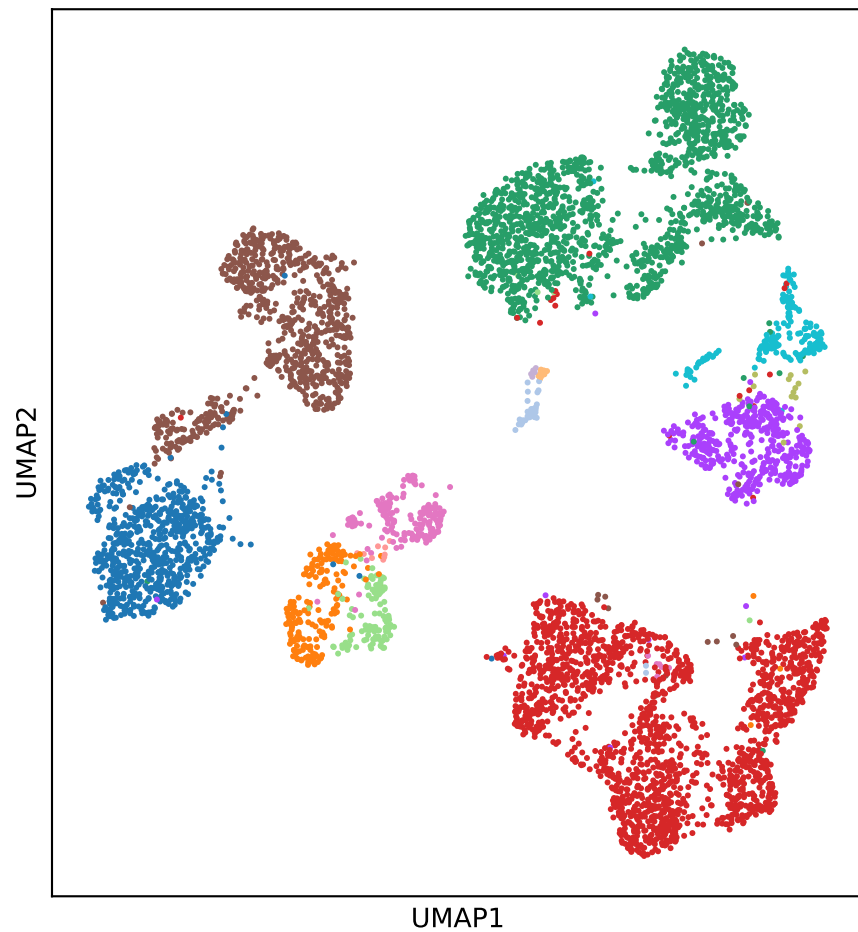
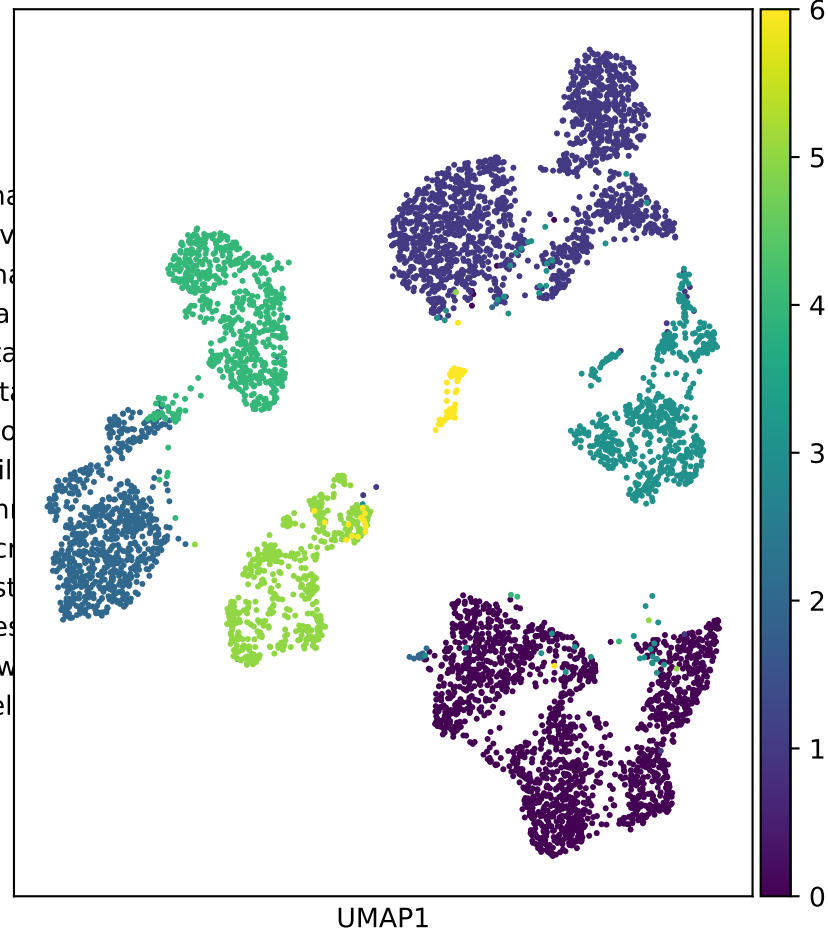


UMAP projection - Labels (train) (Res: 0.06, Iter: 0)

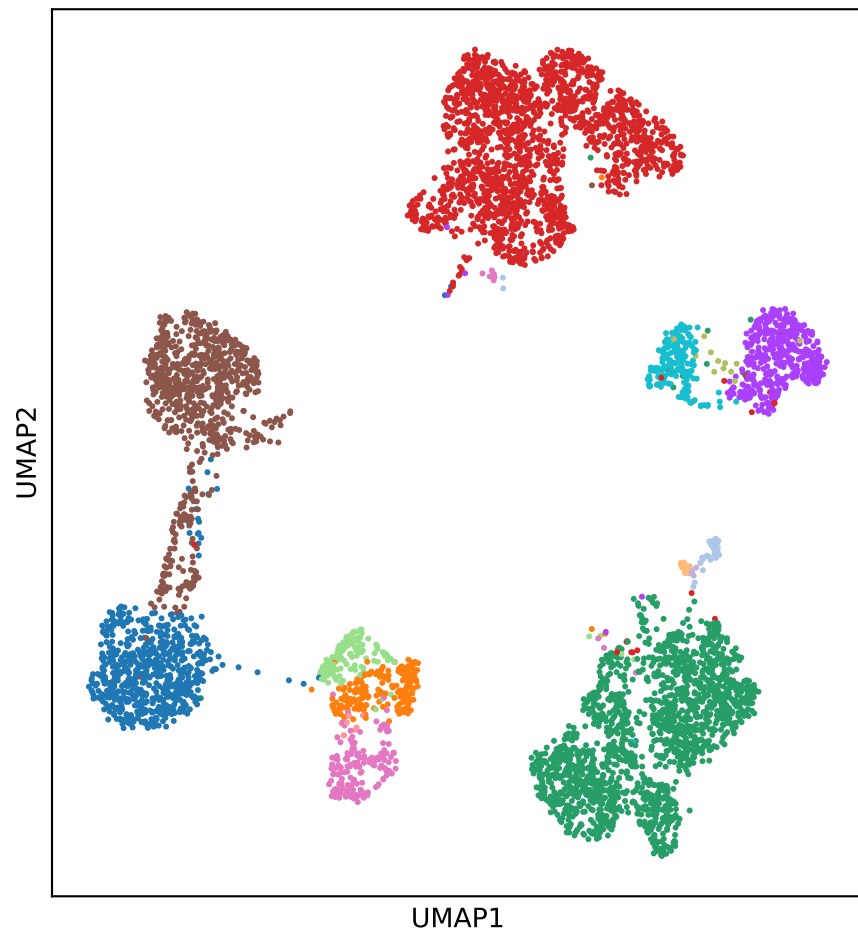


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epill
- gam
- macr
- mast
- quies
- schw
- t_cel

