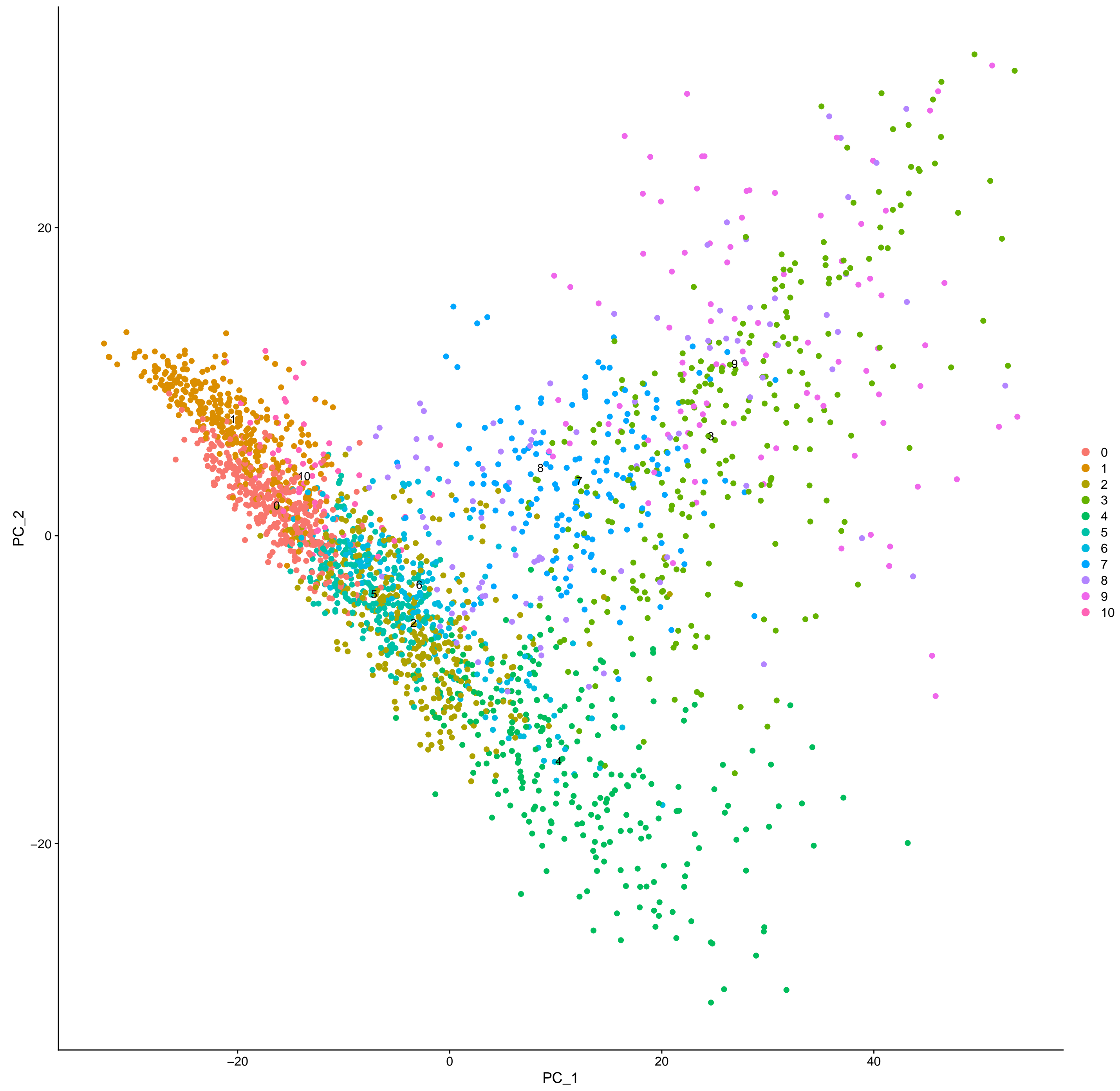
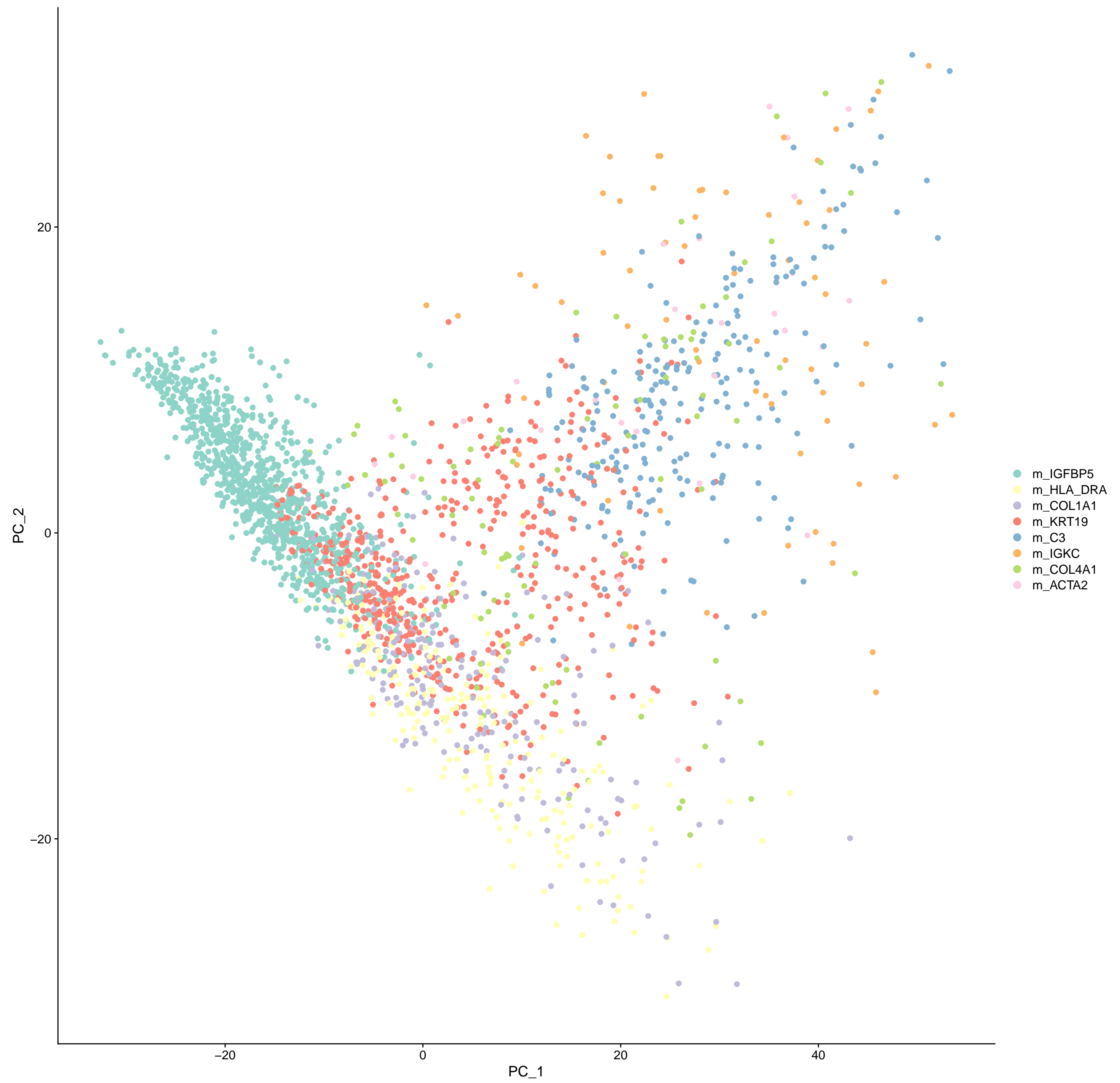
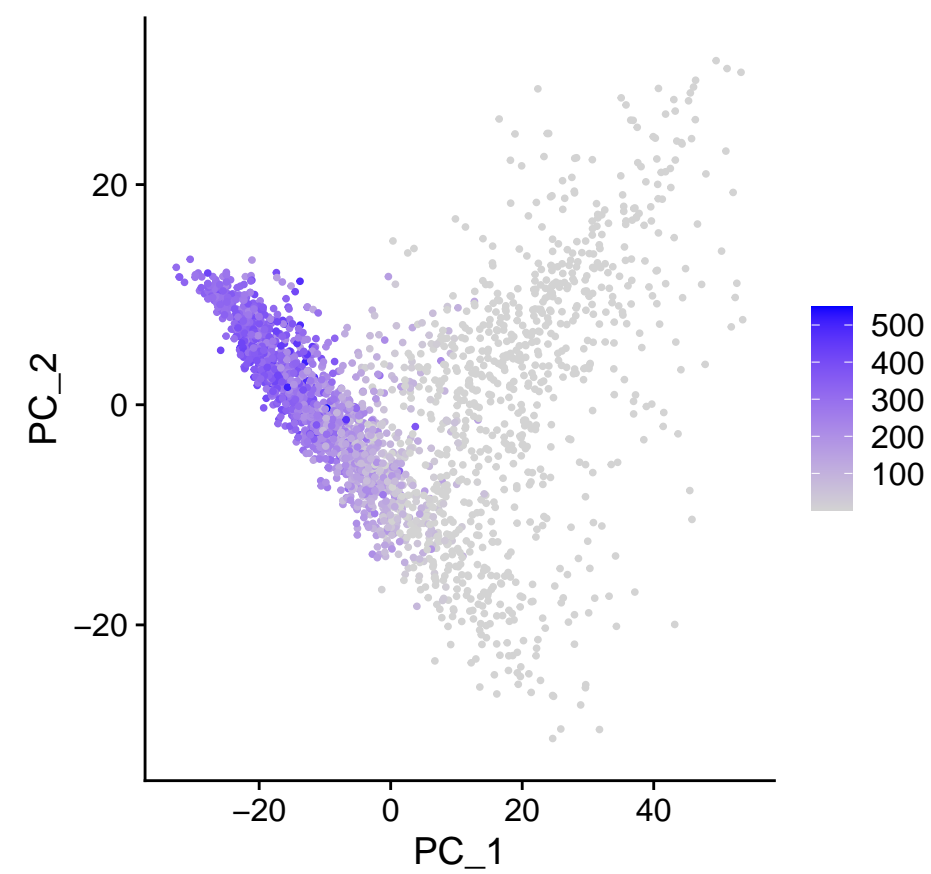
