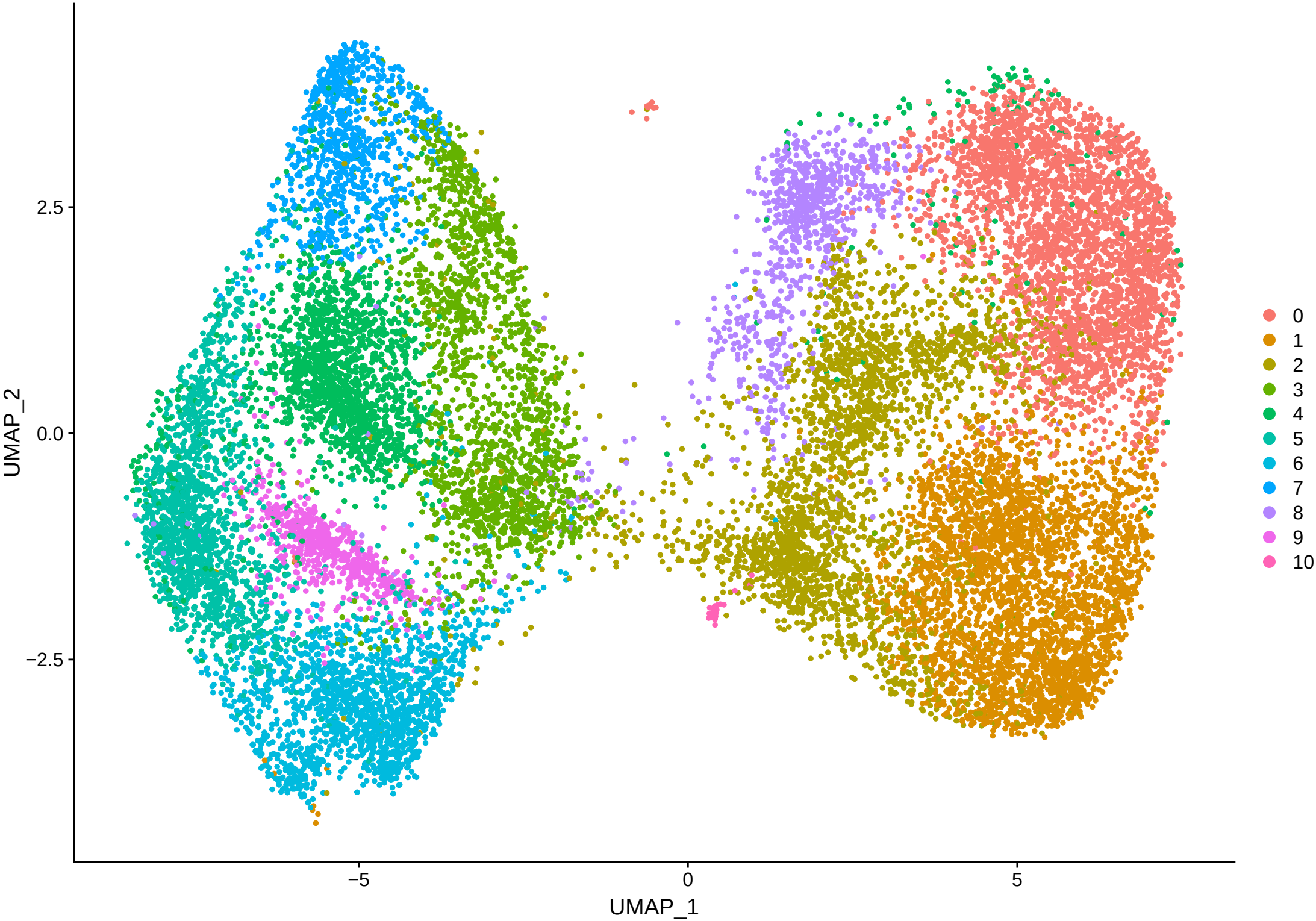
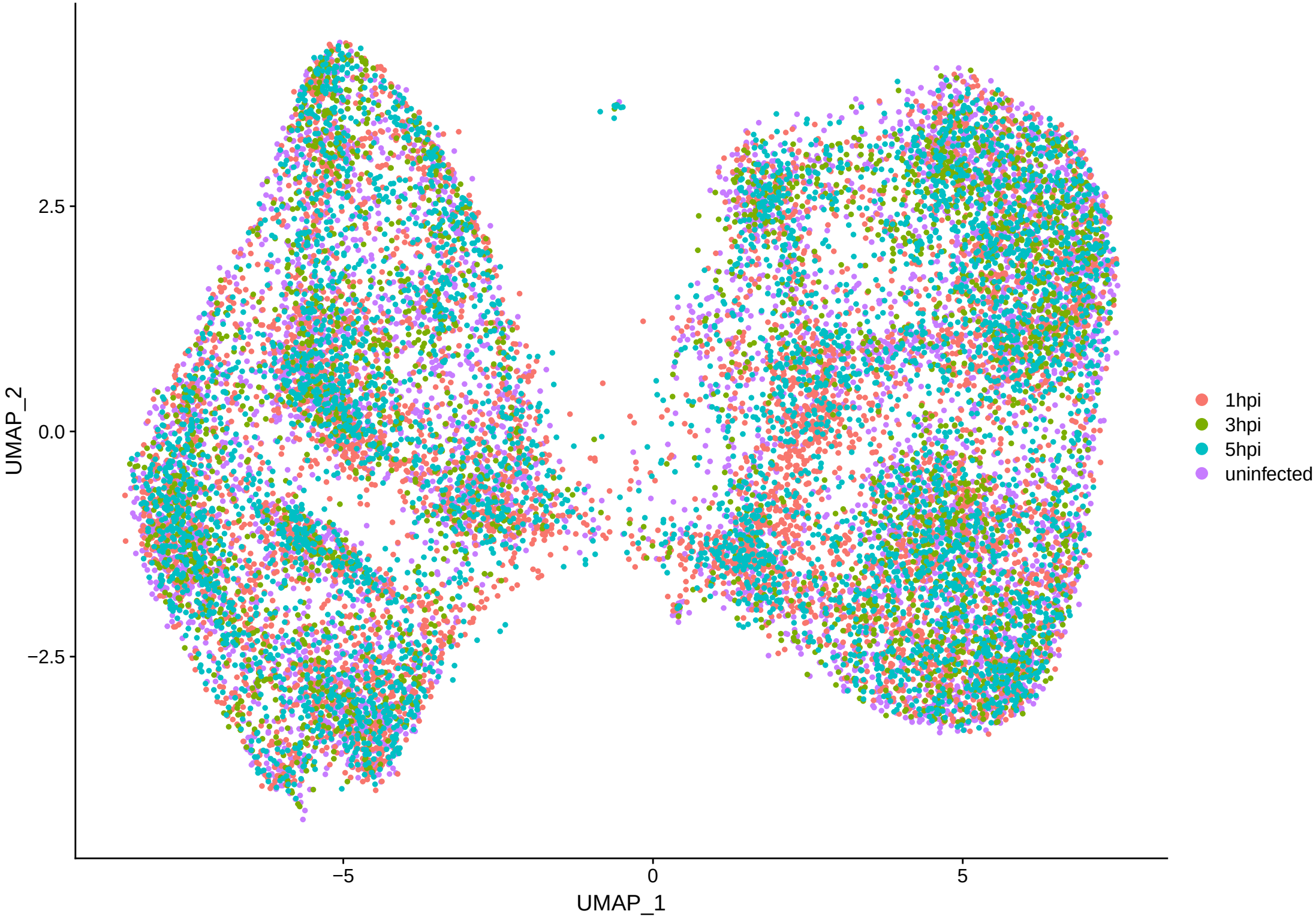
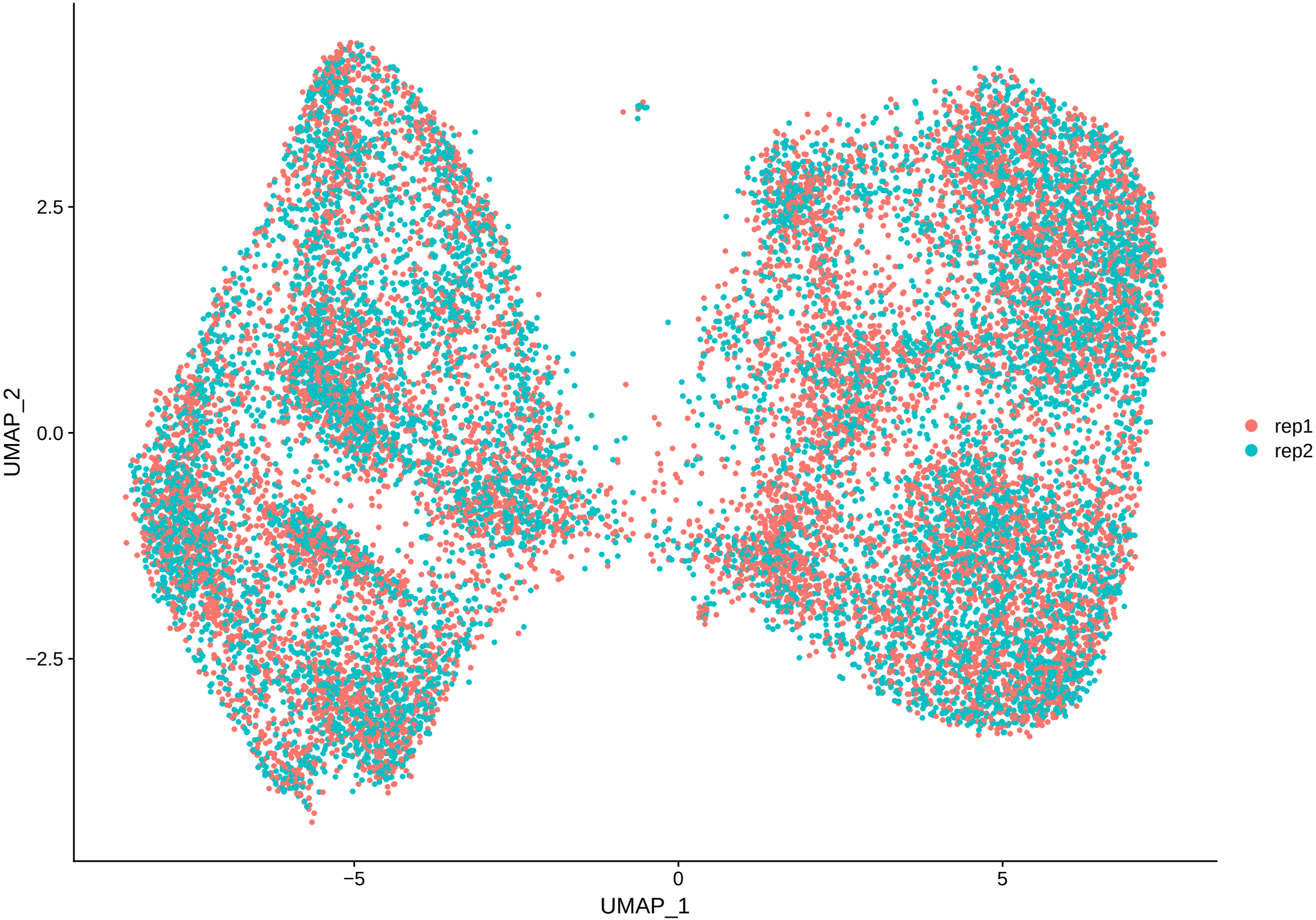
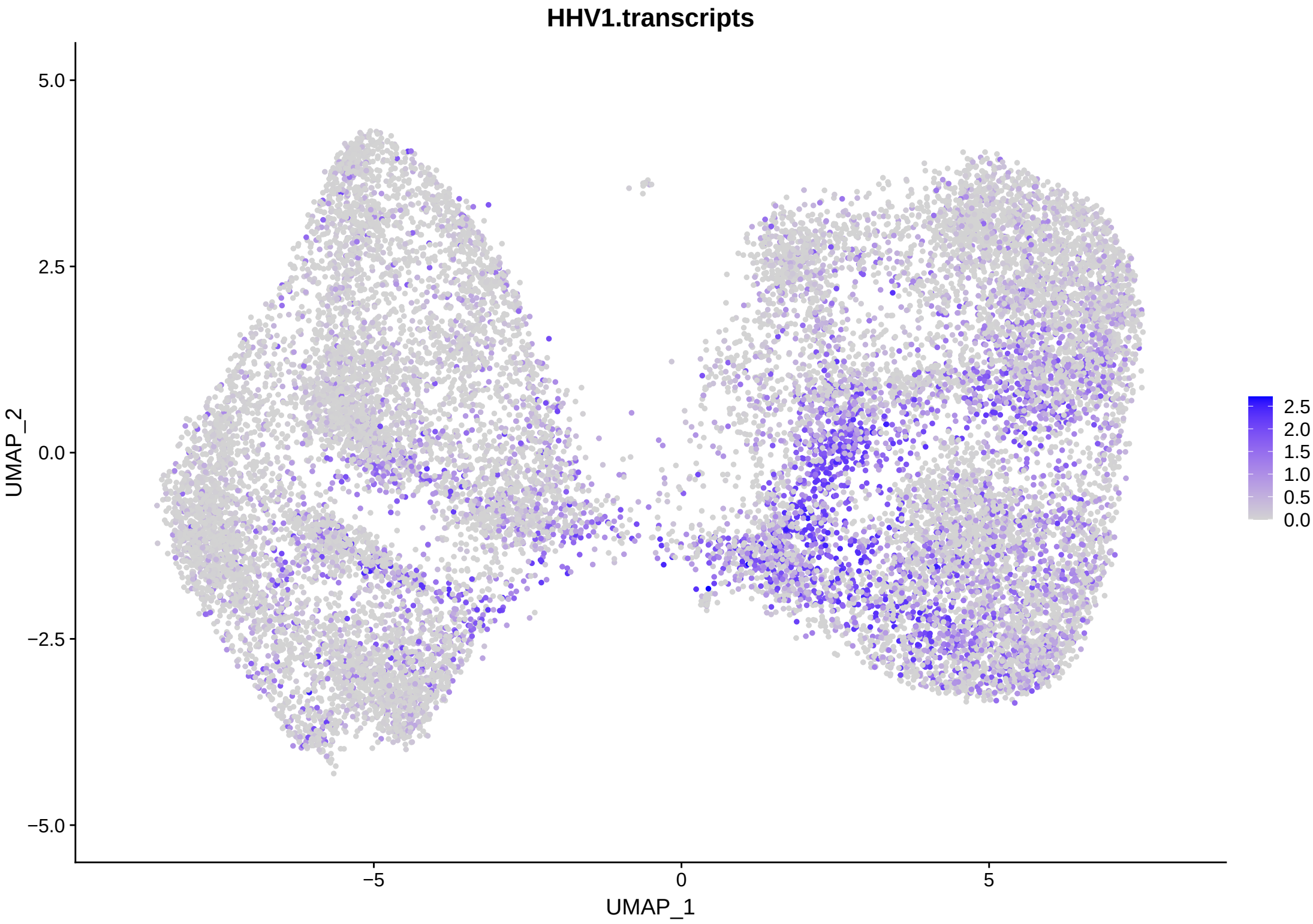

UMAP_2
2.5
0.0
−2.5
−5
0
5
UMAP_1
0
1
2
3
4
5
6
7
8
9
10


2.5
0.0
−2.5
UMAP_2
−5
0
5
UMAP_1
1hpi
3hpi
5hpi
uninfected


2.5
0.0
−2.5
UMAP_2
−5
0
5
UMAP_1
rep1
rep2


HHV1.transcripts
5.0
2.5
0.0
−2.5
−5.0
UMAP_2
−5
0
5
UMAP_1
2.5
2.0
1.5
1.0
0.5
0.0