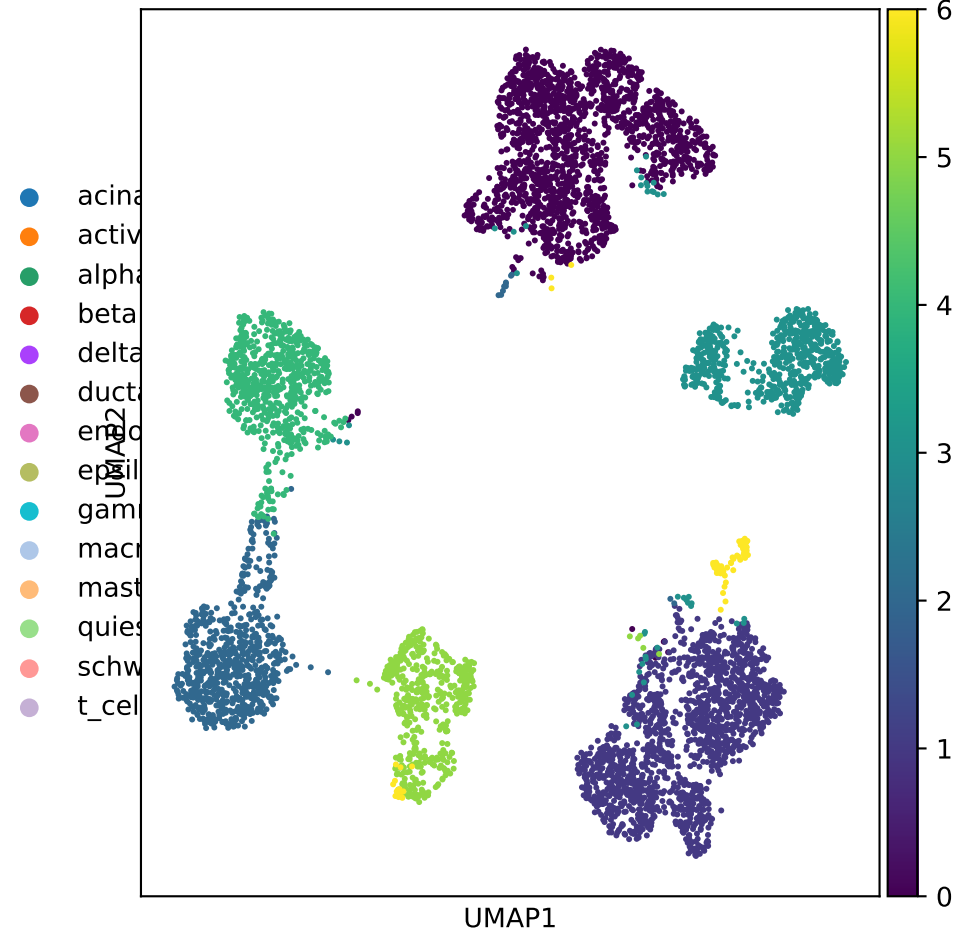
UMAP projection - Predictions (train) (Res: 0.06, Iter: 0)



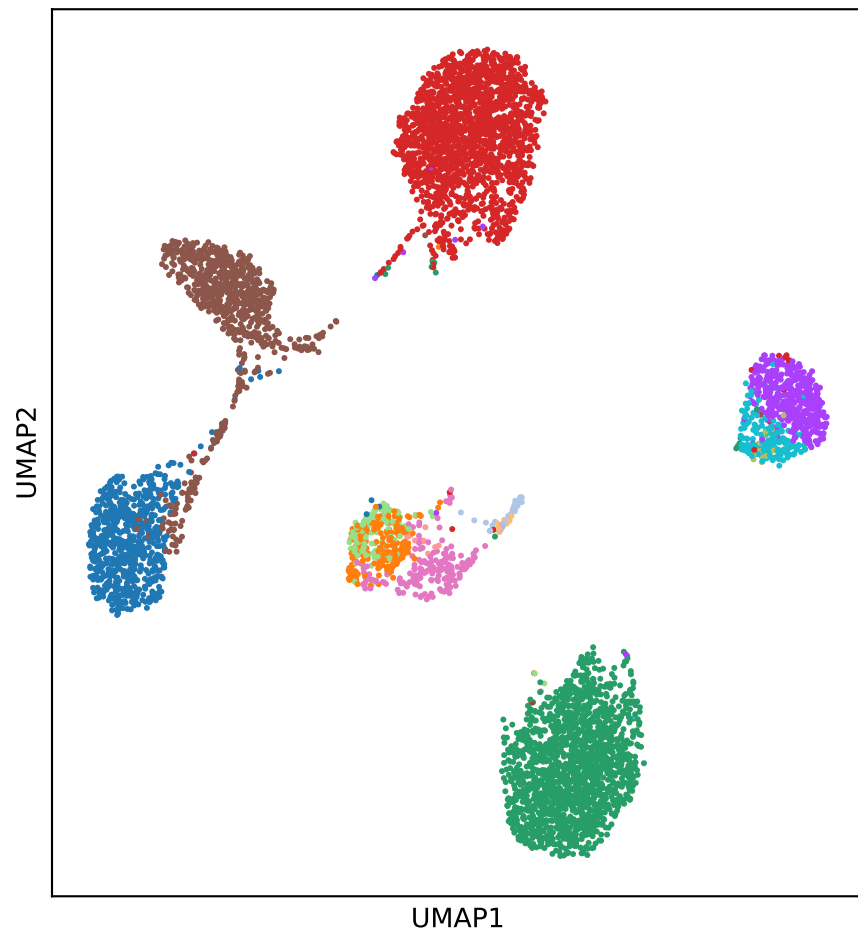
UMAP projection - Labels (train) (Res: 0.06, Iter: 24)



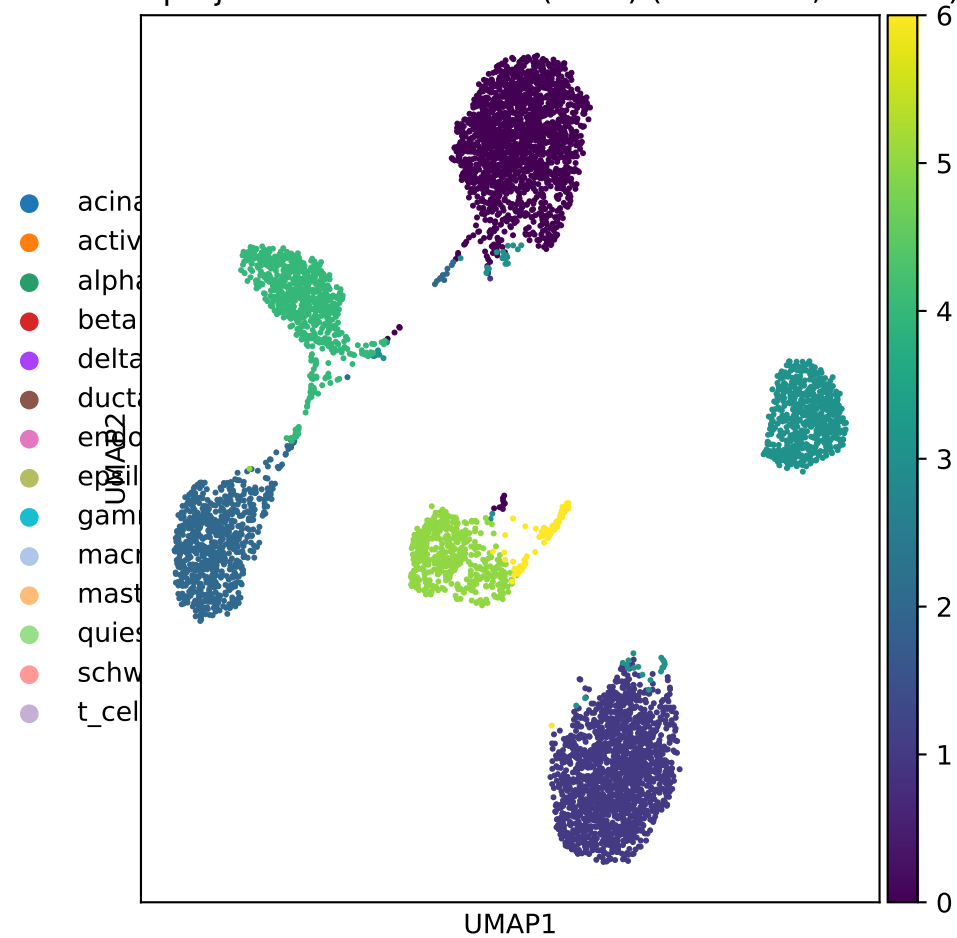
UMAP projection - Predictions (train) (Res: 0.06, Iter: 24)



