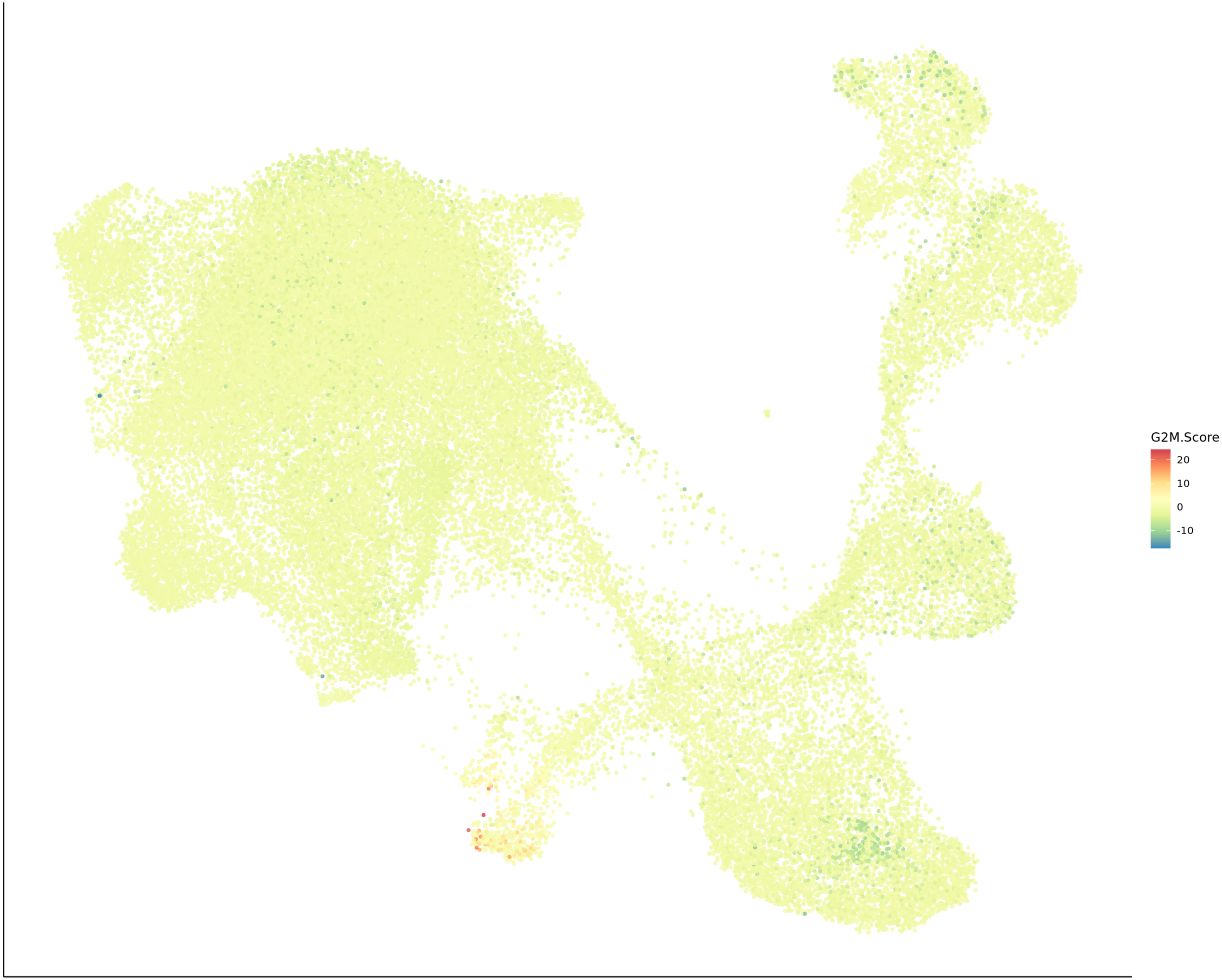
