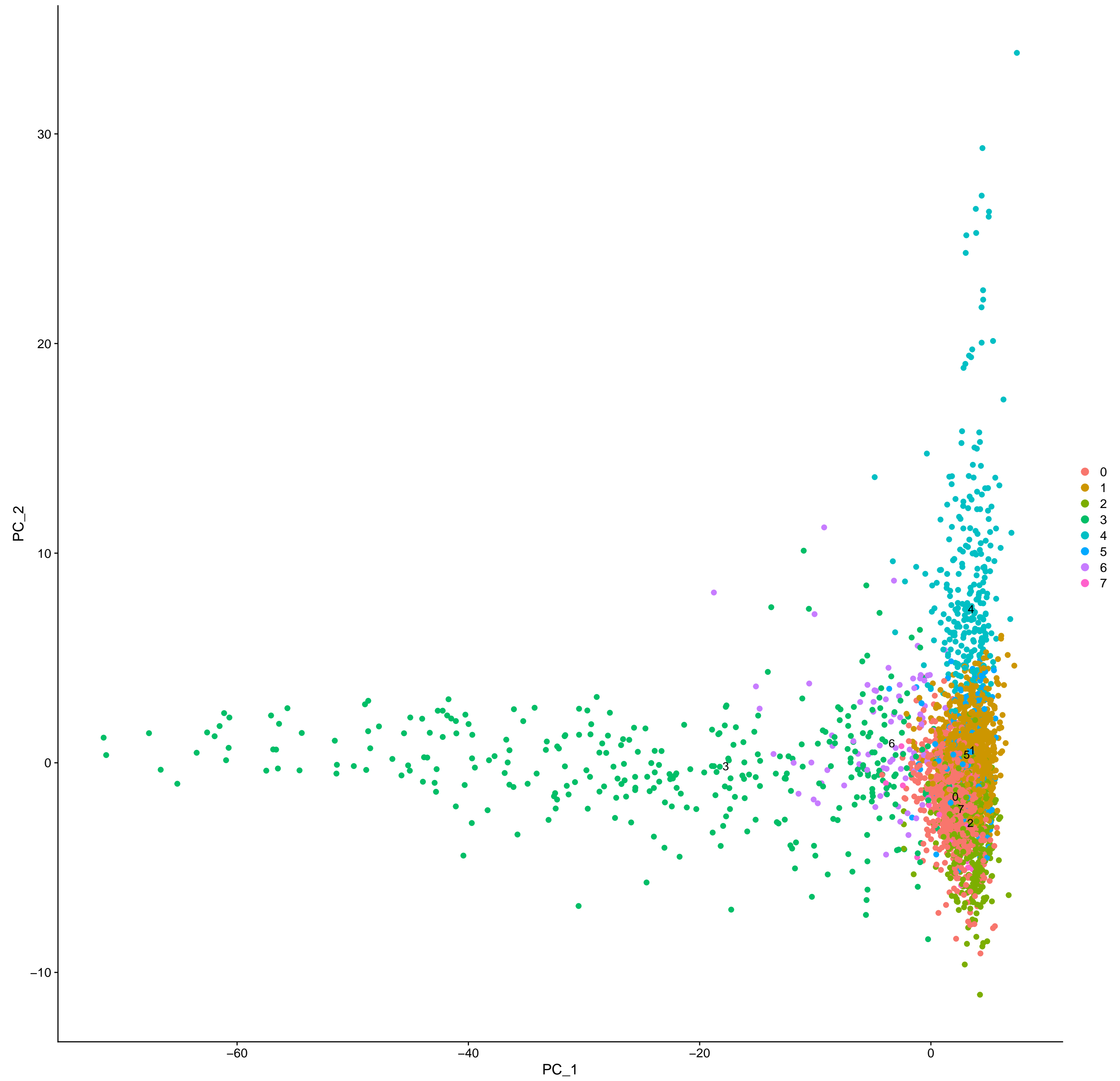
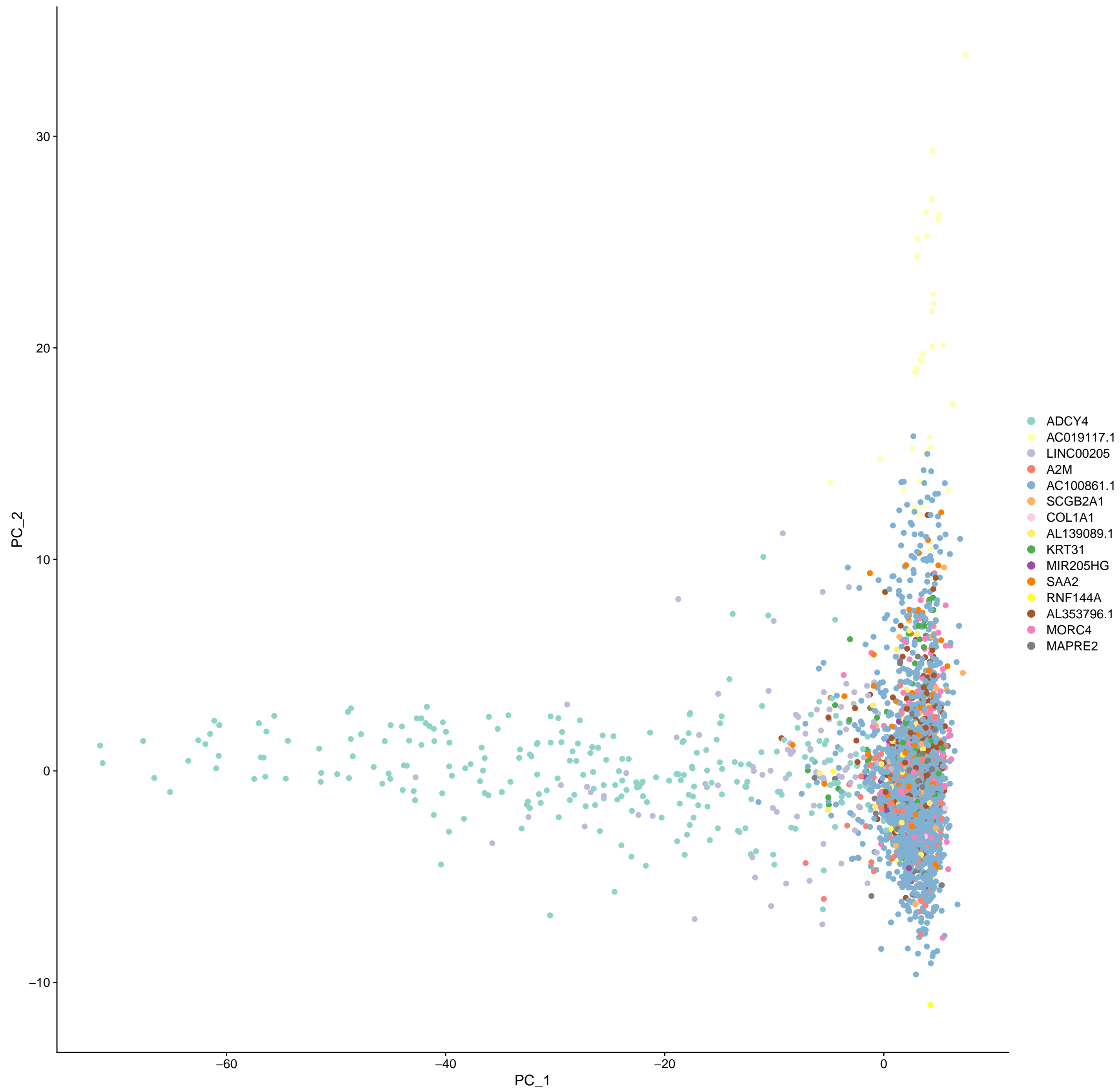


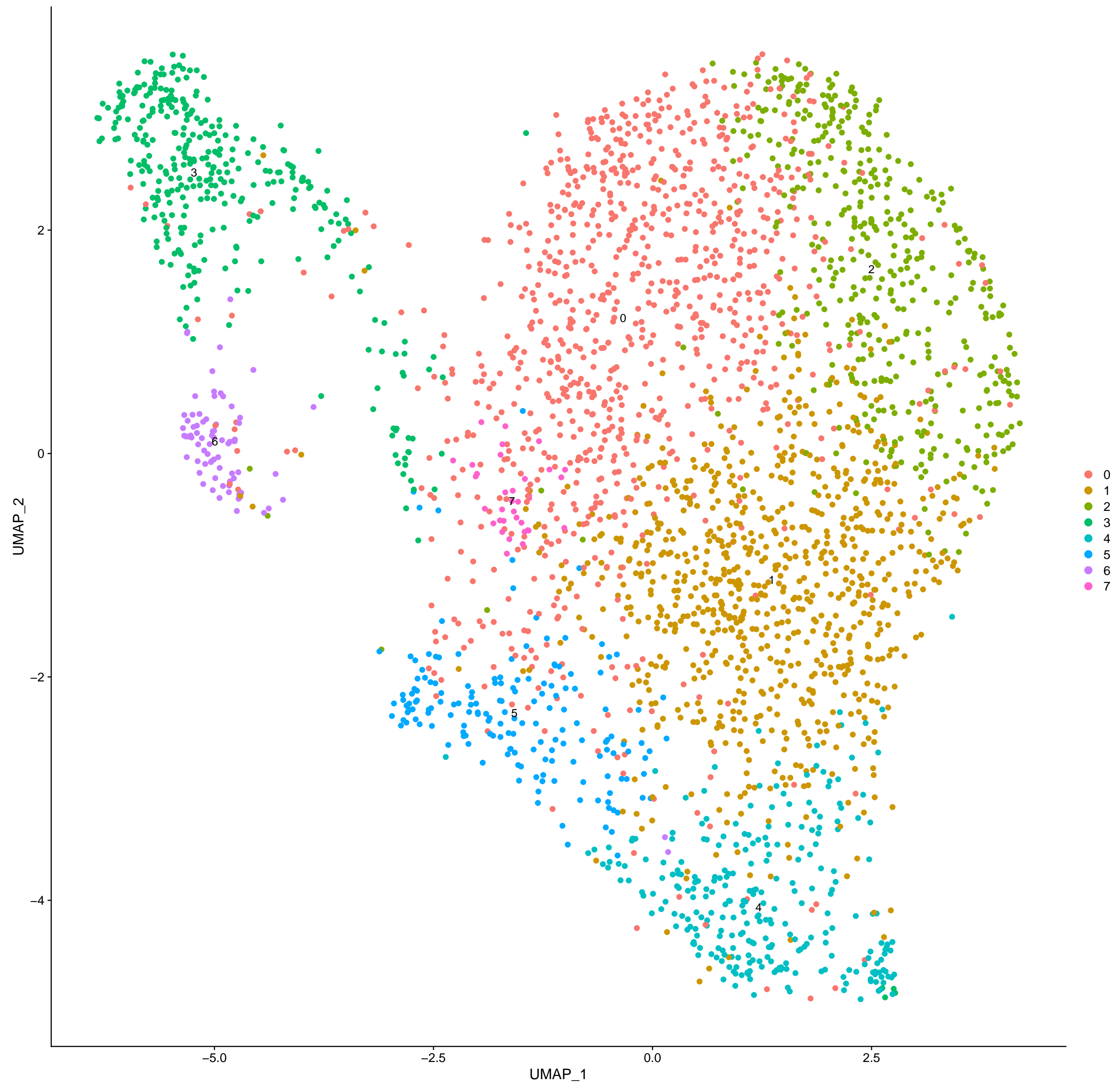
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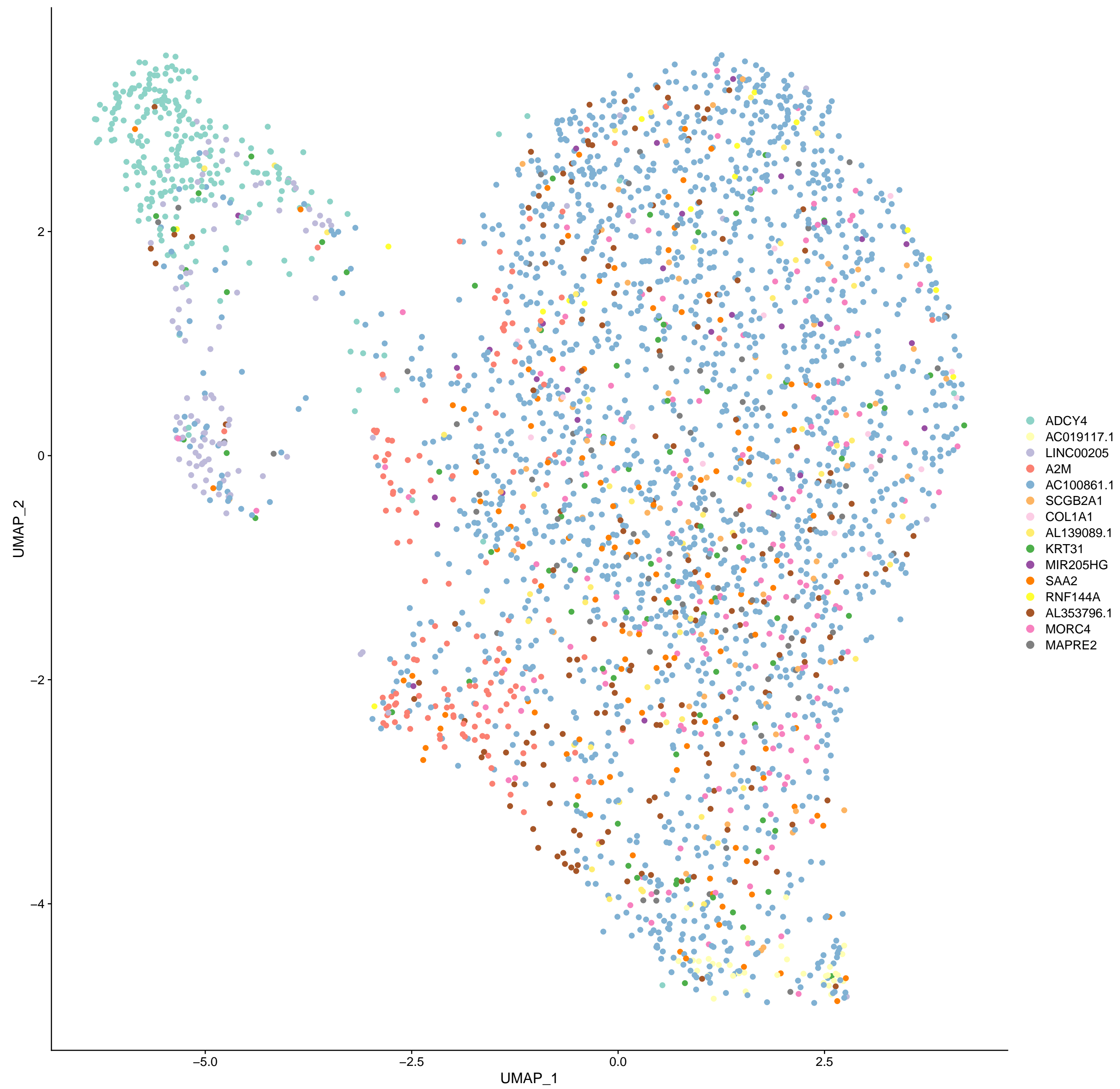