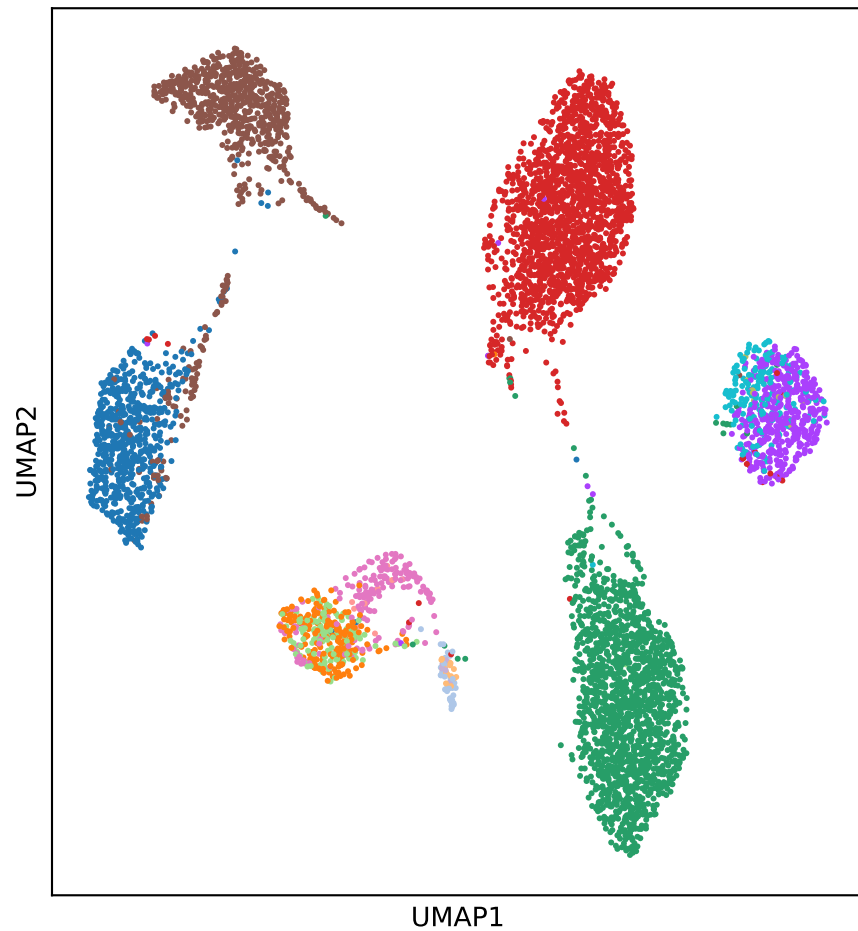
UMAP projection - Labels (train) (Res: 0.06, Iter: 48)



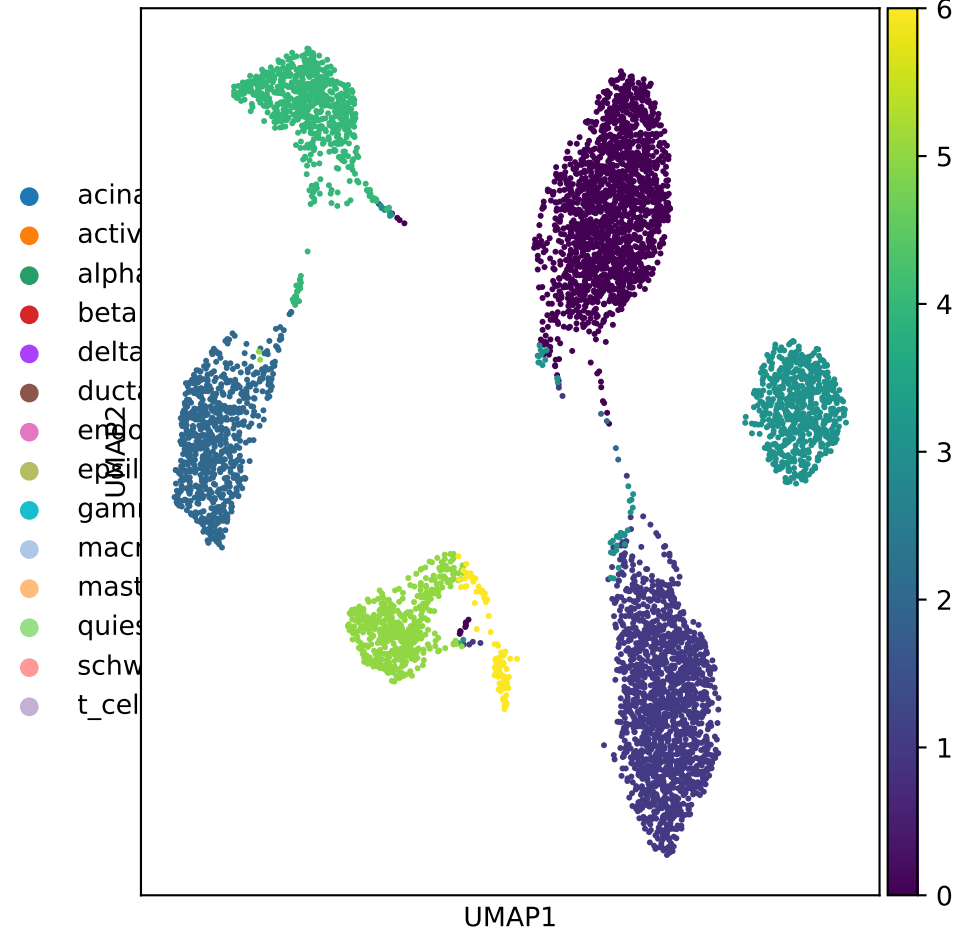
UMAP projection - Predictions (train) (Res: 0.06, Iter: 48)