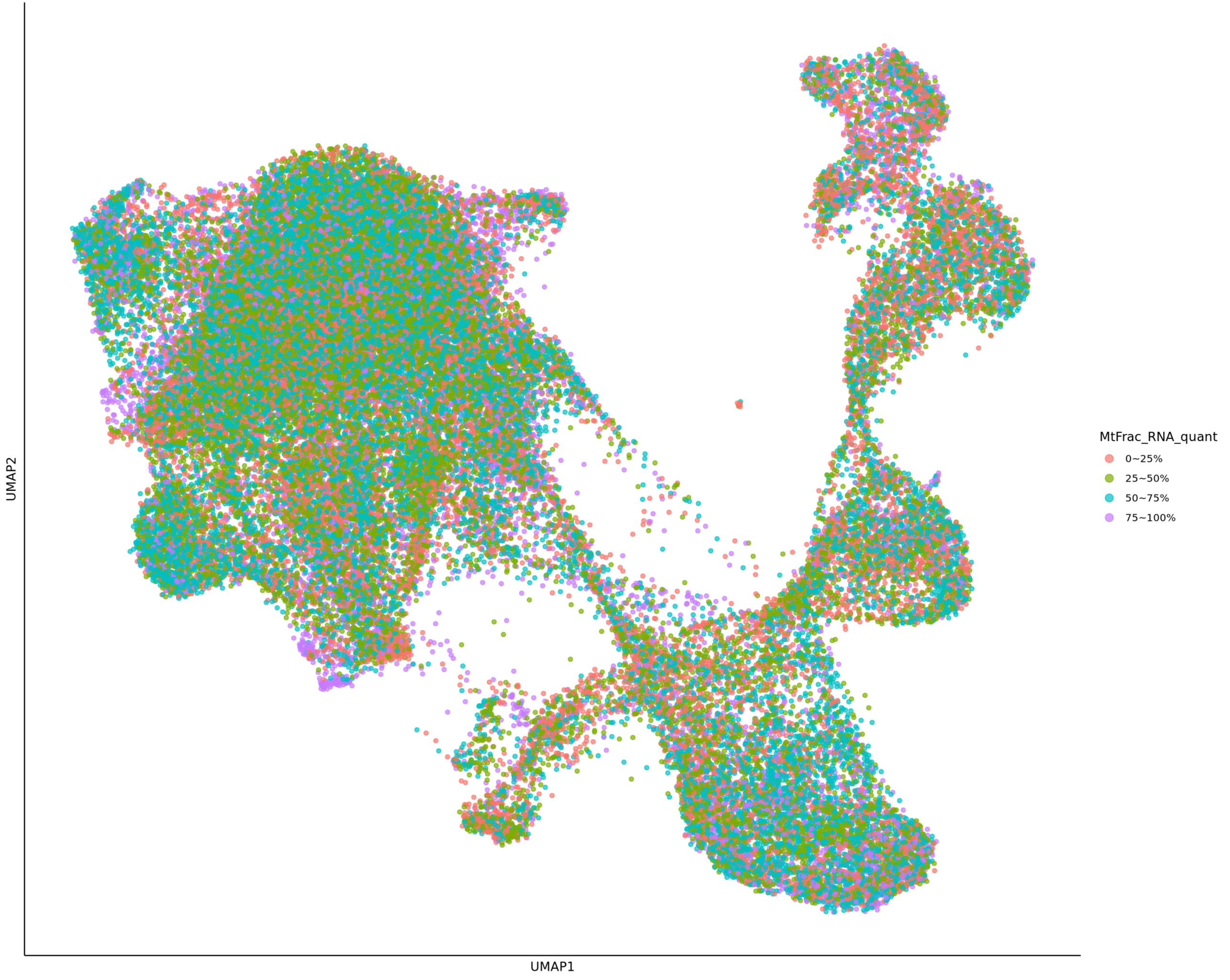