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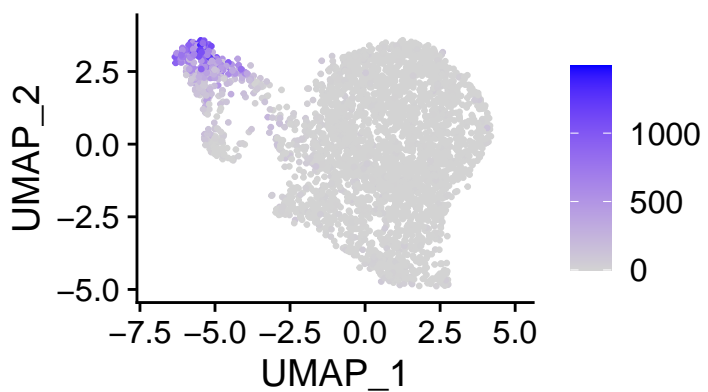
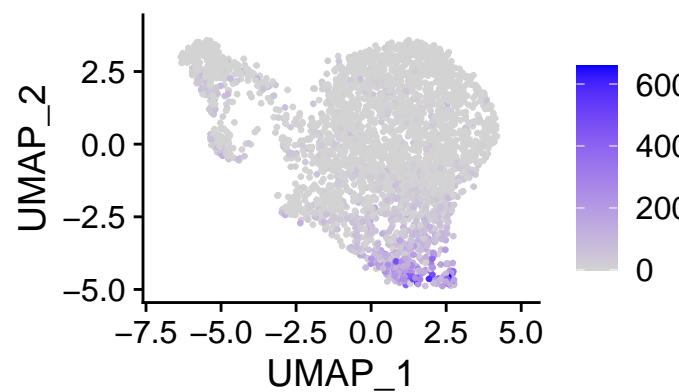
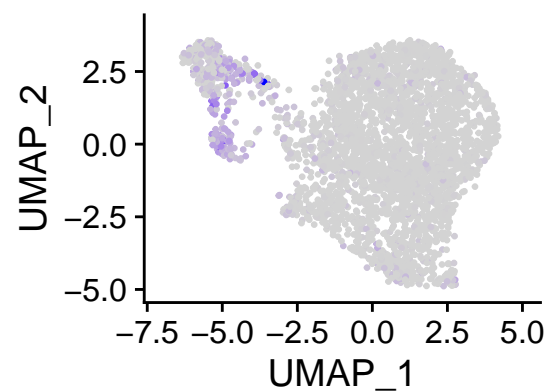