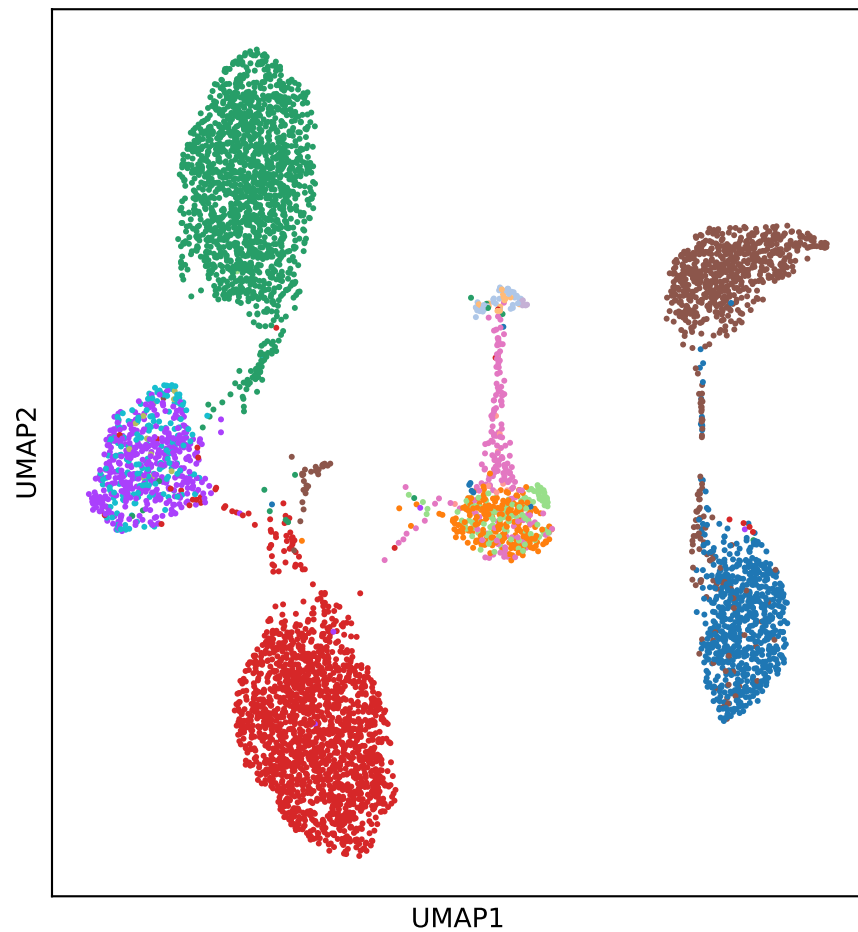
UMAP projection - Labels (train) (Res: 0.06, Iter: 72)



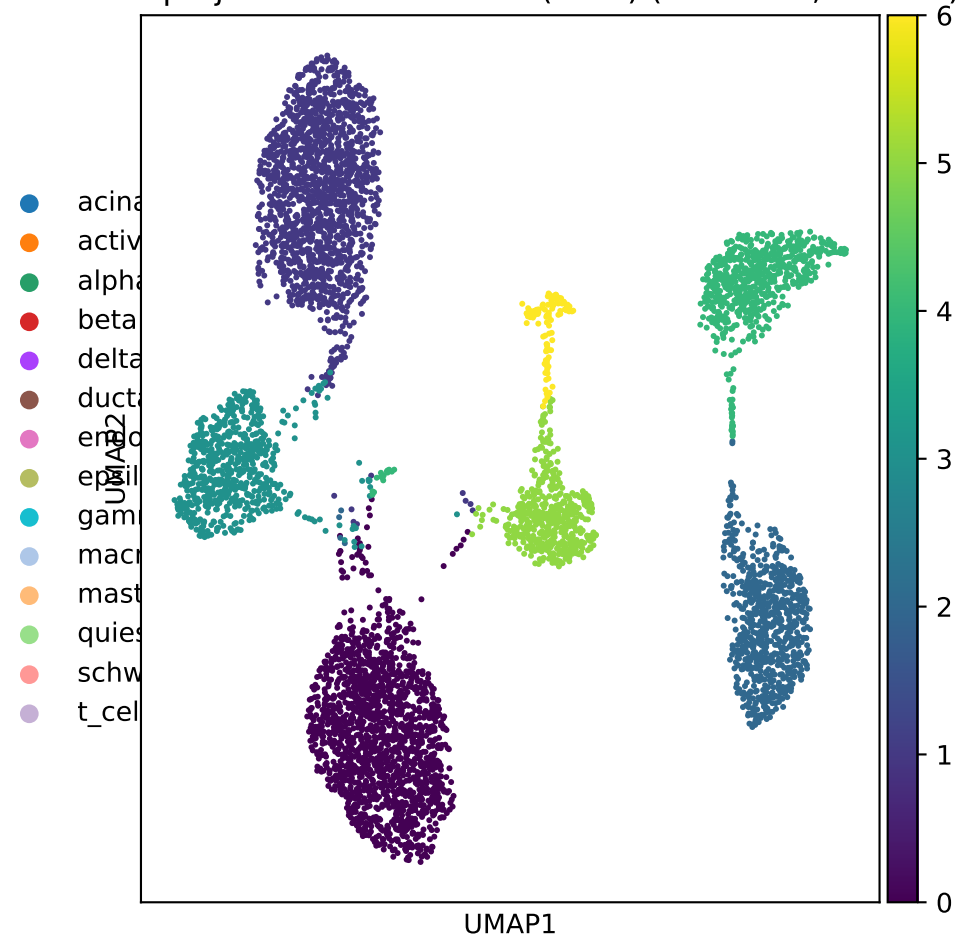
UMAP projection - Predictions (train) (Res: 0.06, Iter: 72)



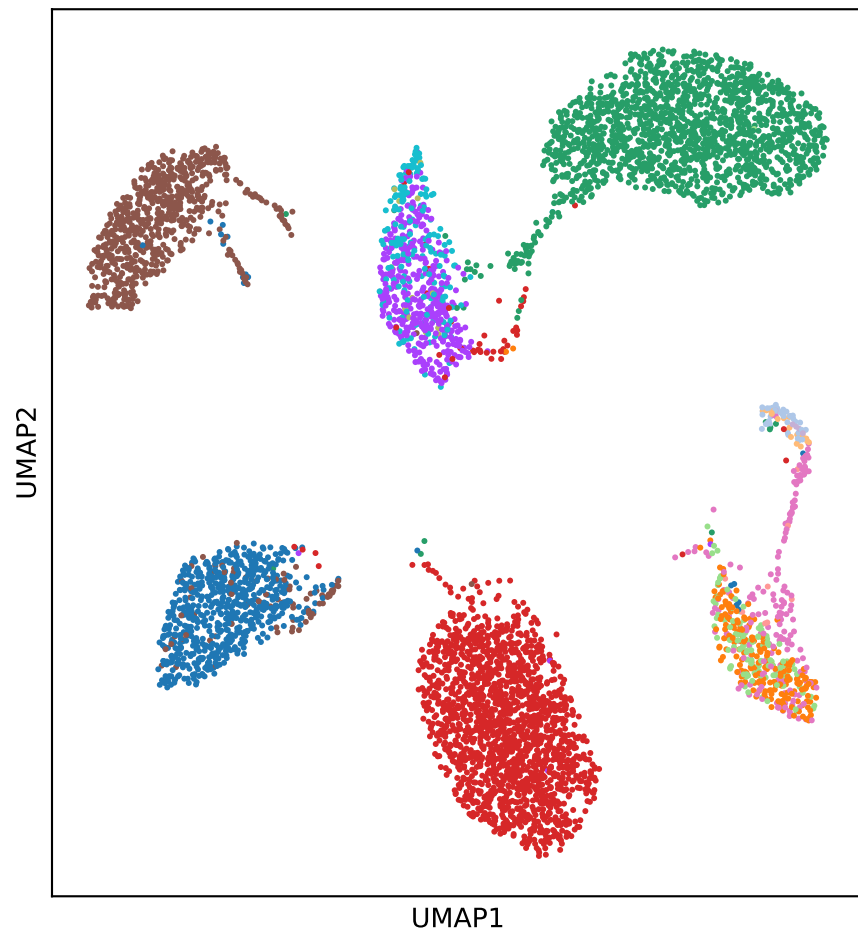
UMAP projection - Labels (train) (Res: 0.06, Iter: 96)