nCount_RNA
80000
60000
40000
20000

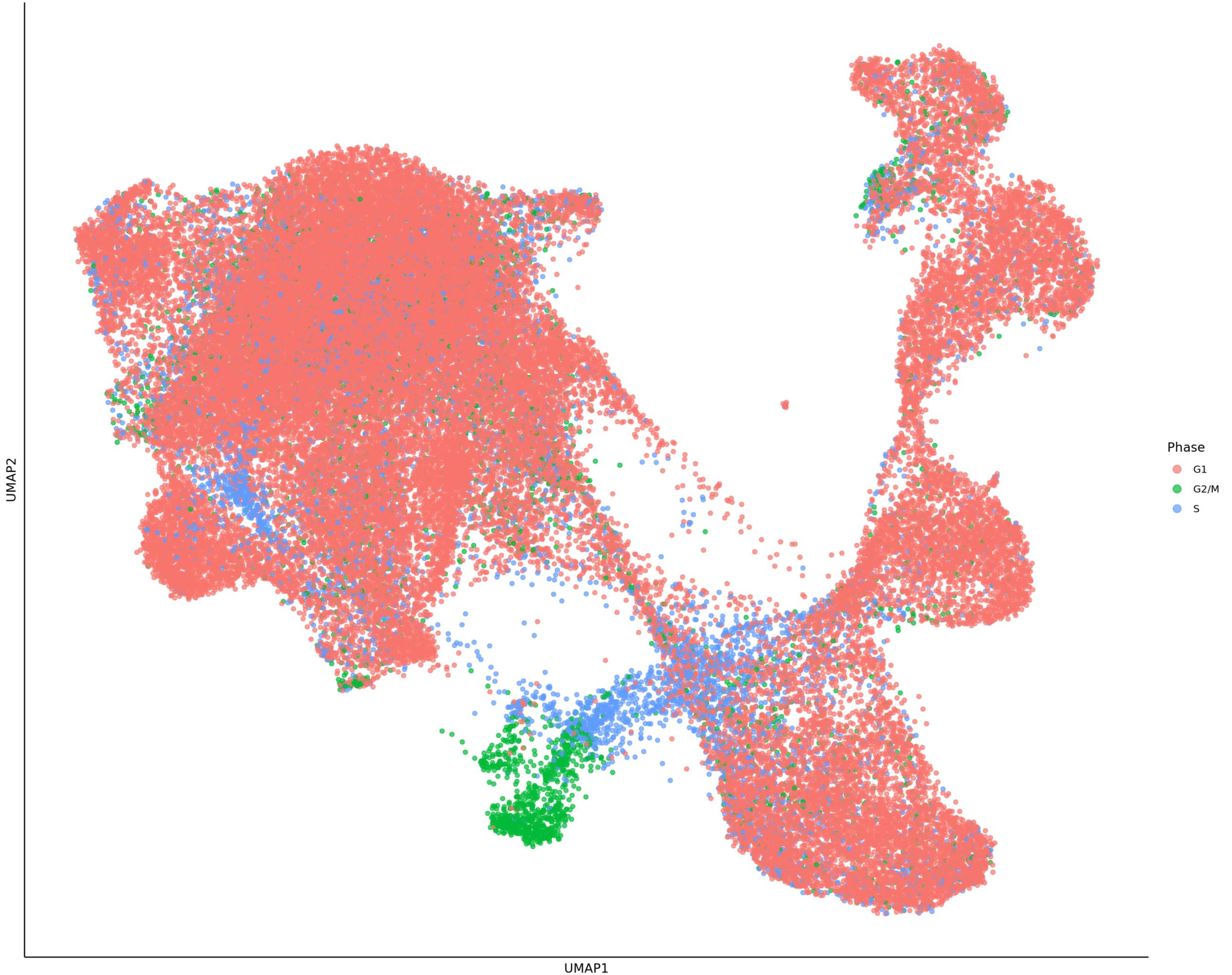

nFeature_RNA
6000
4000
2000


MtFrac_RNA
10
5
0


S.Score
10
5
0
-5


G2M.Score
20
10
0
-10


UMAP2
UMAP1
MtFrac_RNA_quant
0~25%
25~50%
50~75%
75~100%


UMAP2
UMAP1
Phase
G1
G2/M
S