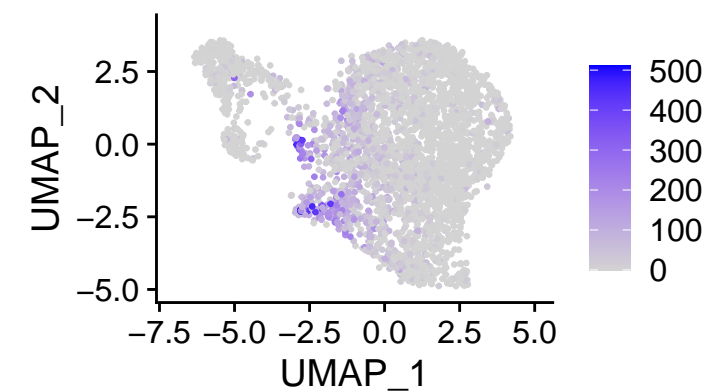
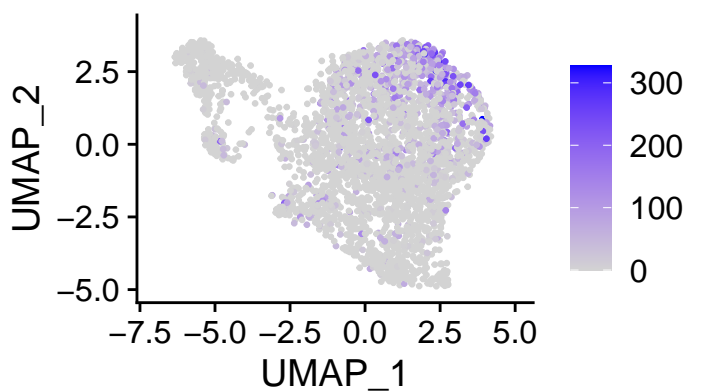
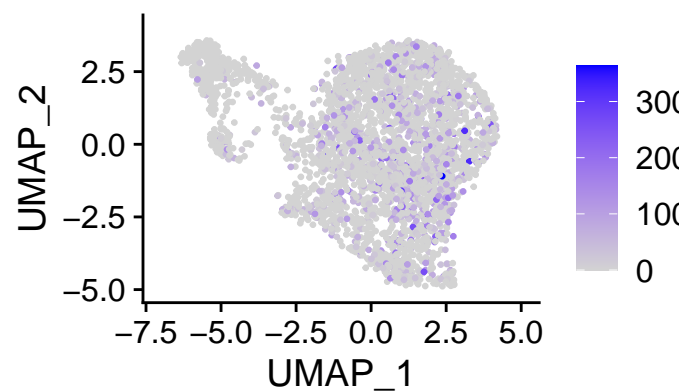
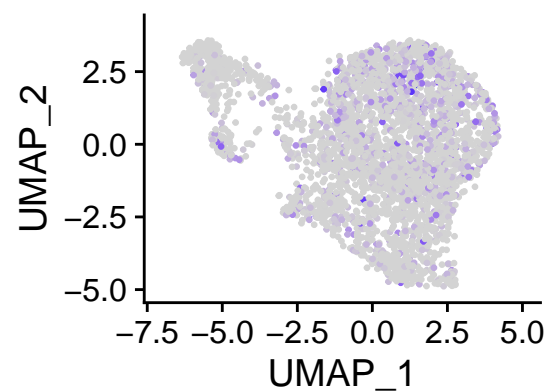
