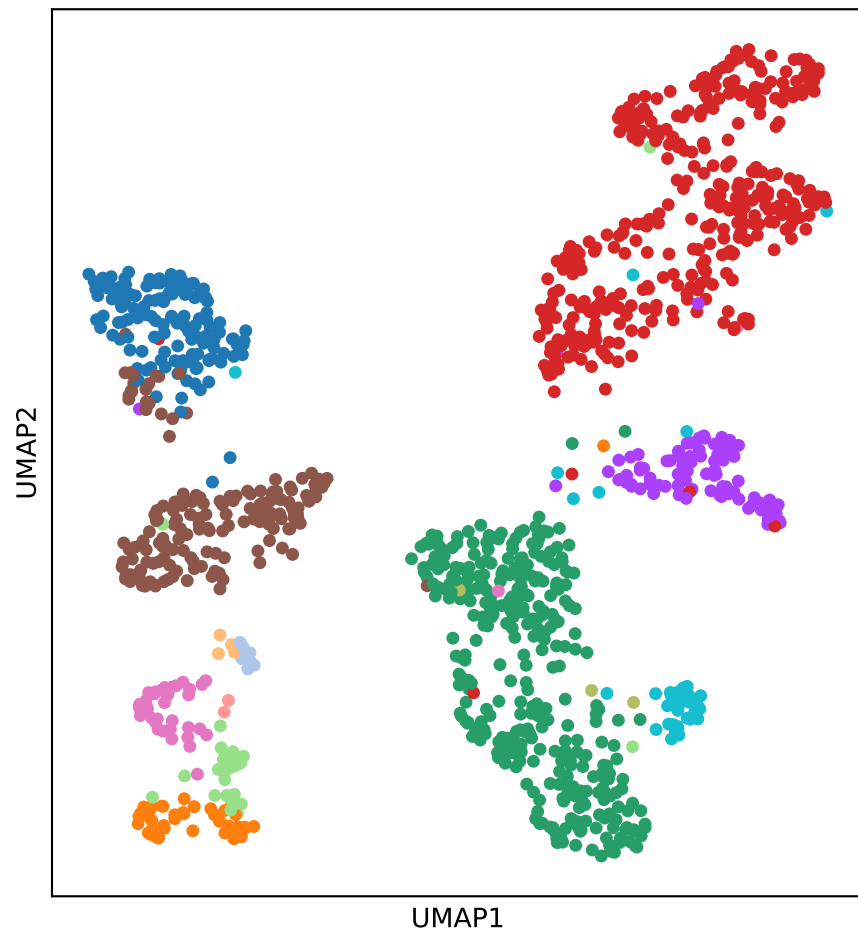
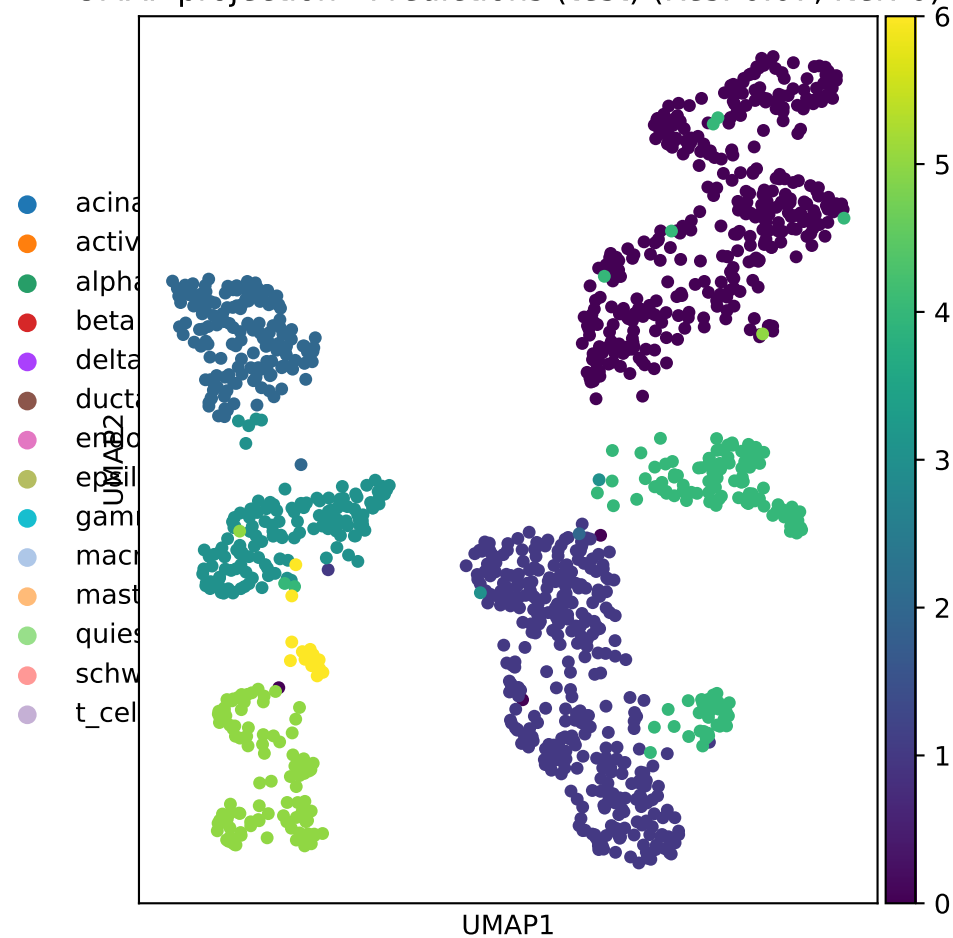


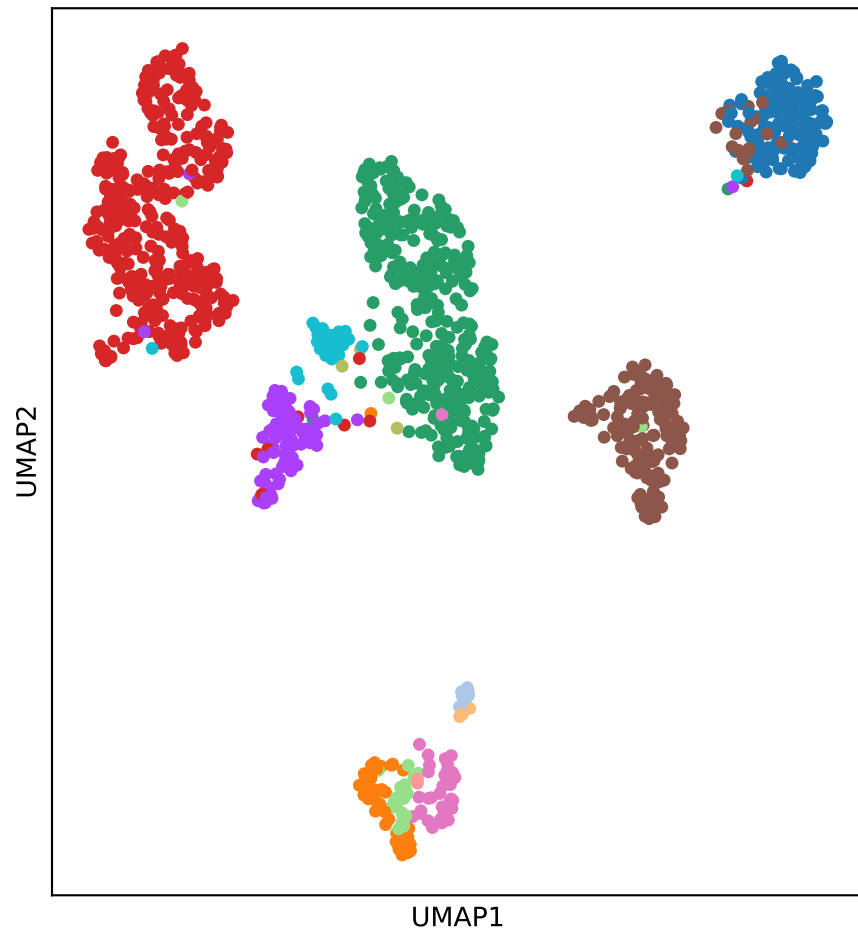
UMAP projection - Labels (test) (Res: 0.07, Iter: 0)