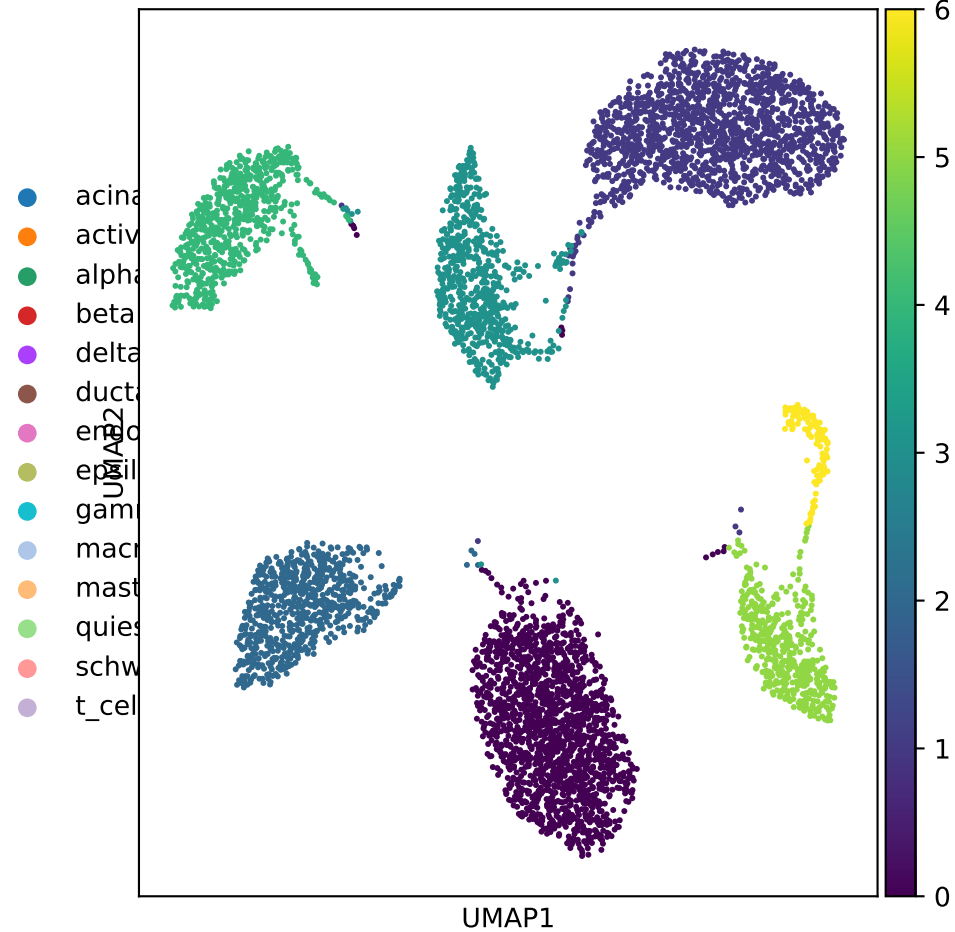
UMAP projection - Predictions (train) (Res: 0.06, Iter: 96)



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

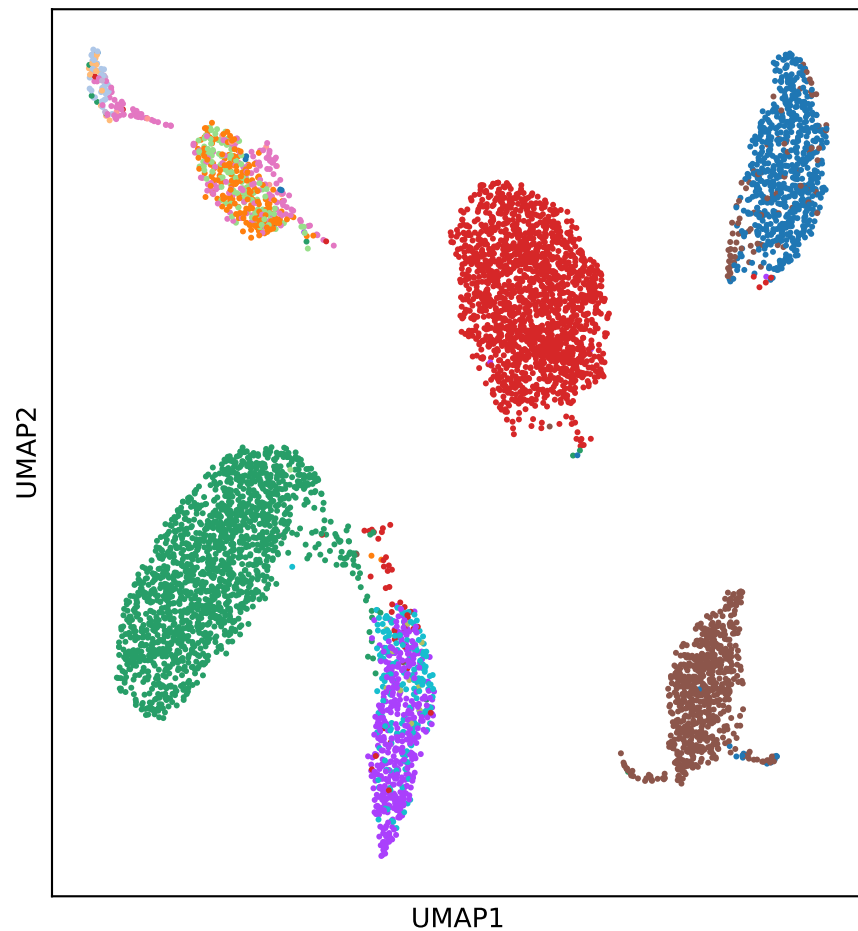


UMAP projection - Predictions (train) (Res: 0.06, Iter: 120)

