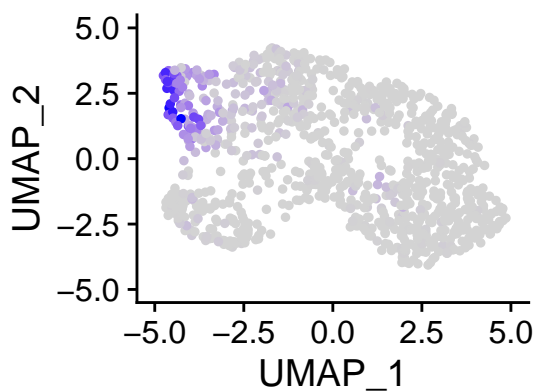
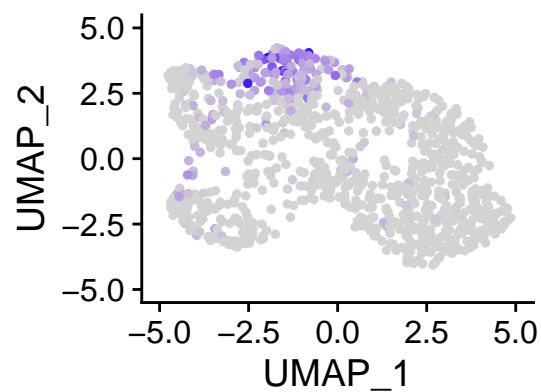
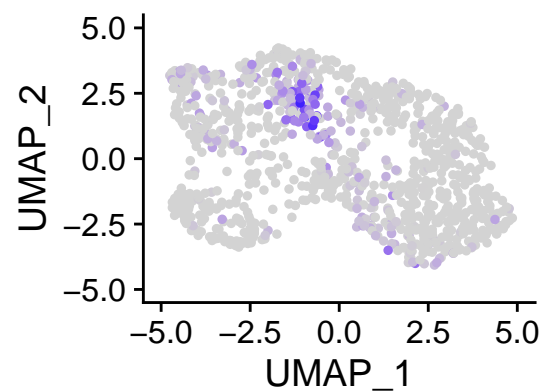
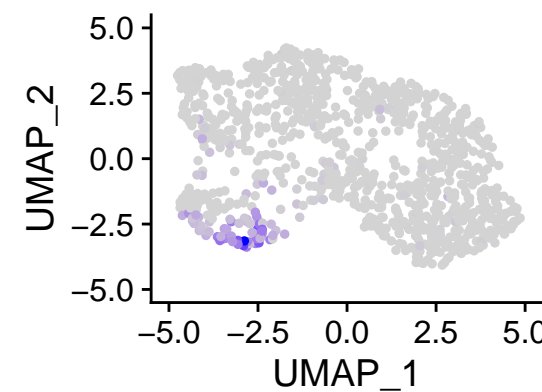
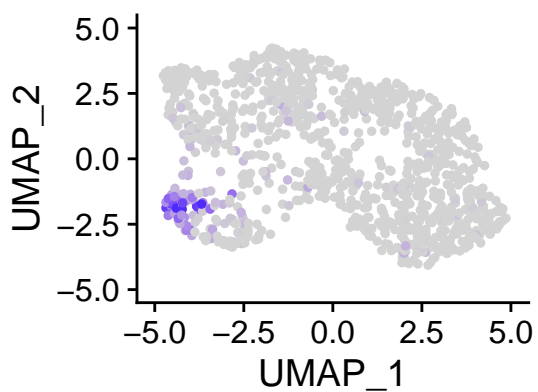
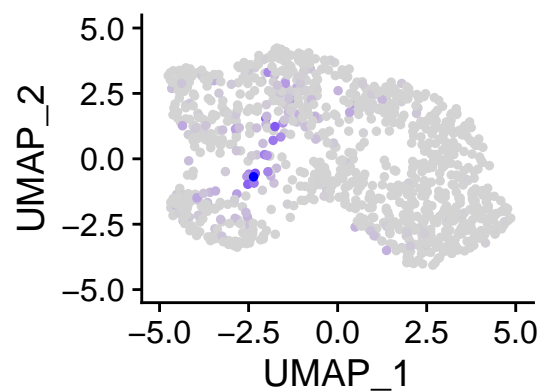