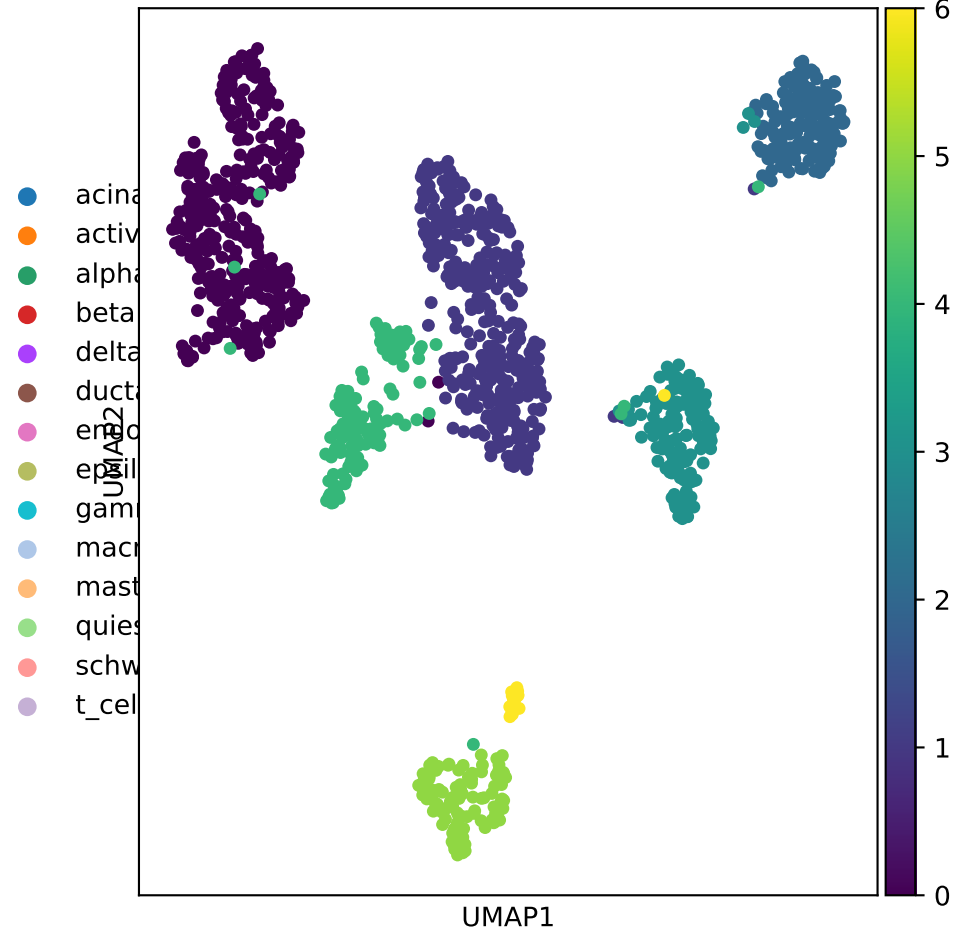
UMAP projection - Predictions (test) (Res: 0.07, Iter: 0)



UMAP projection - Labels (test) (Res: 0.07, Iter: 24)

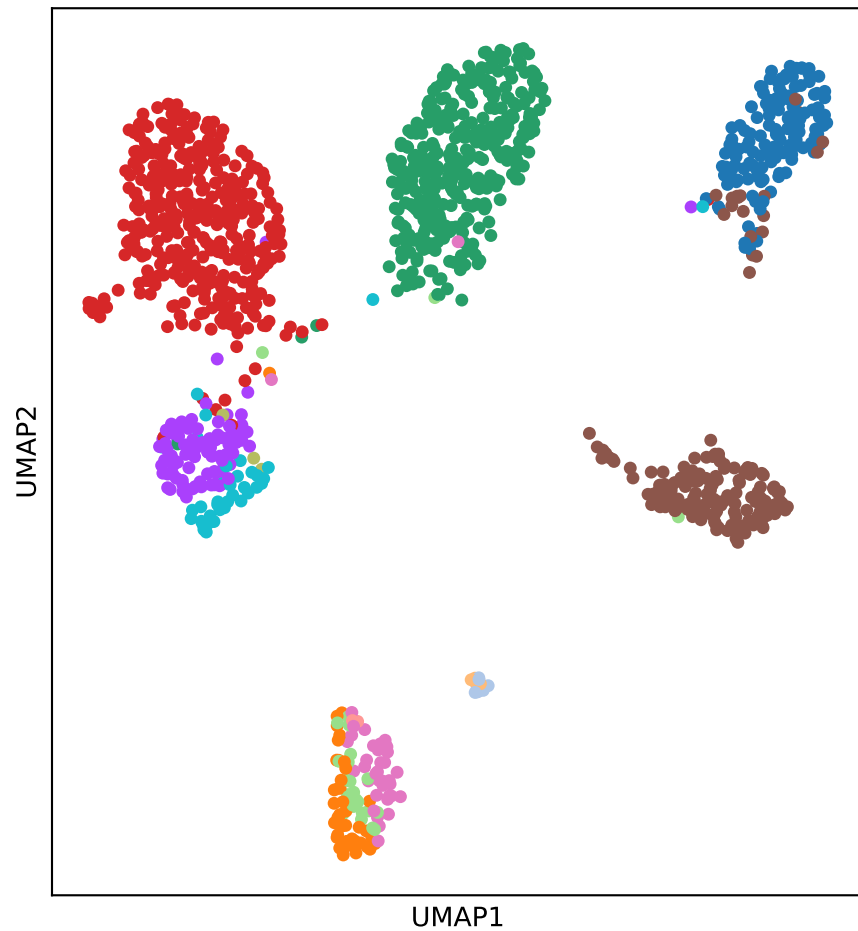


UMAP projection - Predictions (test) (Res: 0.07, Iter: 24)

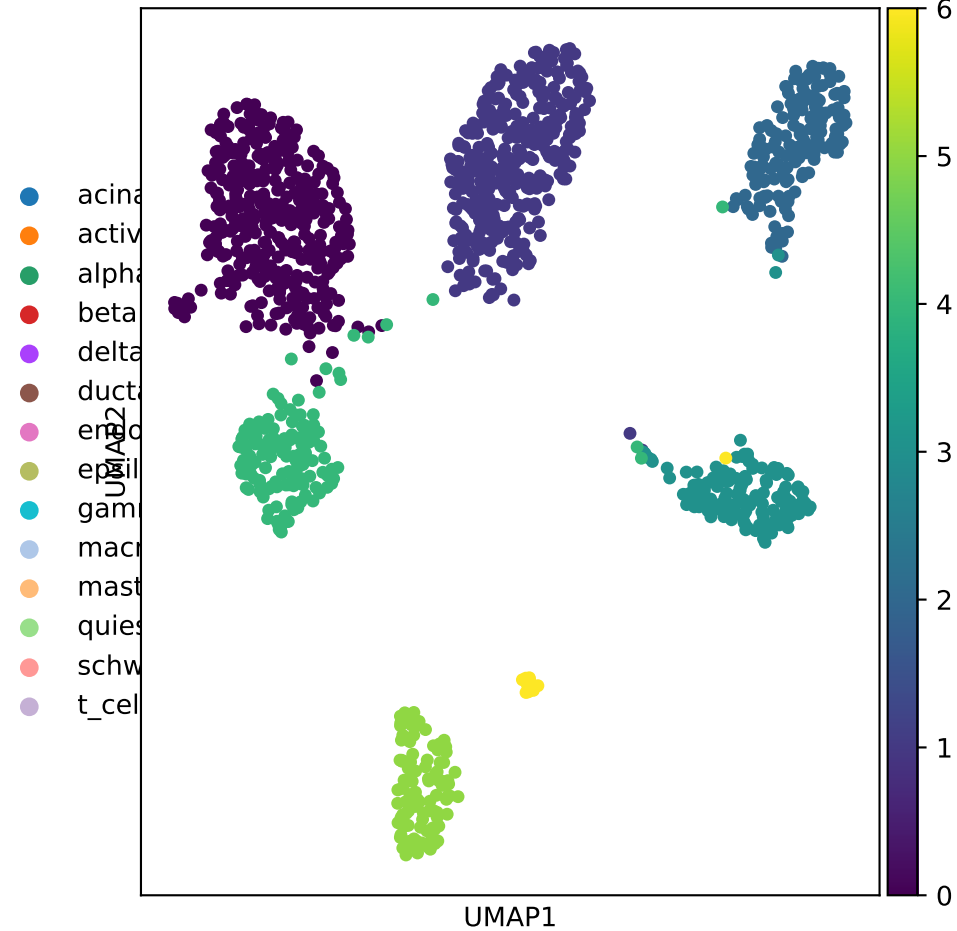


- acina
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- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t\_cel

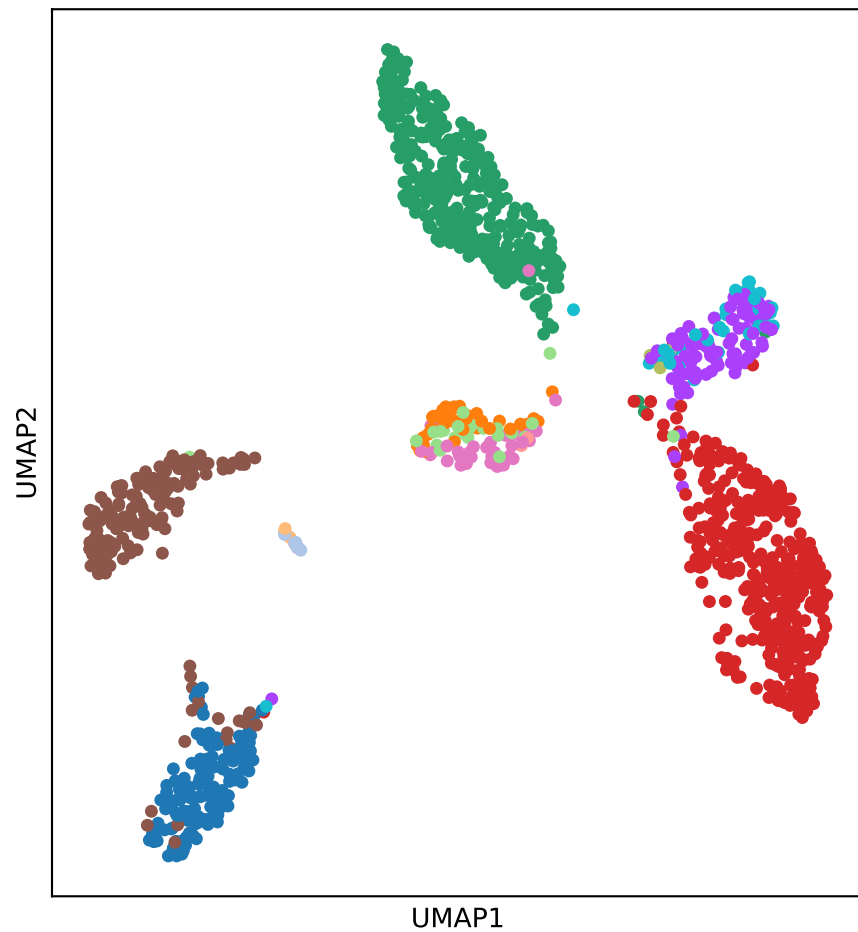
UMAP projection - Labels (test) (Res: 0.07, Iter: 48)



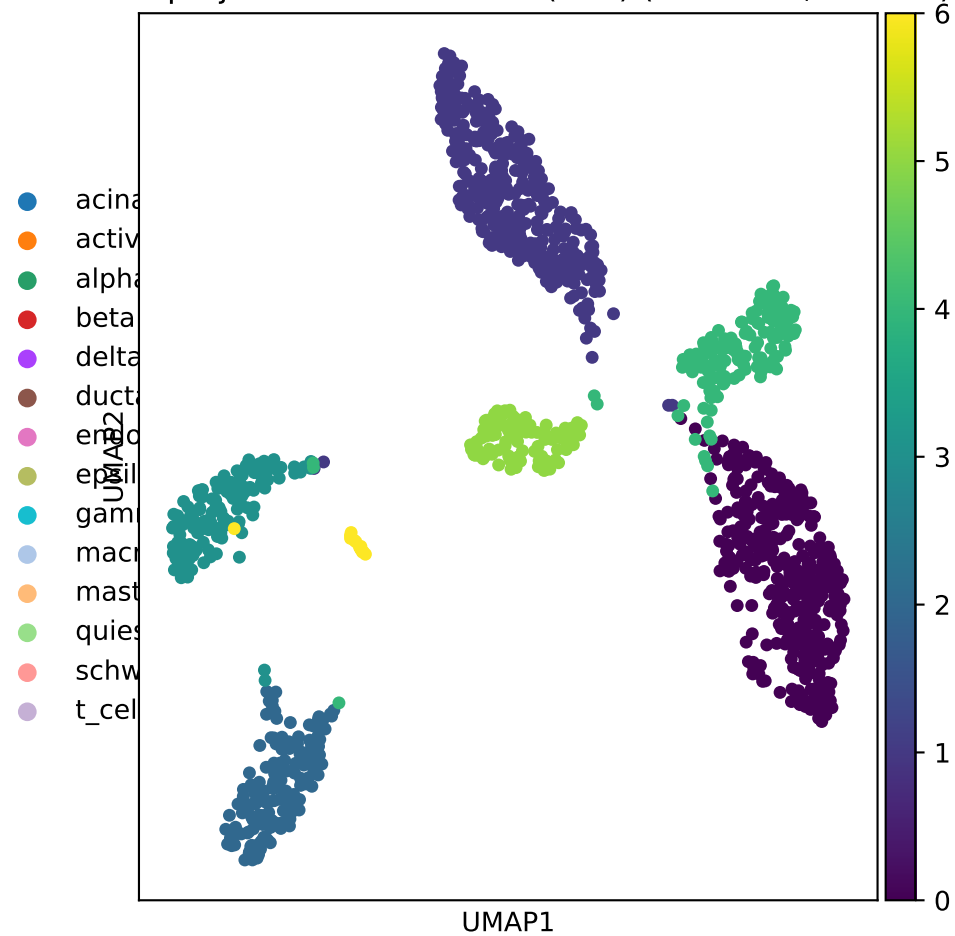
UMAP projection - Predictions (test) (Res: 0.07, Iter: 48)



UMAP projection - Labels (test) (Res: 0.07, Iter: 72)

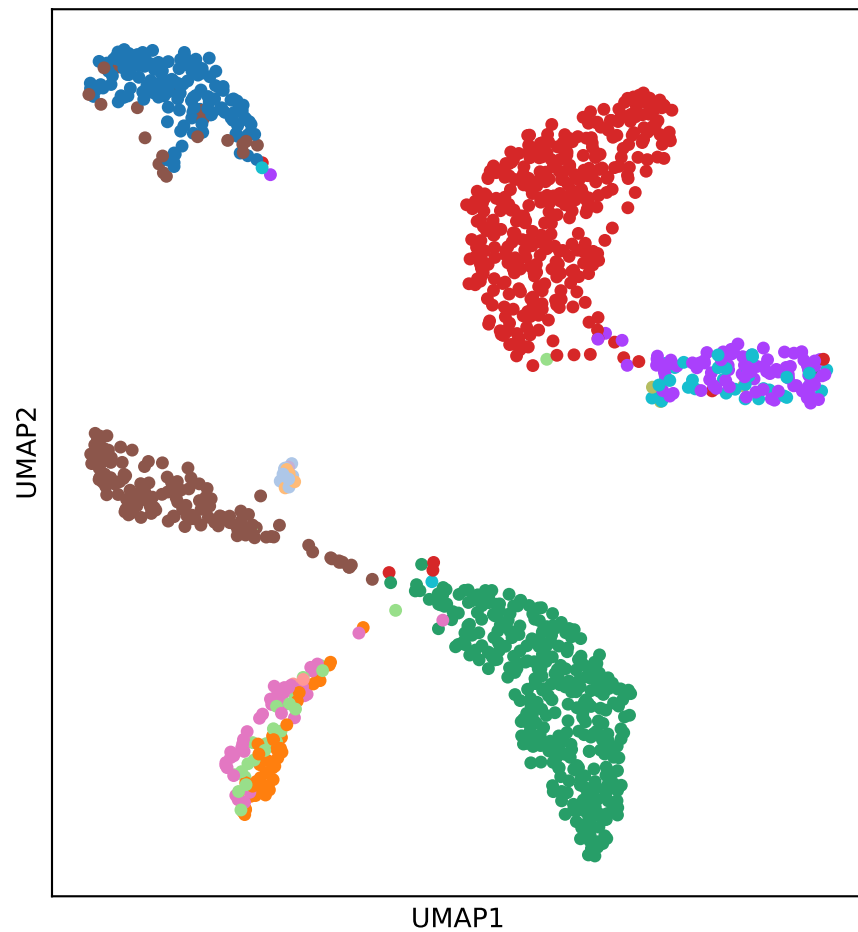


UMAP projection - Predictions (test) (Res: 0.07, Iter: 72)

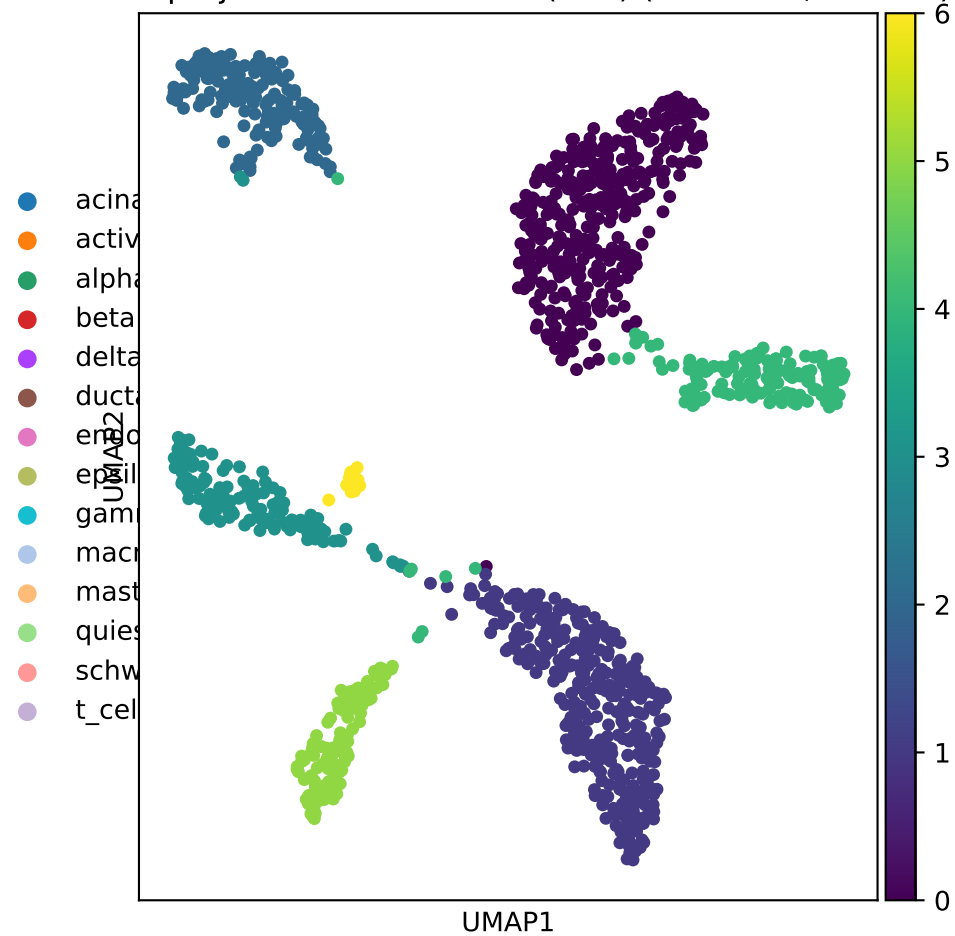


- acinar
- activated
- alpha
- beta
- delta
- ductal
- endocrine
- epithelial
- gamma
- macrophage
- mast cell
- quiescent
- Schwann cell
- T cell

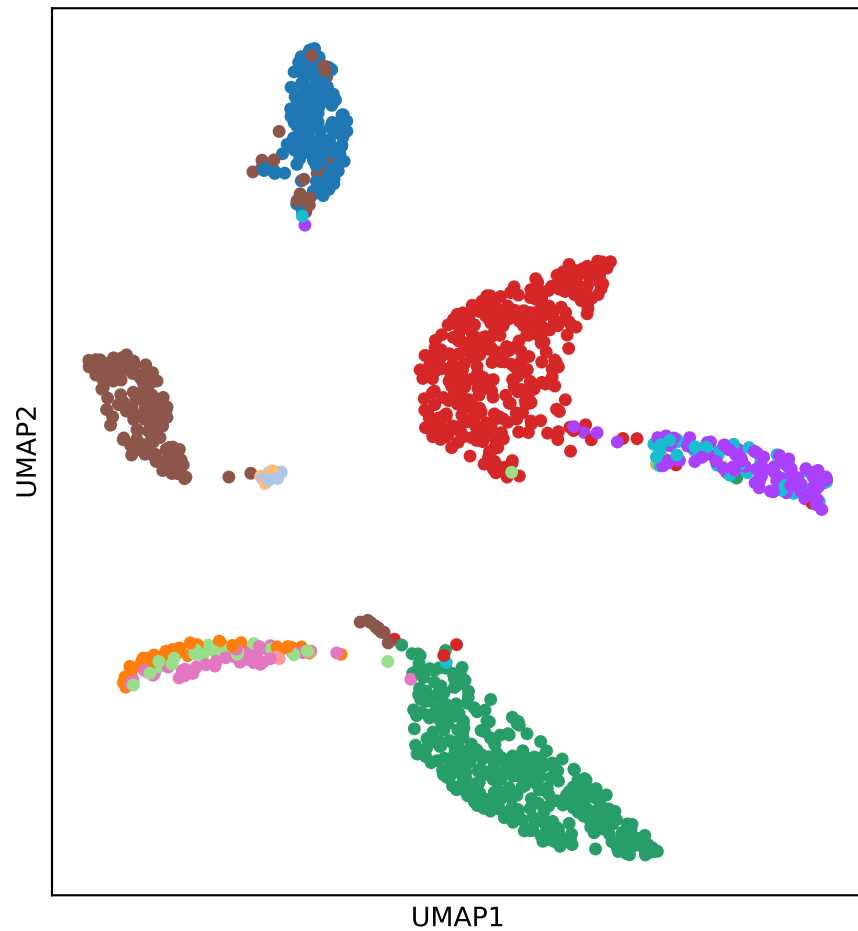
UMAP projection - Labels (test) (Res: 0.07, Iter: 96)



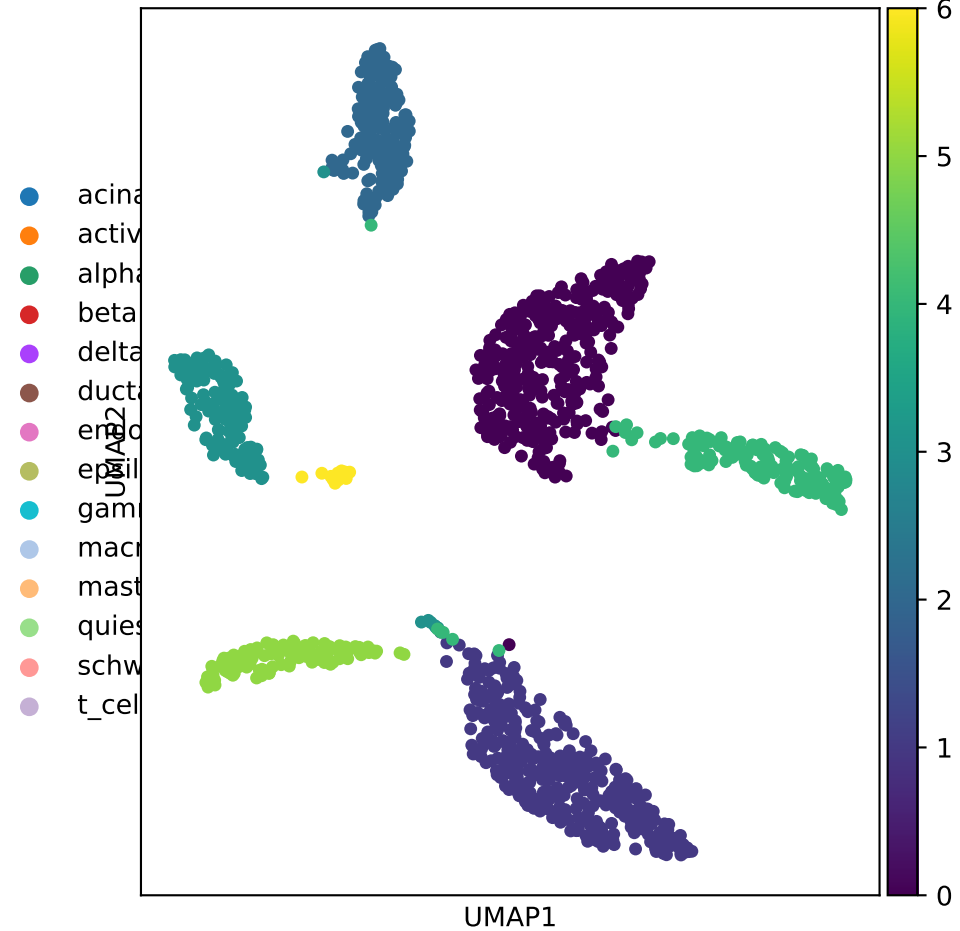
UMAP projection - Predictions (test) (Res: 0.07, Iter: 96)



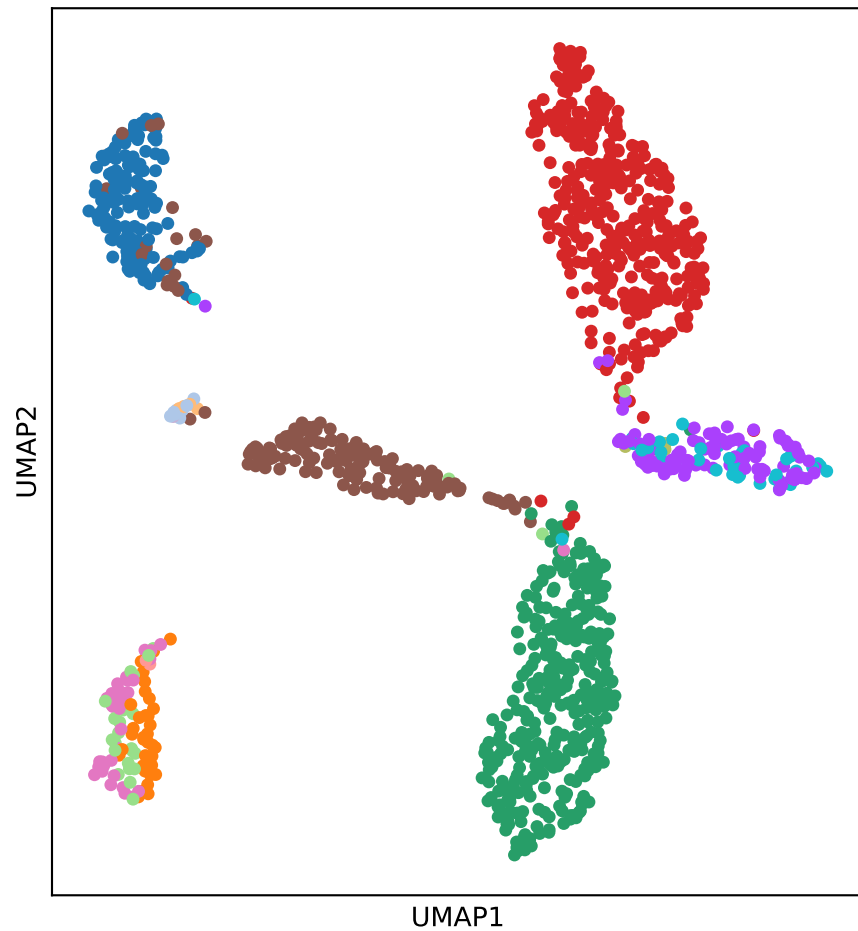
UMAP projection - Labels (test) (Res: 0.07, Iter: 120)



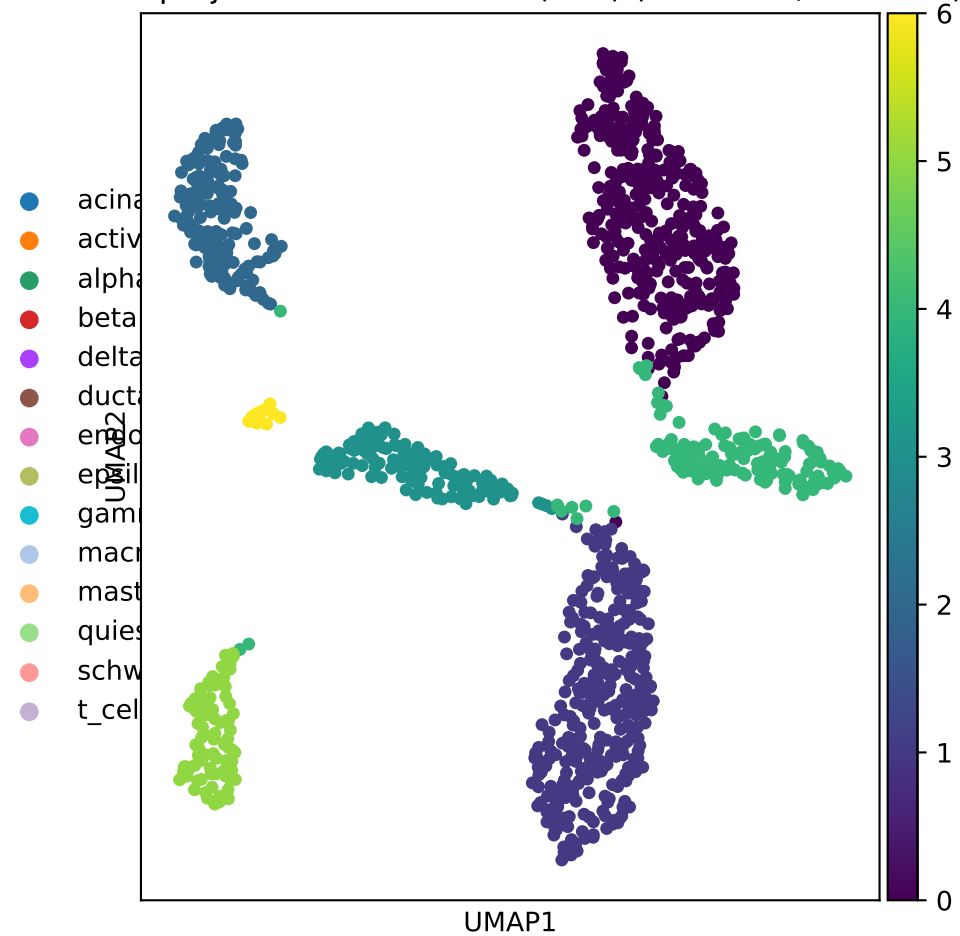
UMAP projection - Predictions (test) (Res: 0.07, Iter: 120)



UMAP projection - Labels (test) (Res: 0.07, Iter: 144)

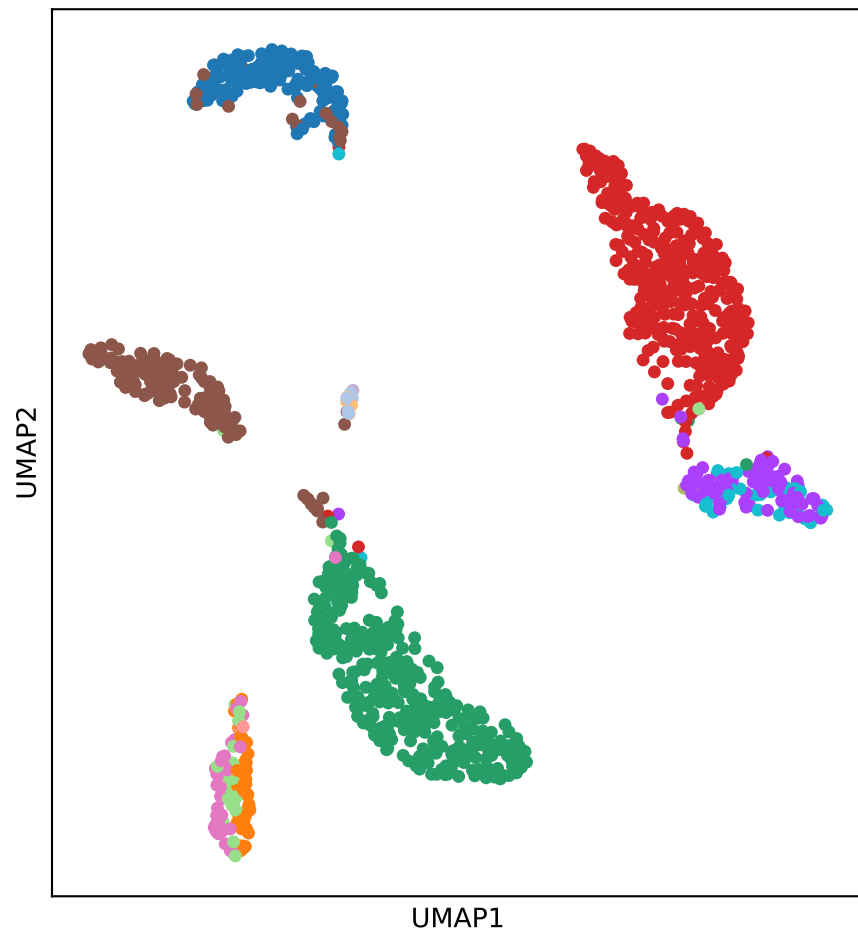


UMAP projection - Predictions (test) (Res: 0.07, Iter: 144)

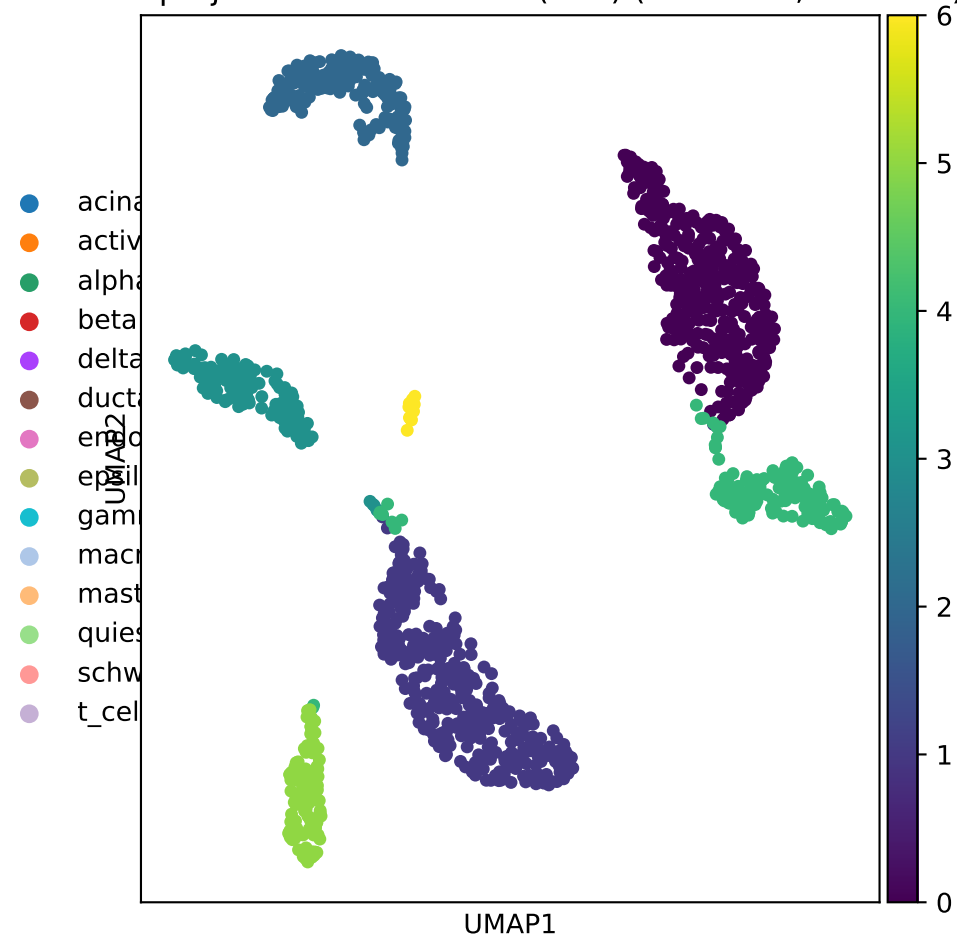


- acina
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- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t\_cel

UMAP projection - Labels (test) (Res: 0.07, Iter: 168)

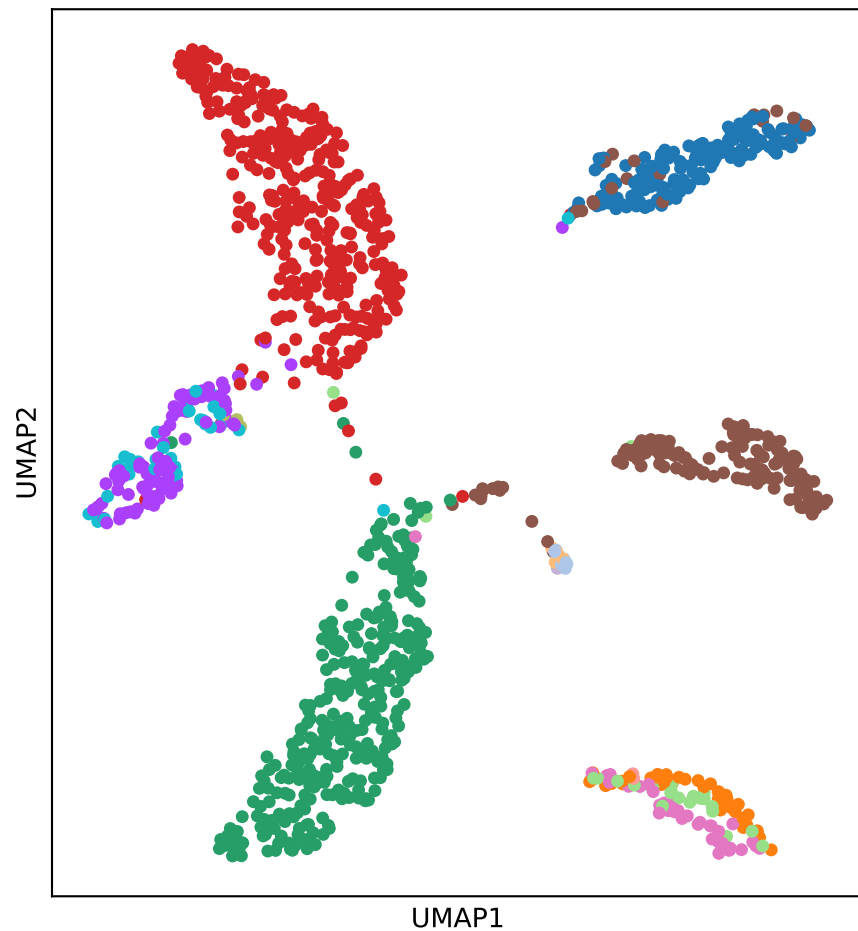


UMAP projection - Predictions (test) (Res: 0.07, Iter: 168)

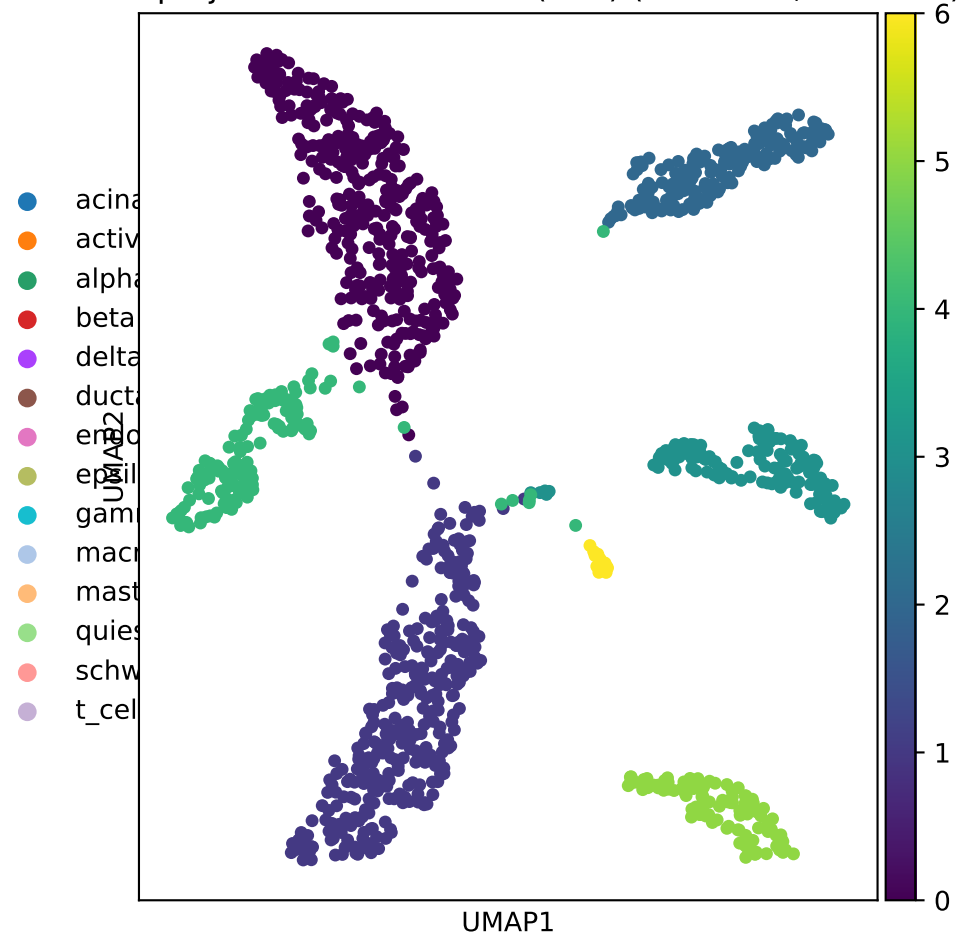




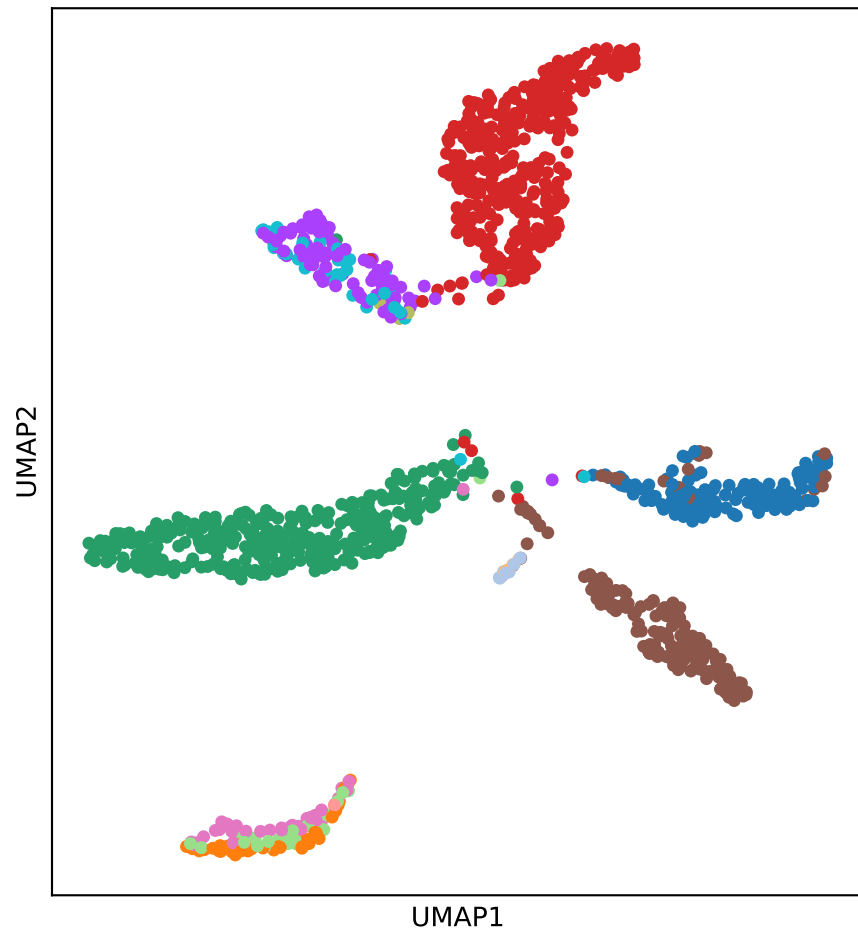
UMAP projection - Labels (test) (Res: 0.07, Iter: 192)



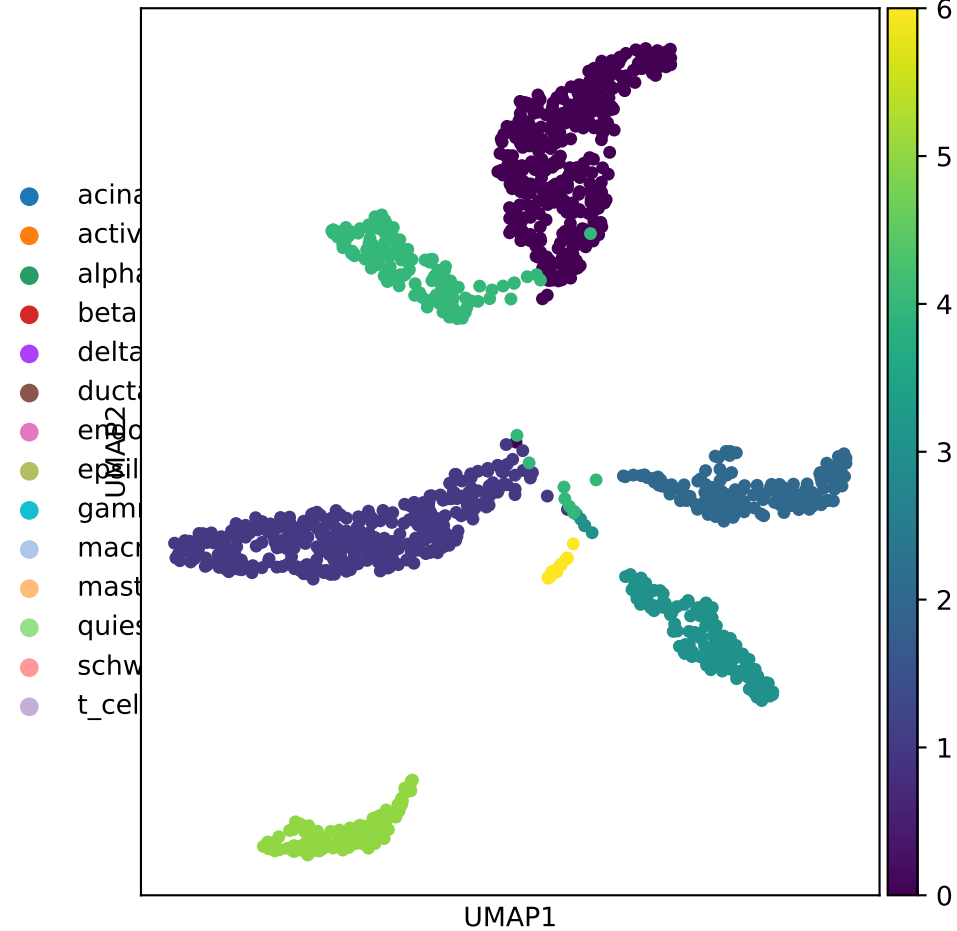
UMAP projection - Predictions (test) (Res: 0.07, Iter: 192)



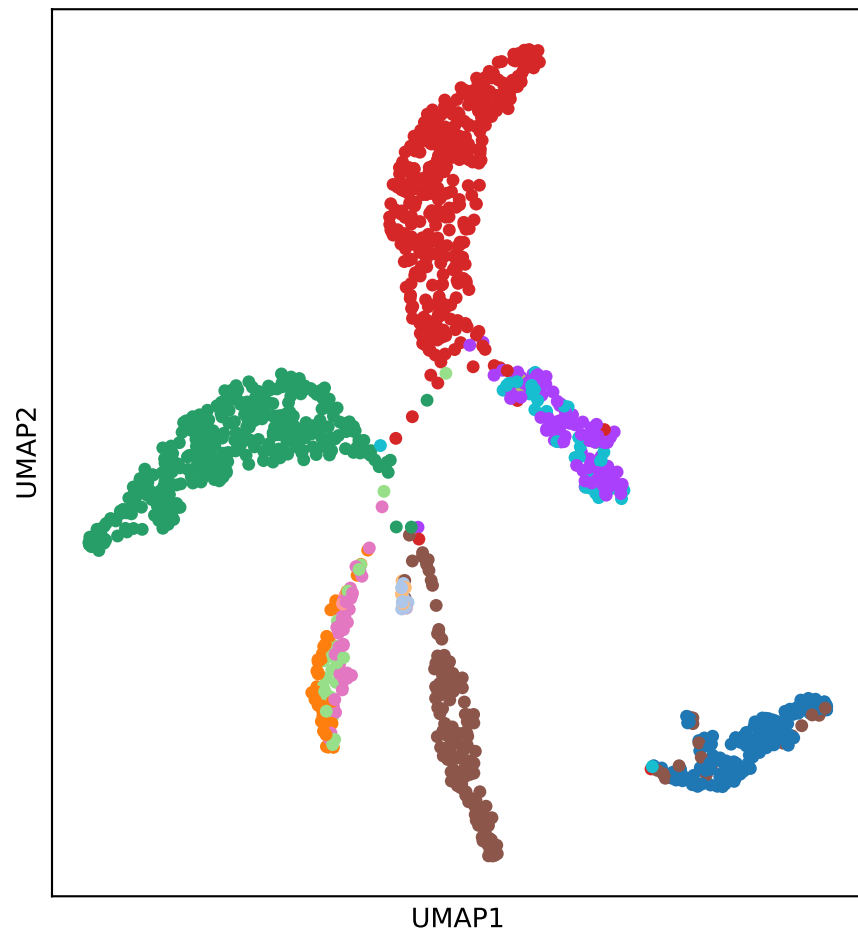
UMAP projection - Labels (test) (Res: 0.07, Iter: 216)



UMAP projection - Predictions (test) (Res: 0.07, Iter: 216)



UMAP projection - Labels (test) (Res: 0.07, Iter: 240)



UMAP projection - Predictions (test) (Res: 0.07, Iter: 240)