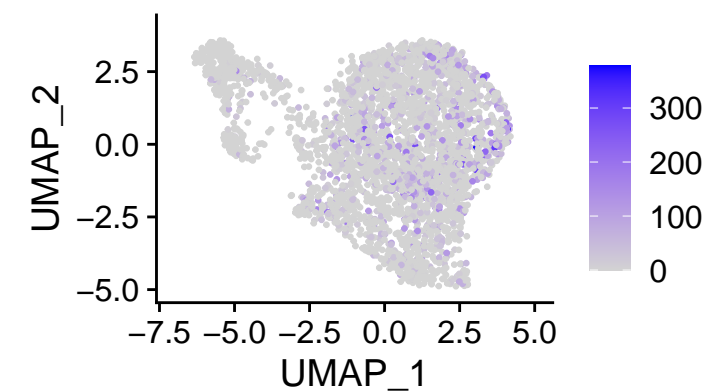
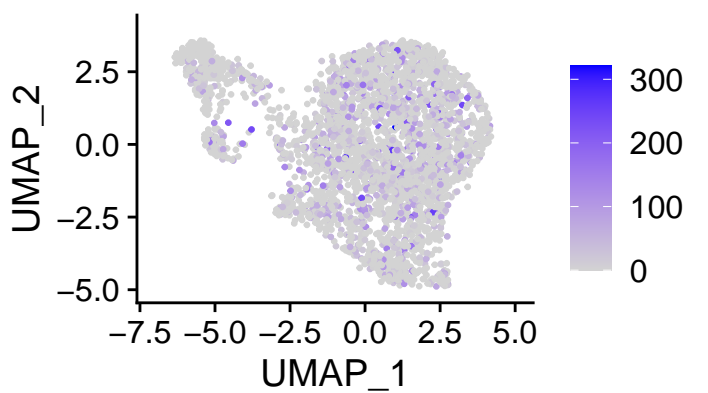
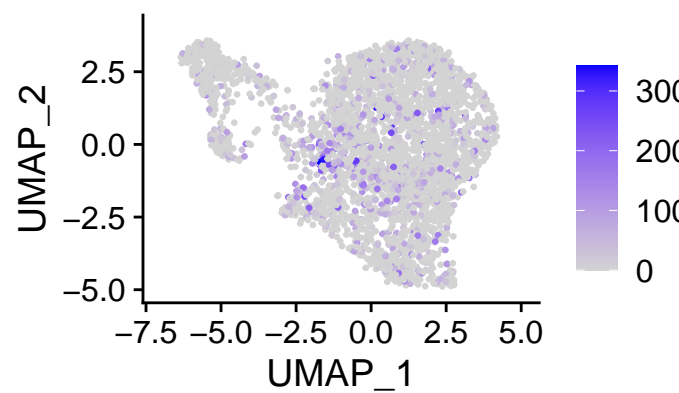
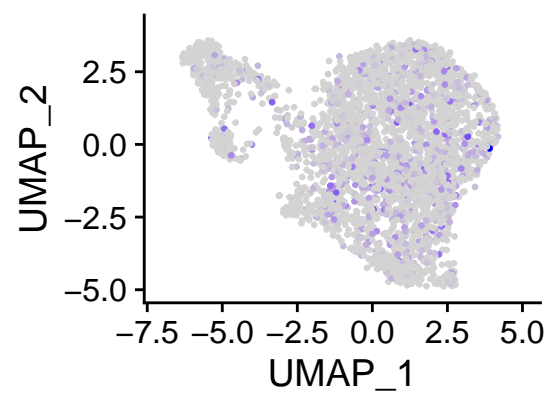
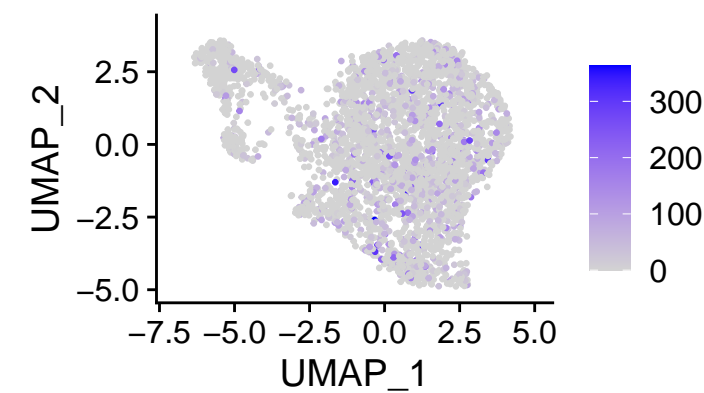
