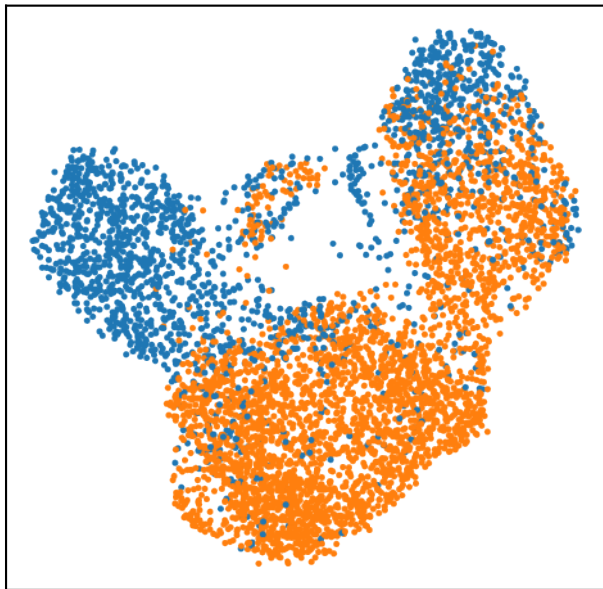


# GC\_celltype\_predictions

UMAP2



- GC\_B\_cells
- Other\_B\_cells

UMAP1