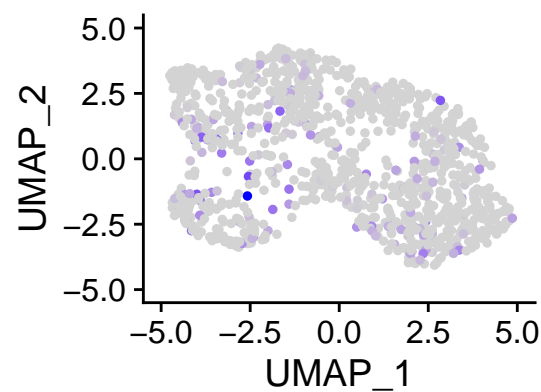
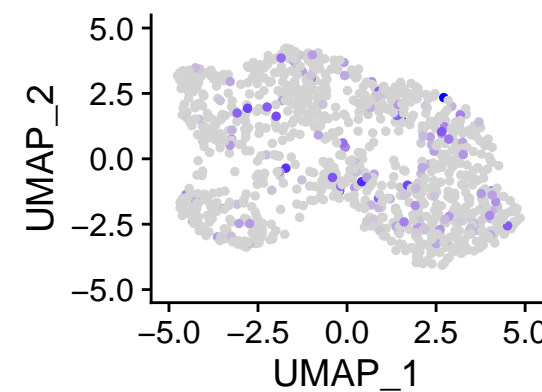
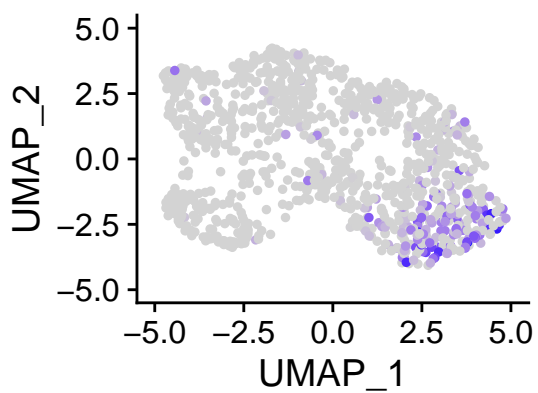
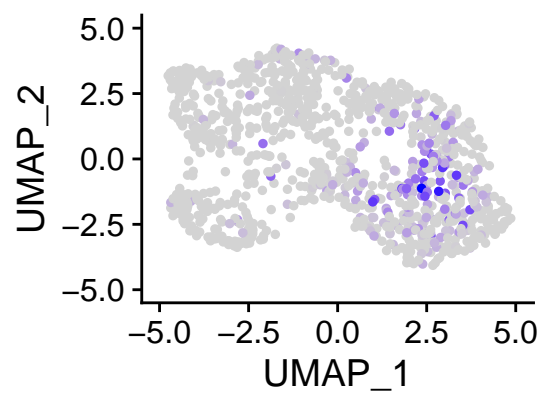
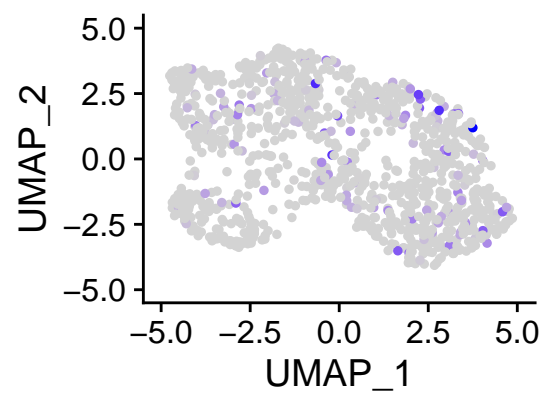