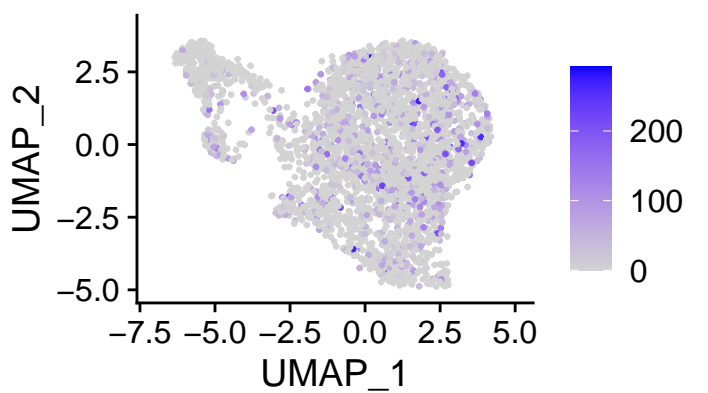
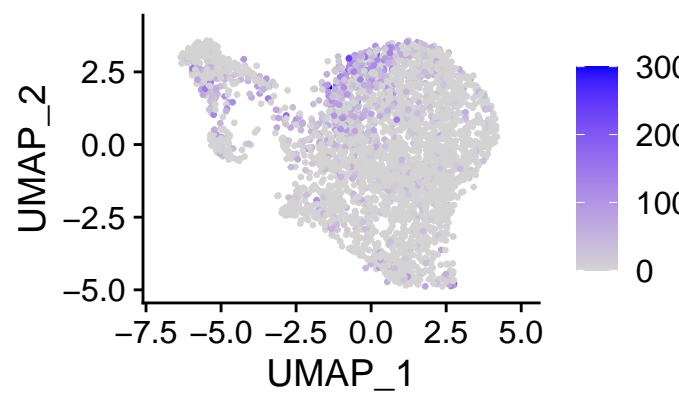
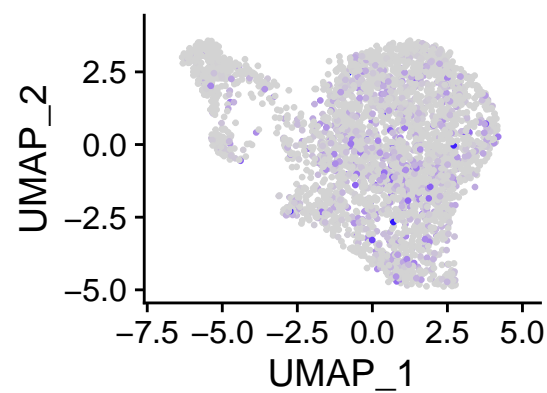
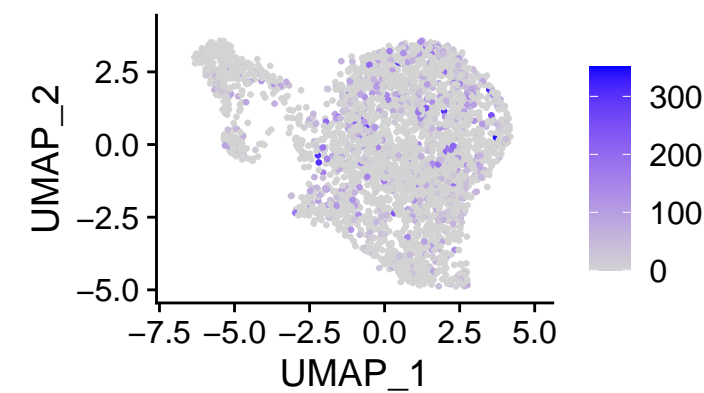
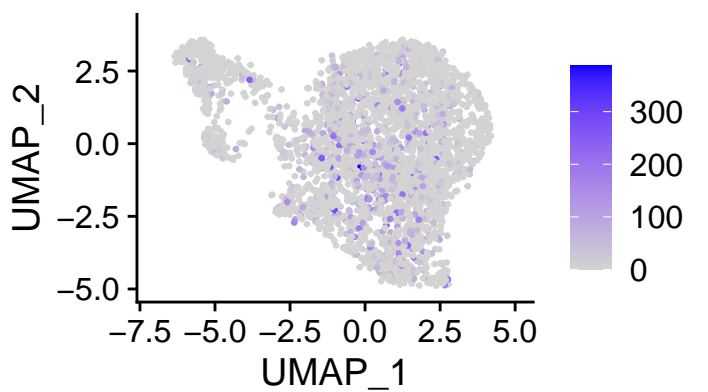
