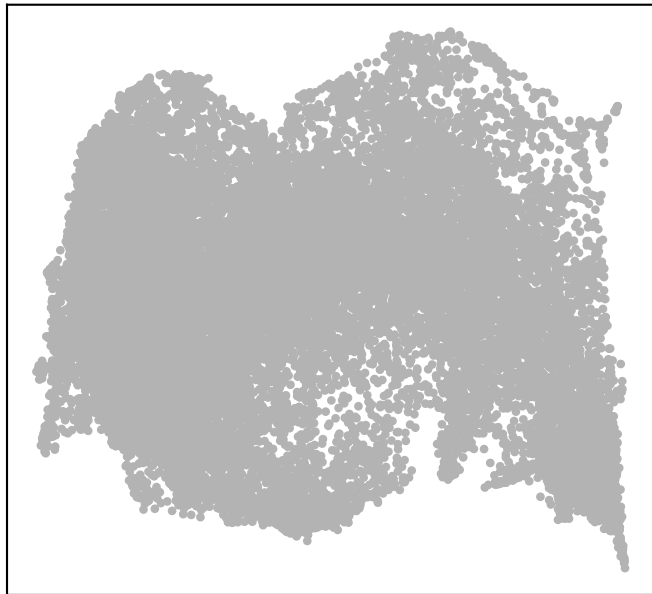


Predicted doublets

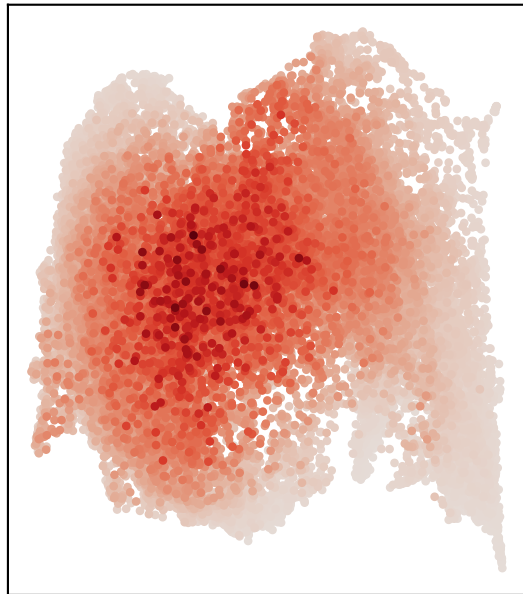
UMAP 2



UMAP 1

UMAP

UMAP 2



UMAP 1

0.3

0.2

0.1