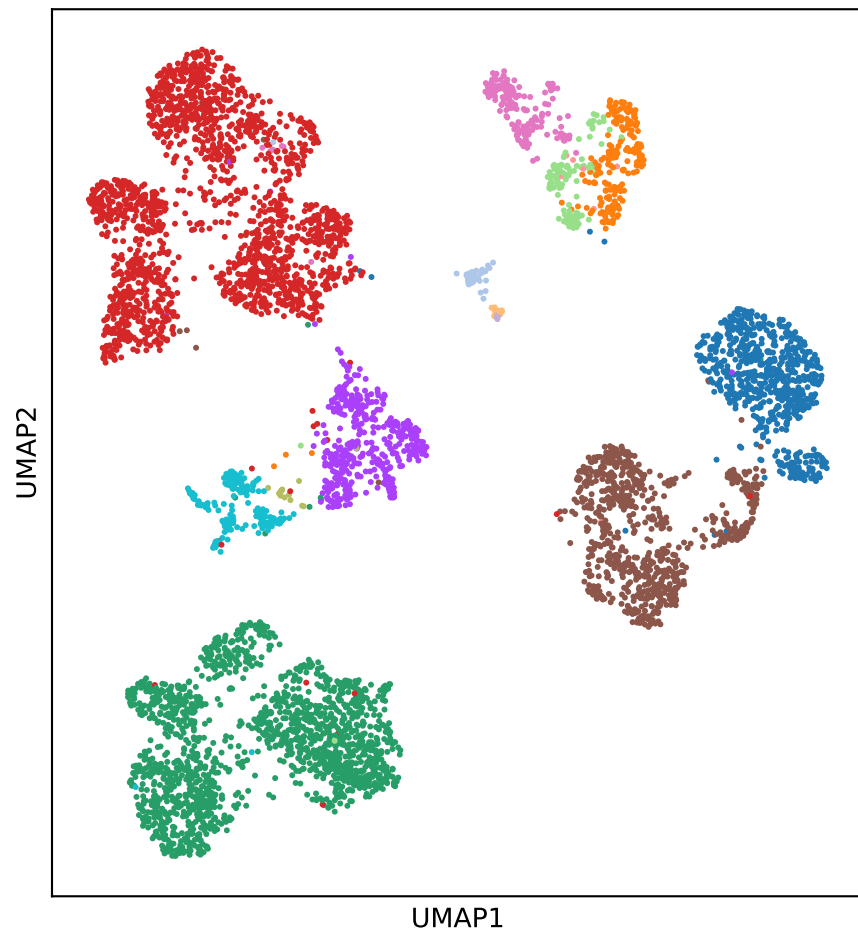
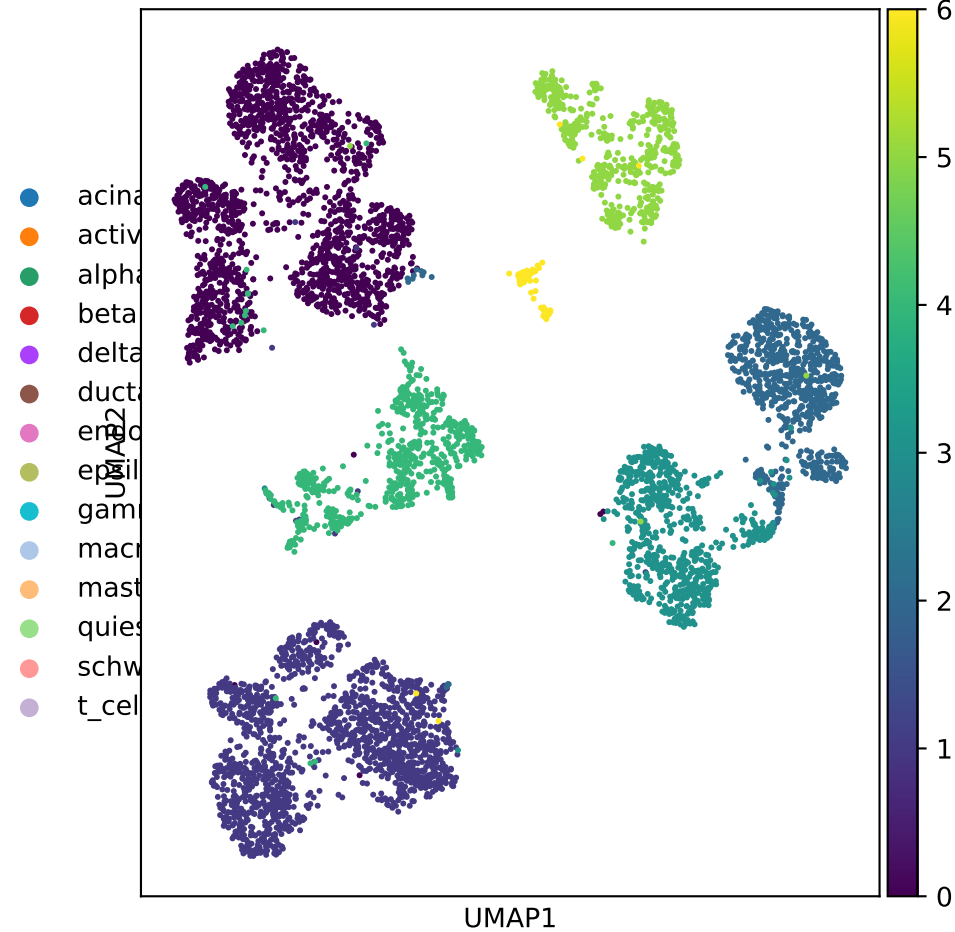