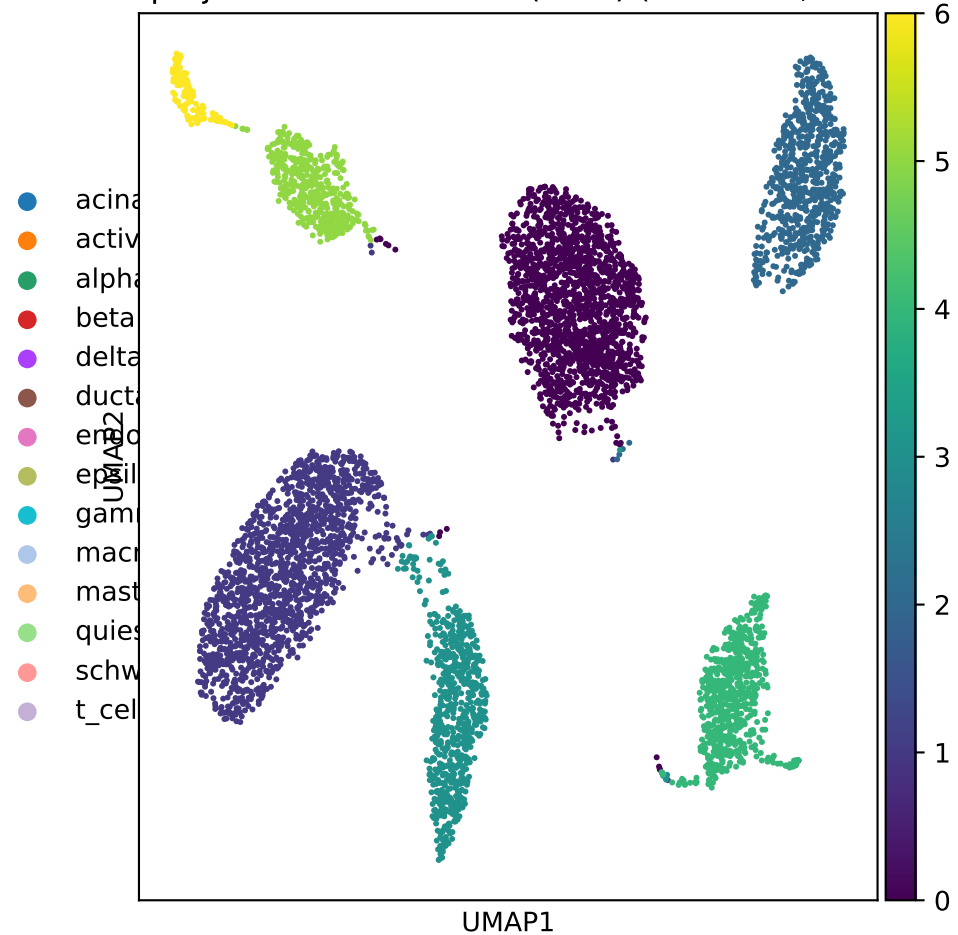


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

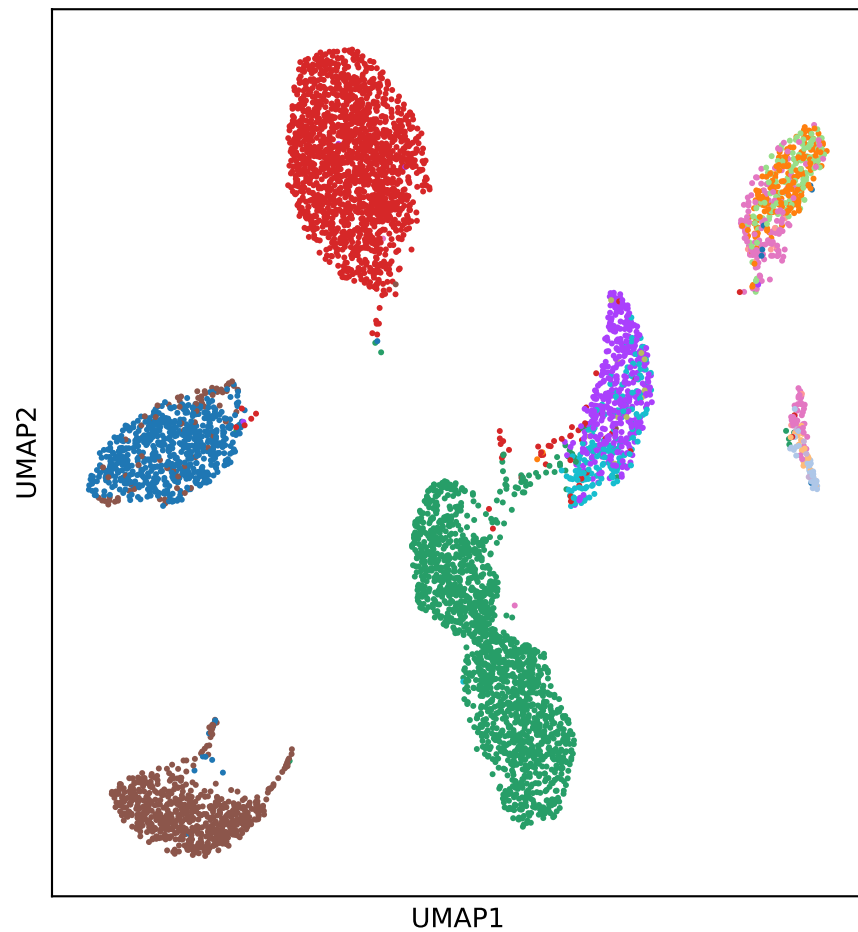
UMAP projection - Labels (train) (Res: 0.06, Iter: 144)



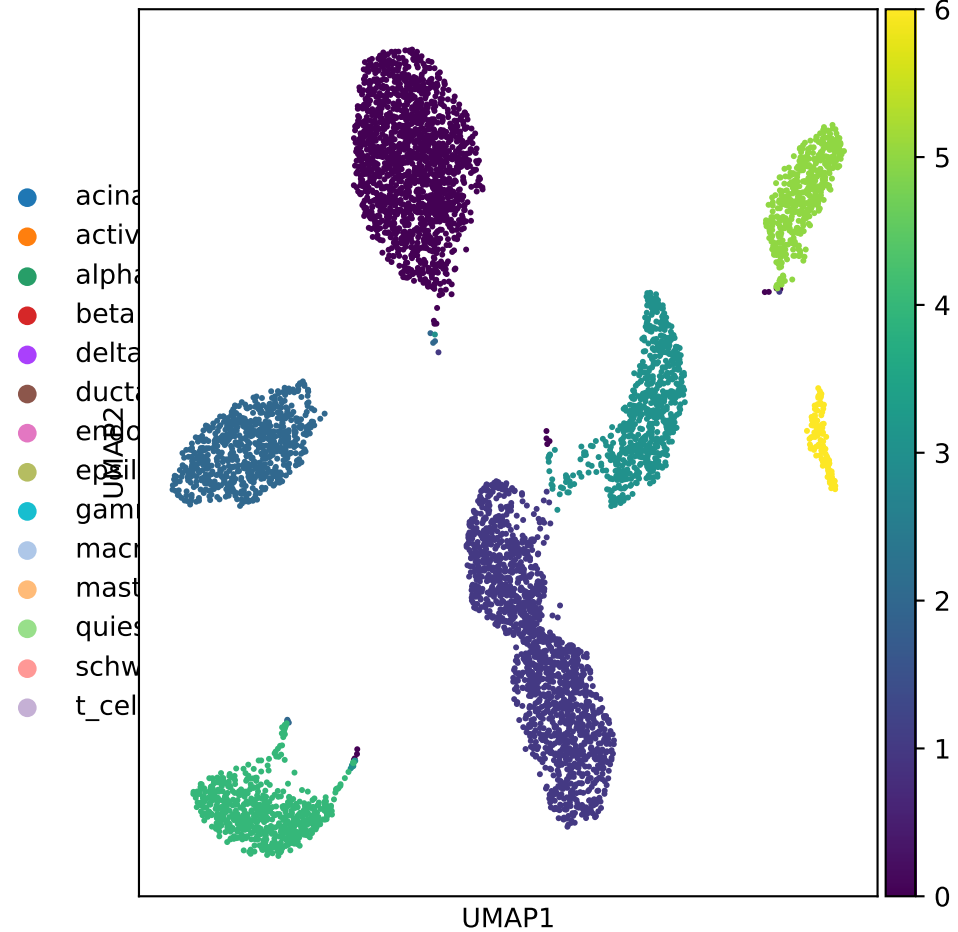
UMAP projection - Predictions (train) (Res: 0.06, Iter: 144)



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

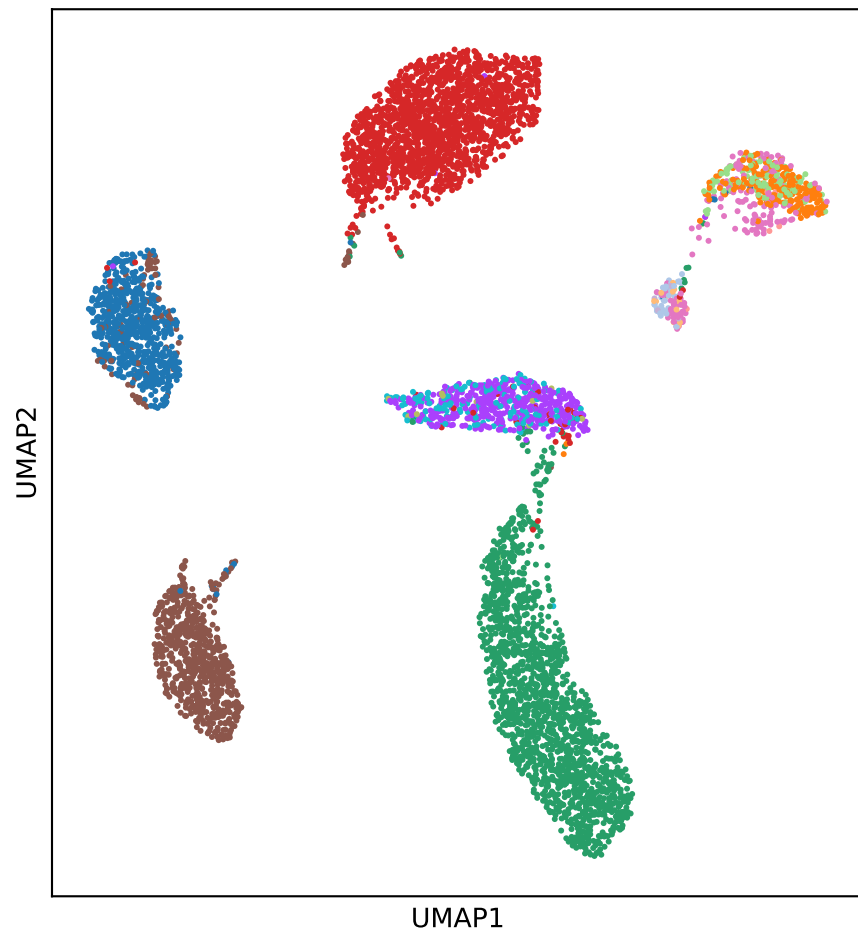


UMAP projection - Predictions (train) (Res: 0.06, Iter: 168)