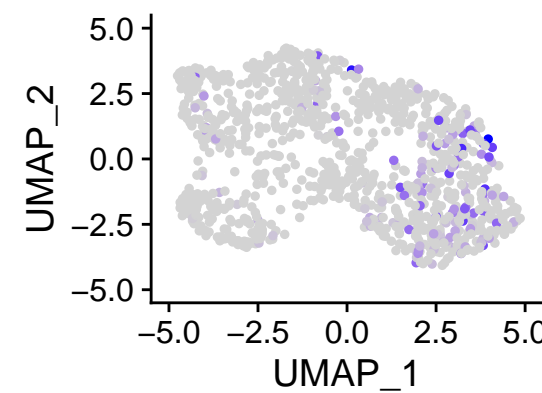
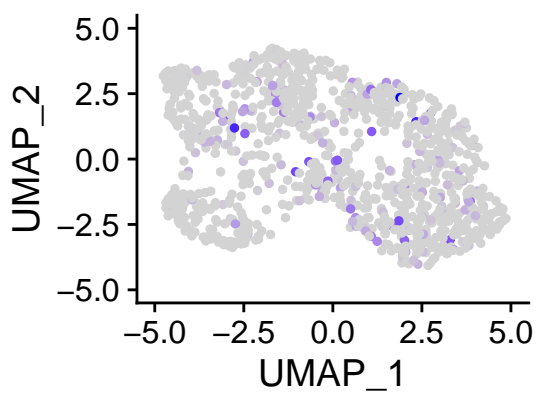
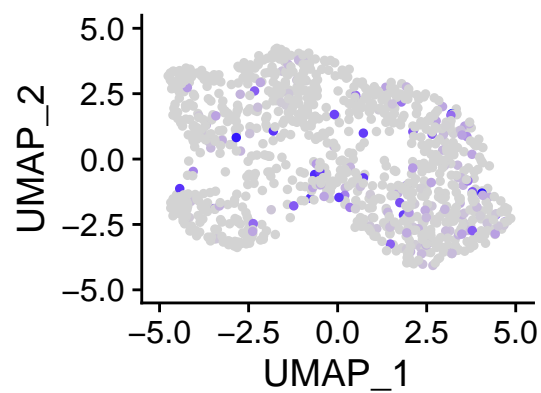
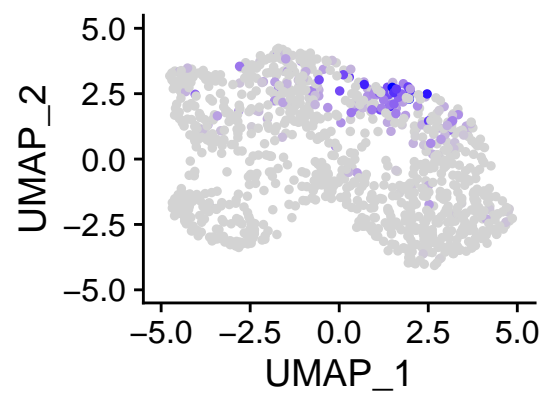
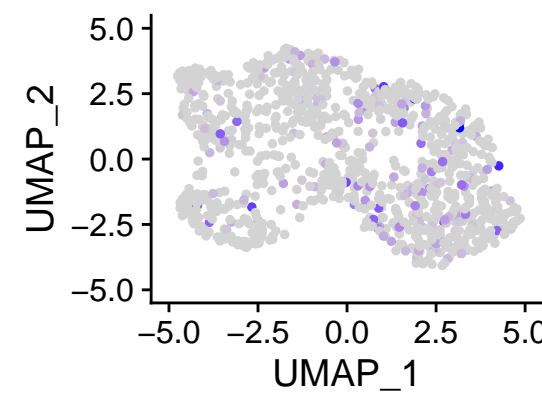
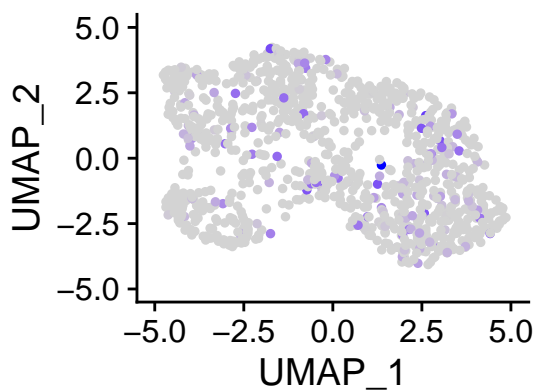