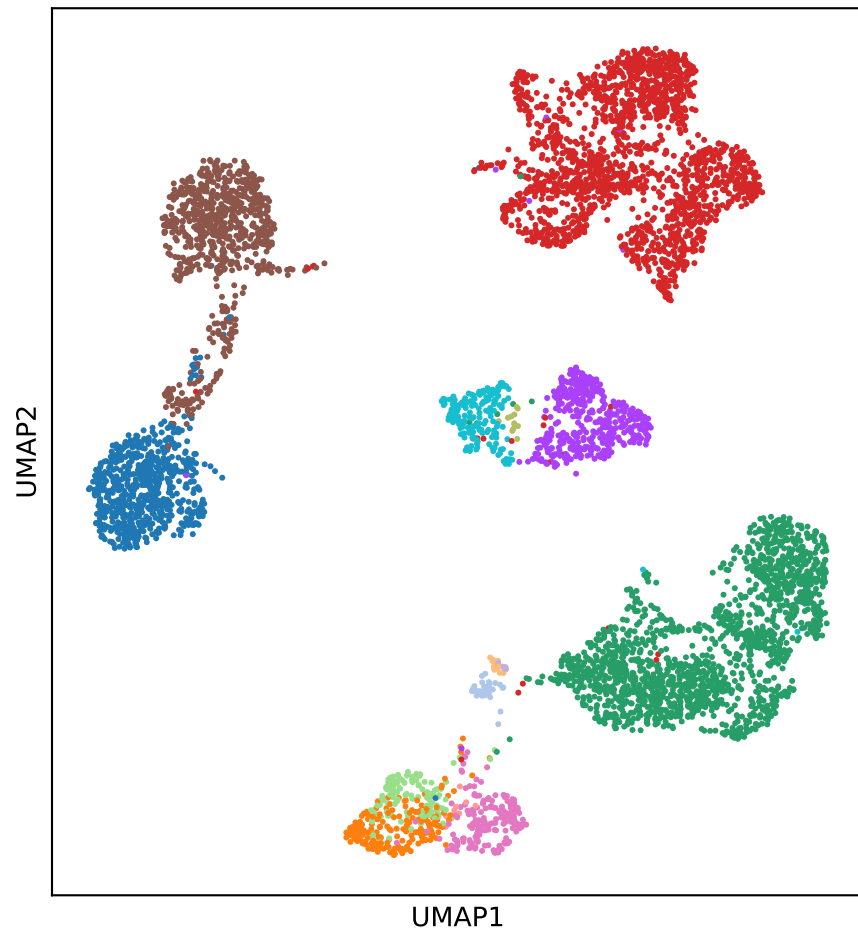
UMAP projection - Labels (train) (Res: 0.06, Iter: 0)



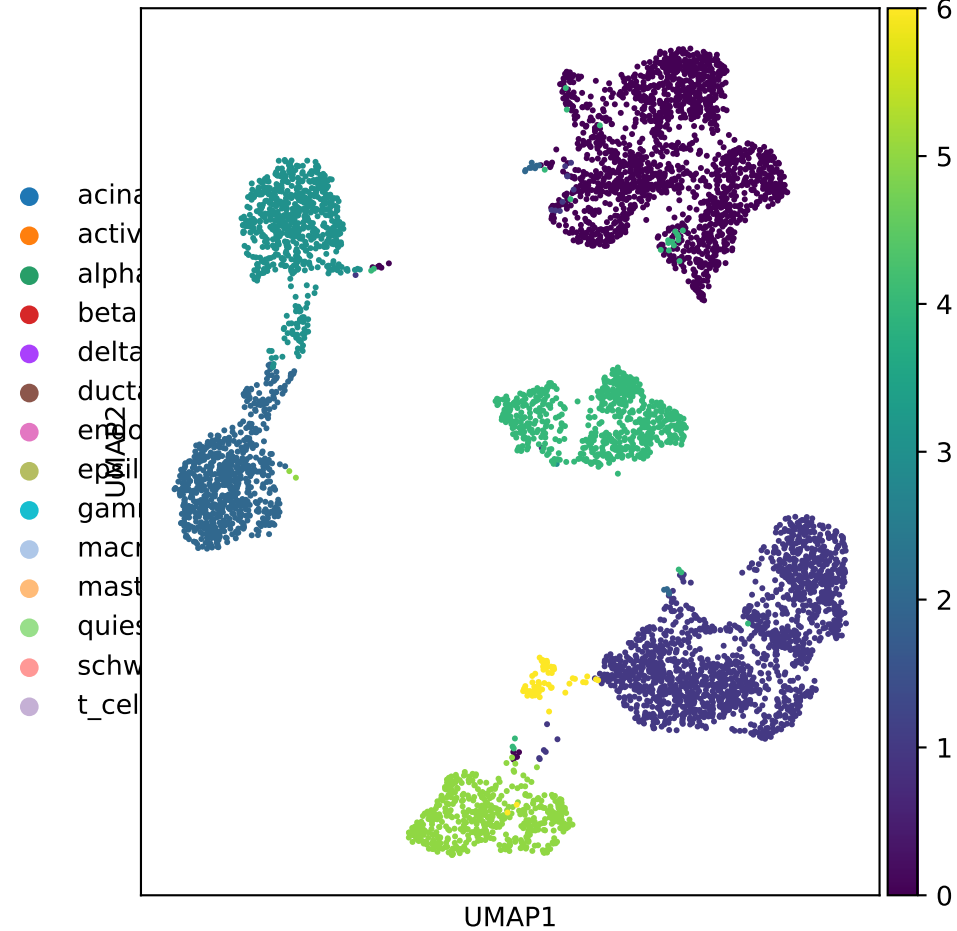
UMAP projection - Predictions (train) (Res: 0.06, Iter: 0)



UMAP projection - Labels (train) (Res: 0.06, Iter: 24)