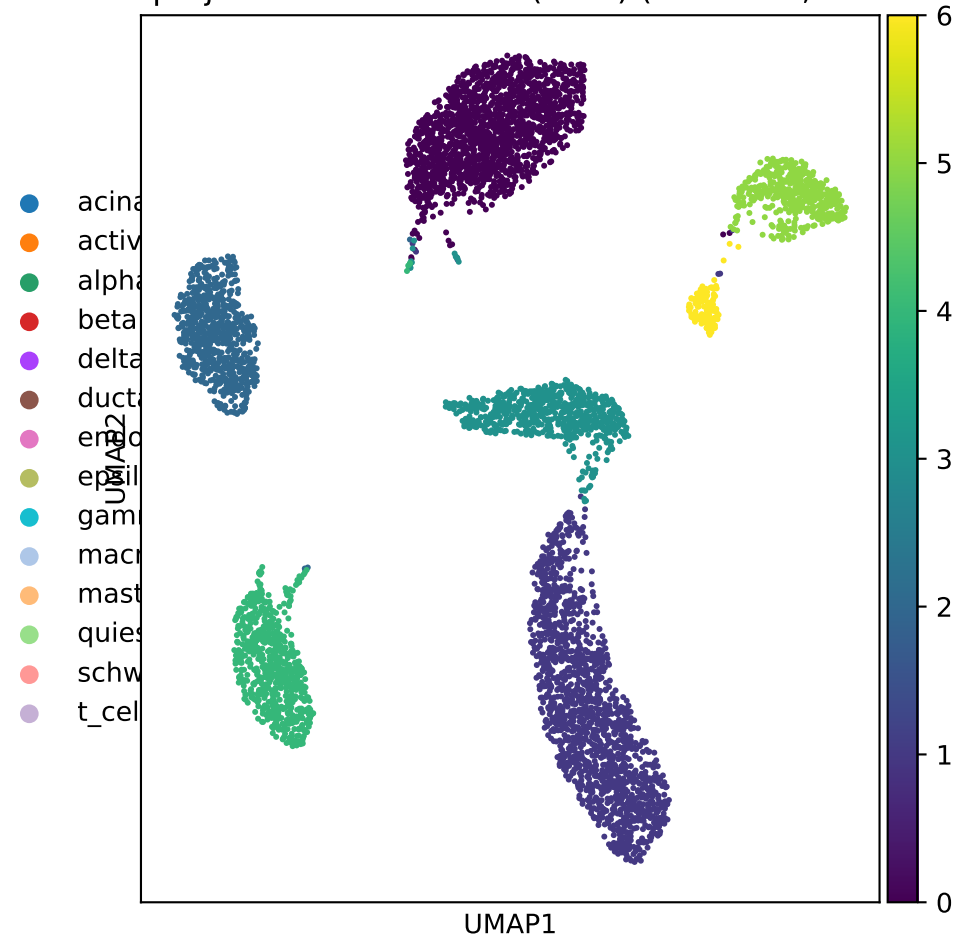


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

UMAP projection - Labels (train) (Res: 0.06, Iter: 192)

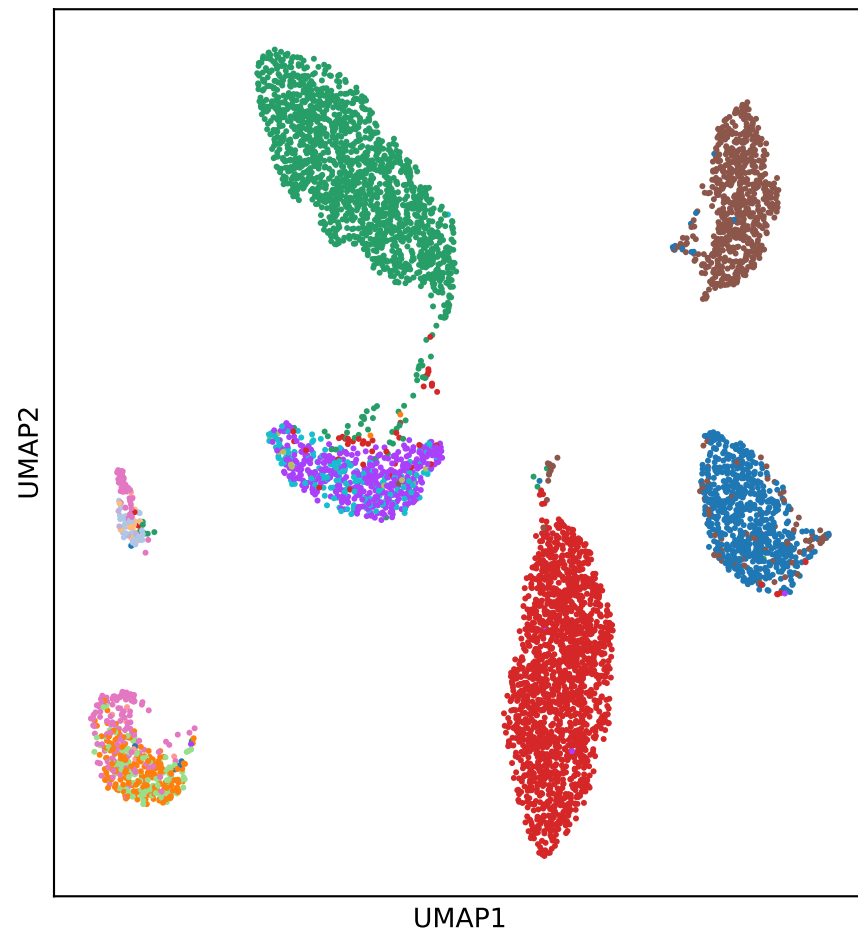


UMAP projection - Predictions (train) (Res: 0.06, Iter: 192)