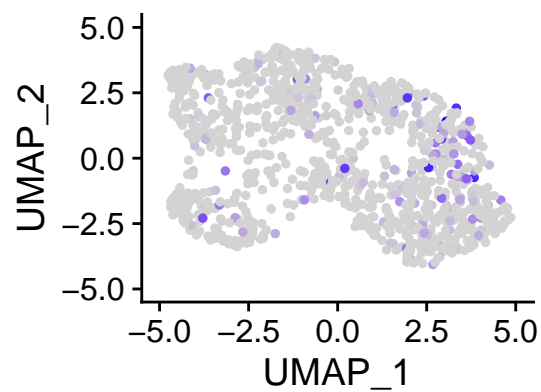
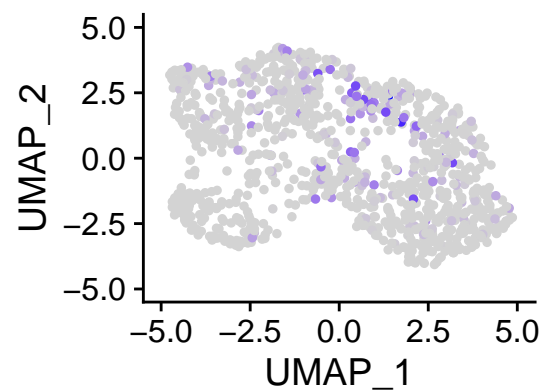
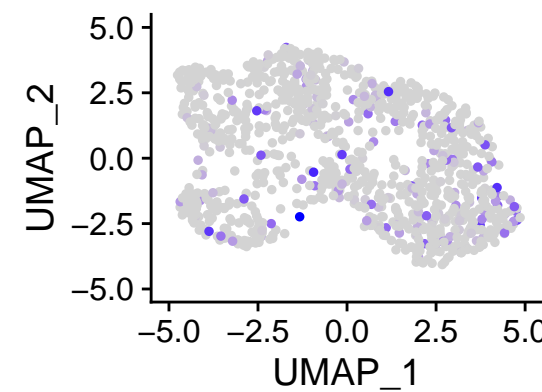
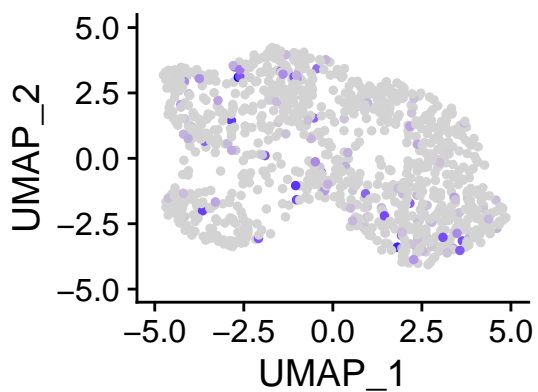
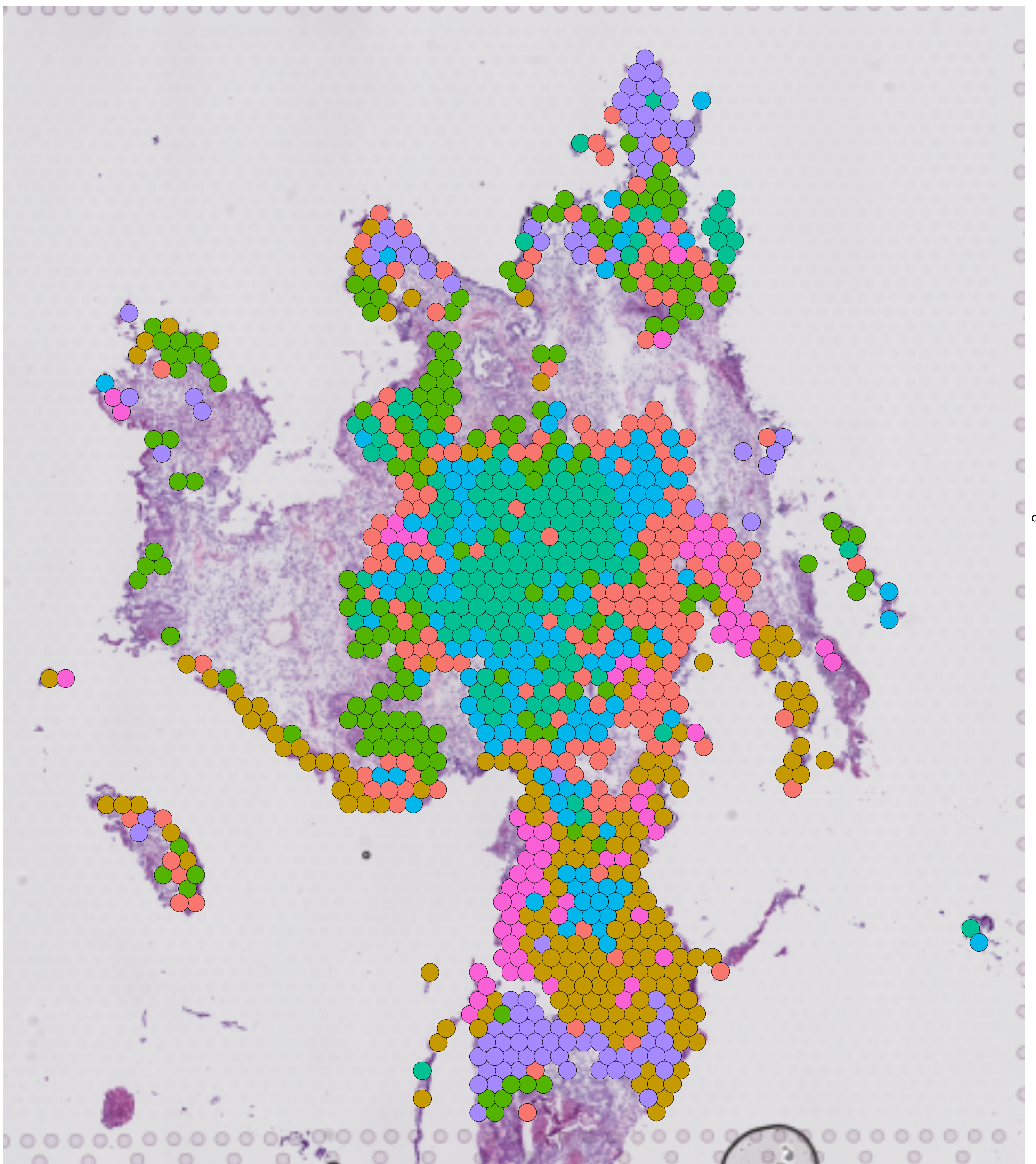
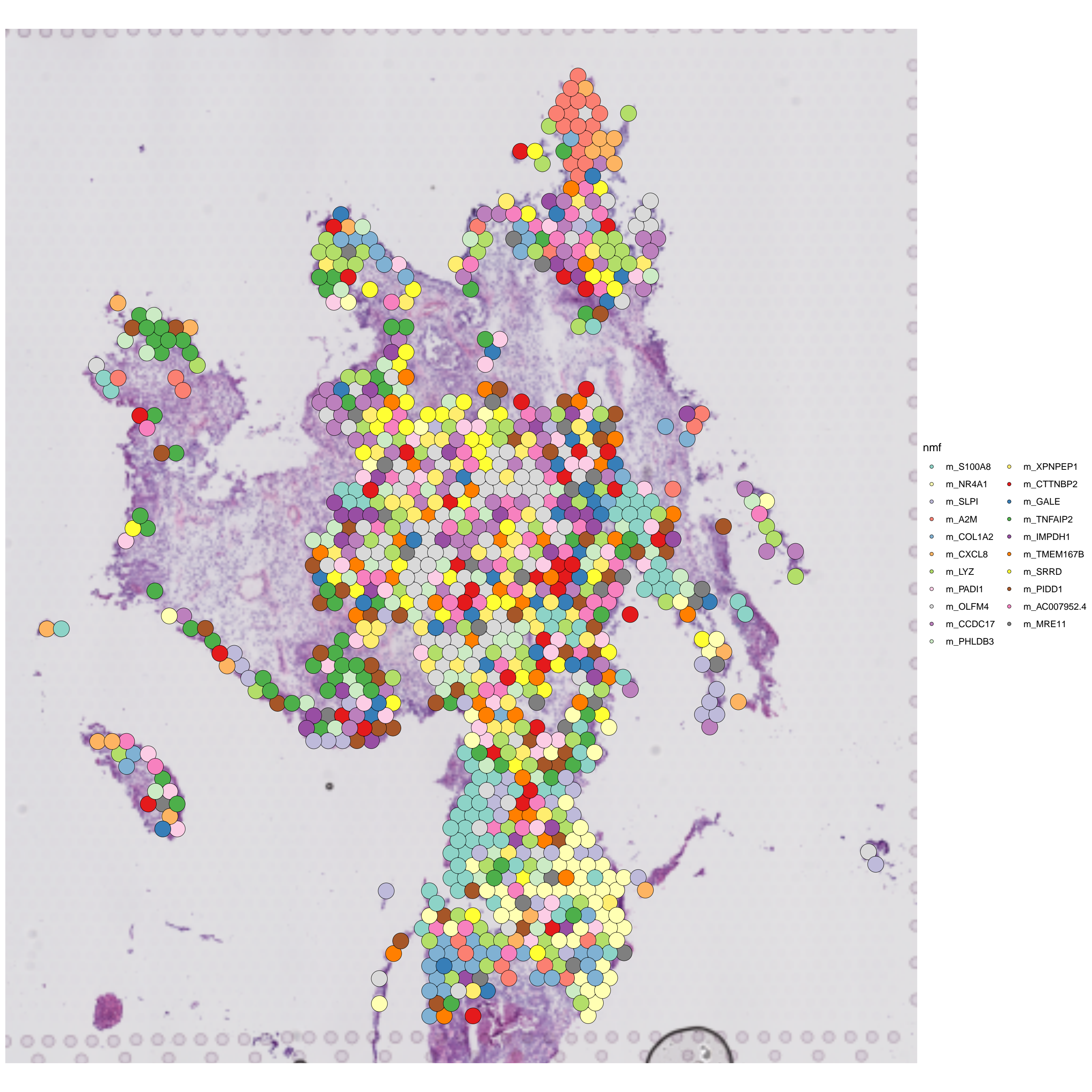


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cluster

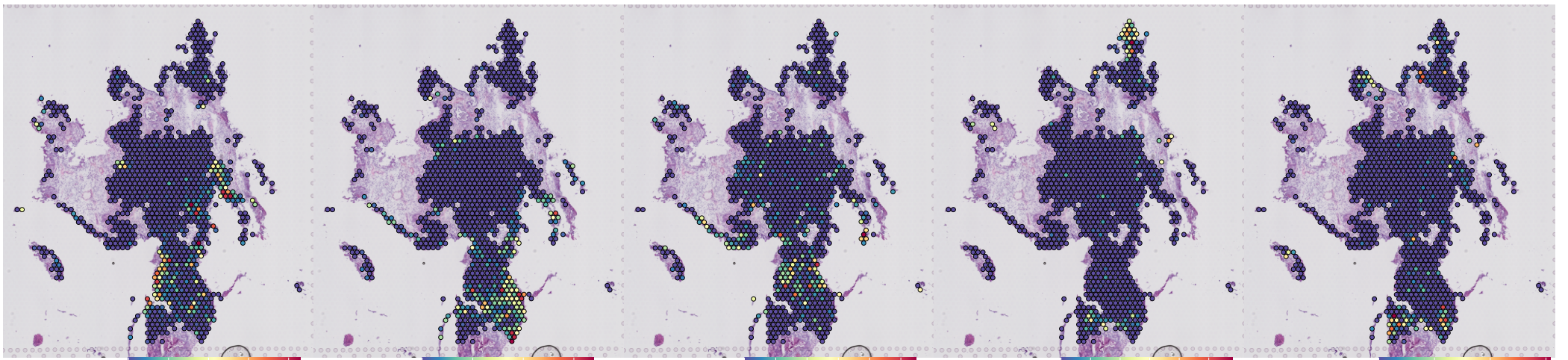
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- 3
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- 6



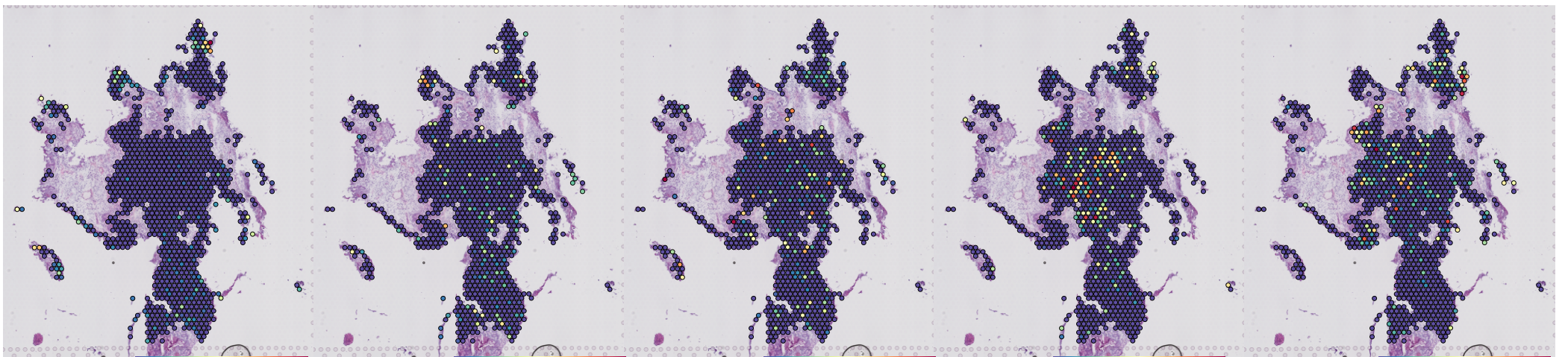
nmf

- | | |
|------------|----------------|
| ● m_S100A8 | ● m_XPNPEP1 |
| ● m_NR4A1 | ● m_CTTNBP2 |
| ● m_SLPI | ● m_GALE |
| ● m_A2M | ● m_TNFAIP2 |
| ● m_COL1A2 | ● m_IMPDI1 |
| ● m_CXCL8 | ● m_TMED167B |
| ● m_LYZ | ● m_SRRD |
| ● m_PADI1 | ● m_PIDD1 |
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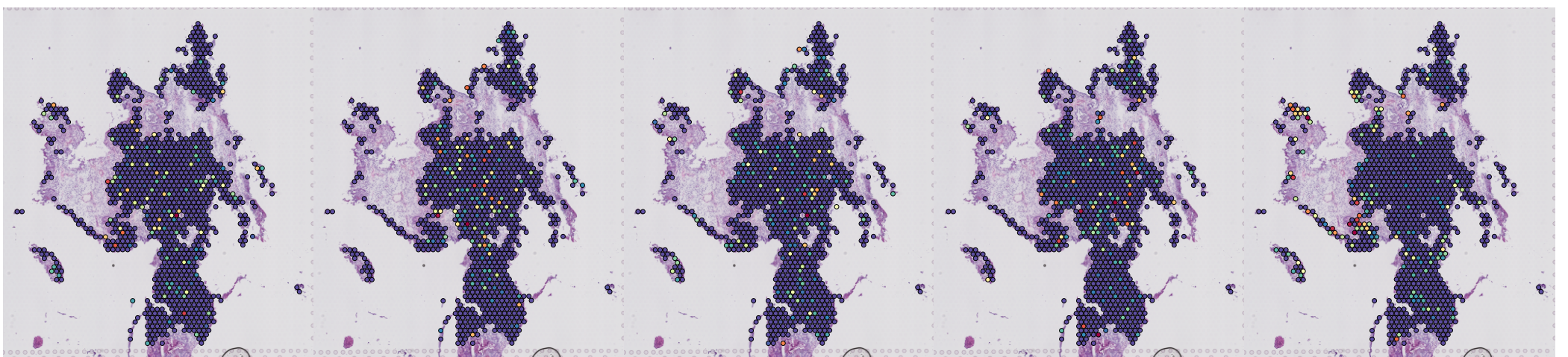
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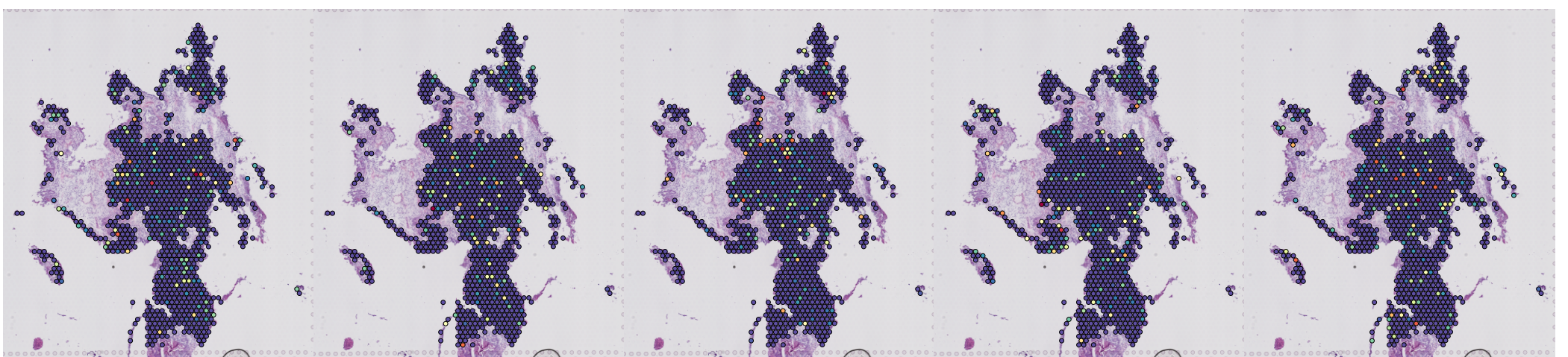
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m_IMP DH1 0 100 200 300 400 500 m_TM EM167B 0 200 400 600 m_SRRD 0 100 200 300 400 500 m_PIDD1 0 200 400 600 m_AC007952.4 100 200 300 400 500



m_MRE11 0 100 200 300 400