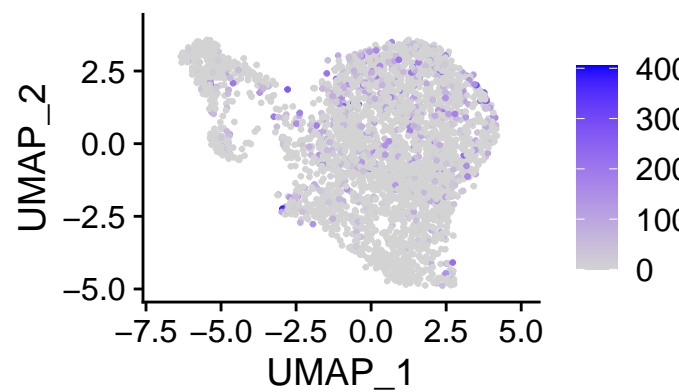
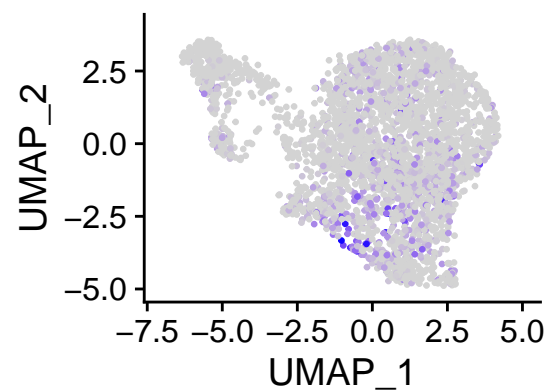
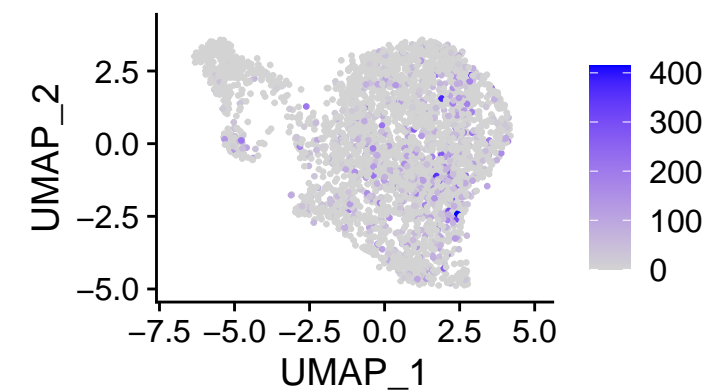
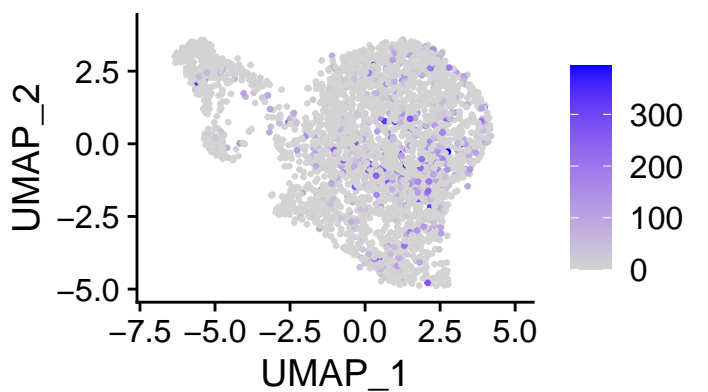


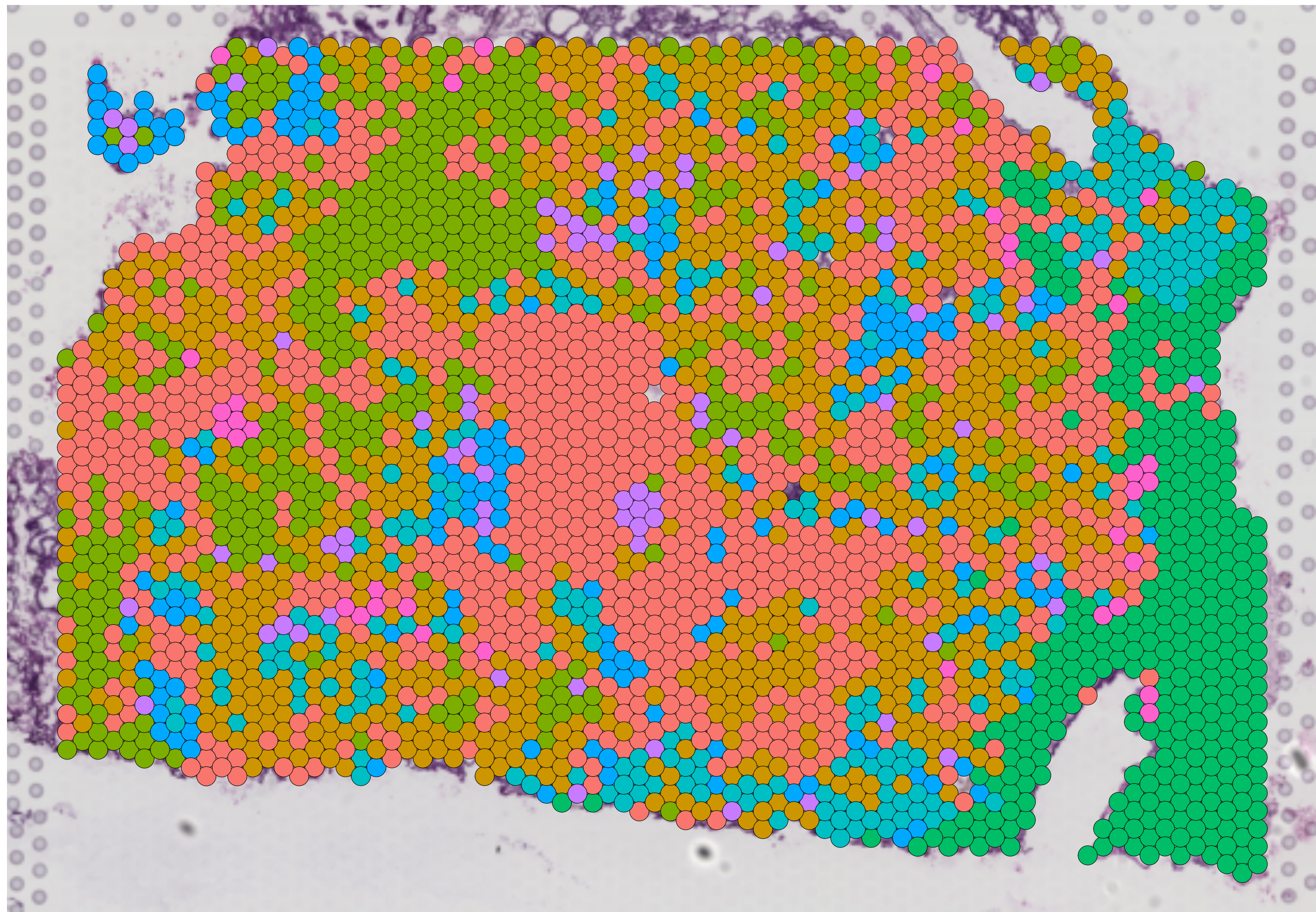
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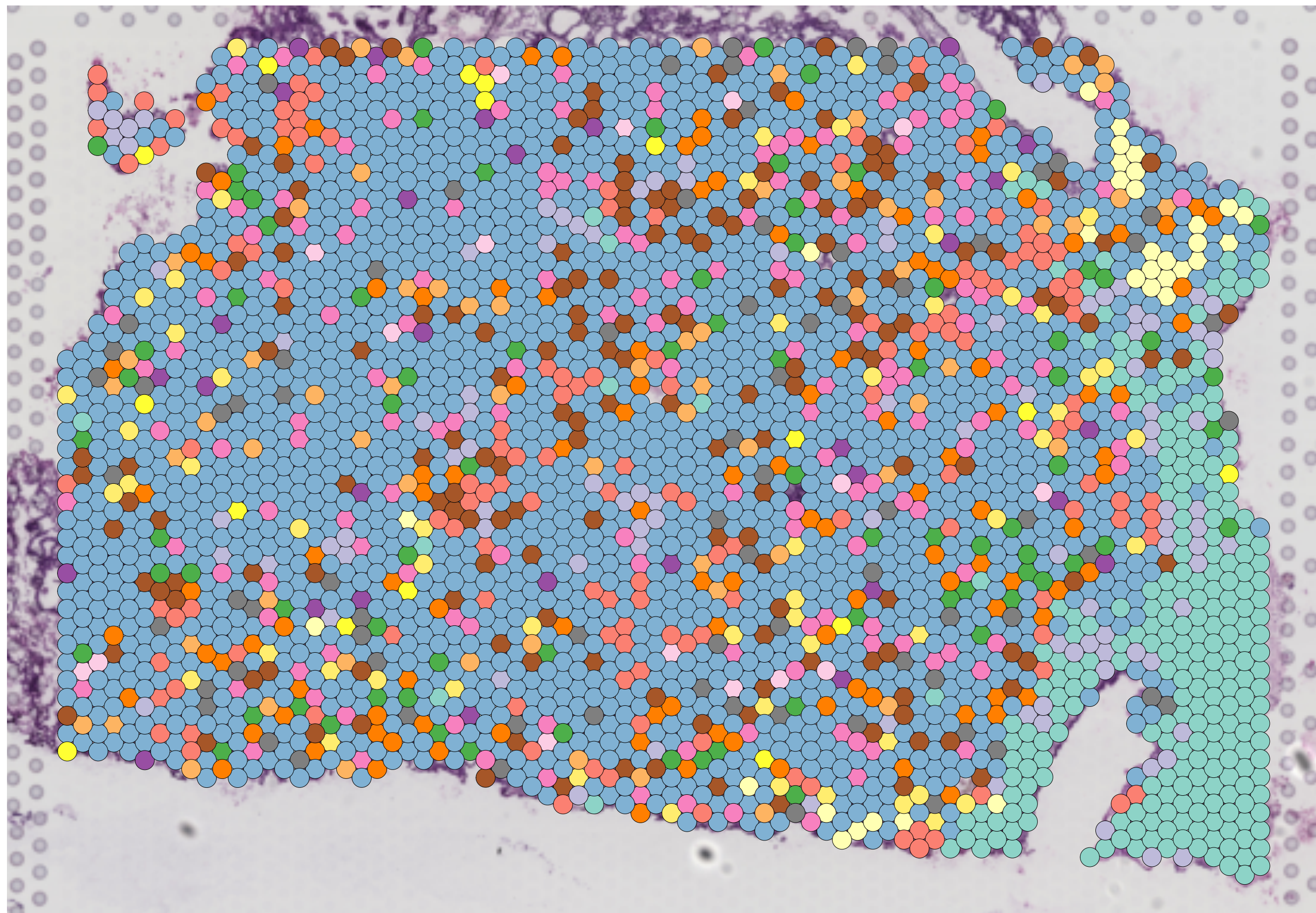


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seurat_clusters

- 0
- 1
- 2
- 3
- 4
- 5
- 6
- 7



nmf

- ADCY4
- AC019117.1
- LINC00205
- A2M
- AC100861.1
- SCGB2A1
- COL1A1
- AL139089.1
- KRT31
- MIR205HG
- SAA2
- RNF144A
- AL353796.1
- MORC4
- MAPRE2