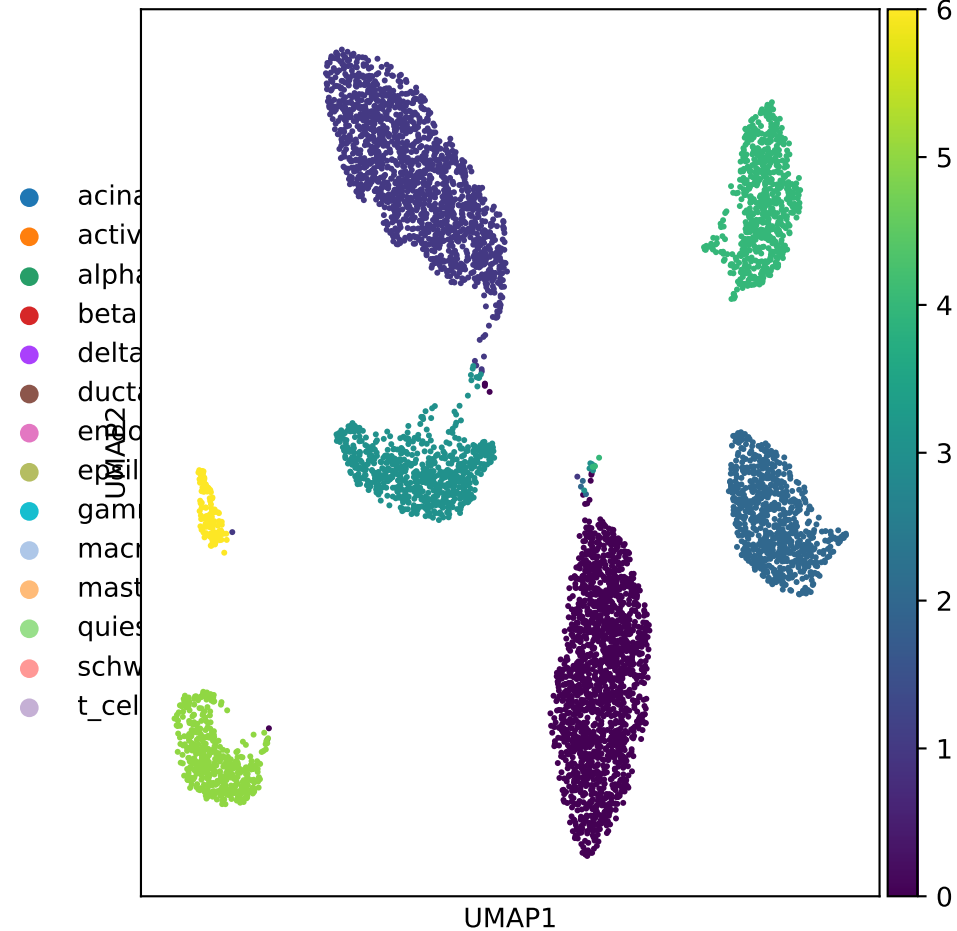


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

UMAP projection - Labels (train) (Res: 0.06, Iter: 216)

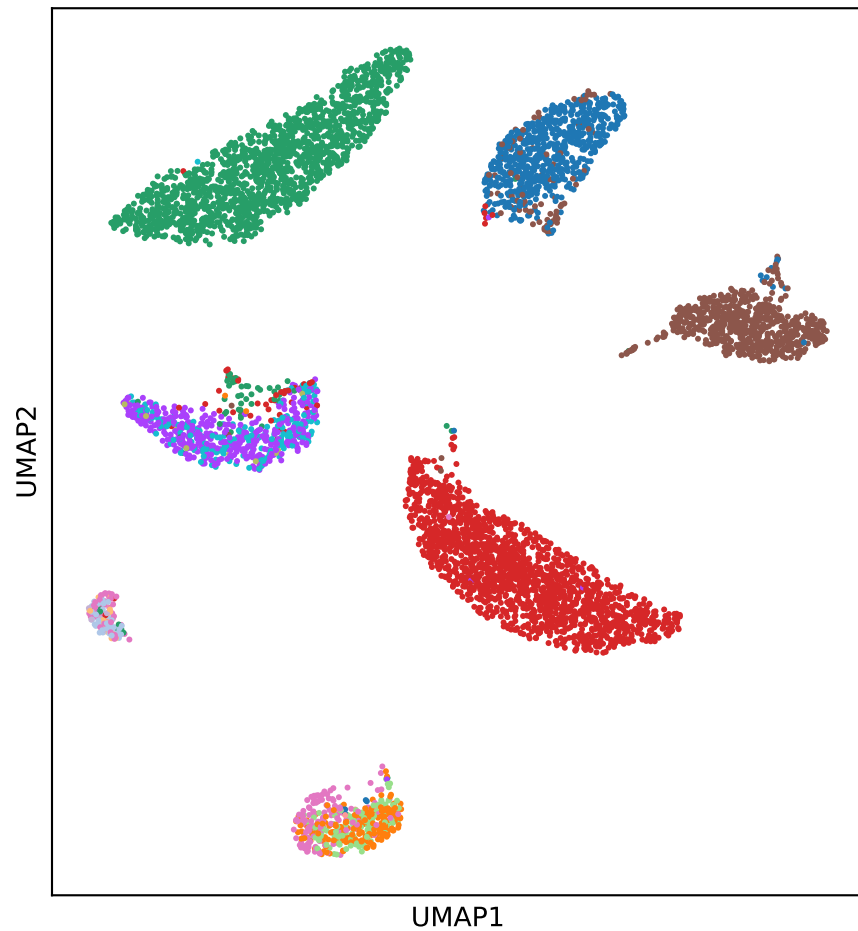


UMAP projection - Predictions (train) (Res: 0.06, Iter: 216)

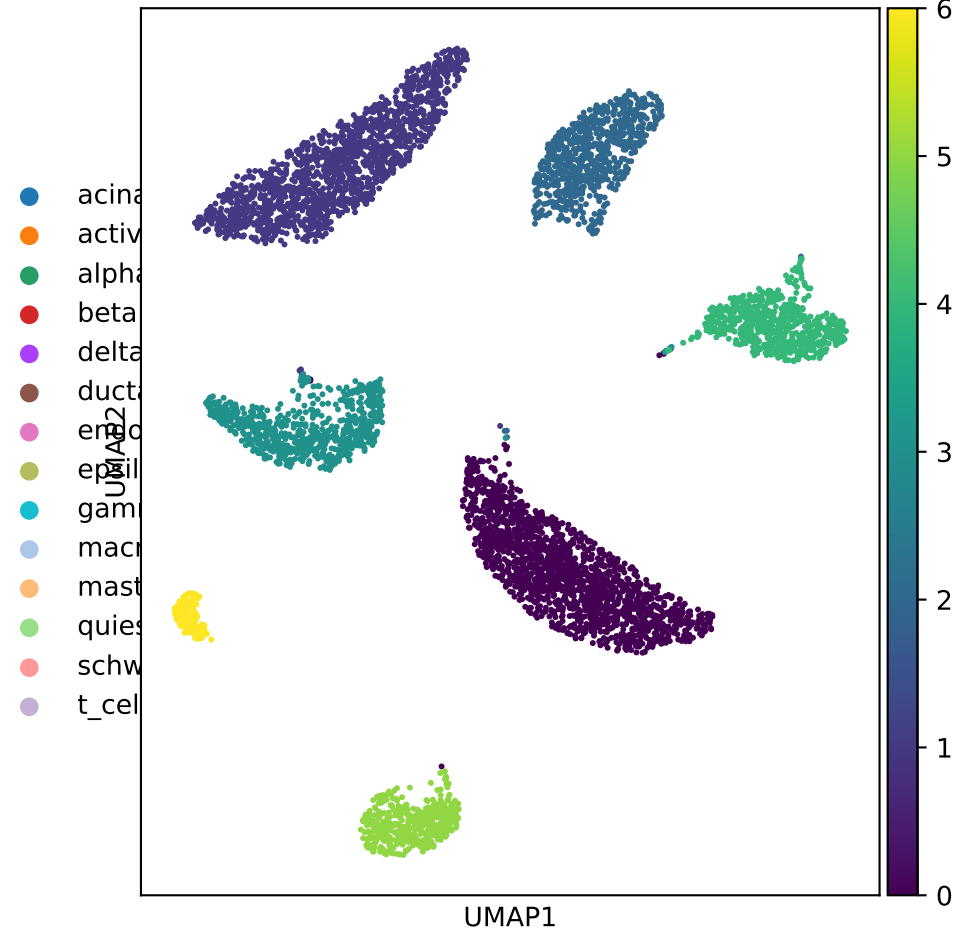


- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

UMAP projection - Labels (train) (Res: 0.06, Iter: 240)



UMAP projection - Predictions (train) (Res: 0.06, Iter: 240)



- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel