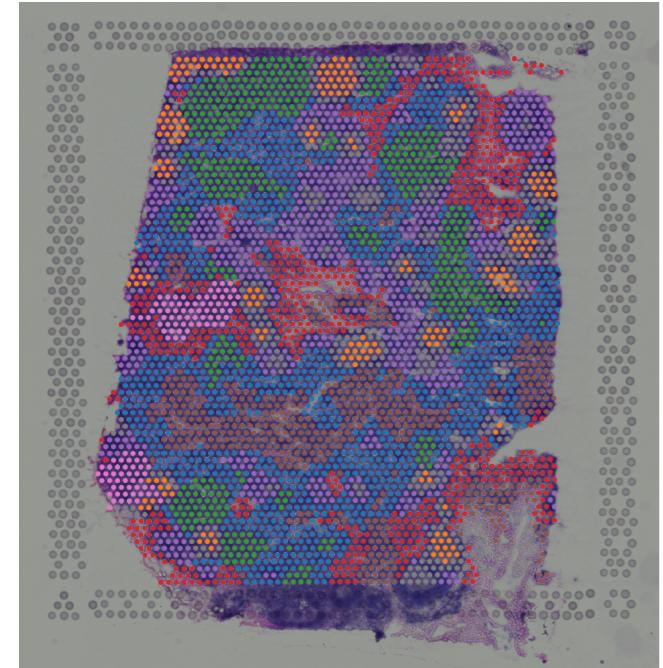
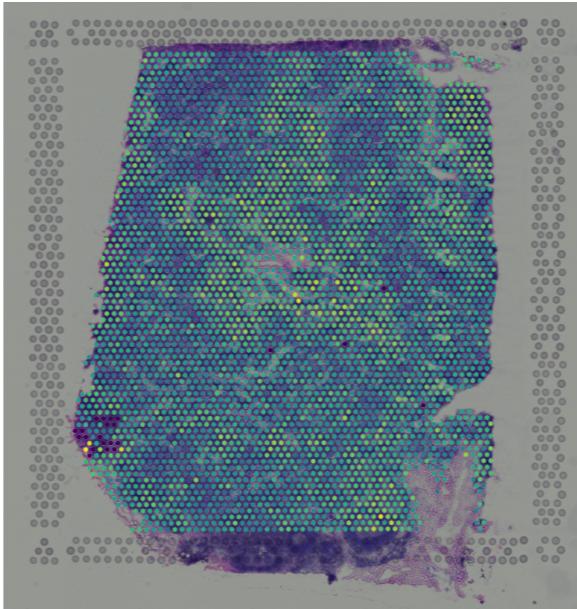


Spatial Transcriptomic Analysis of Human Lymph Node

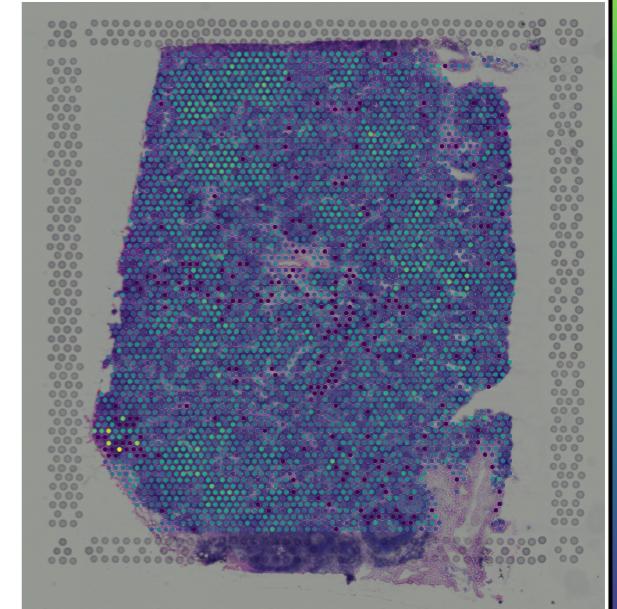
A) Spatial Domains



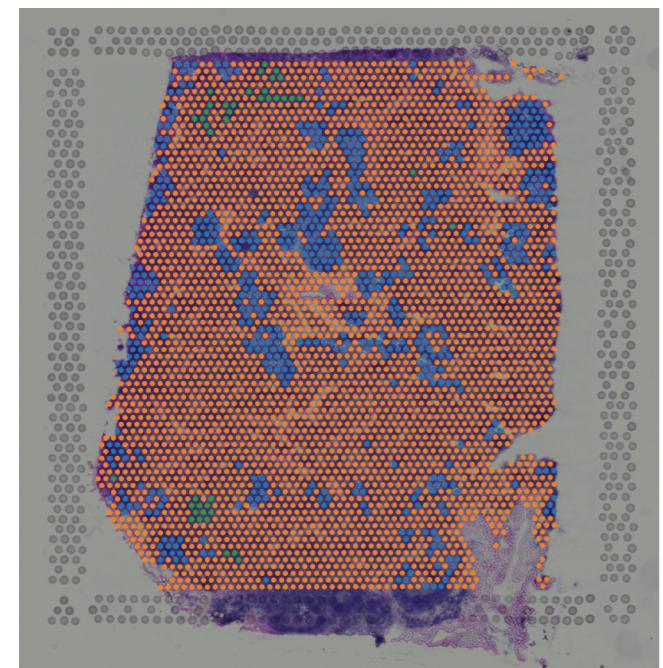
B) MS4A1 Expression (B cells)



C) CD3E Expression (T cells)

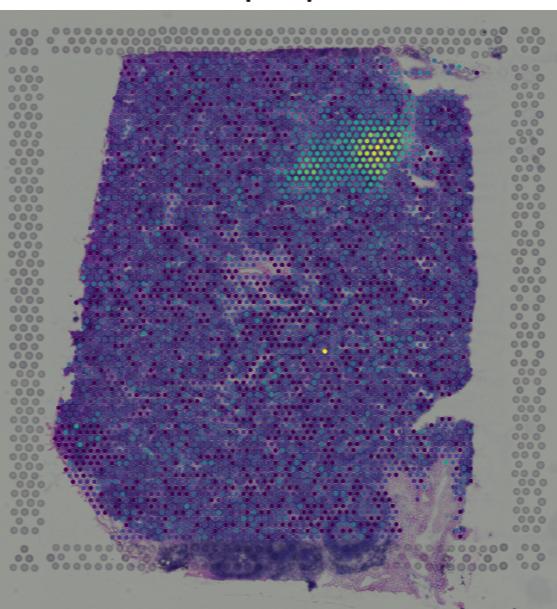


D) Lymph Node Regions



- B_cell_zone
- Medullary_cords
- T_cell_zone

E) ISG15 (Top Spatial Gene)



F) B Cell Proportion