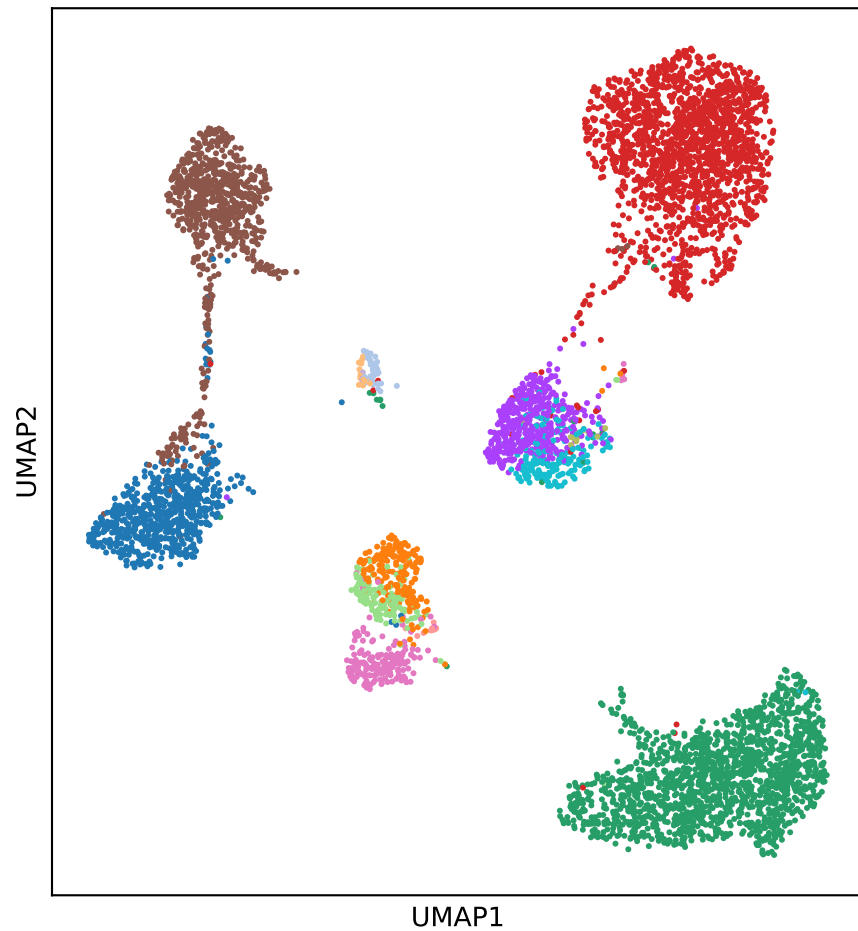


UMAP projection - Predictions (train) (Res: 0.06, Iter: 24)

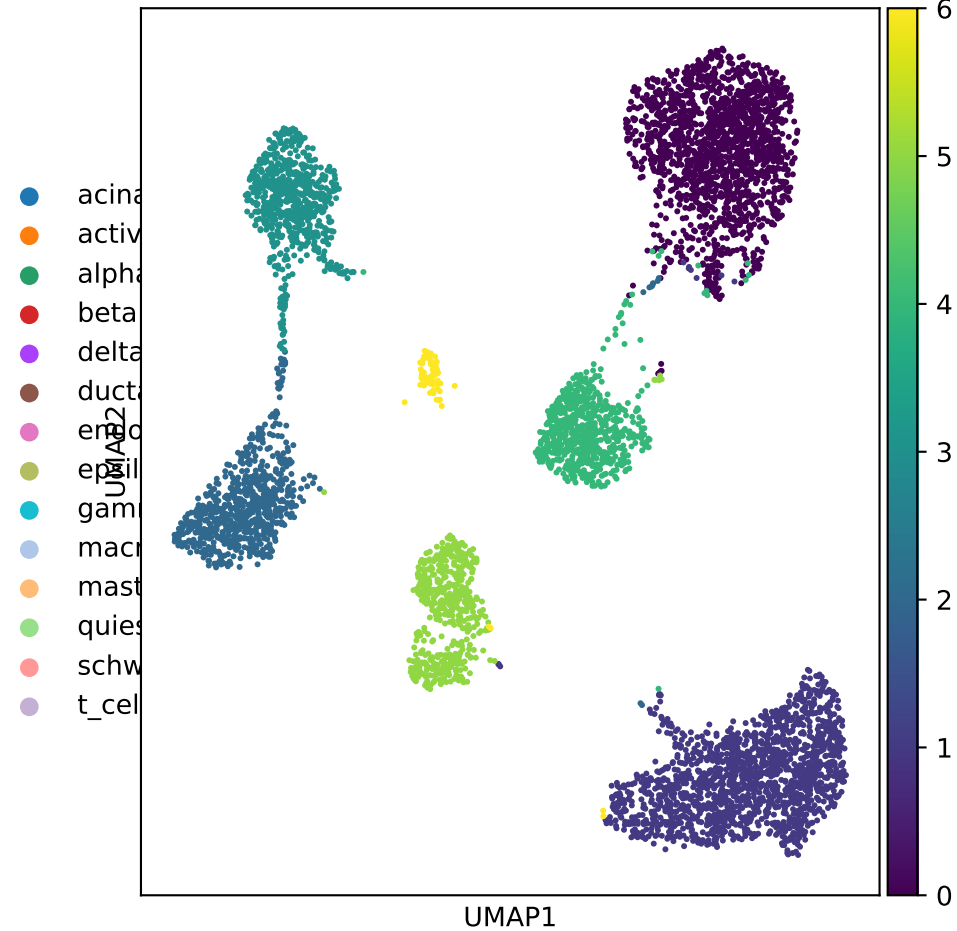


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t\_cel

UMAP projection - Labels (train) (Res: 0.06, Iter: 48)

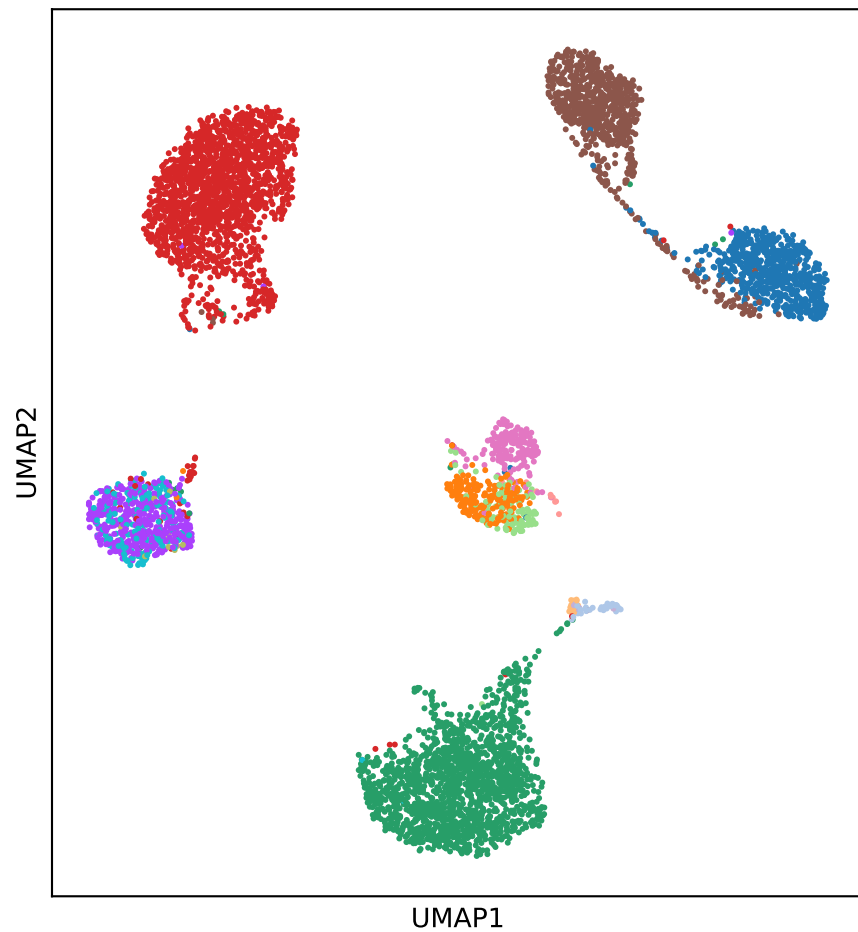


UMAP projection - Predictions (train) (Res: 0.06, Iter: 48)

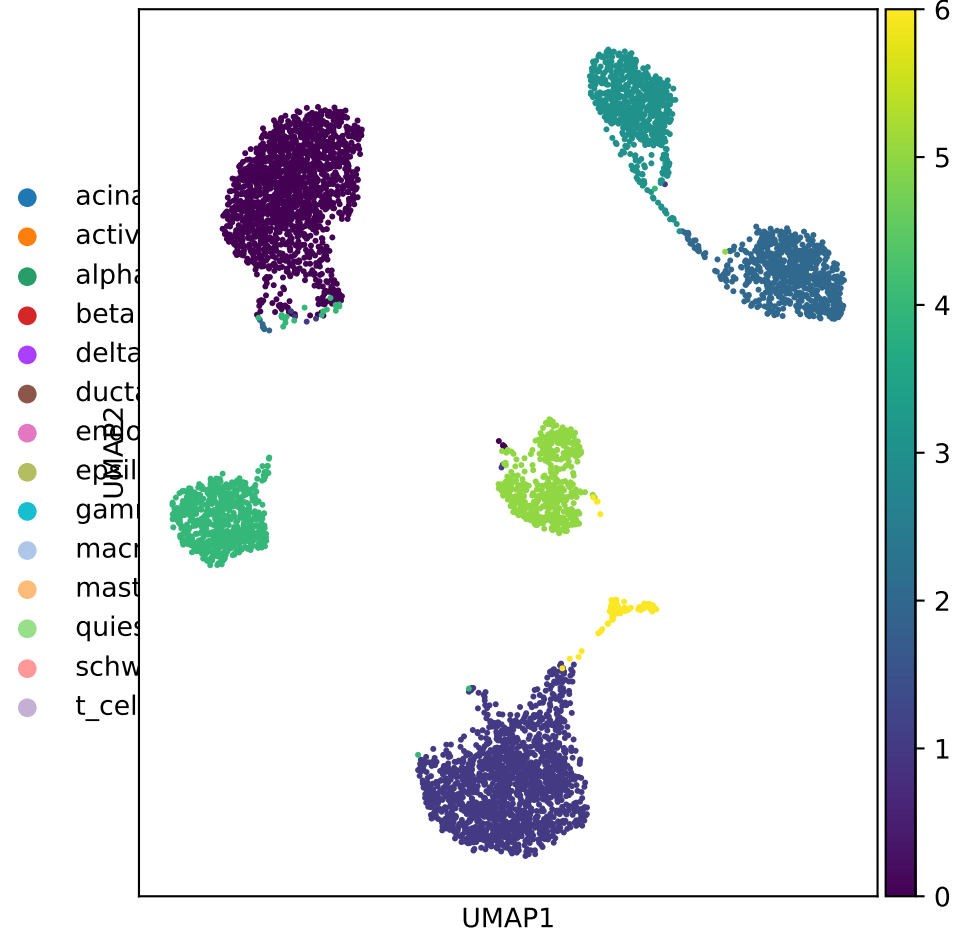


- acinar
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- alpha
- beta
- delta
- ductal
- endocrine
- epithelial
- gamma
- macrophage
- mast cell
- quiescent
- schwann
- t\_cell

UMAP projection - Labels (train) (Res: 0.06, Iter: 72)

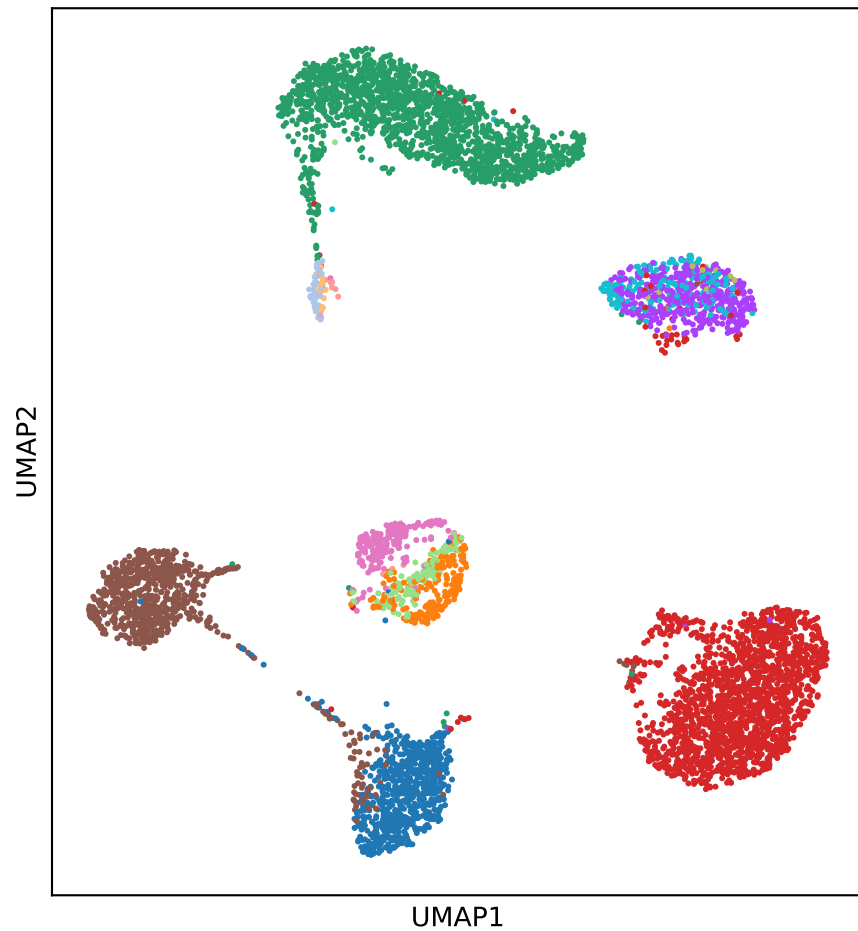


UMAP projection - Predictions (train) (Res: 0.06, Iter: 72)

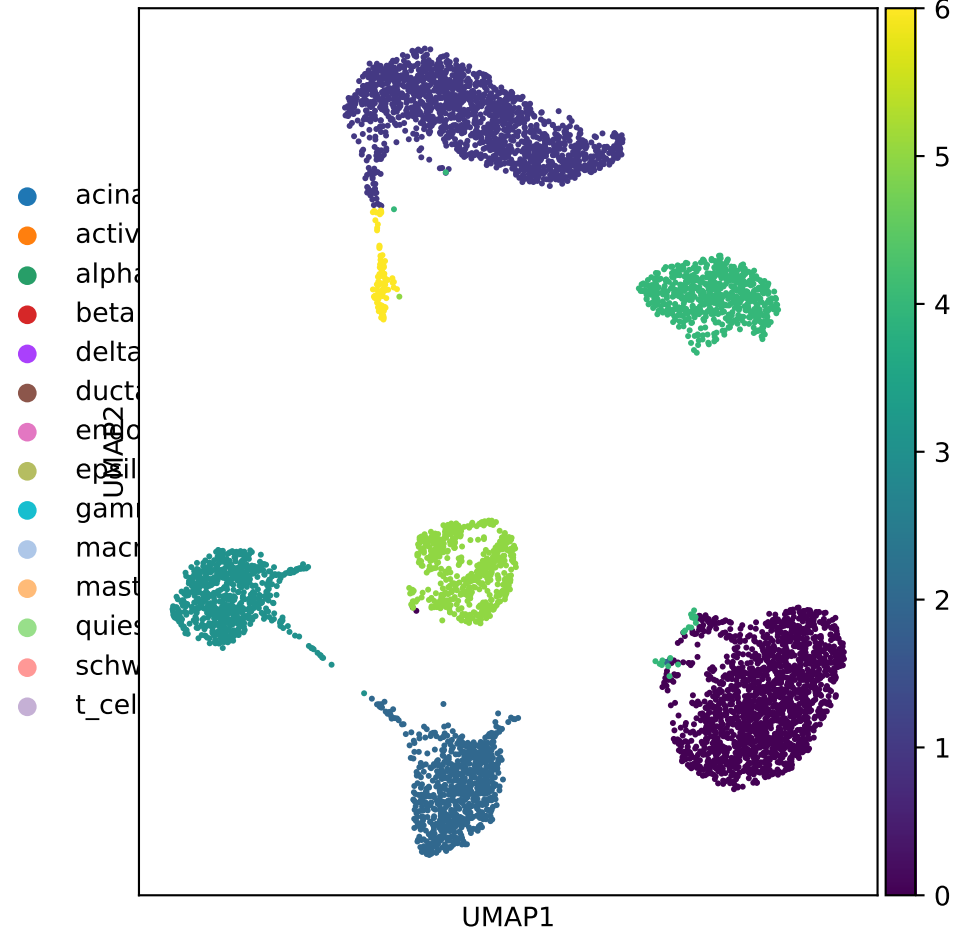


- acina
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- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t\_cel

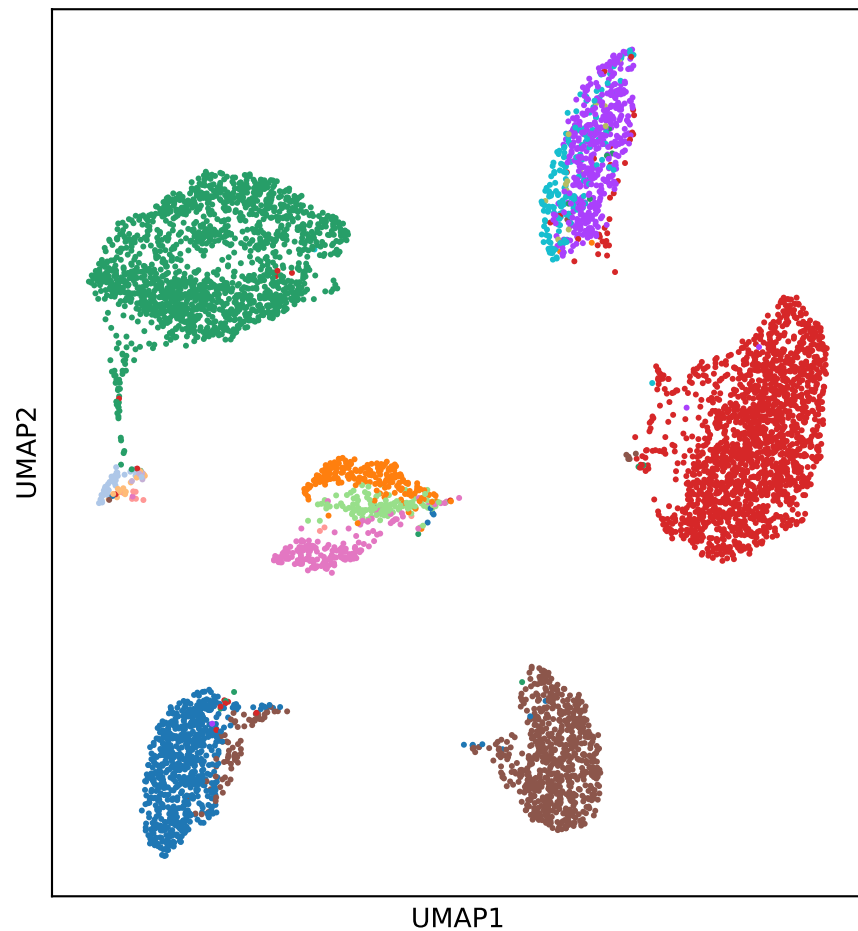
UMAP projection - Labels (train) (Res: 0.06, Iter: 96)



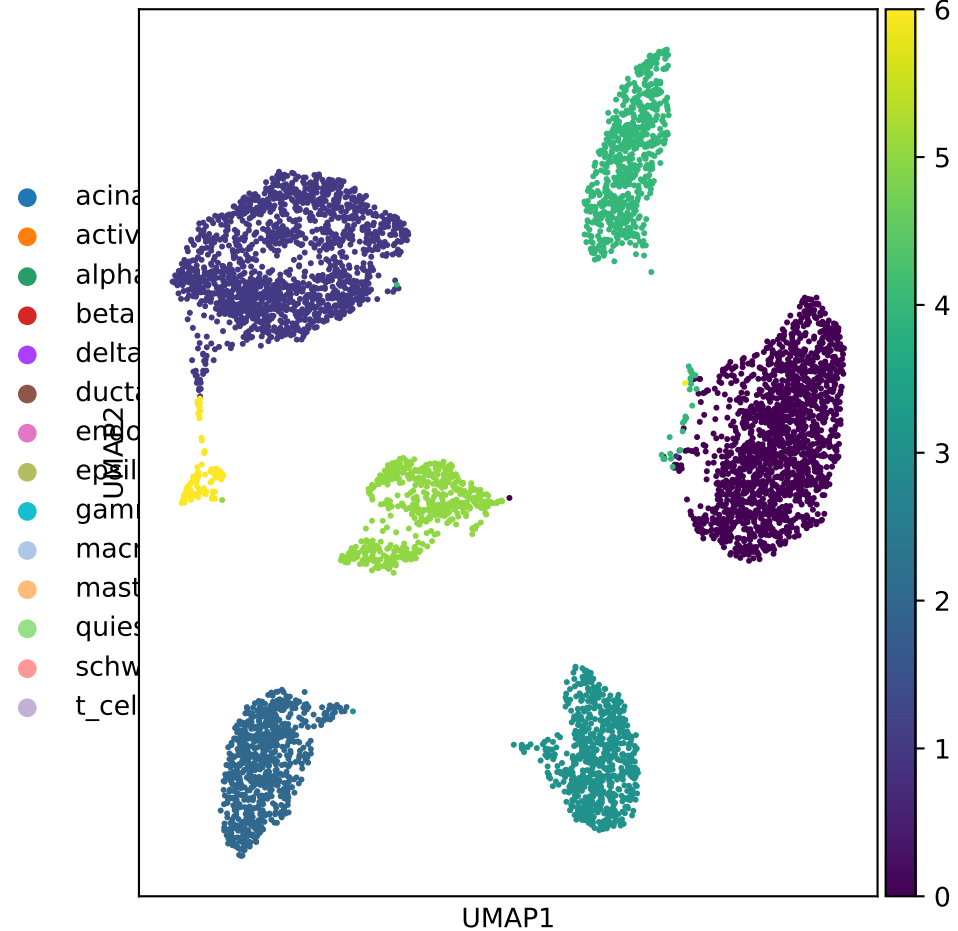
UMAP projection - Predictions (train) (Res: 0.06, Iter: 96)



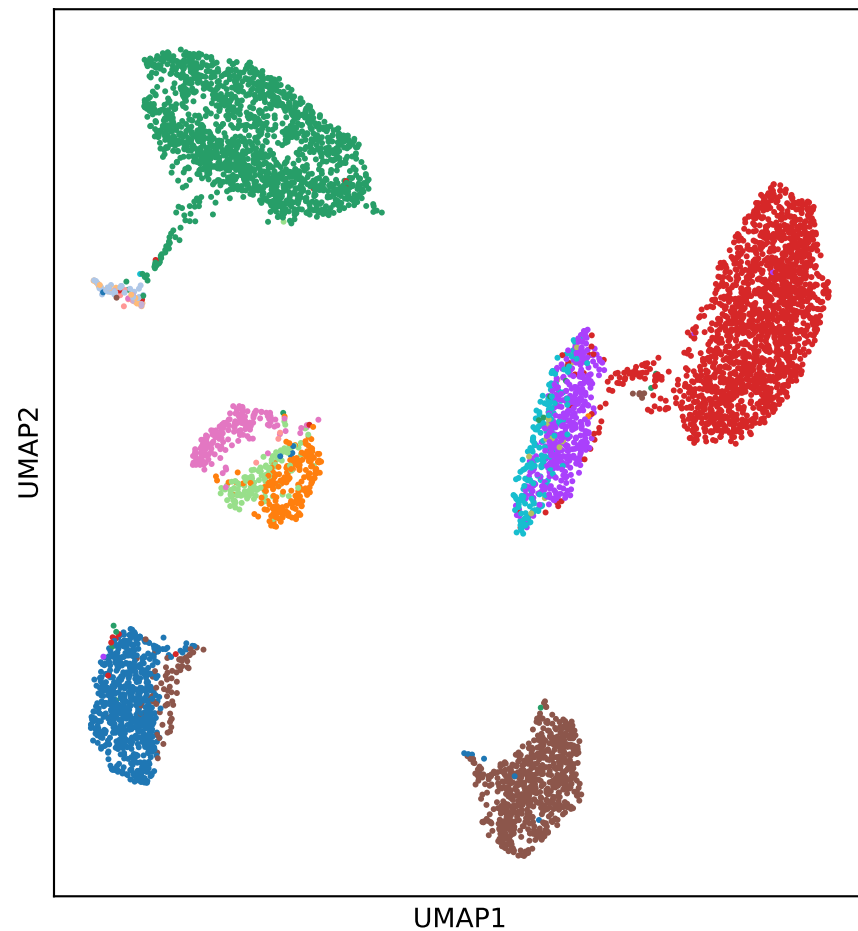
UMAP projection - Labels (train) (Res: 0.06, Iter: 120)



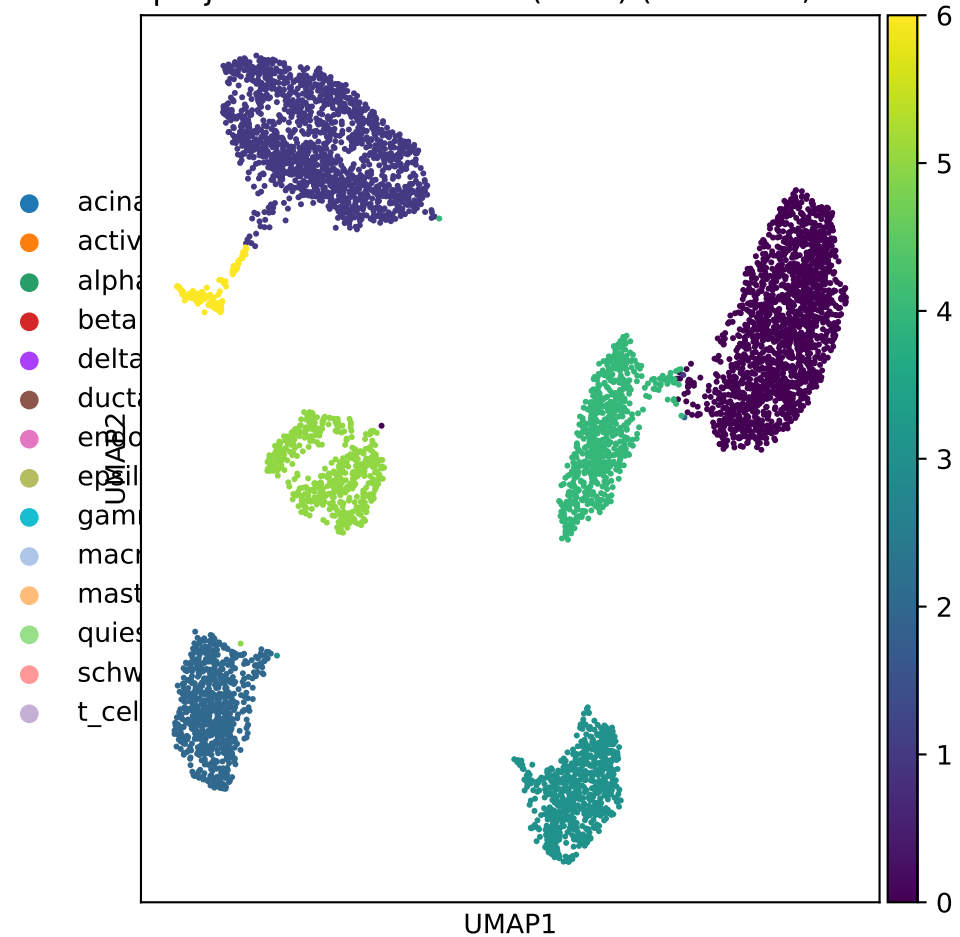
UMAP projection - Predictions (train) (Res: 0.06, Iter: 120)



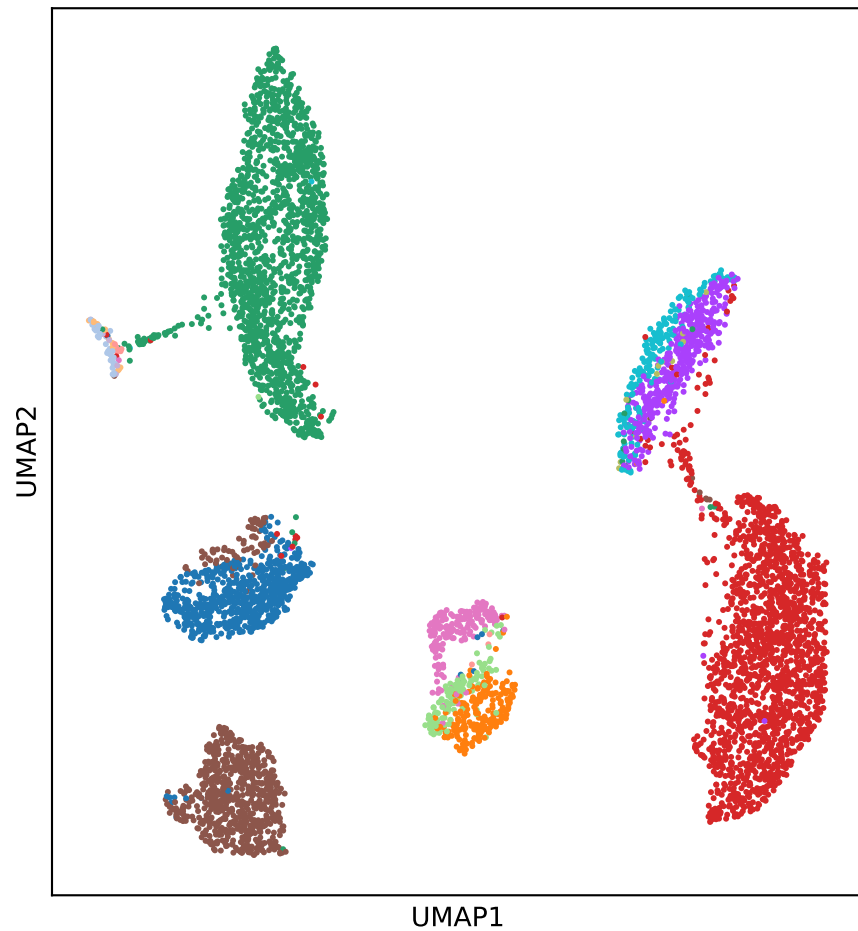
UMAP projection - Labels (train) (Res: 0.06, Iter: 144)



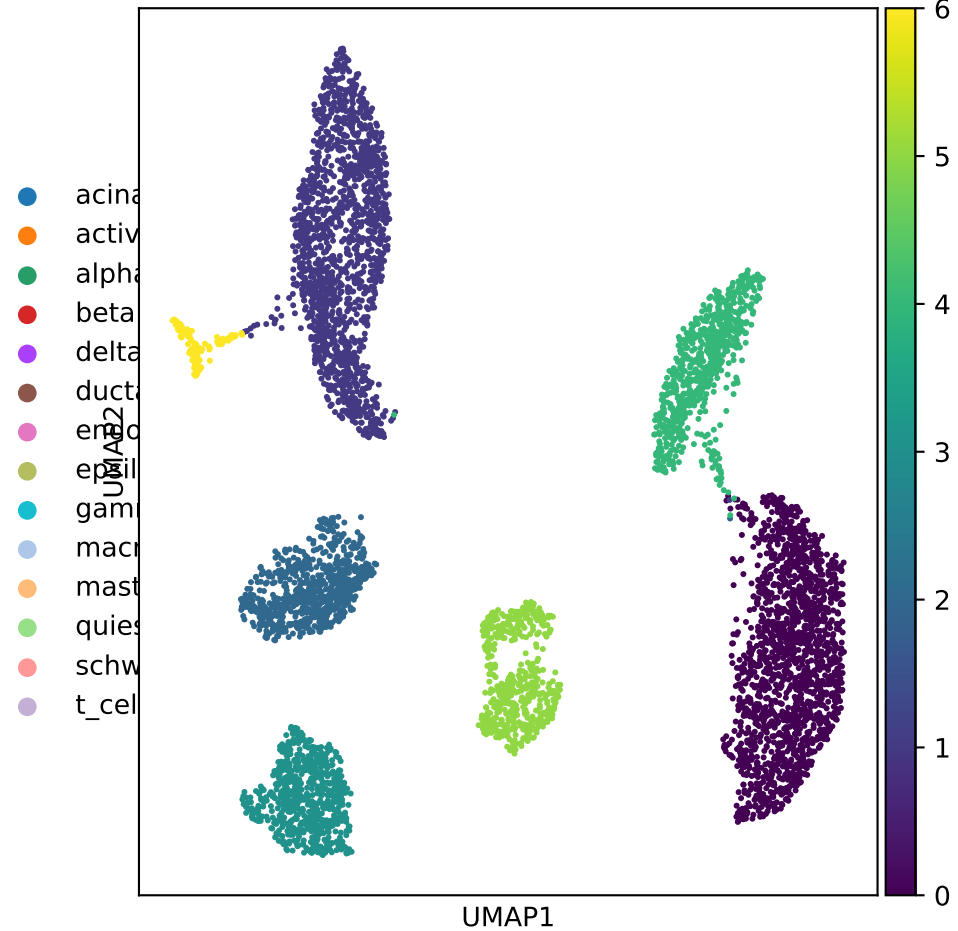
UMAP projection - Predictions (train) (Res: 0.06, Iter: 144)



UMAP projection - Labels (train) (Res: 0.06, Iter: 168)



UMAP projection - Predictions (train) (Res: 0.06, Iter: 168)



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