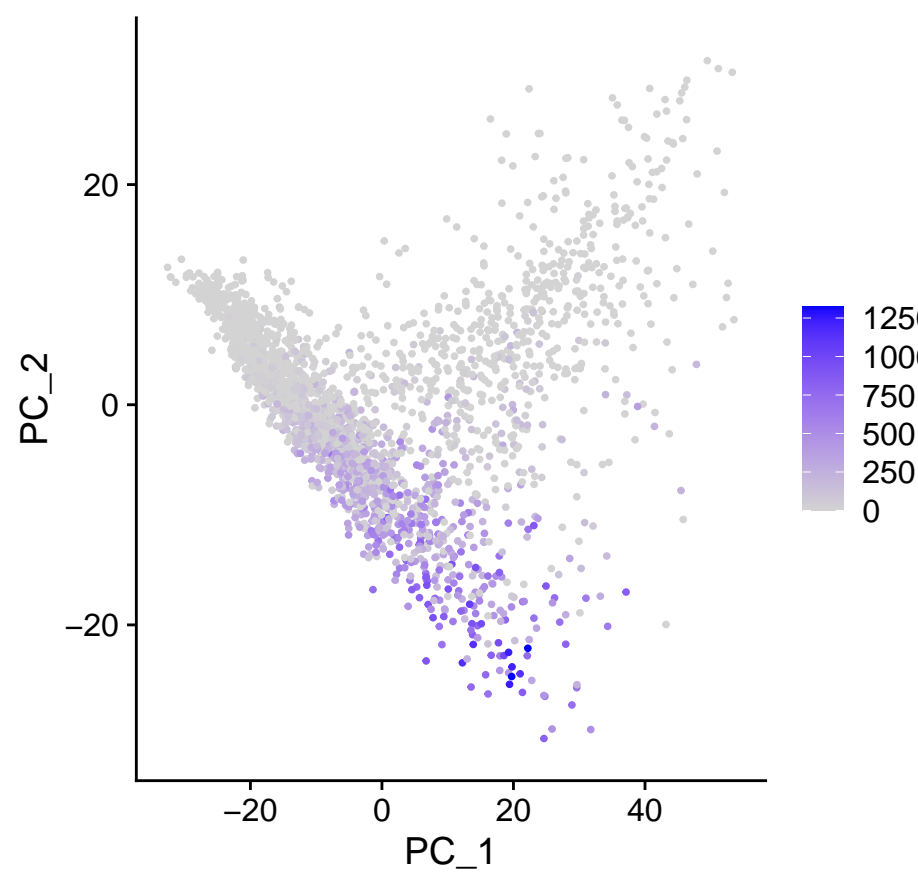
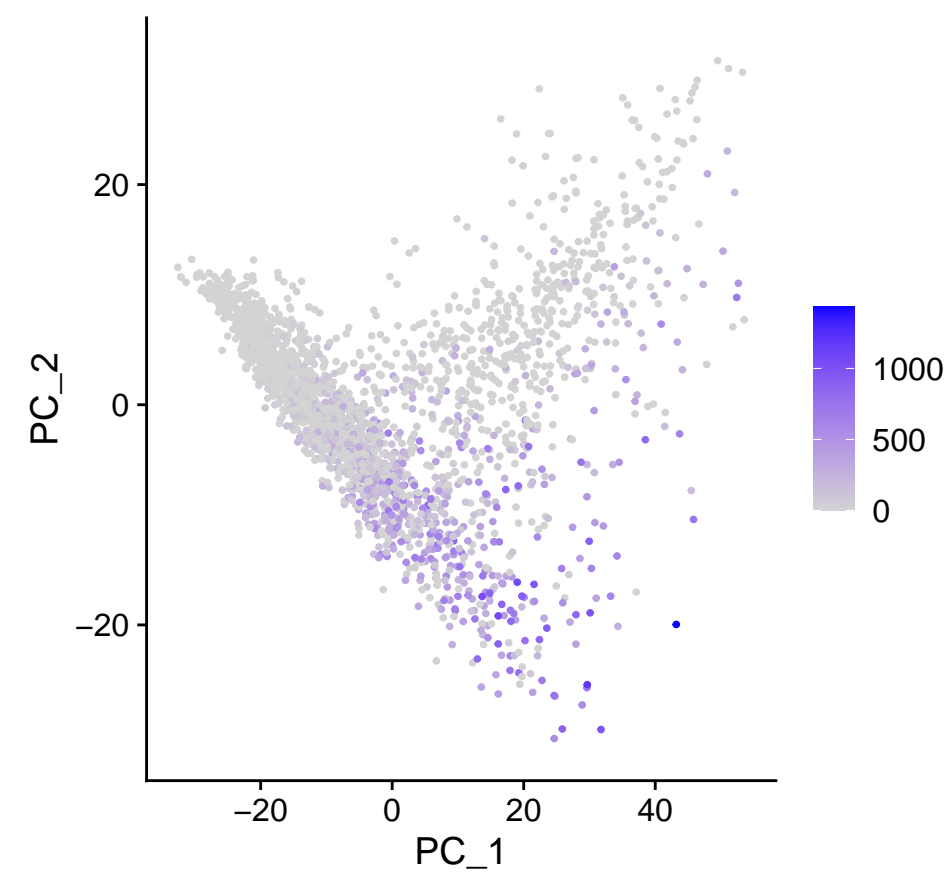
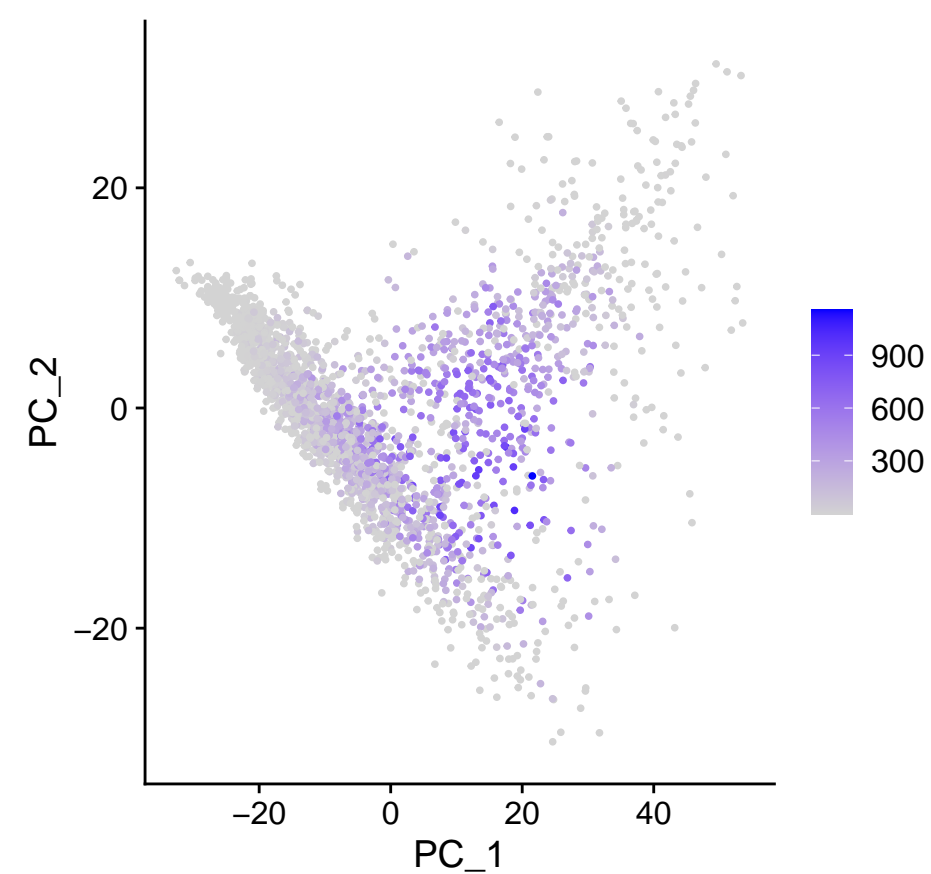
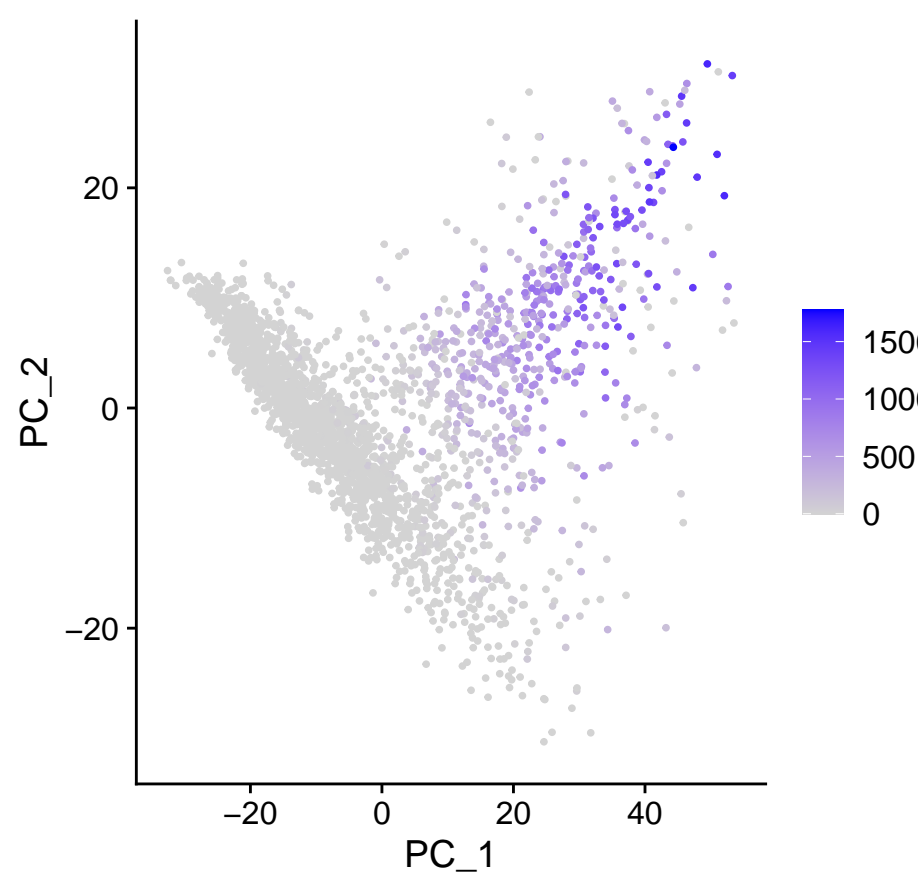
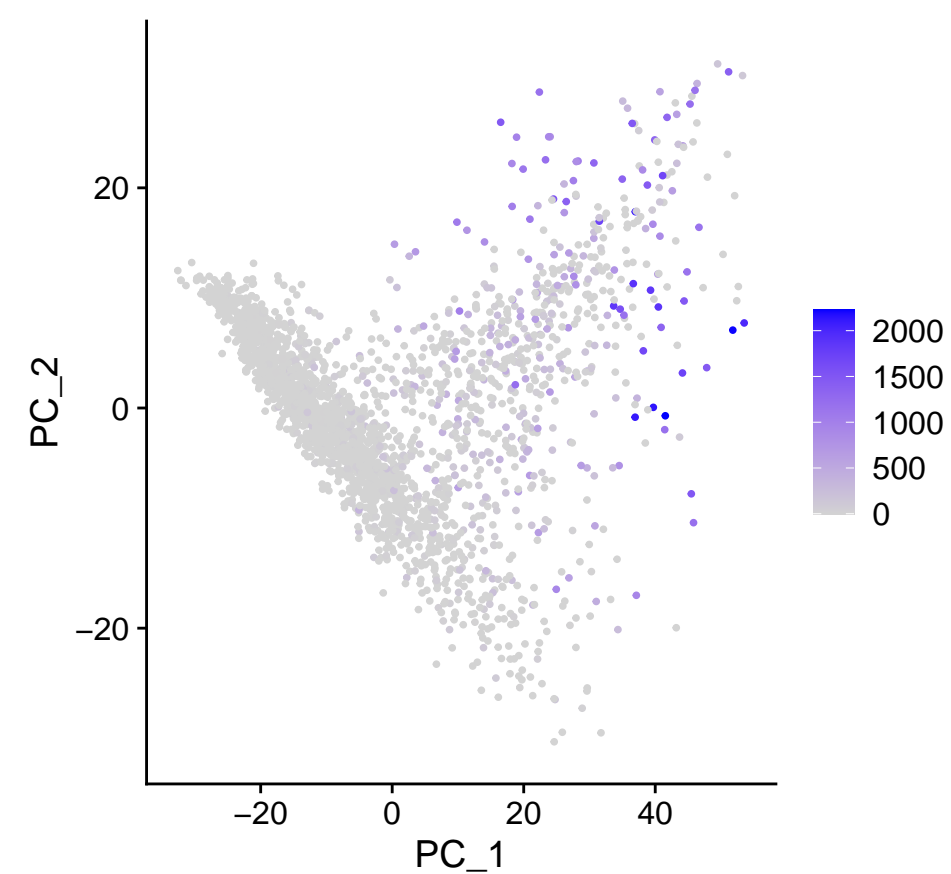
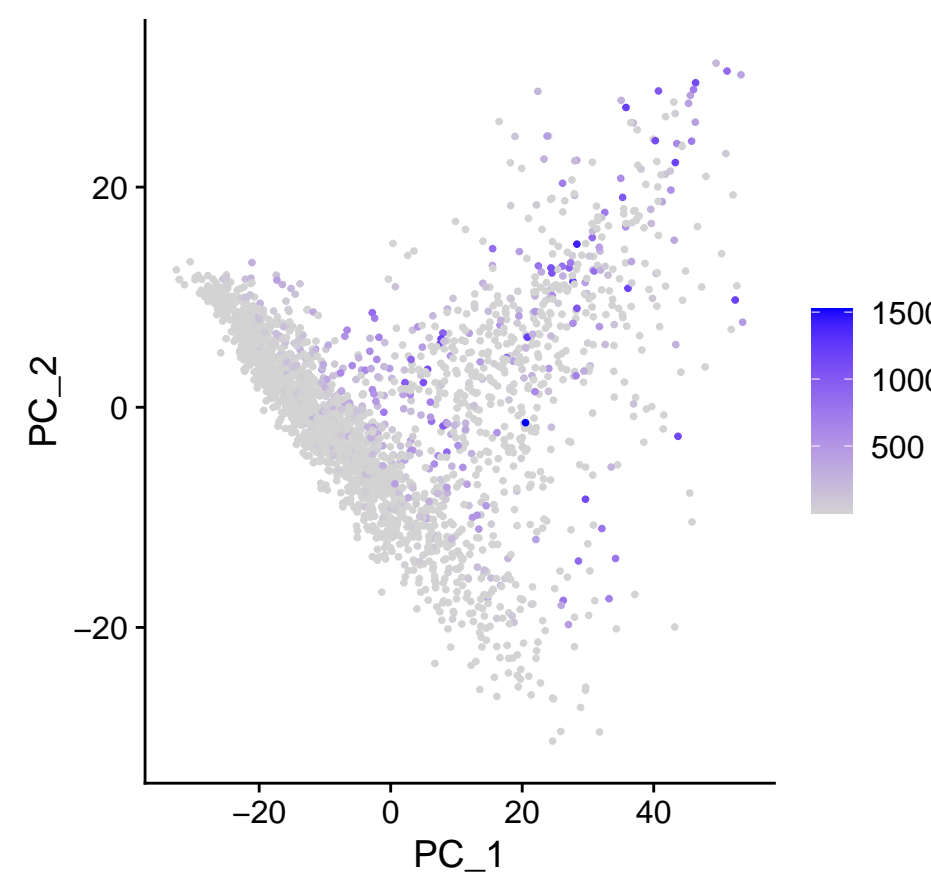
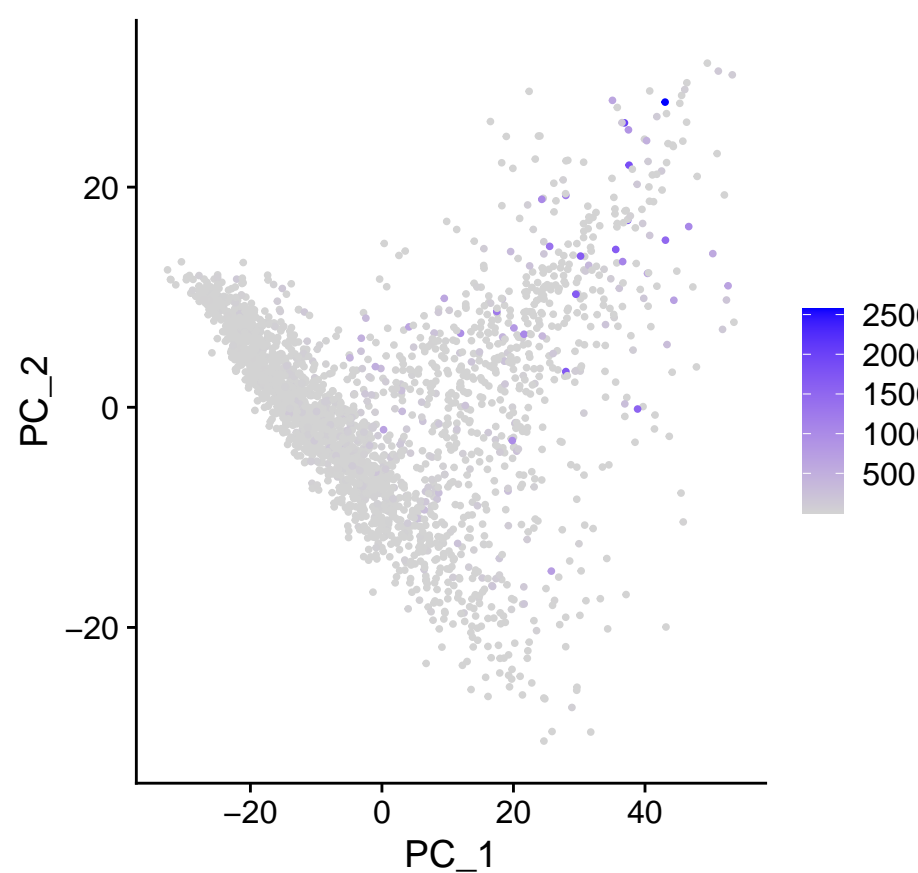


cluster



nmf



m_IGFBP5**m_HLA_DRA****m_COL1A1****m_KRT19****m_C3****m_IGKC****m_COL4A1****m_ACTA2**

cluster

5.0

2.5

0.0

-2.5

-5.0

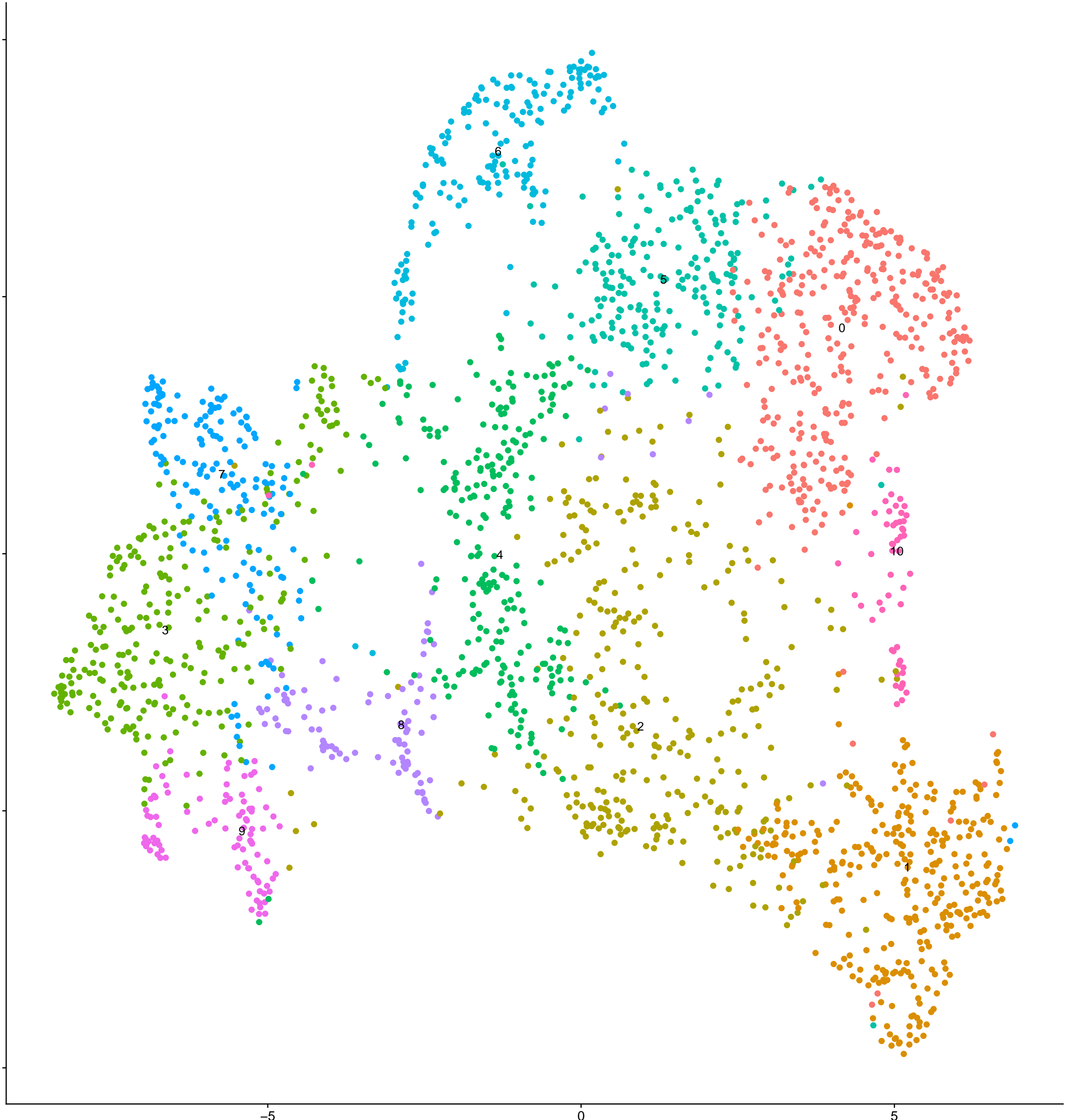
- 0
- 1
- 2
- 3
- 4
- 5
- 6
- 7
- 8
- 9
- 10

UMAP_1

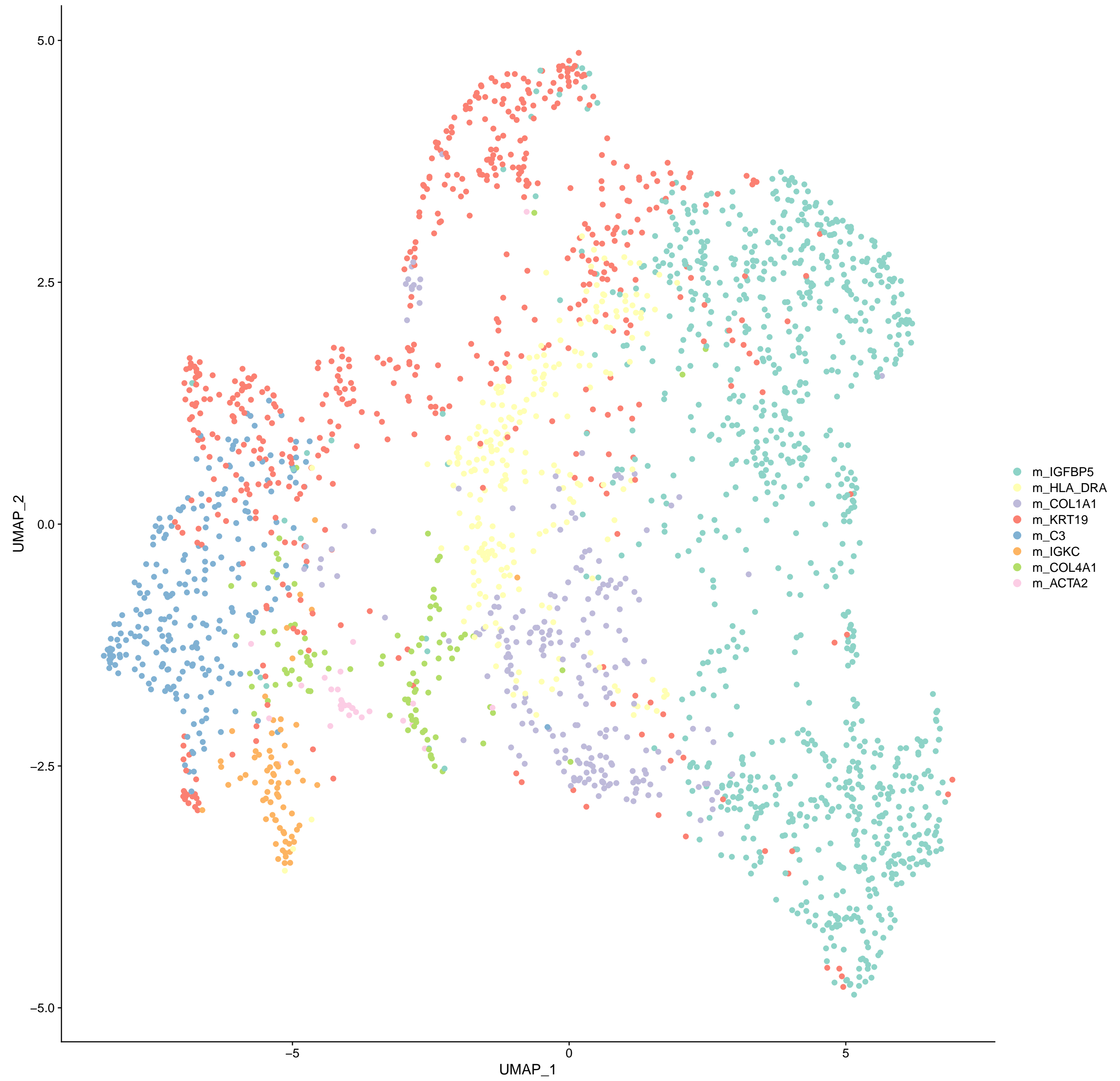
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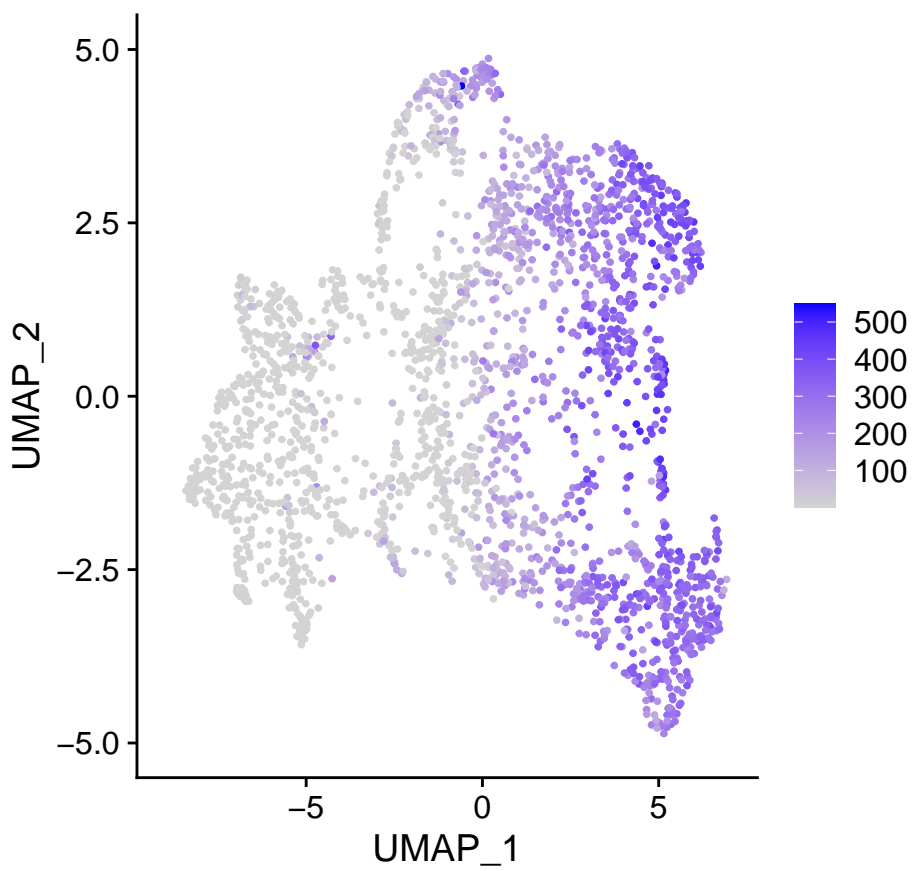
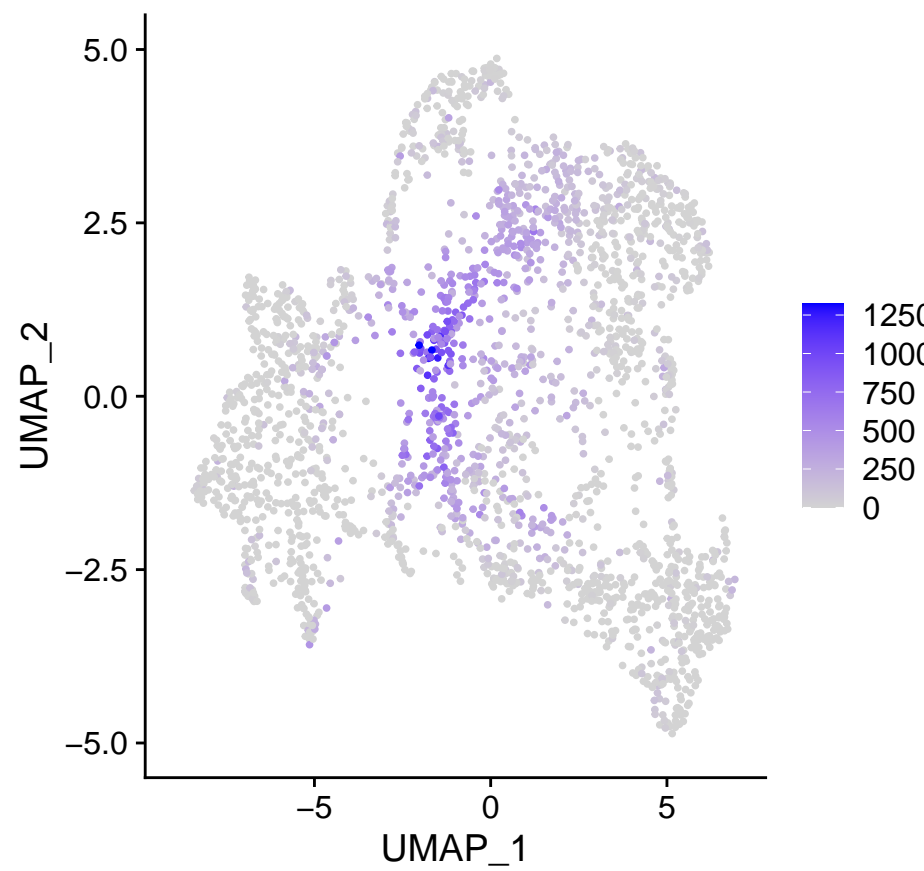
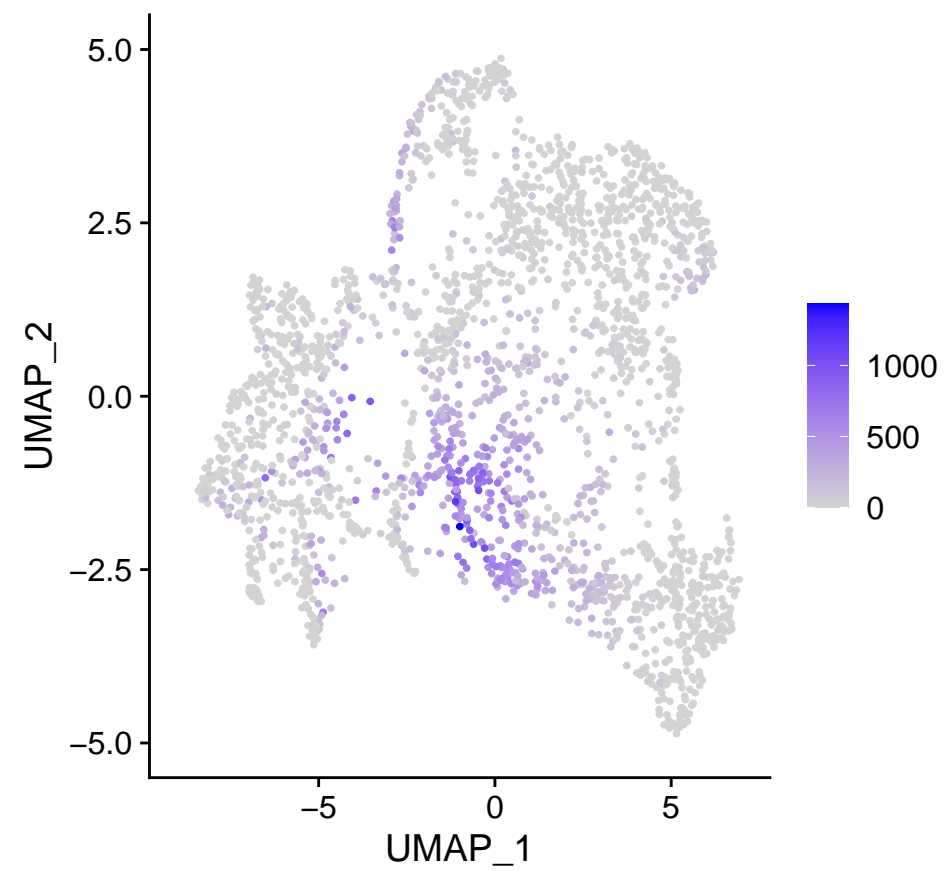
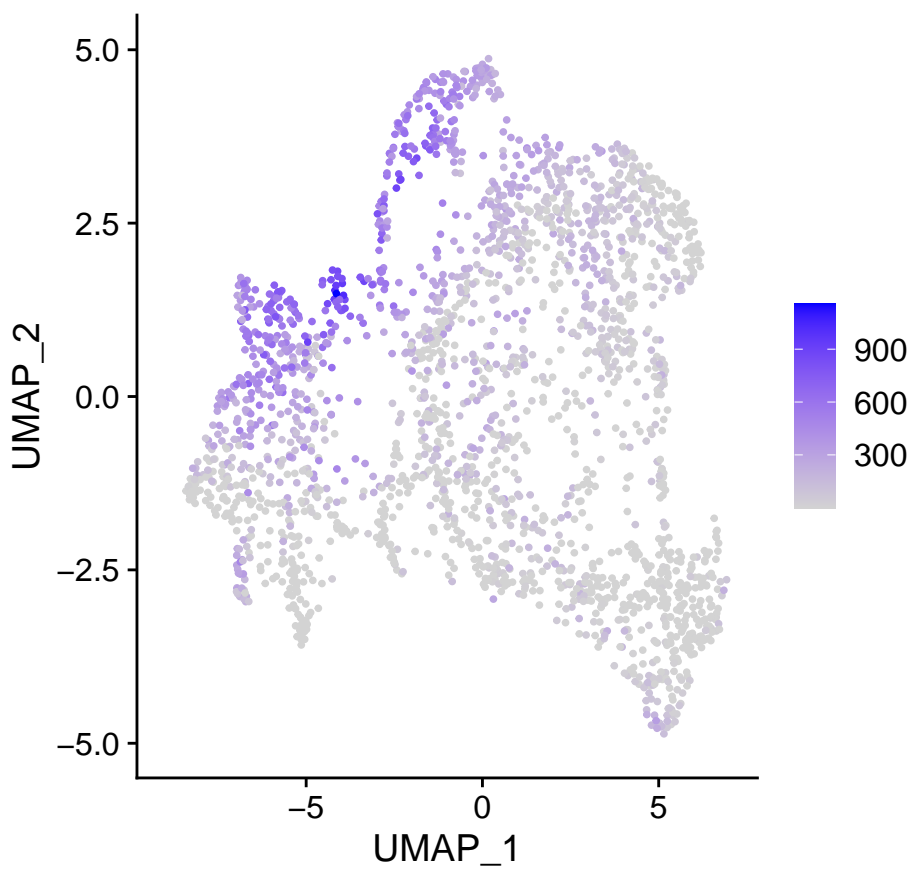
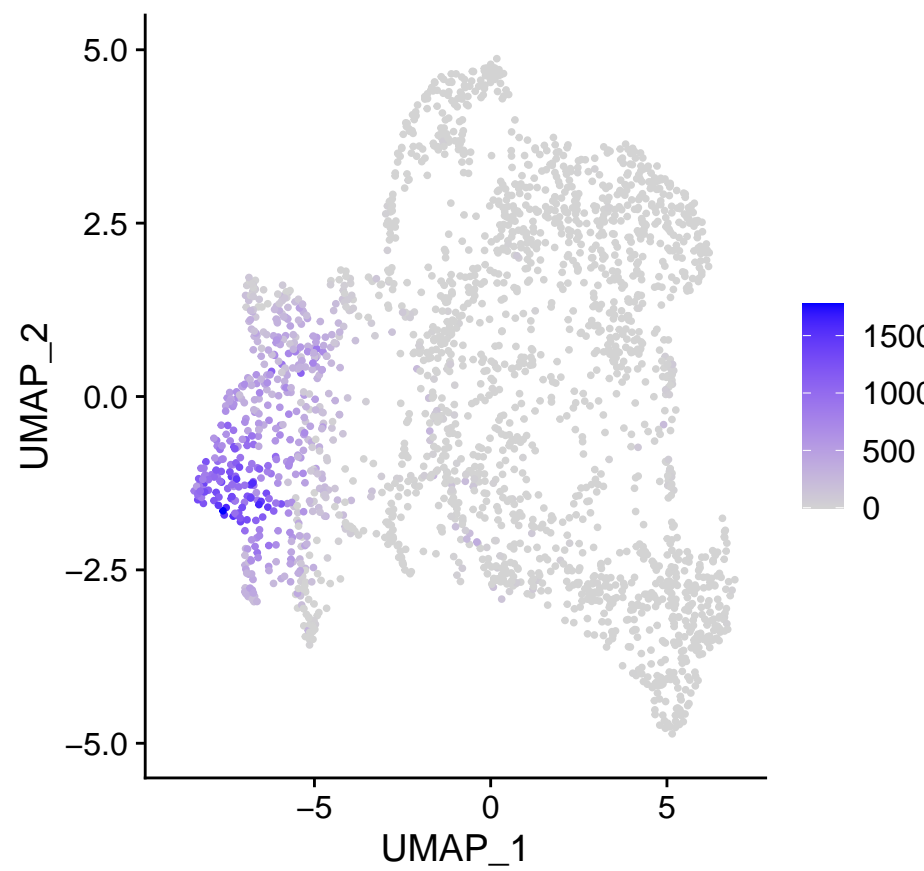
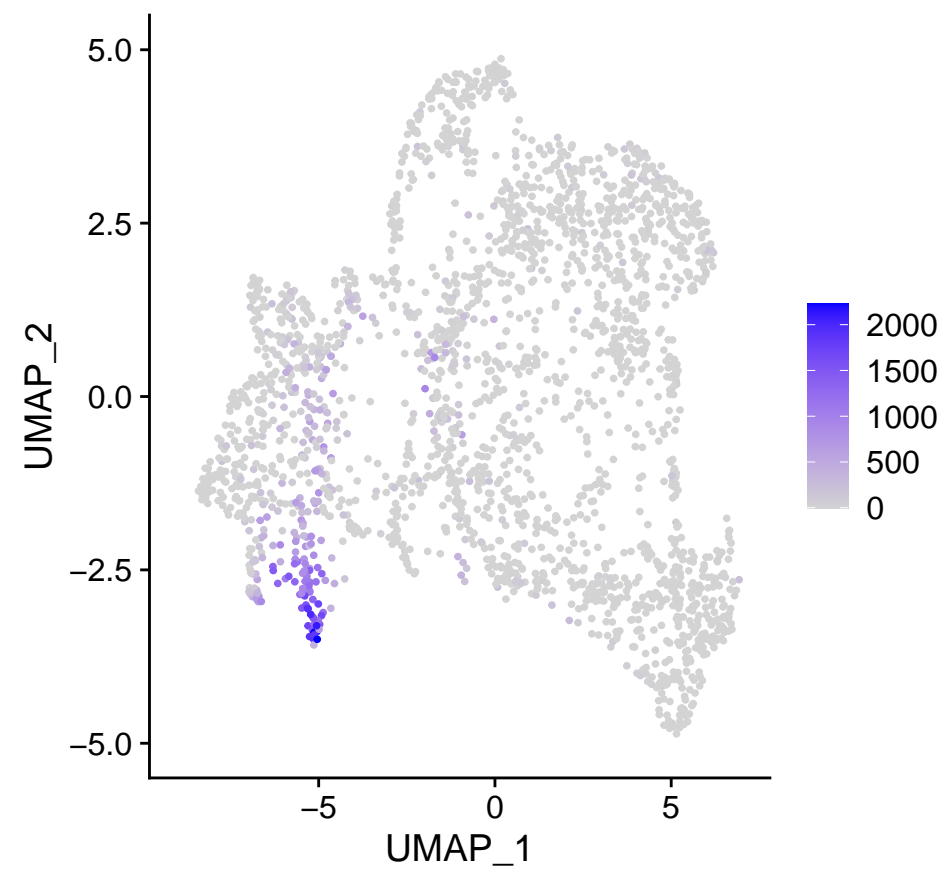
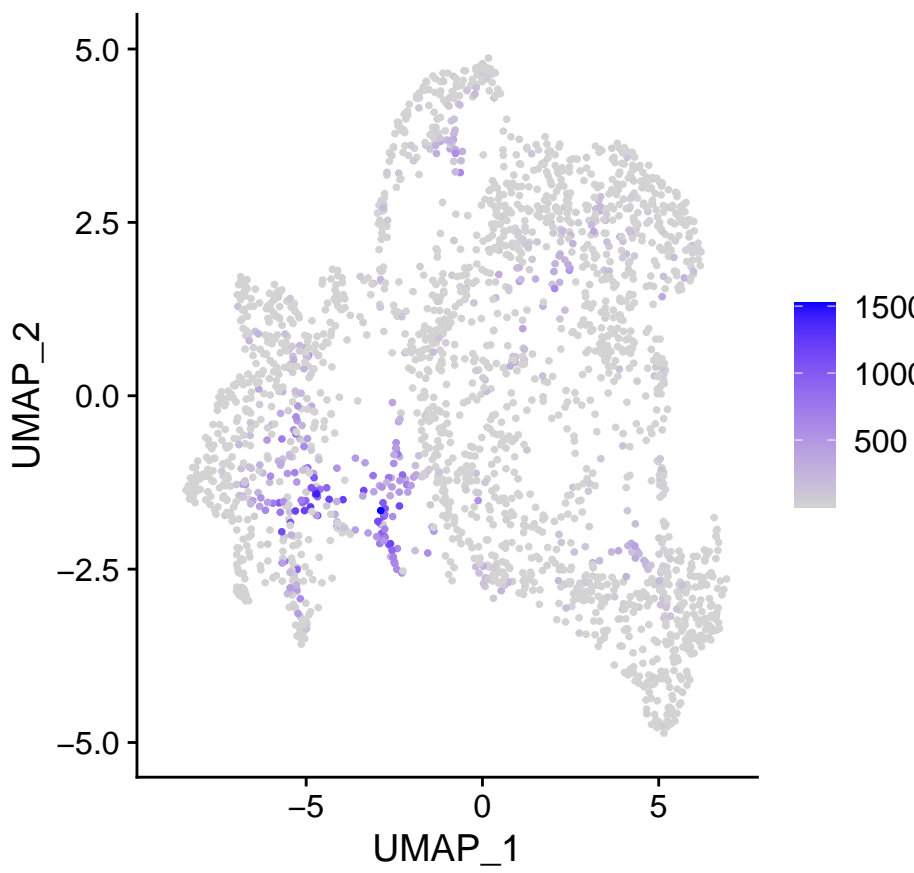
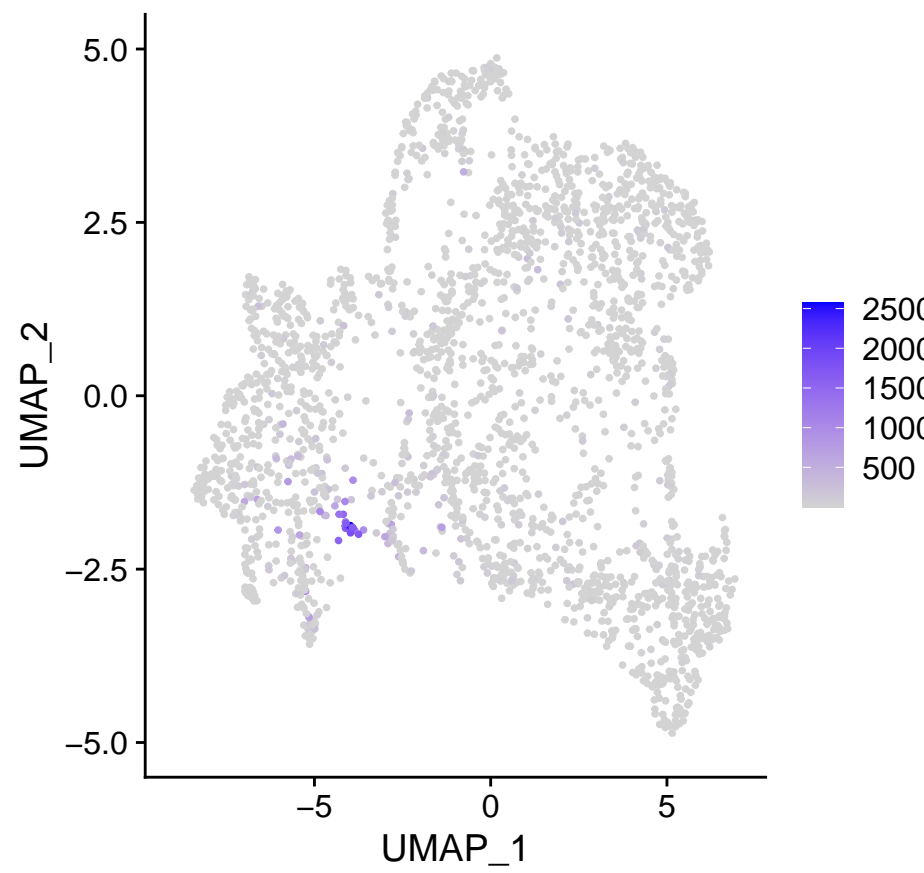
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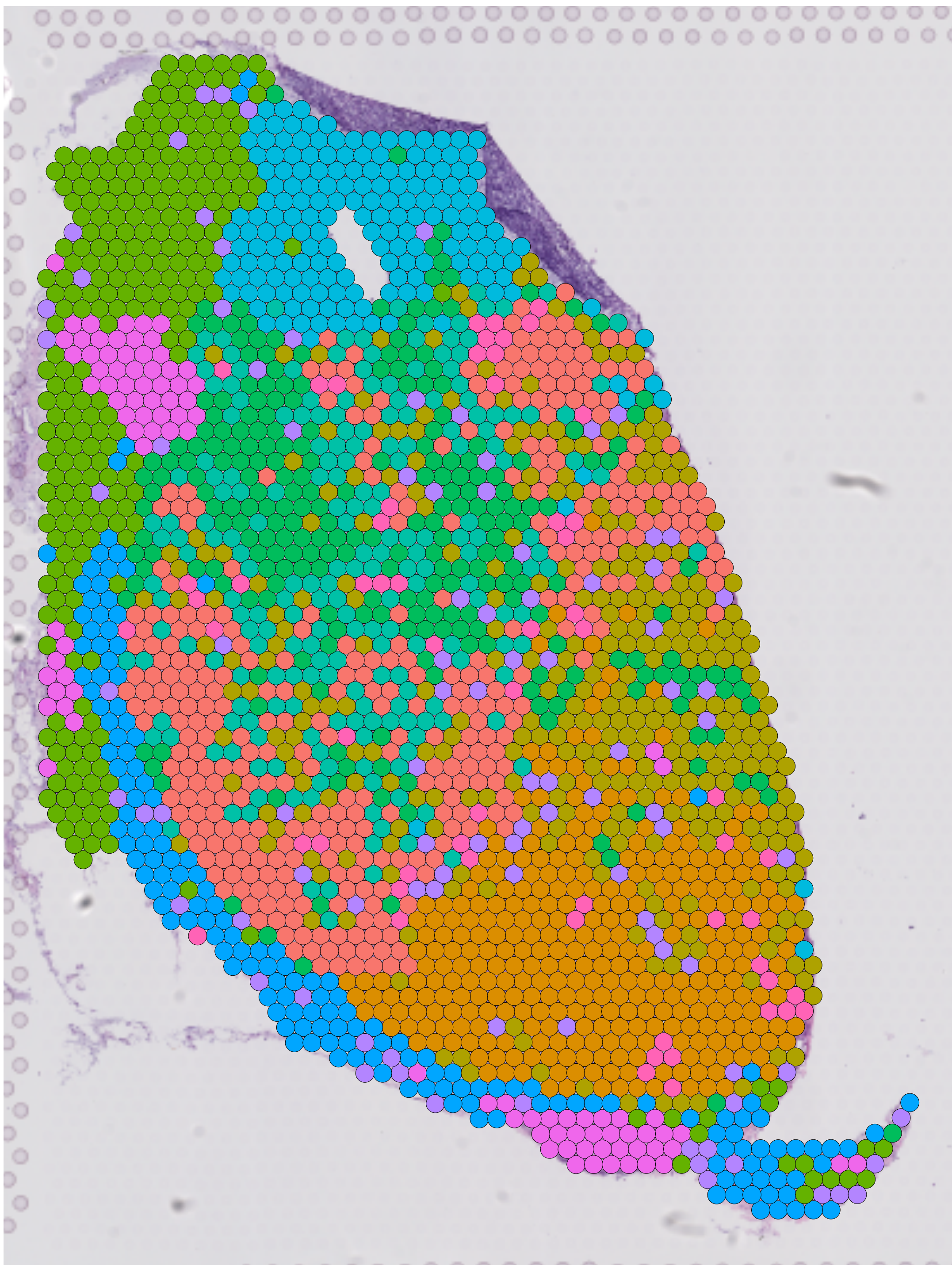
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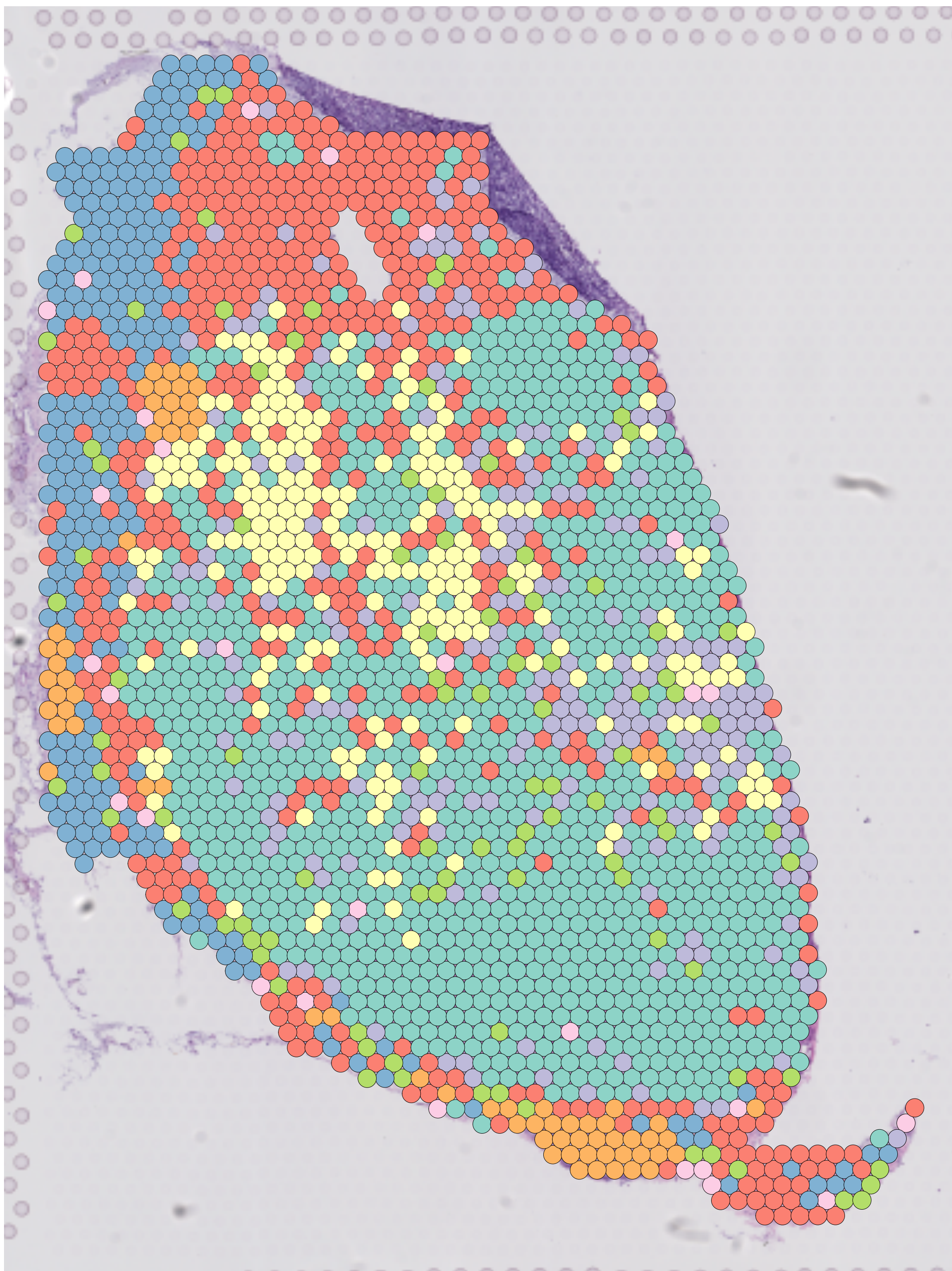


nmf



m_IGFBP5**m_HLA_DRA****m_COL1A1****m_KRT19****m_C3****m_IGKC****m_COL4A1****m_ACTA2**





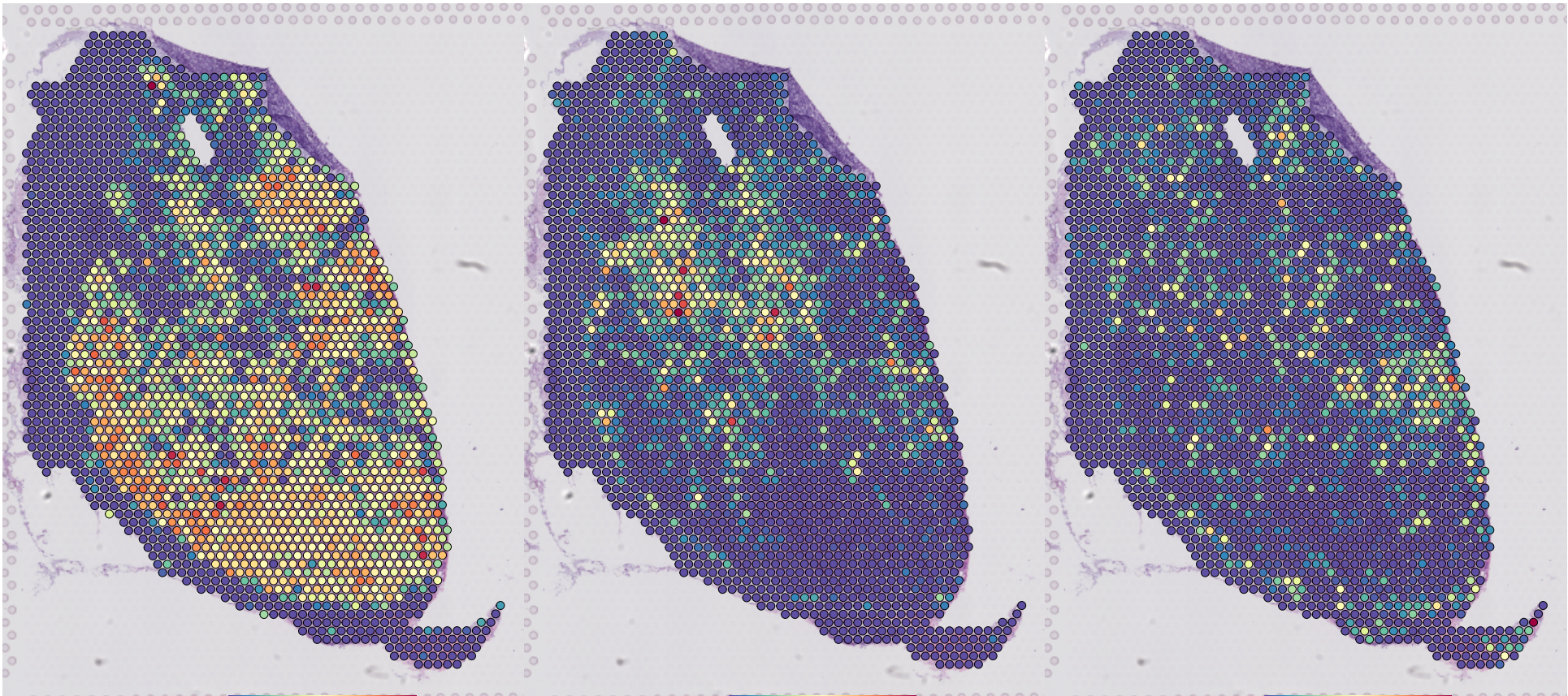
nmf

- m_IGFBP5
- m_HLA_DRA
- m_COL1A1
- m_KRT19
- m_C3
- m_IGKC
- m_COL4A1
- m_ACTA2

m_IGFBP5
100 200 300 400 500

m_HLA_DRA
0 250 500 750 1000 250

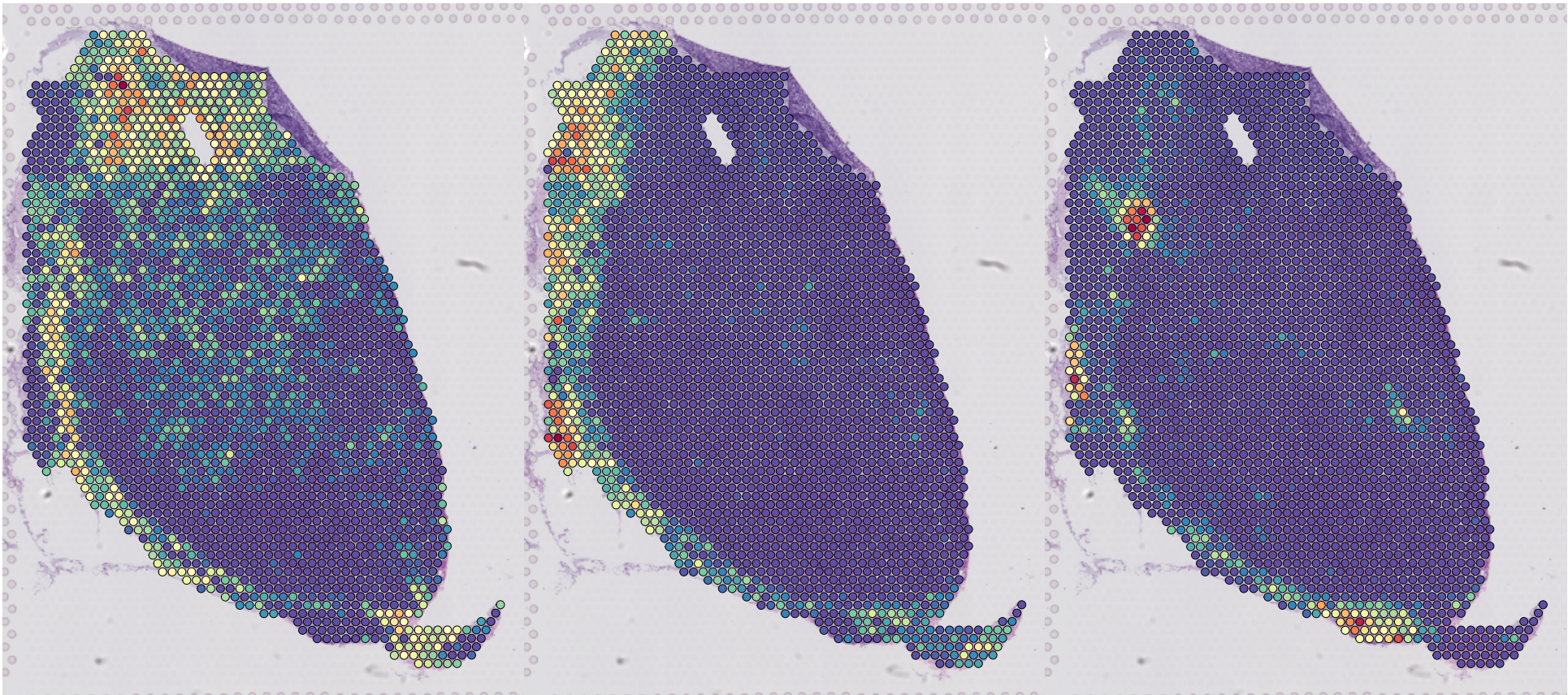
m_COL1A1
0 500 1000



m_KRT19
300 600 900

m_C3
0 500 1000 1500

m_IGKC
0 500 1000 1500 2000



m_COL4A1
500 1000 1500

m_ACTA2
500 1000 1500 2000 2500

