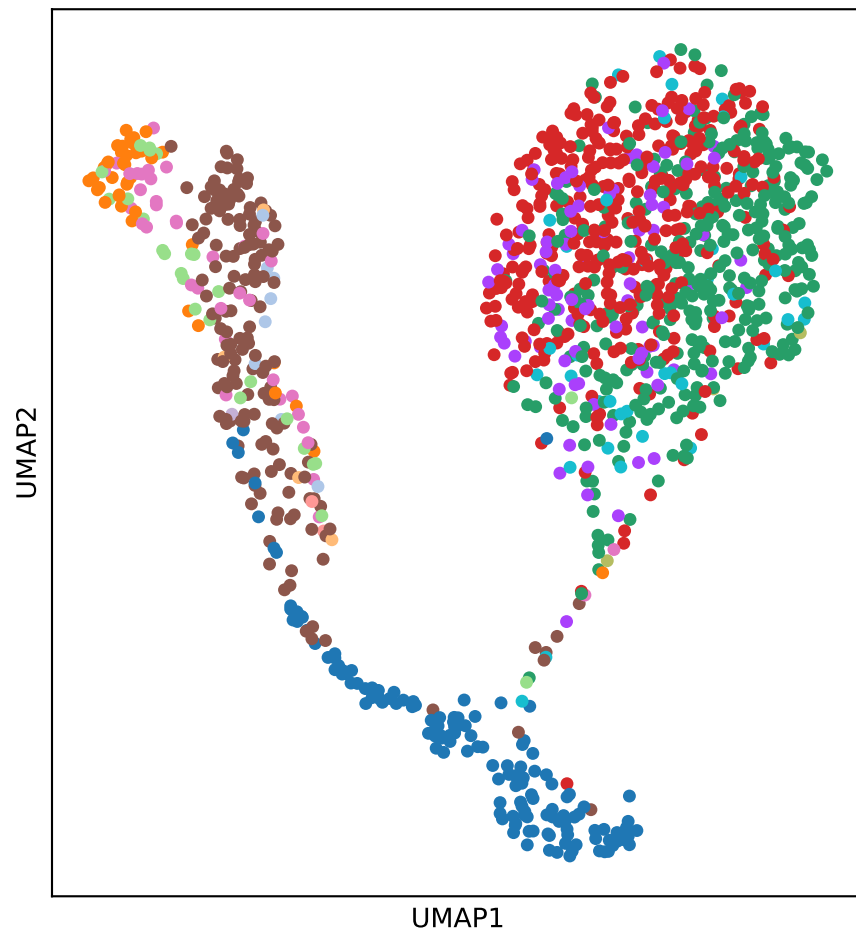
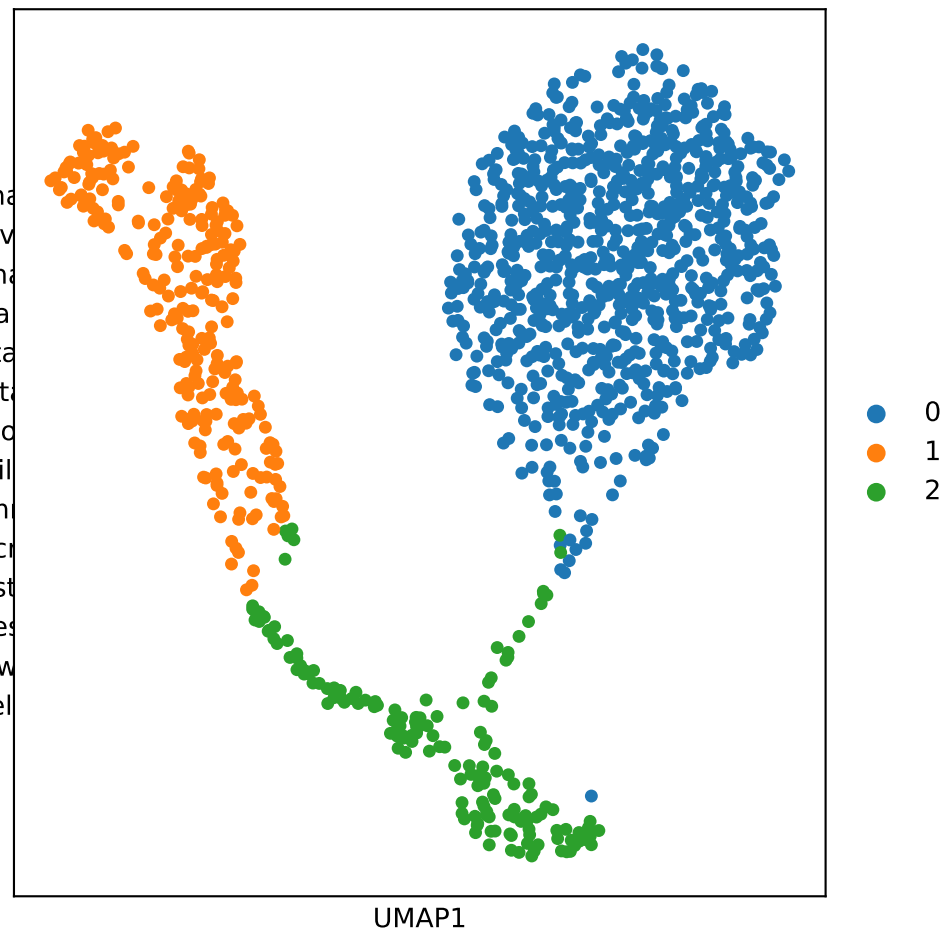


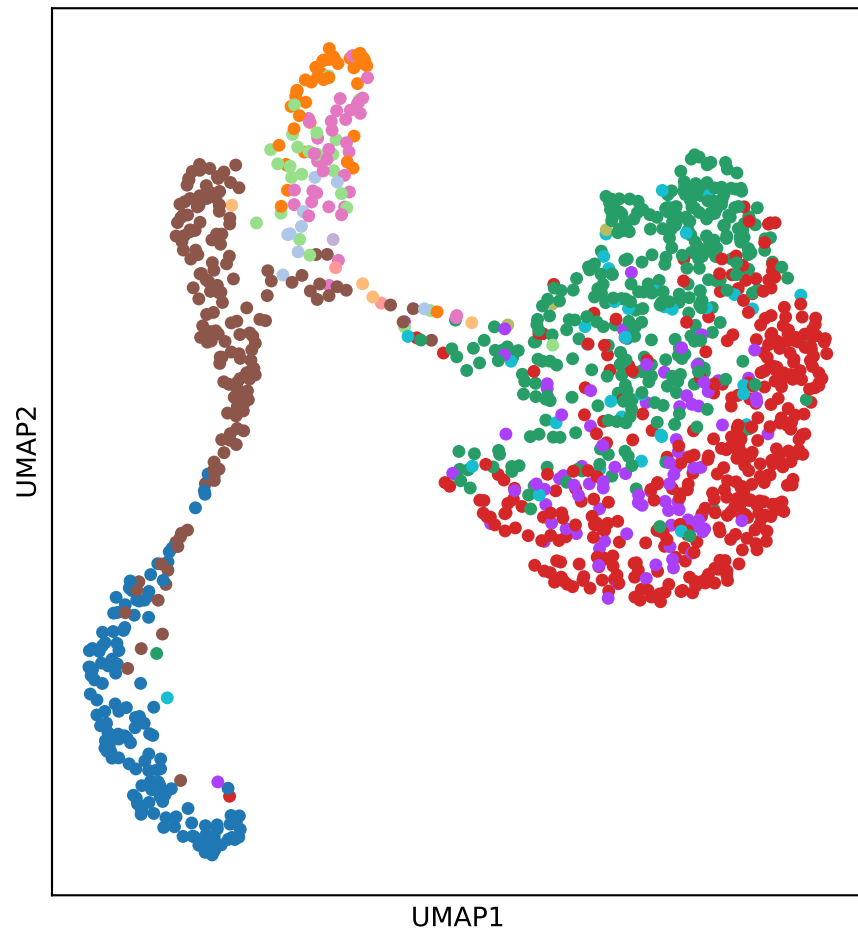
UMAP projection - Labels (val) (Res: 0.1, Iter: 0)



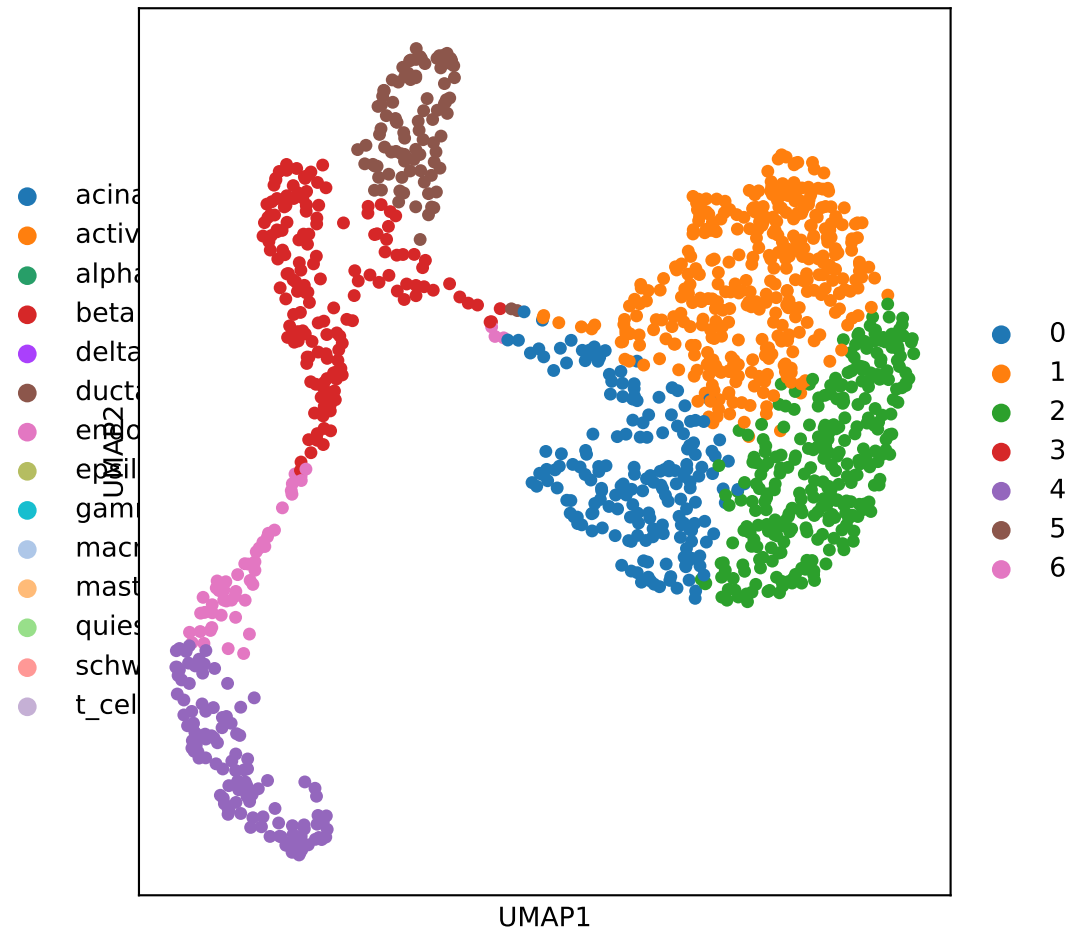
UMAP projection - Predictions (val) (Res: 0.1, Iter: 0)



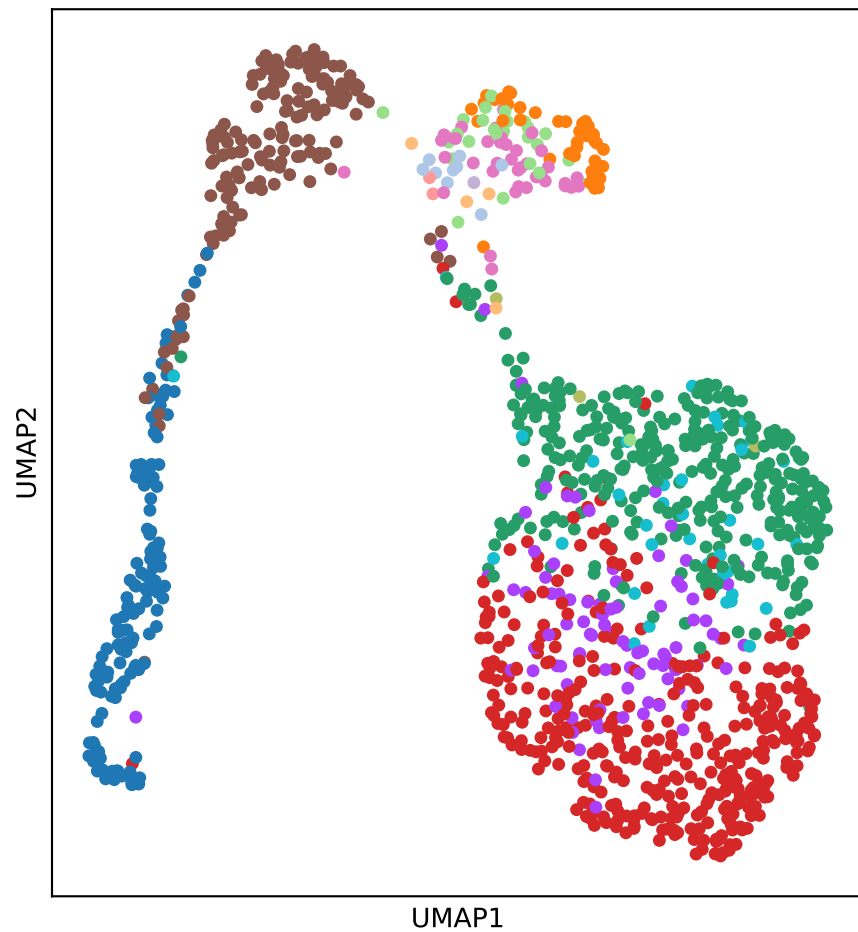
UMAP projection - Labels (val) (Res: 0.2, Iter: 1)



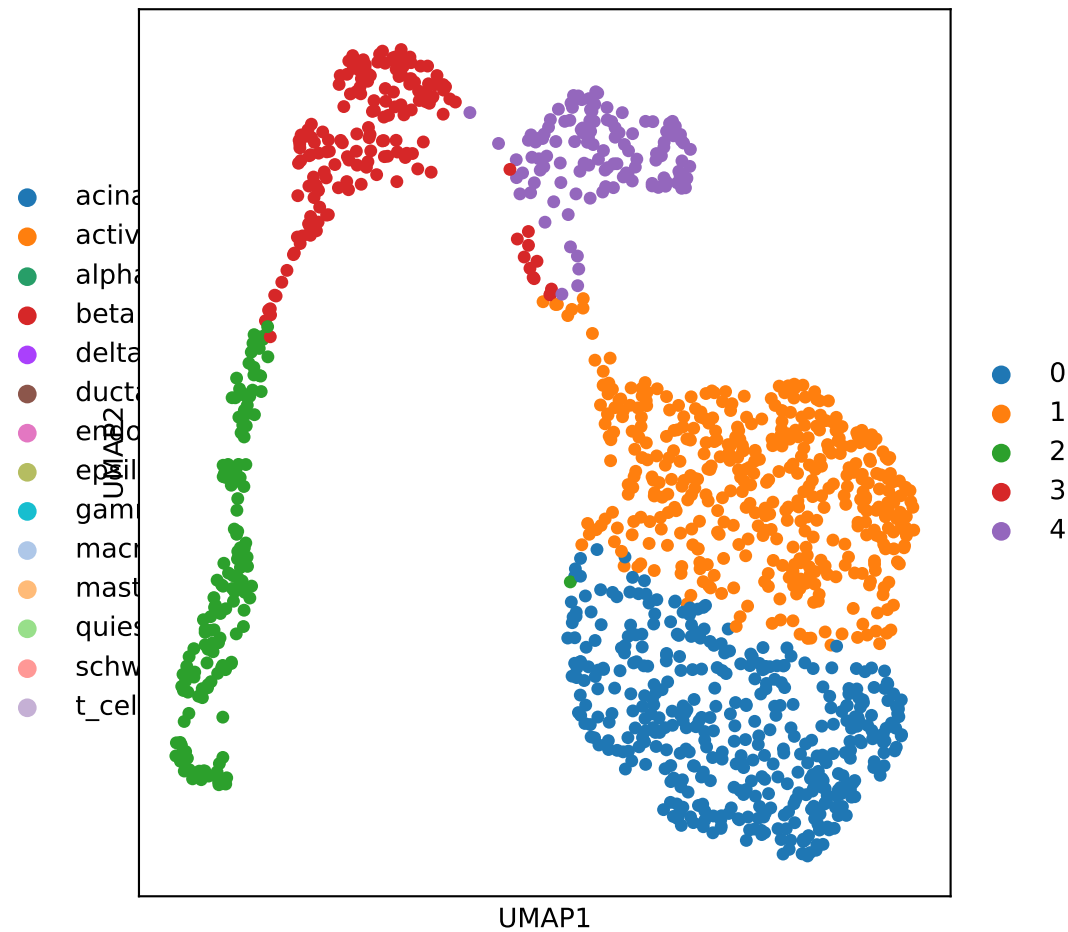
UMAP projection - Predictions (val) (Res: 0.2, Iter: 1)



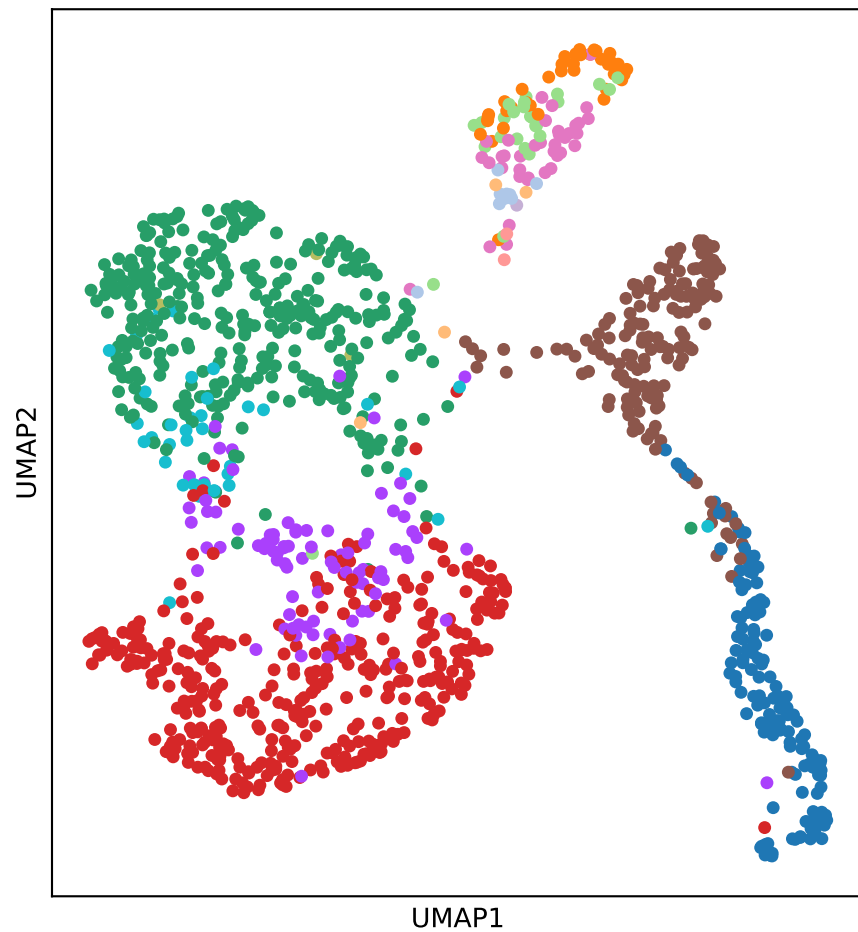
UMAP projection - Labels (val) (Res: 0.05, Iter: 2)



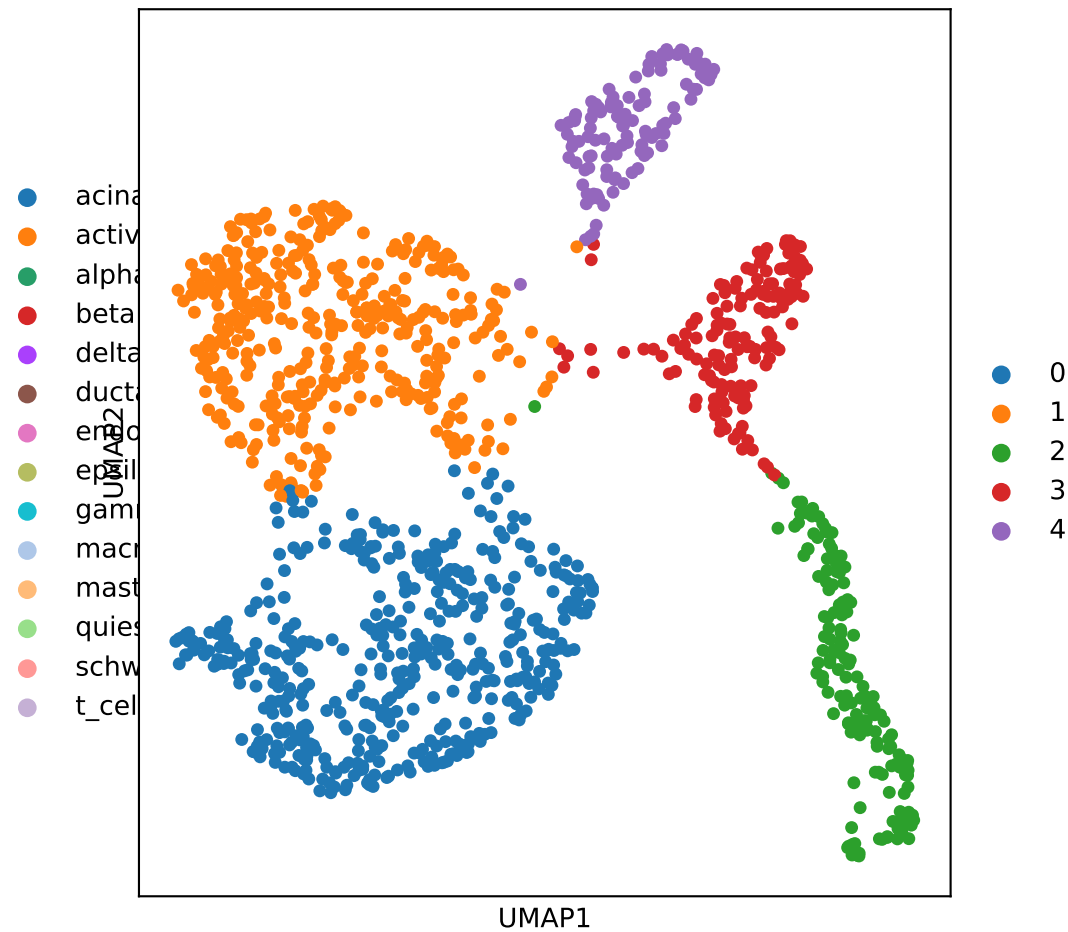
UMAP projection - Predictions (val) (Res: 0.05, Iter: 2)



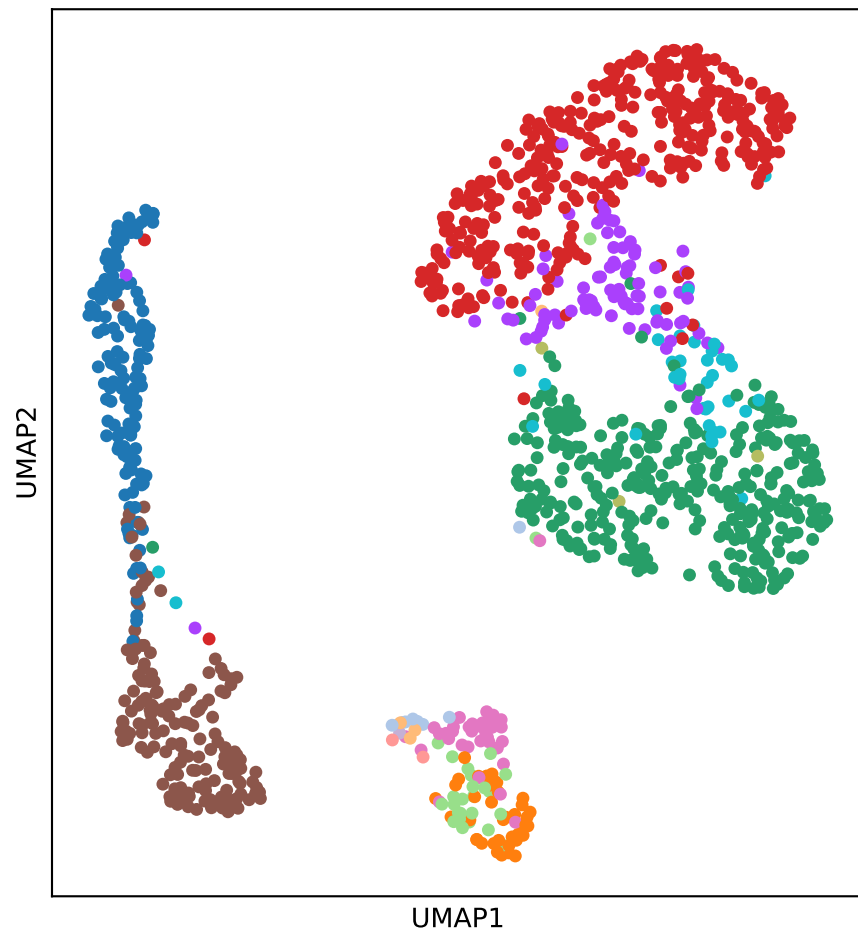
UMAP projection - Labels (val) (Res: 0.03, Iter: 3)



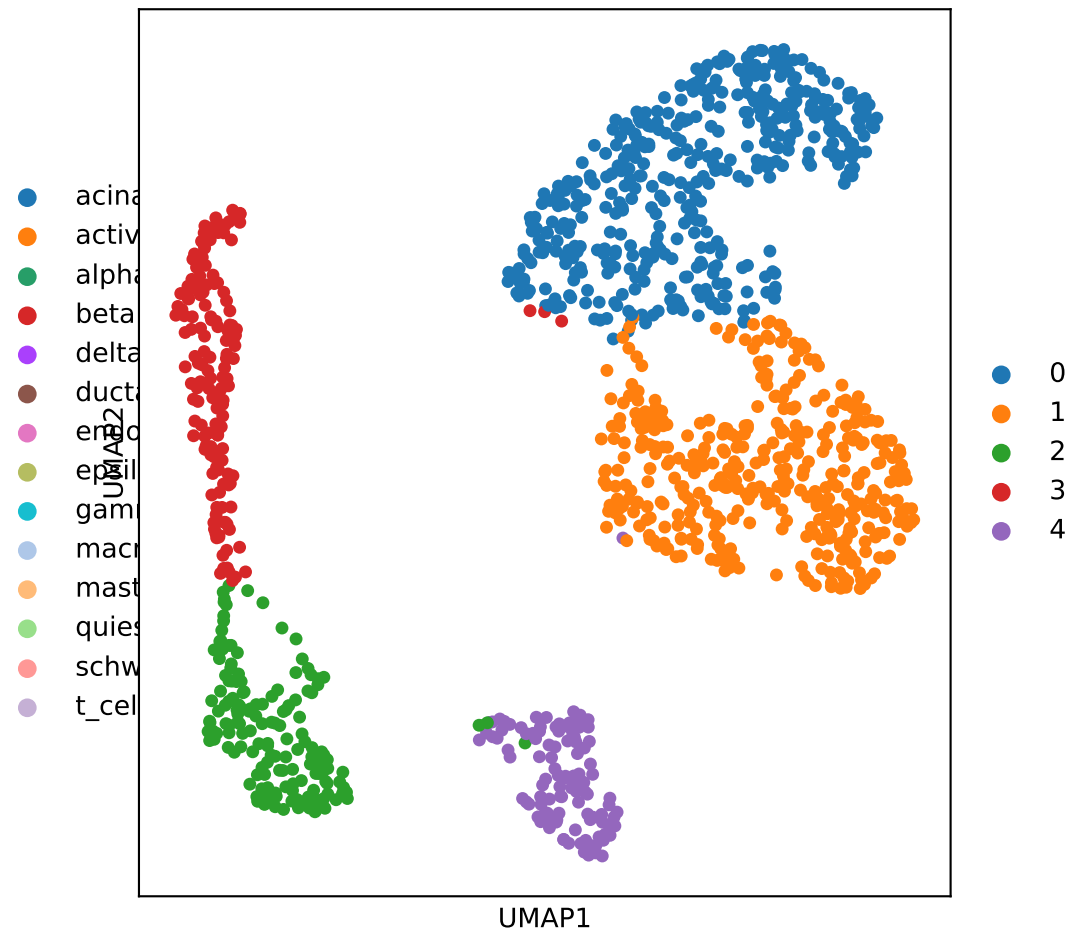
UMAP projection - Predictions (val) (Res: 0.03, Iter: 3)



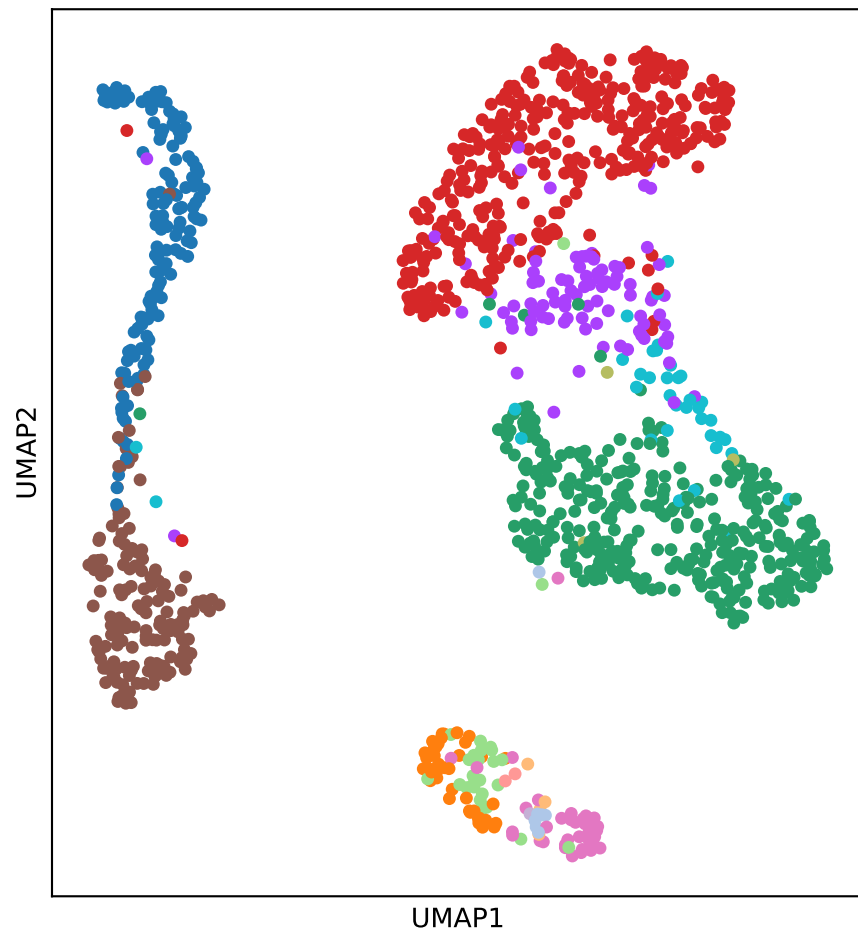
UMAP projection - Labels (val) (Res: 0.05, Iter: 4)



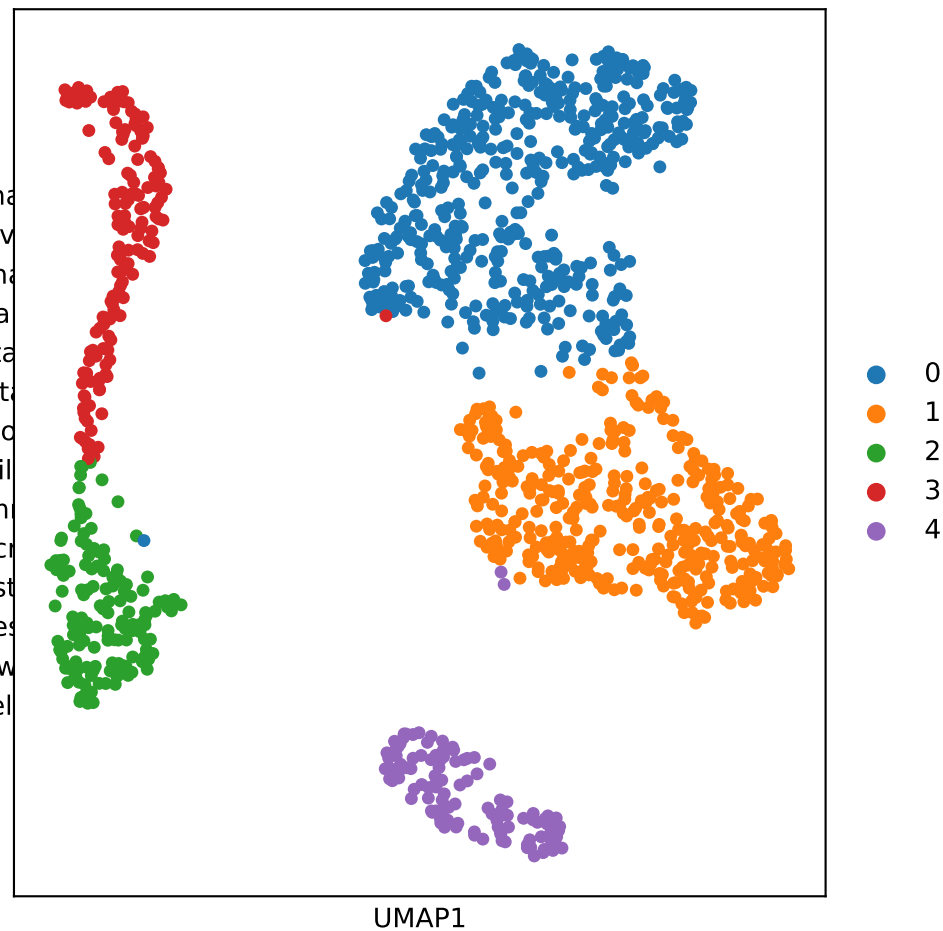
UMAP projection - Predictions (val) (Res: 0.05, Iter: 4)



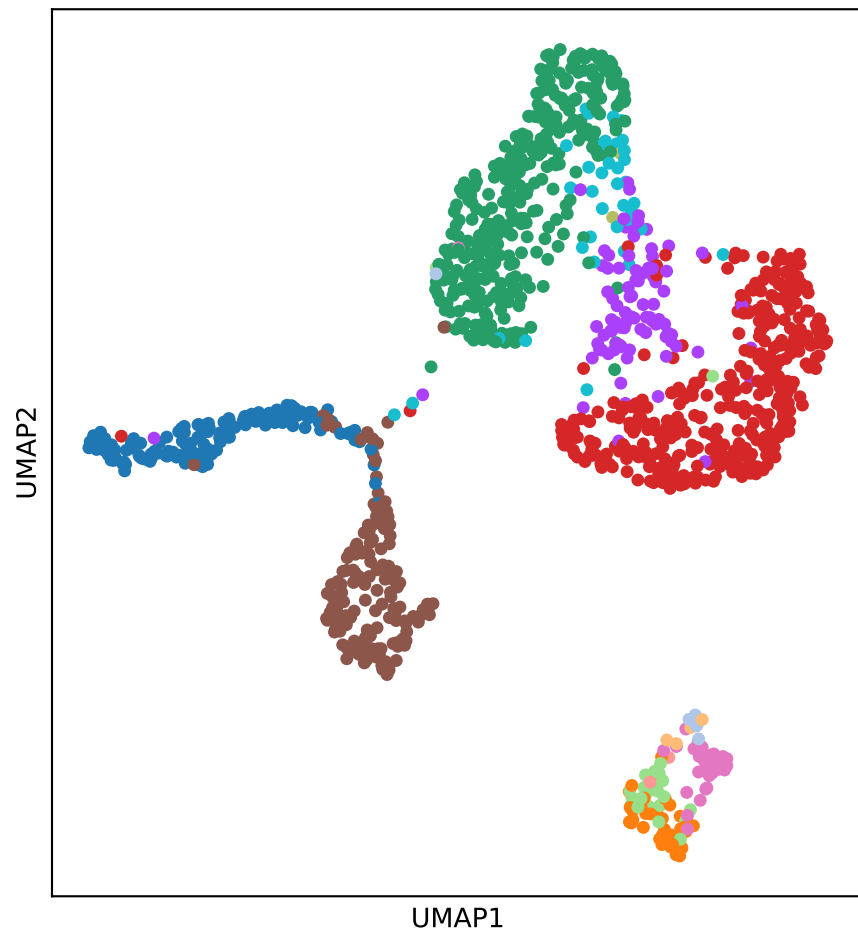
UMAP projection - Labels (val) (Res: 0.03, Iter: 5)



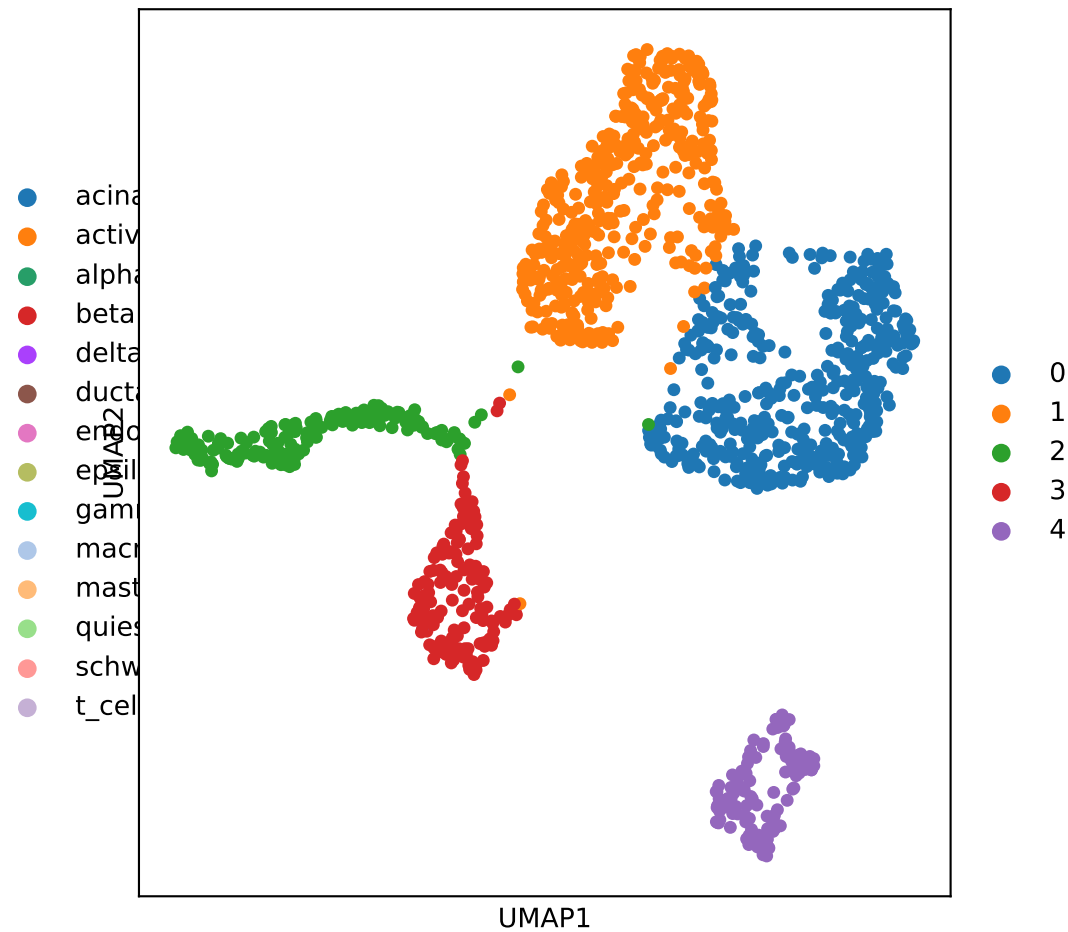
UMAP projection - Predictions (val) (Res: 0.03, Iter: 5)



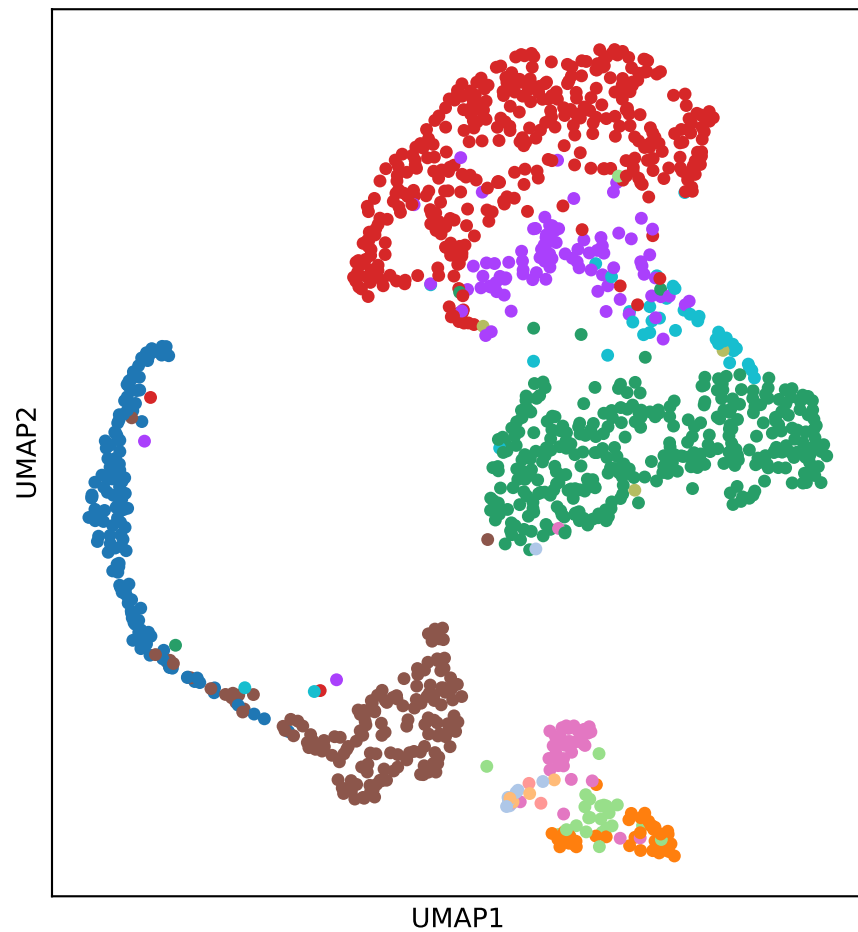
UMAP projection - Labels (val) (Res: 0.03, Iter: 6)



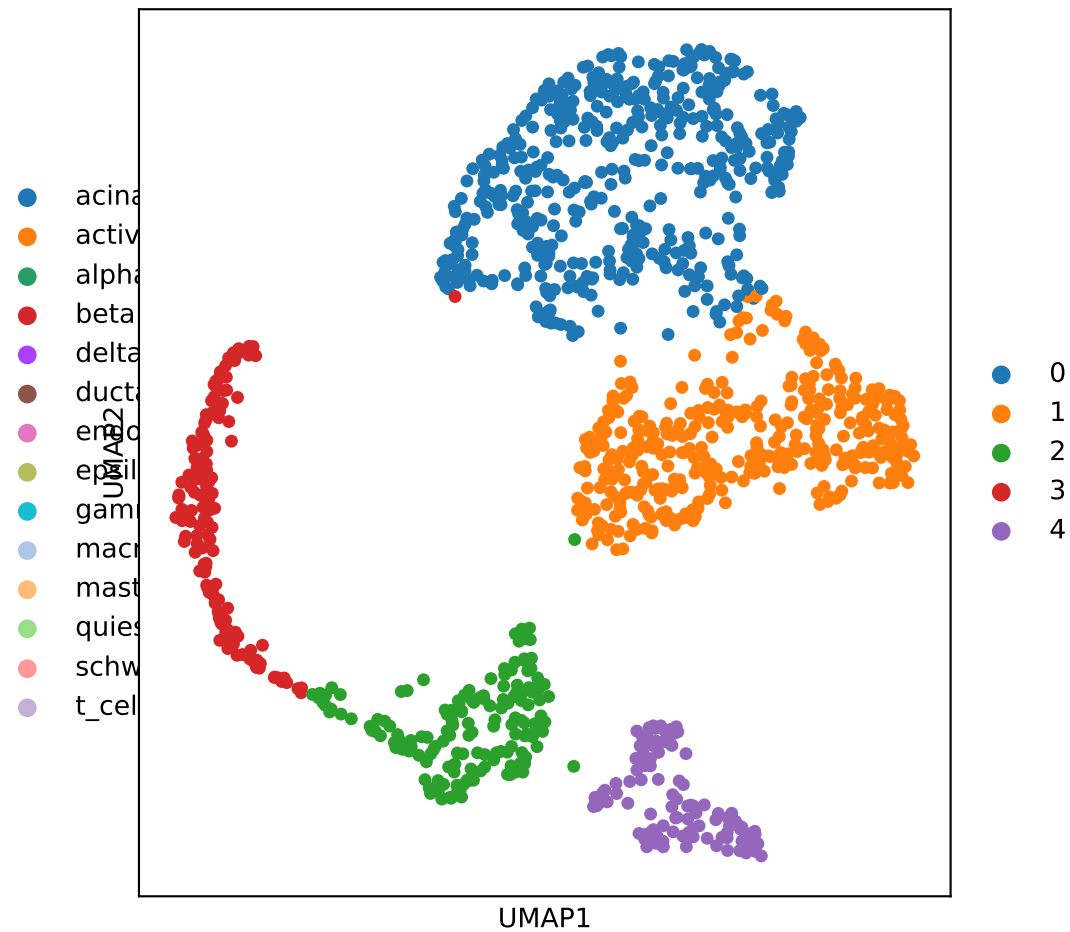
UMAP projection - Predictions (val) (Res: 0.03, Iter: 6)



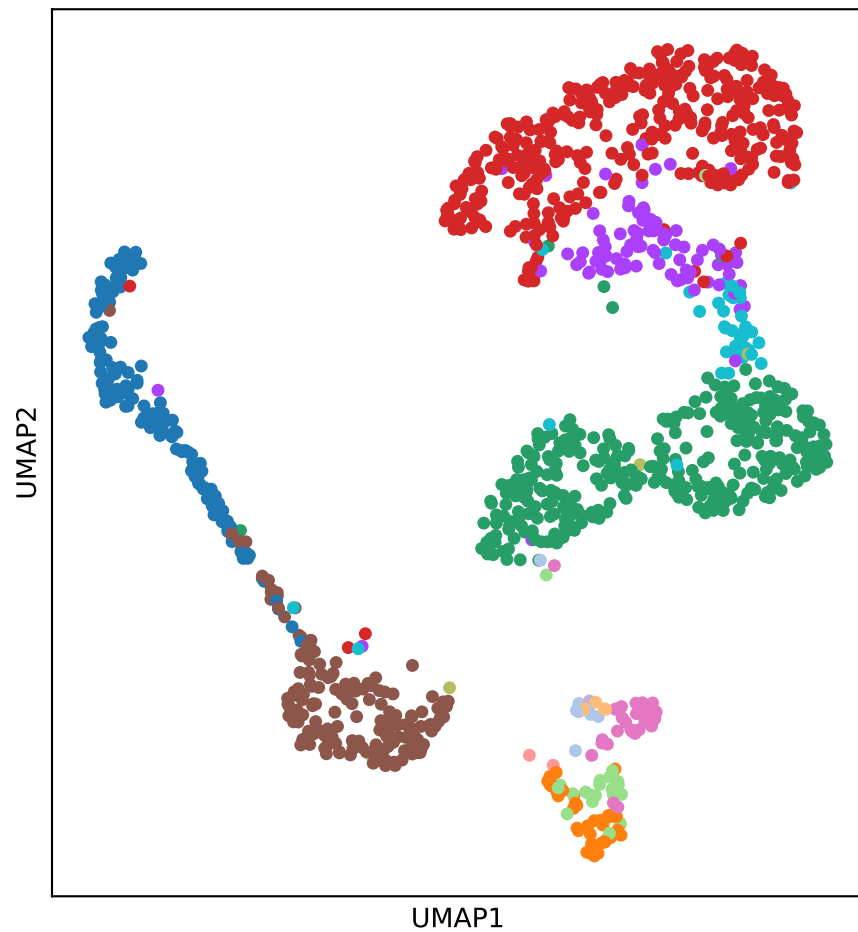
UMAP projection - Labels (val) (Res: 0.06, Iter: 7)



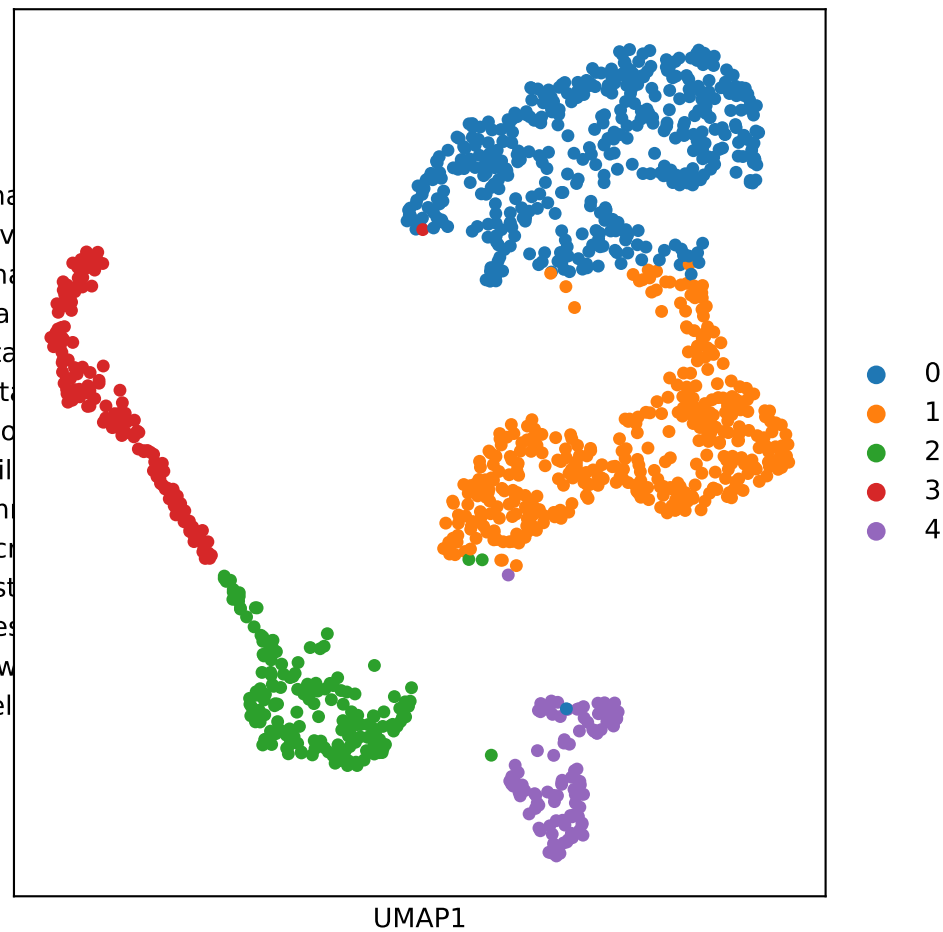
UMAP projection - Predictions (val) (Res: 0.06, Iter: 7)



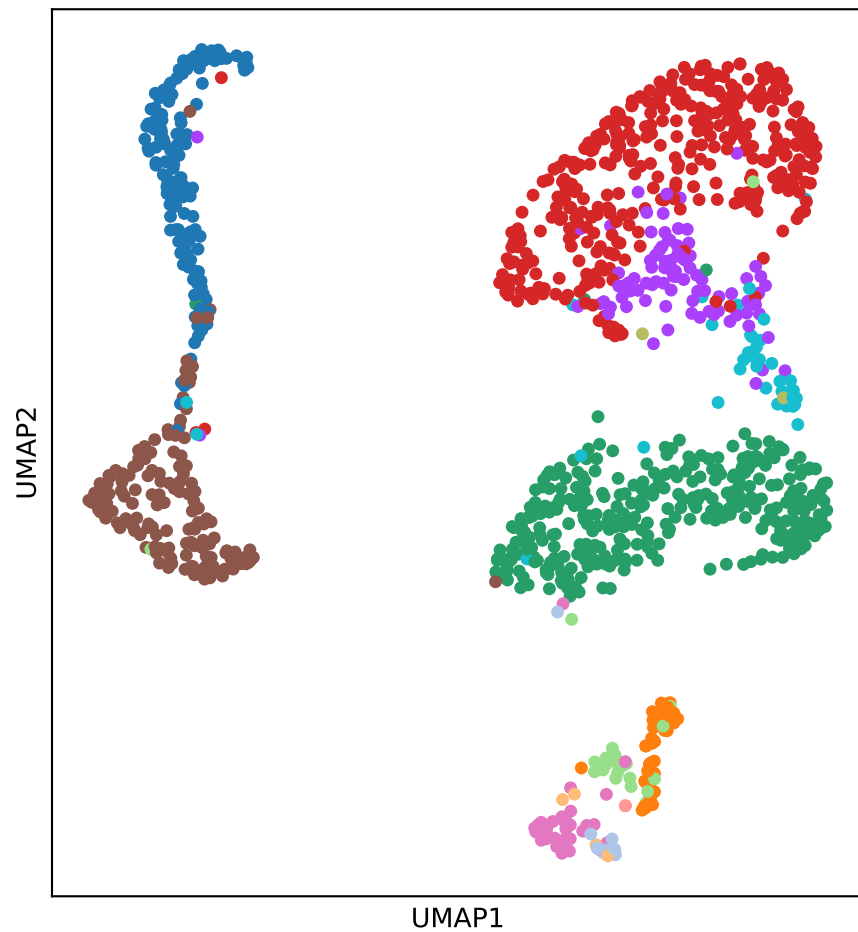
UMAP projection - Labels (val) (Res: 0.03, Iter: 8)



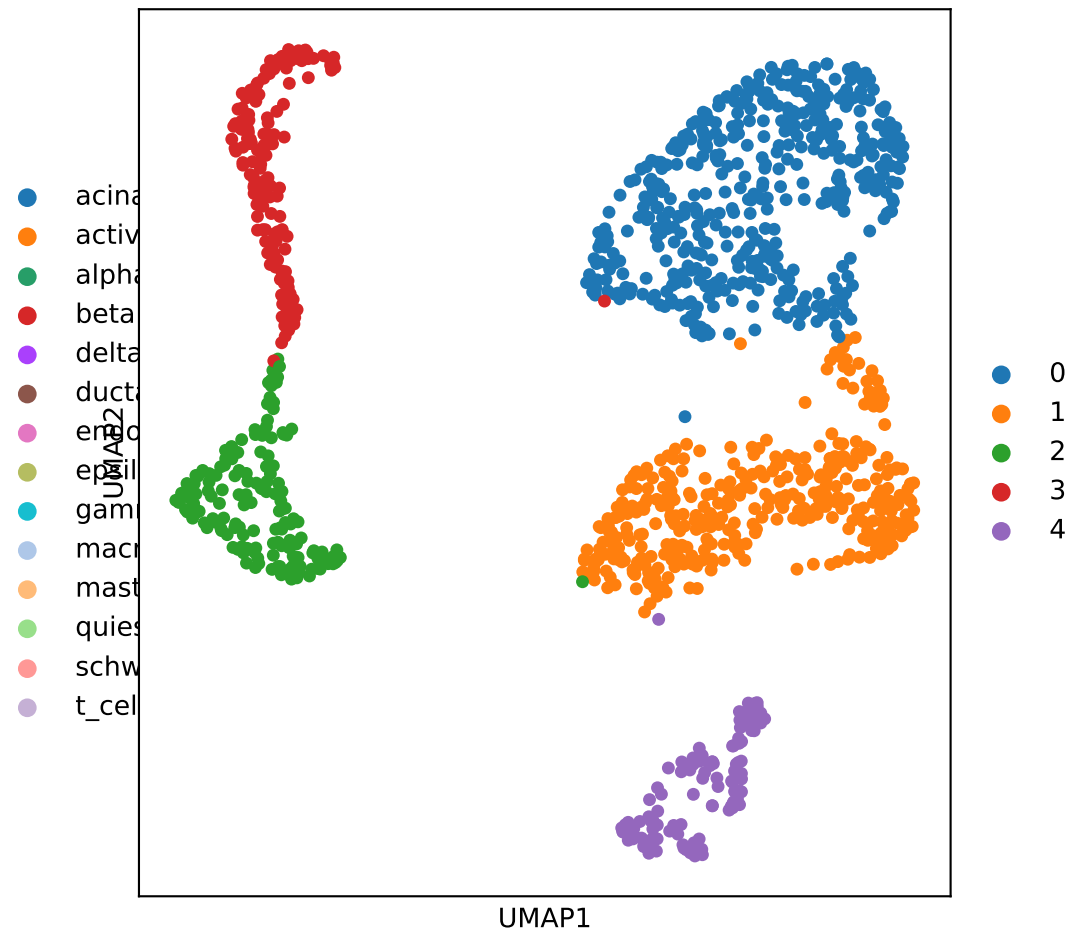
UMAP projection - Predictions (val) (Res: 0.03, Iter: 8)



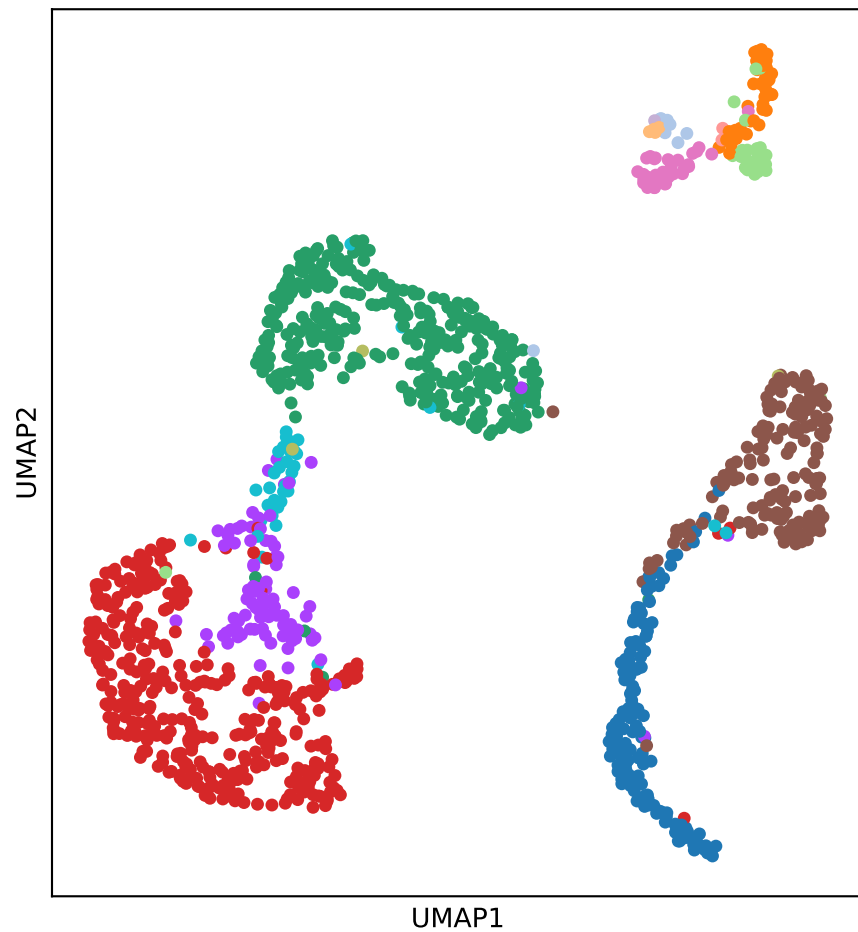
UMAP projection - Labels (val) (Res: 0.04, Iter: 9)



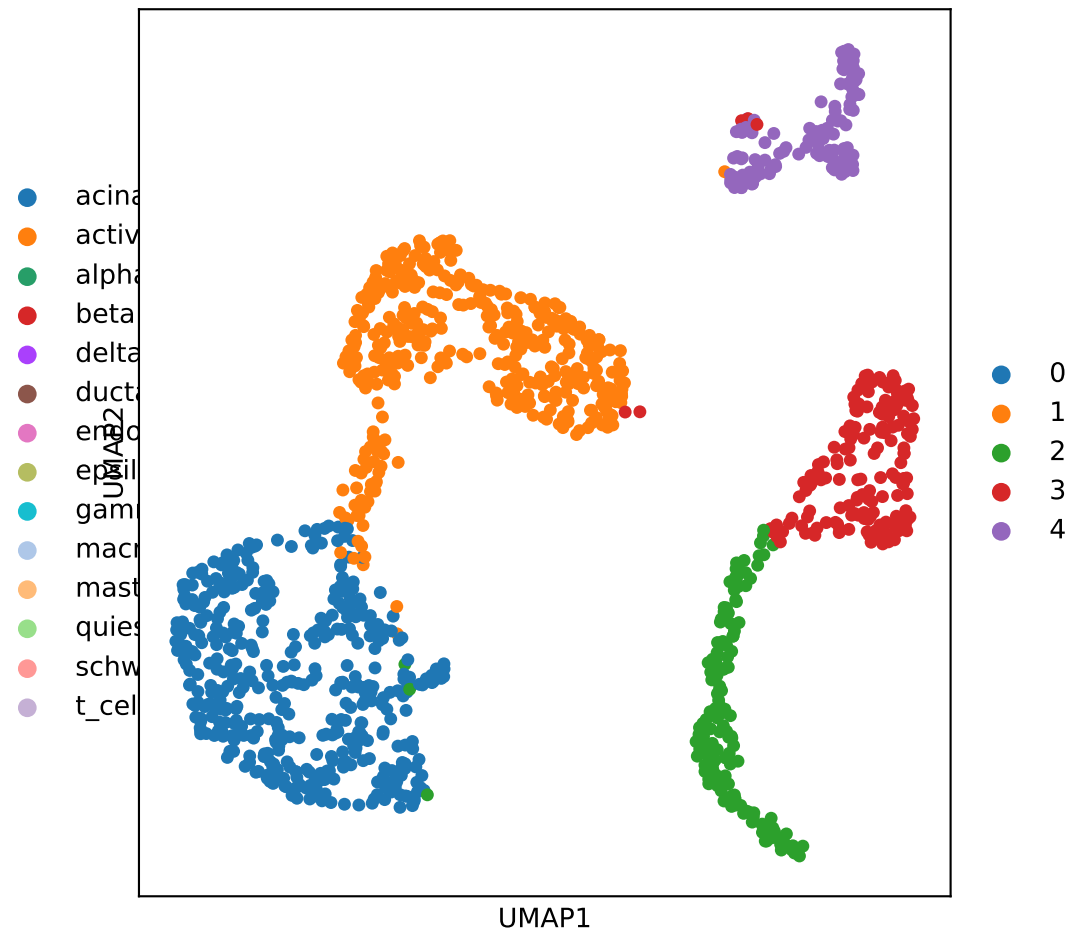
UMAP projection - Predictions (val) (Res: 0.04, Iter: 9)



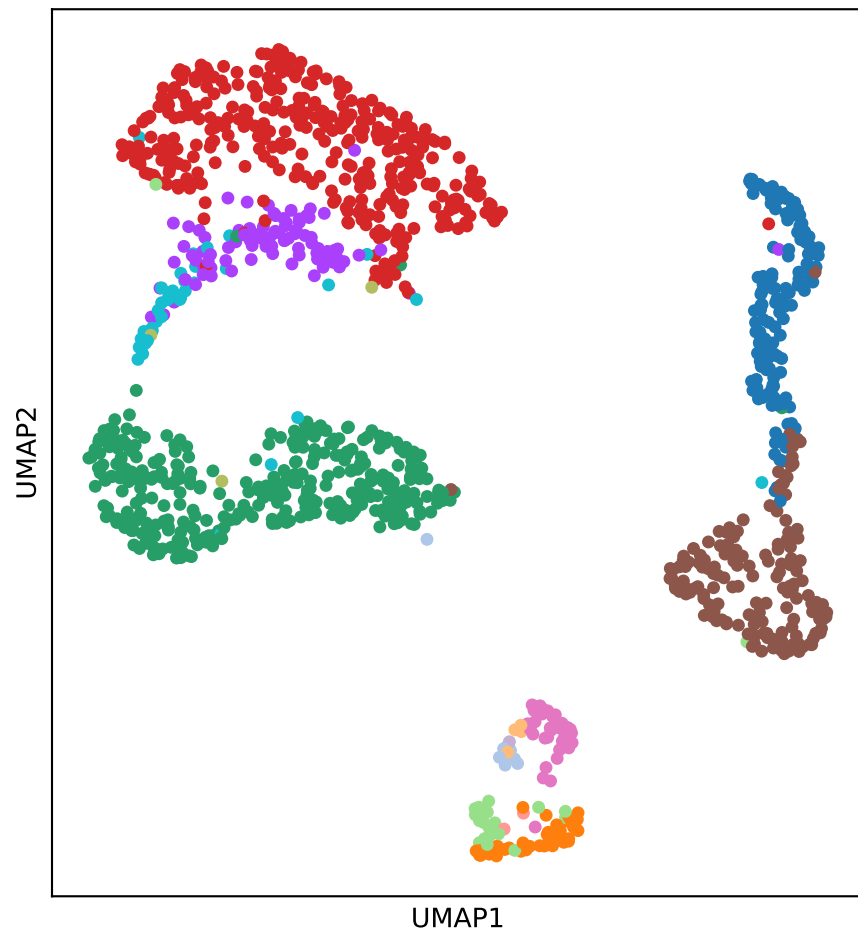
UMAP projection - Labels (val) (Res: 0.07, Iter: 10)



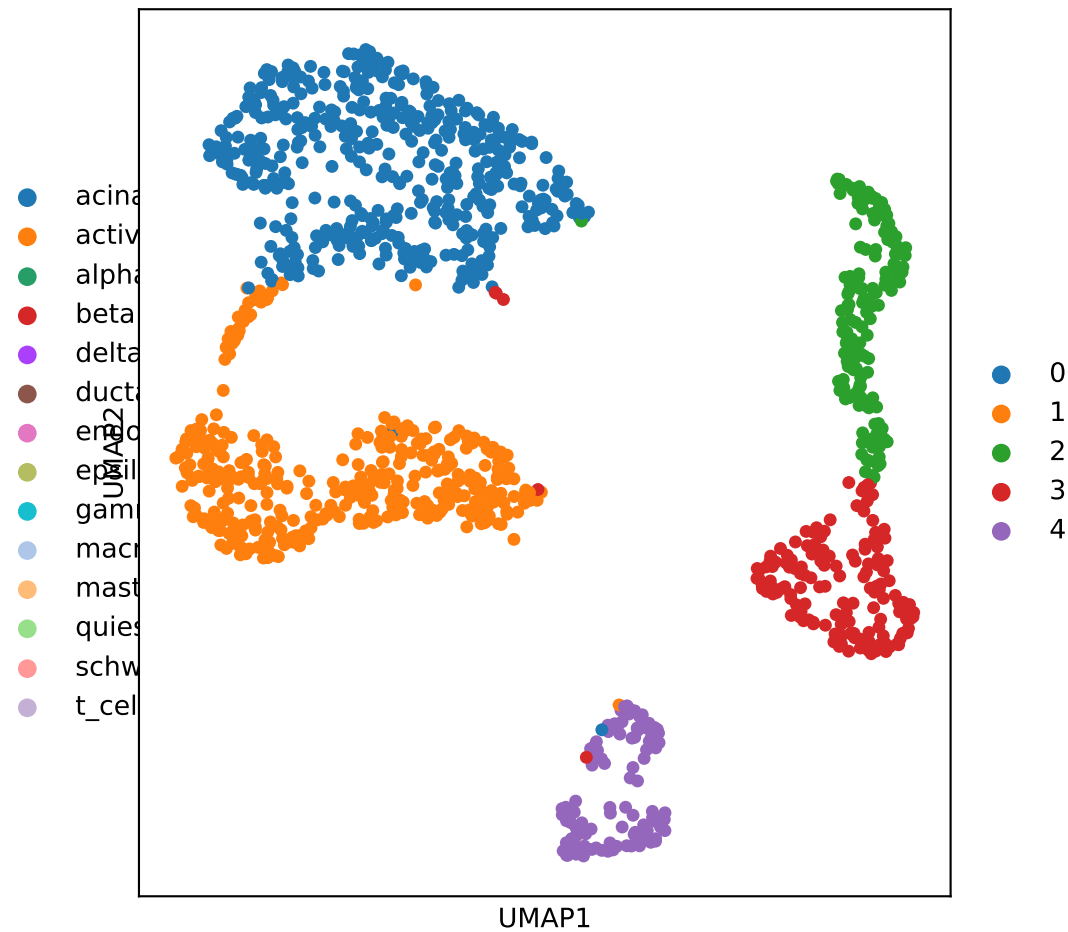
UMAP projection - Predictions (val) (Res: 0.07, Iter: 10)



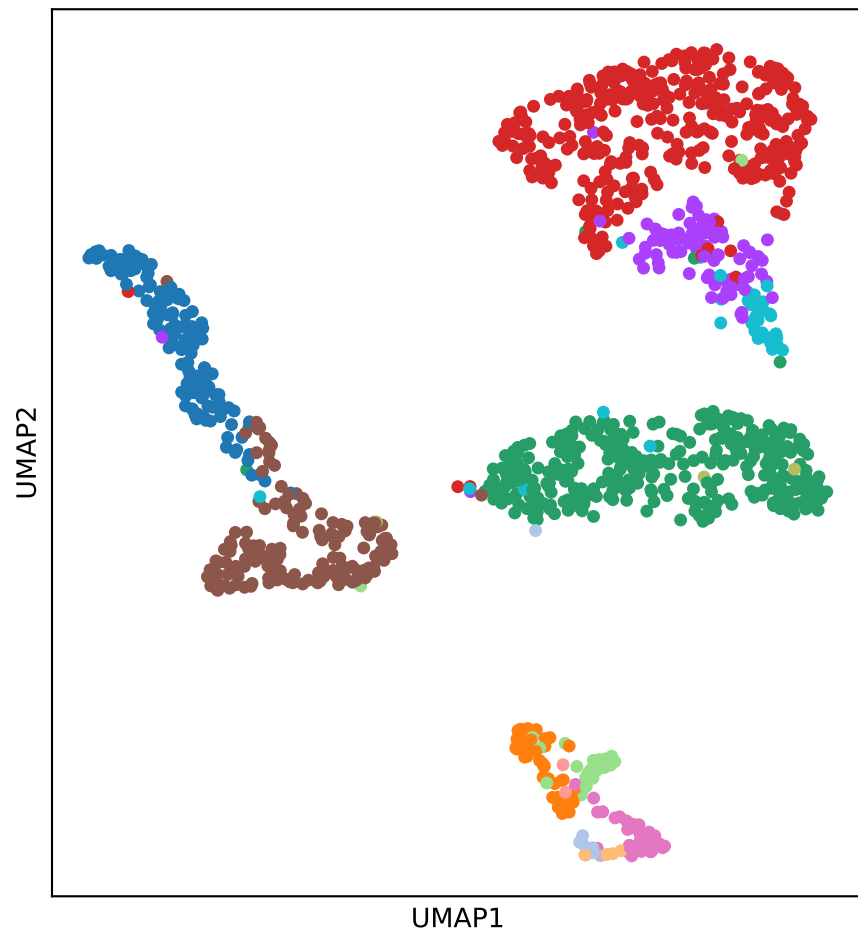
UMAP projection - Labels (val) (Res: 0.06, Iter: 11)



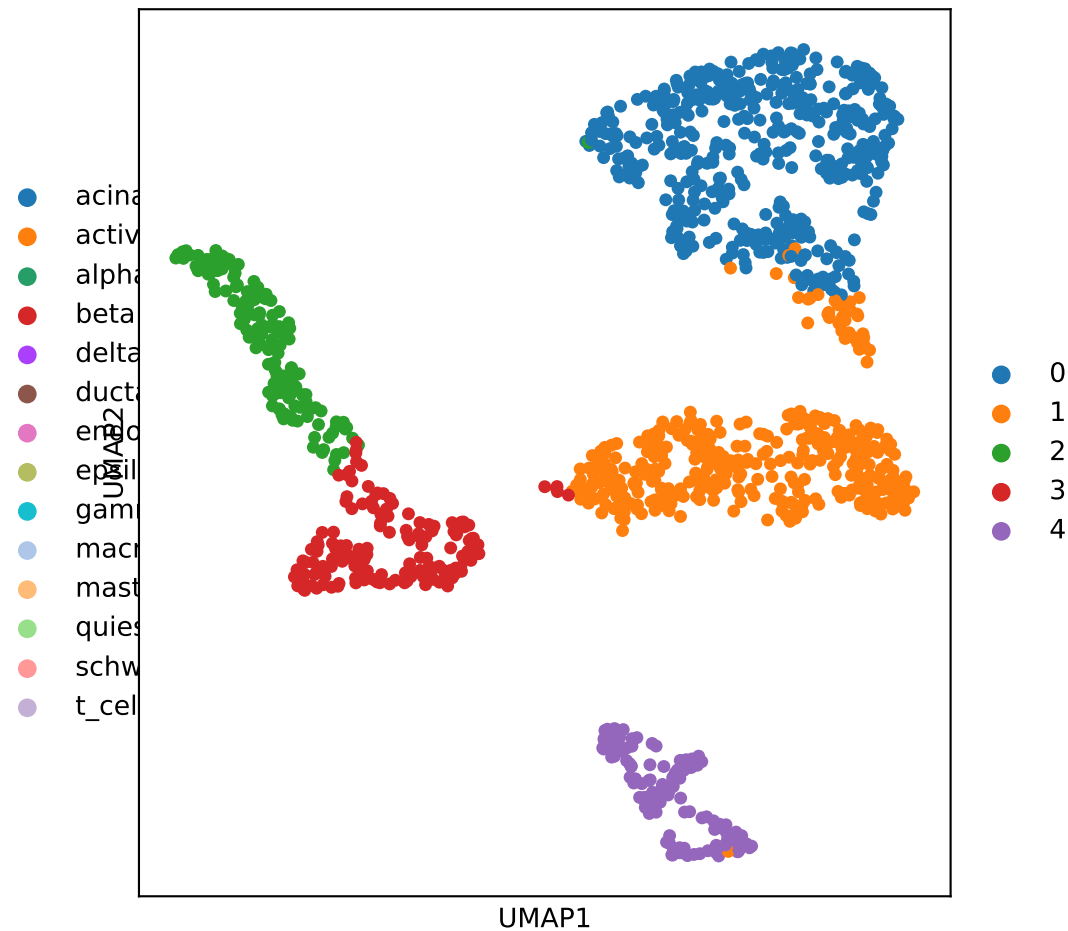
UMAP projection - Predictions (val) (Res: 0.06, Iter: 11)



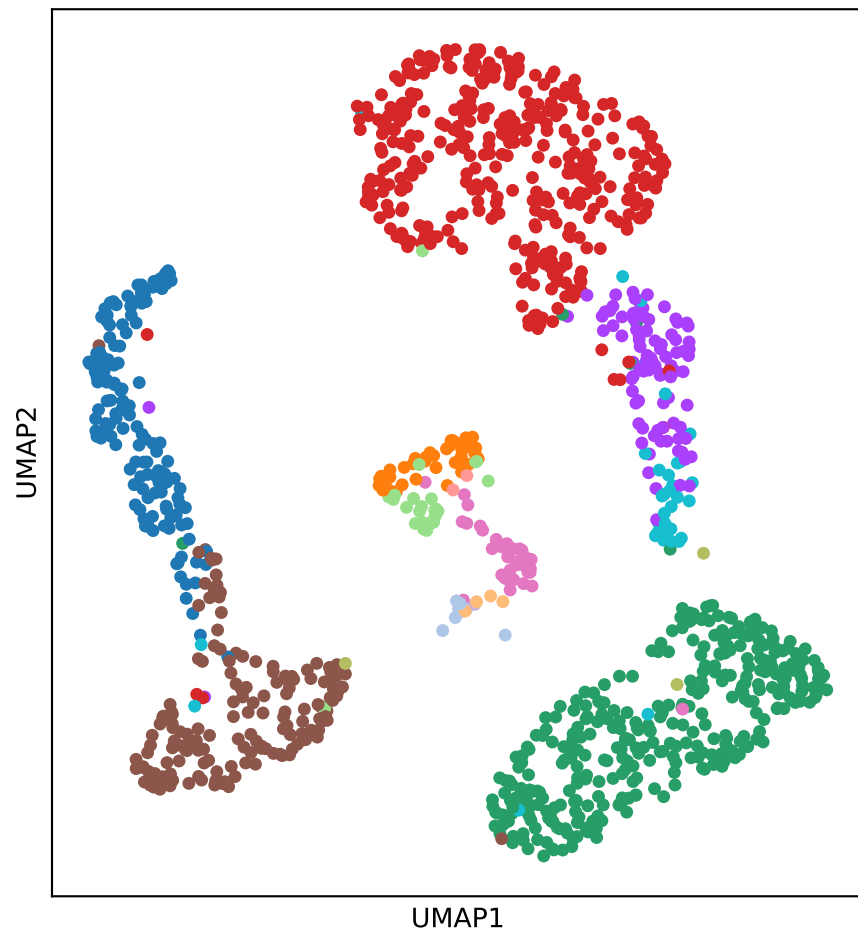
UMAP projection - Labels (val) (Res: 0.06, Iter: 12)



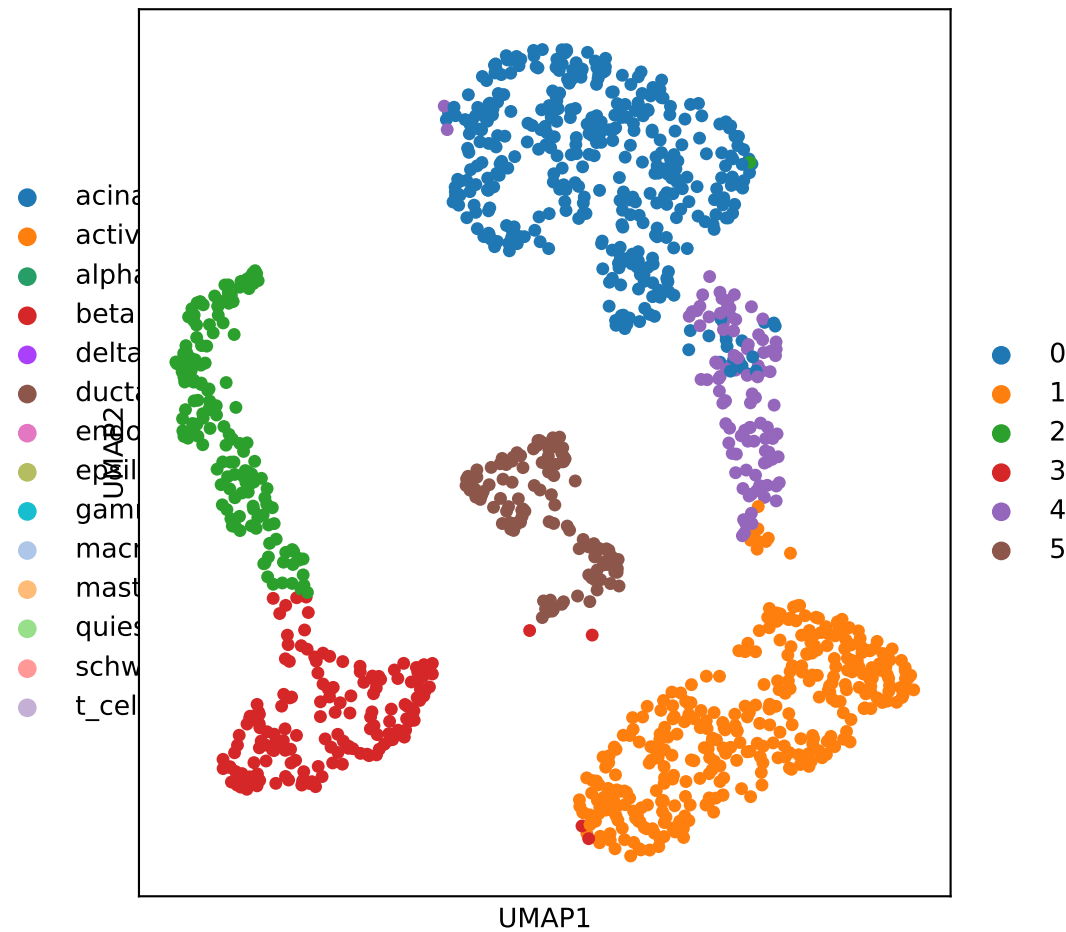
UMAP projection - Predictions (val) (Res: 0.06, Iter: 12)



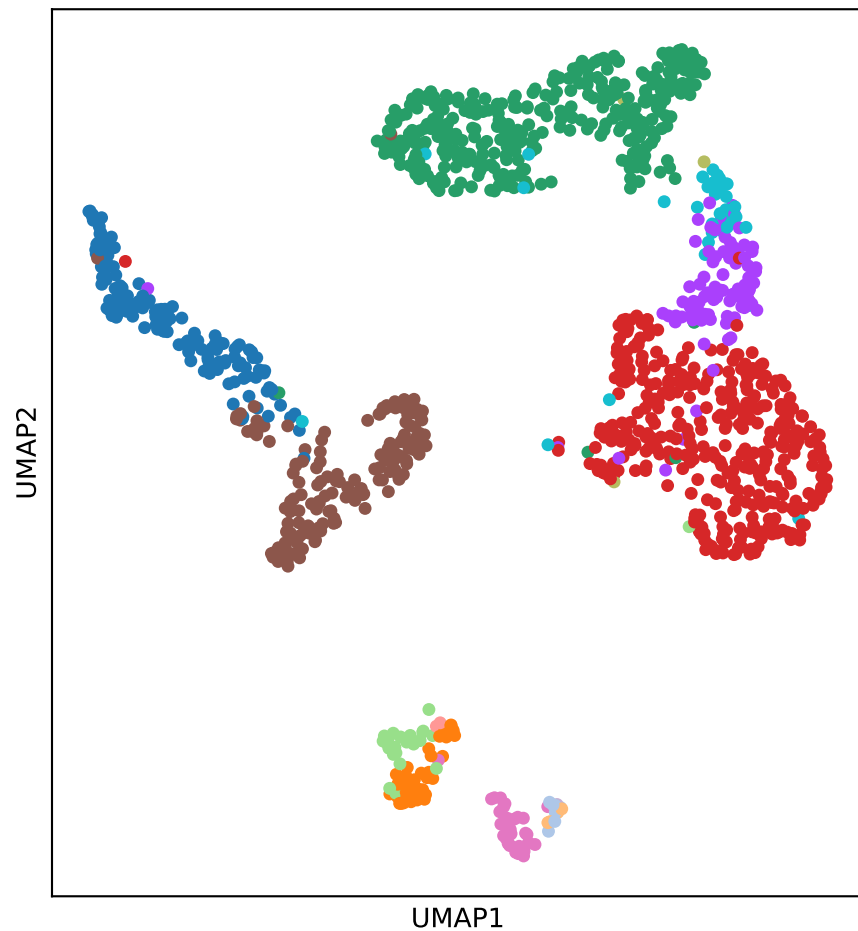
UMAP projection - Labels (val) (Res: 0.1, Iter: 13)



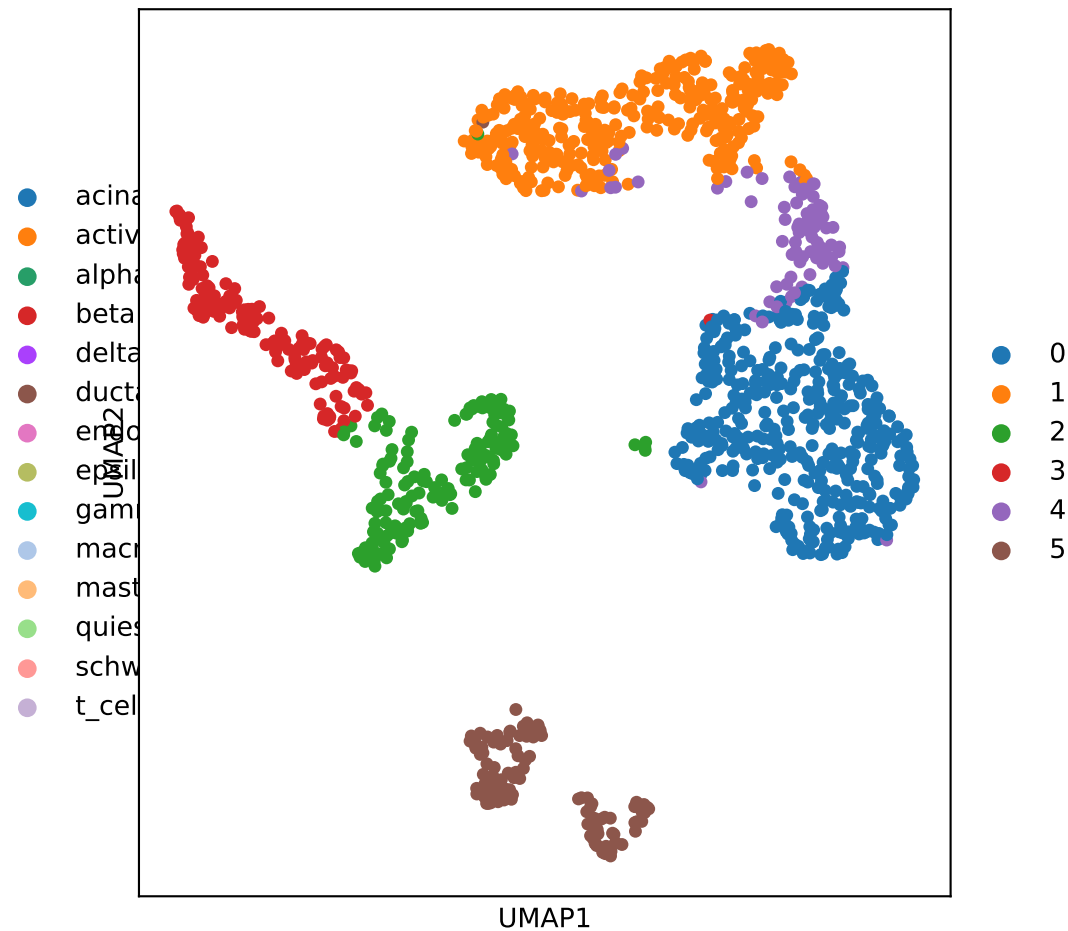
UMAP projection - Predictions (val) (Res: 0.1, Iter: 13)



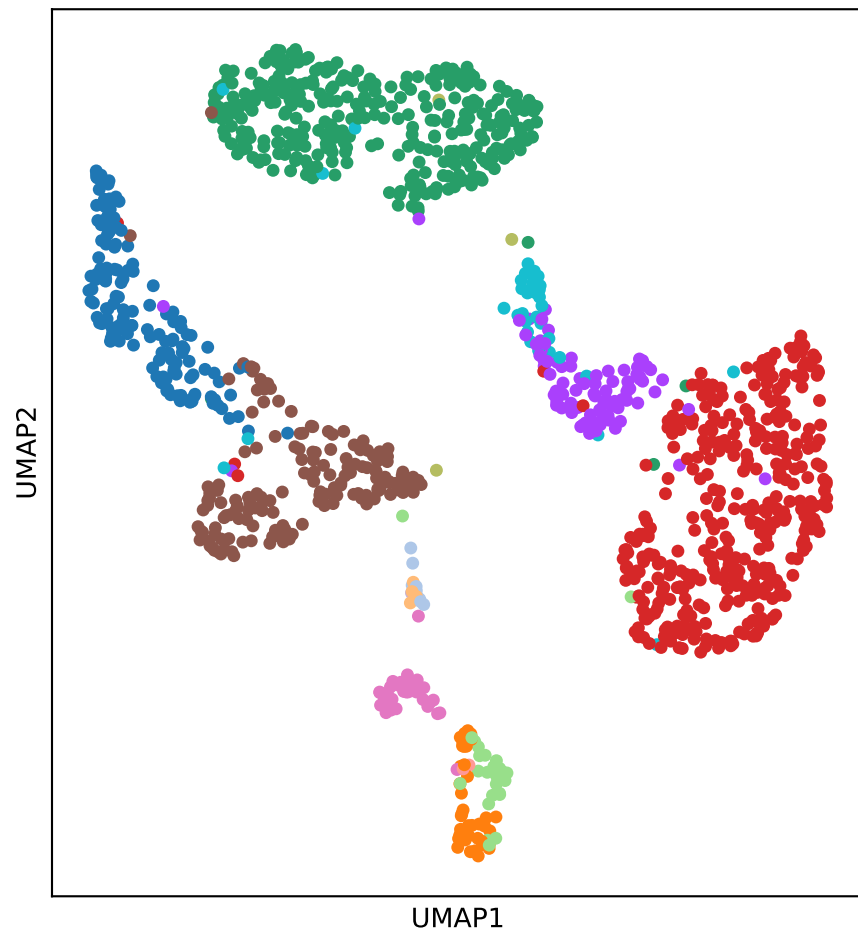
UMAP projection - Labels (val) (Res: 0.1, Iter: 14)



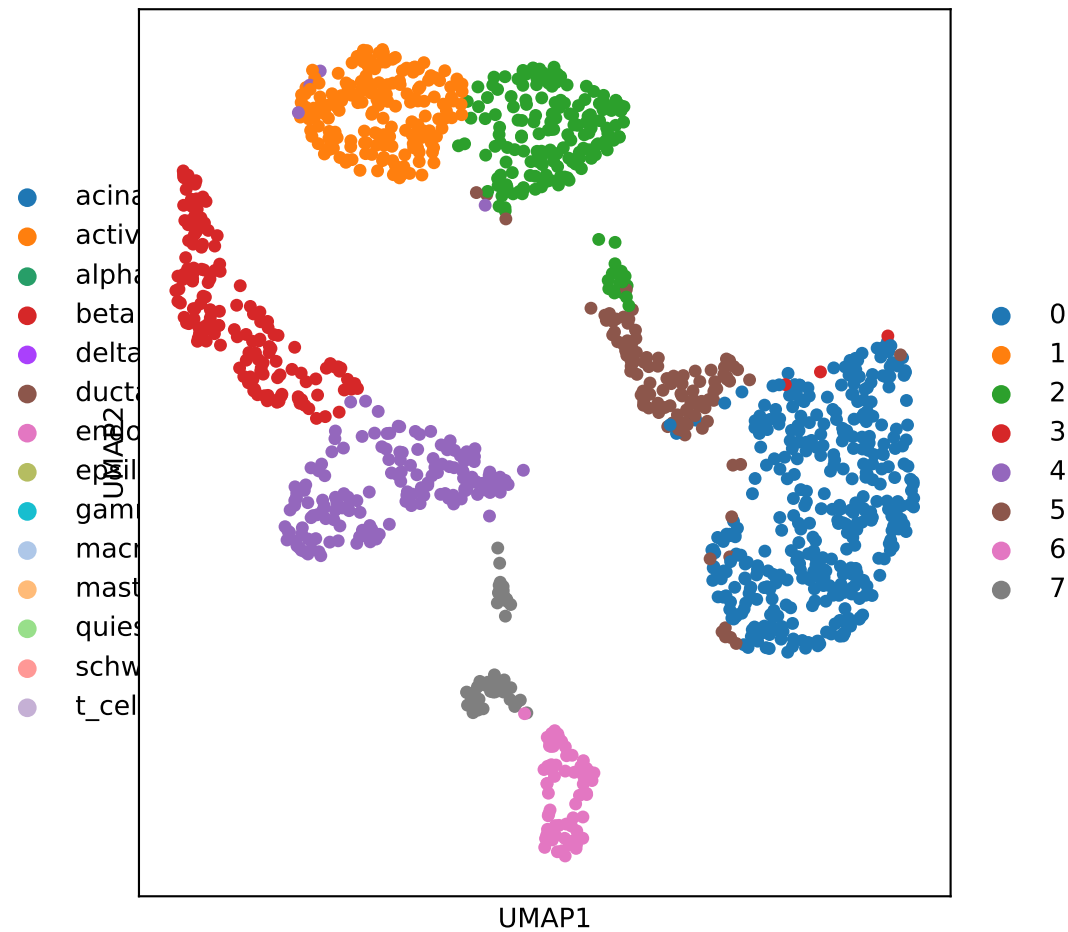
UMAP projection - Predictions (val) (Res: 0.1, Iter: 14)



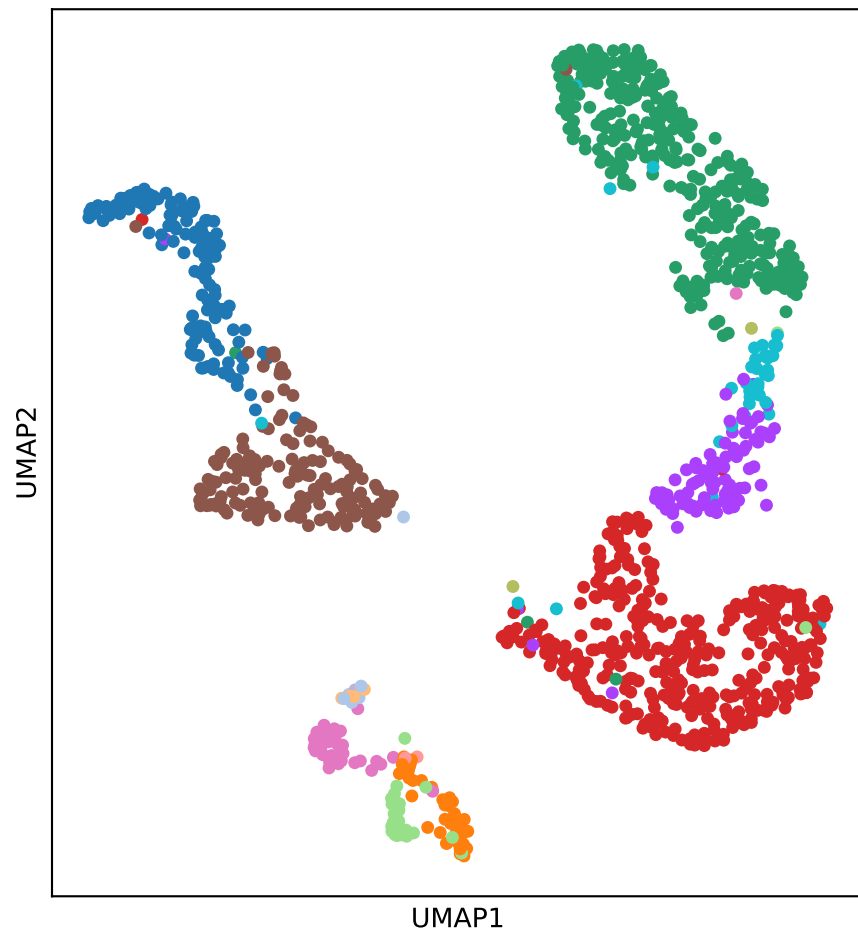
UMAP projection - Labels (val) (Res: 0.1, Iter: 15)



UMAP projection - Predictions (val) (Res: 0.1, Iter: 15)

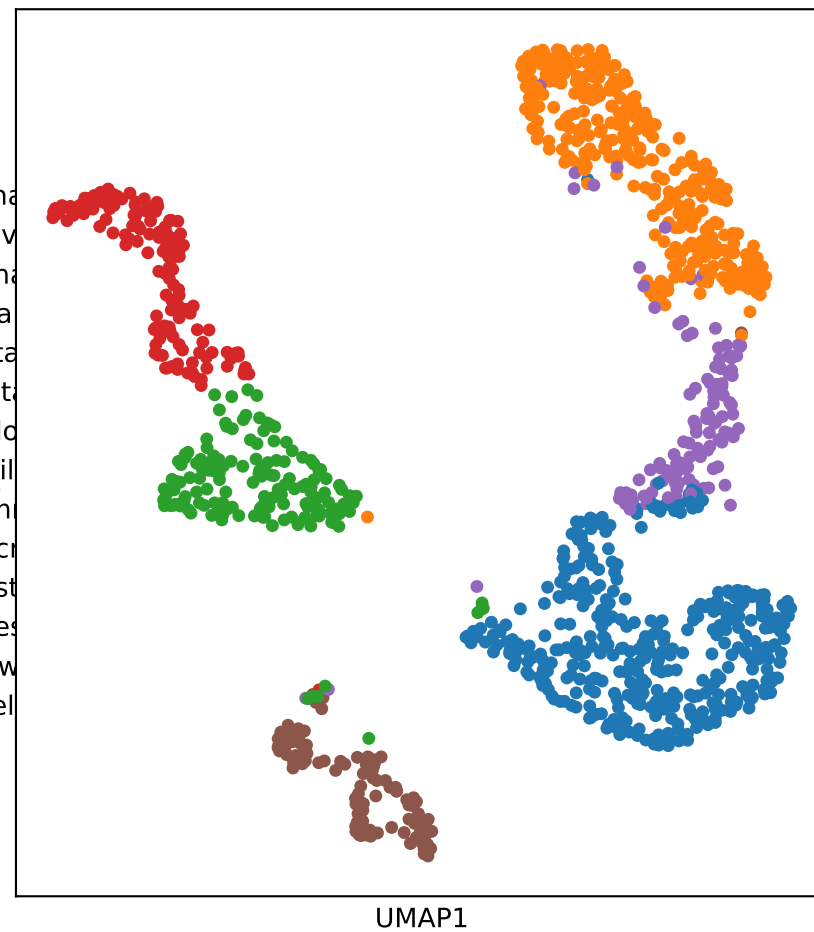


UMAP projection - Labels (val) (Res: 0.08, Iter: 16)



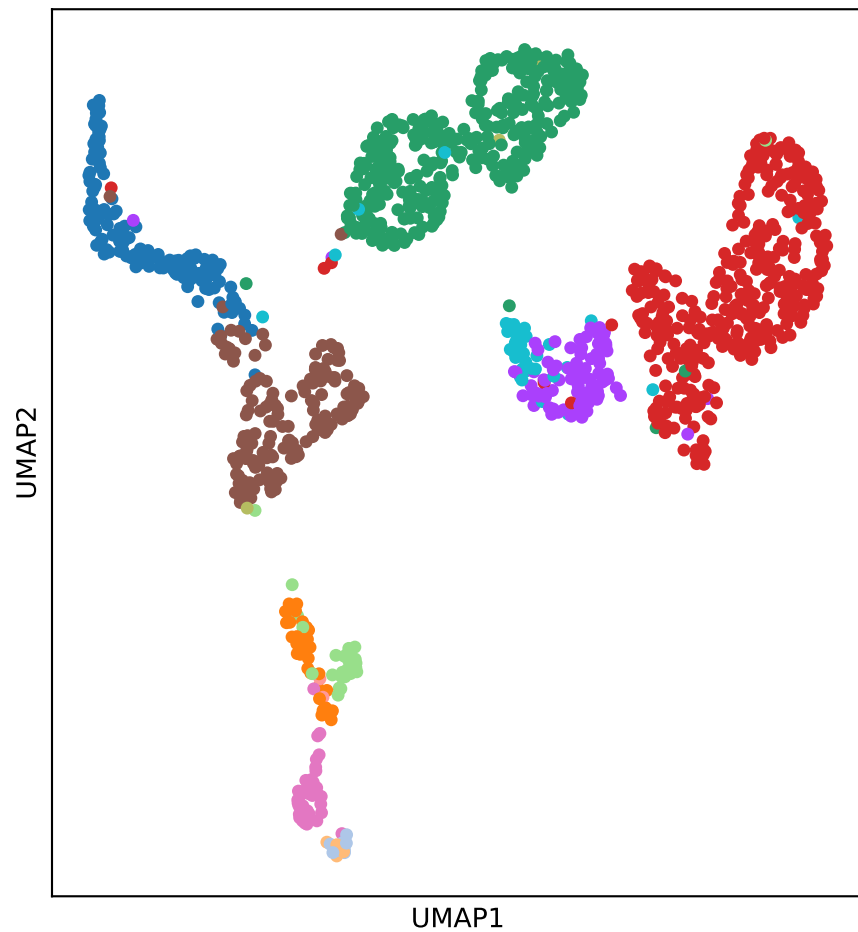
- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

UMAP projection - Predictions (val) (Res: 0.08, Iter: 16)

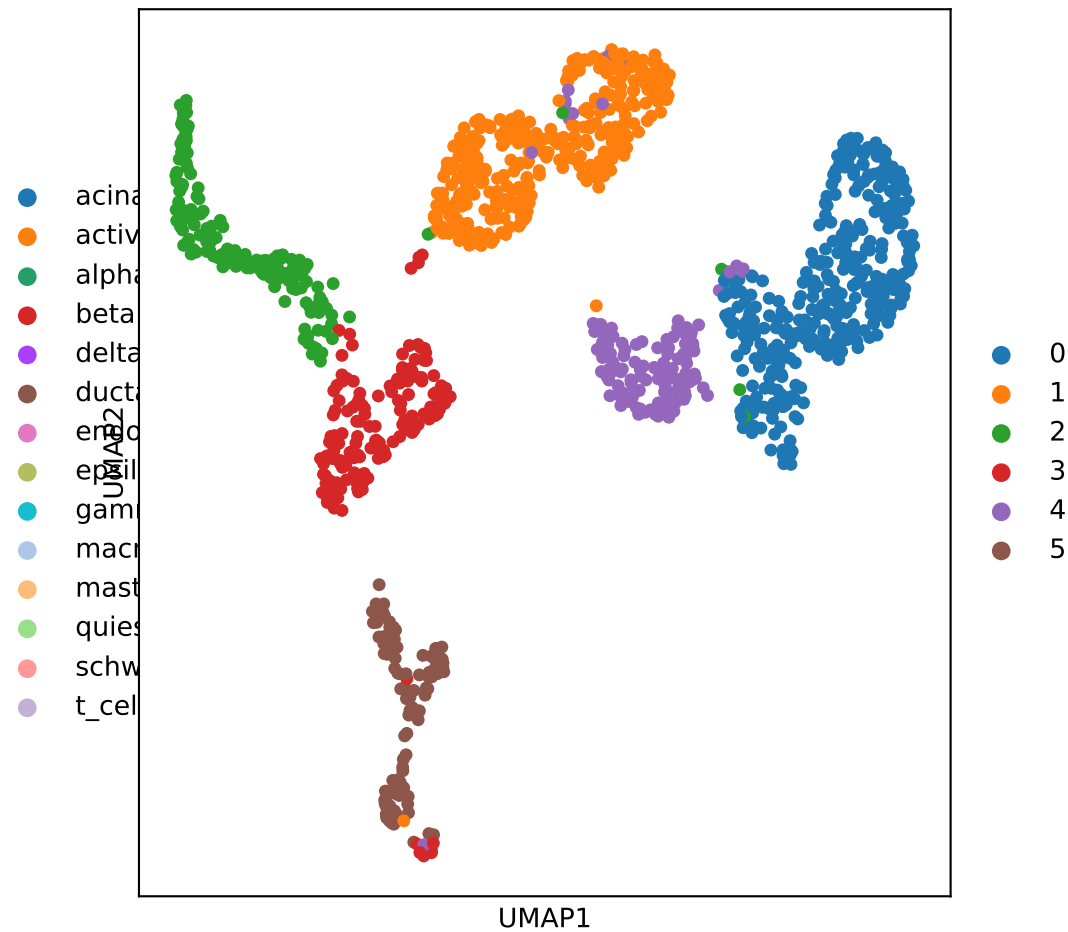


- 0
- 1
- 2
- 3
- 4
- 5

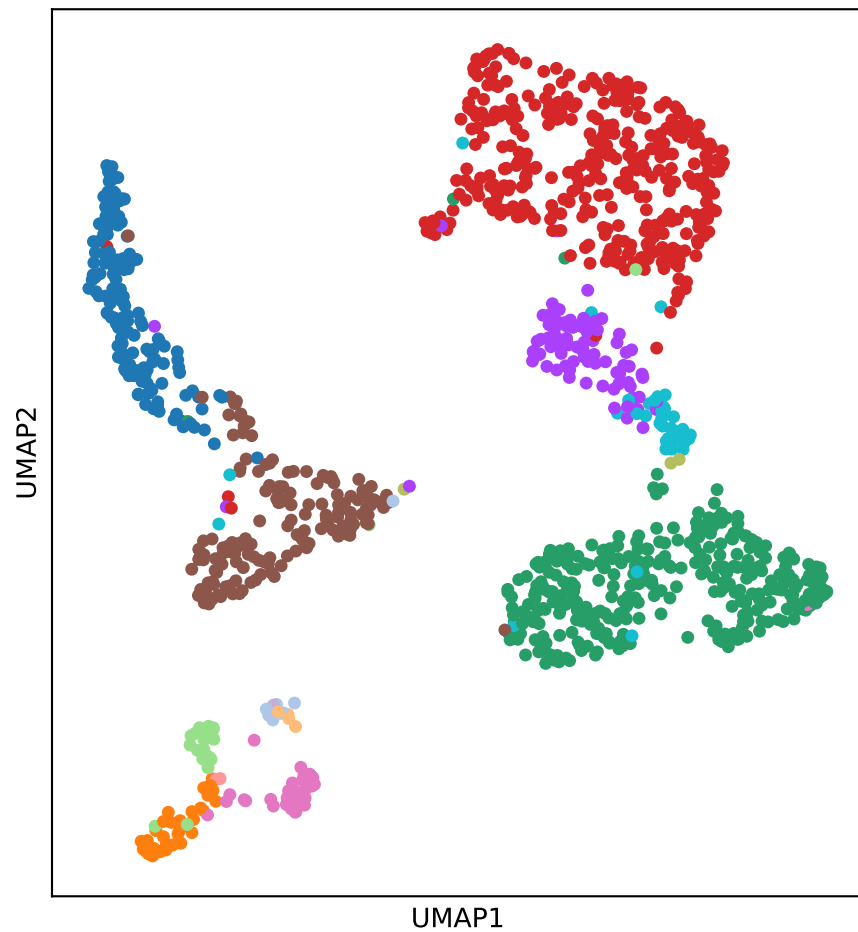
UMAP projection - Labels (val) (Res: 0.08, Iter: 17)



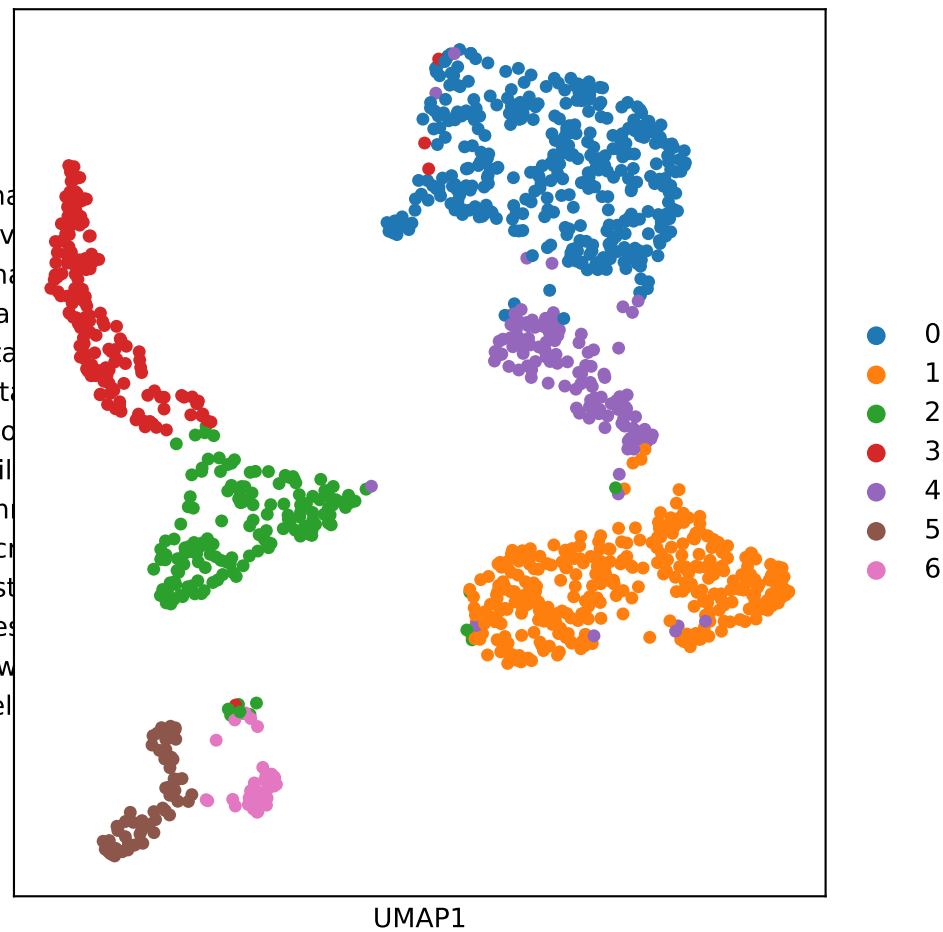
UMAP projection - Predictions (val) (Res: 0.08, Iter: 17)



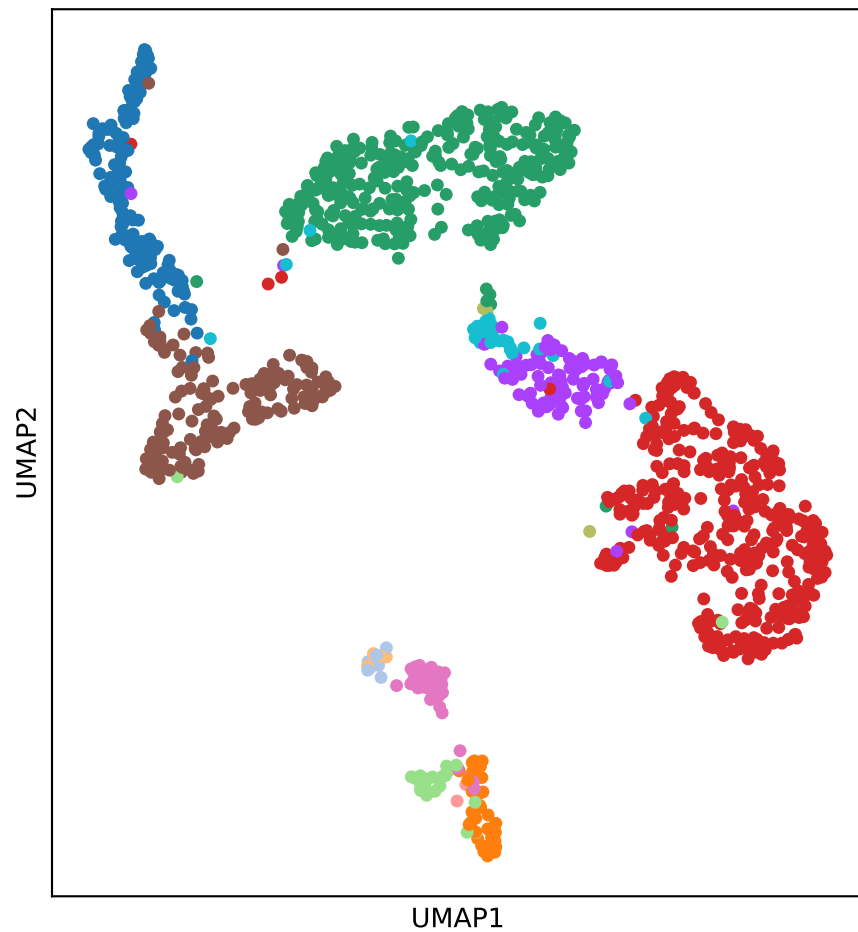
UMAP projection - Labels (val) (Res: 0.07, Iter: 18)



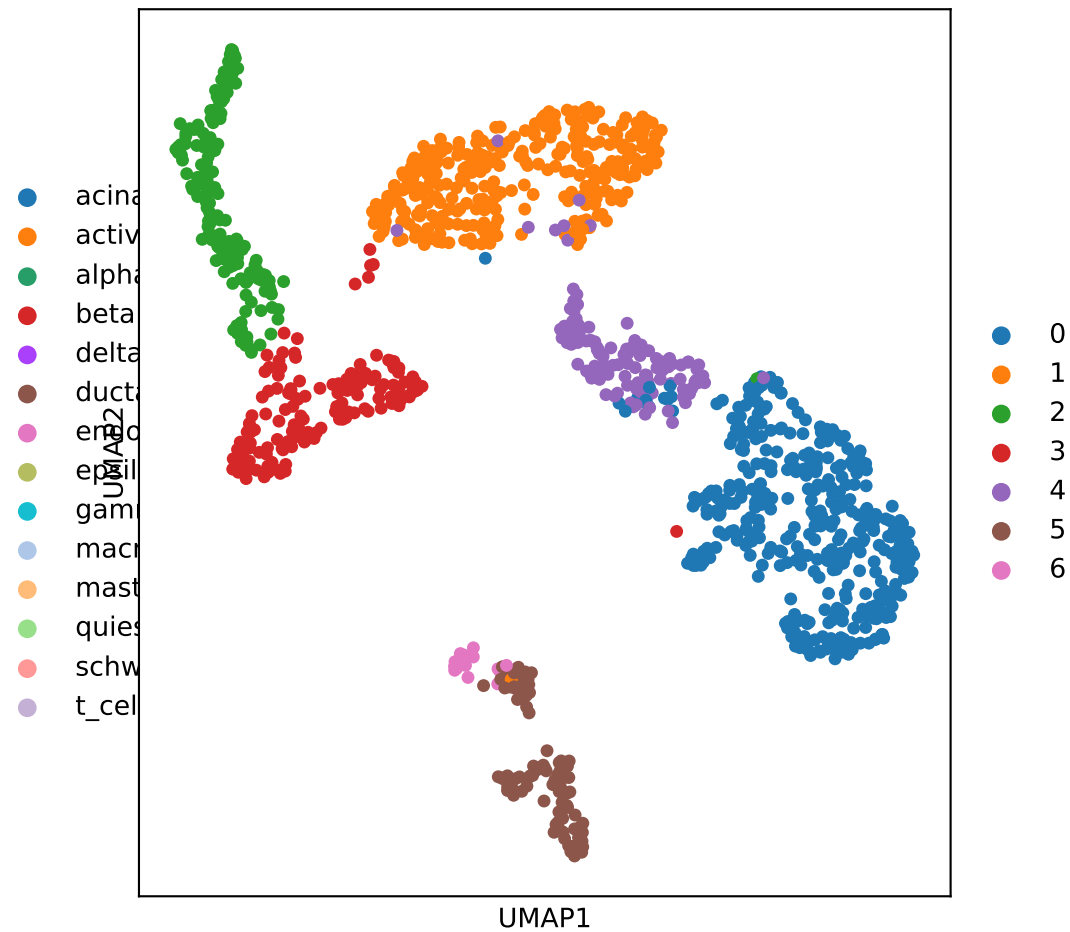
UMAP projection - Predictions (val) (Res: 0.07, Iter: 18)



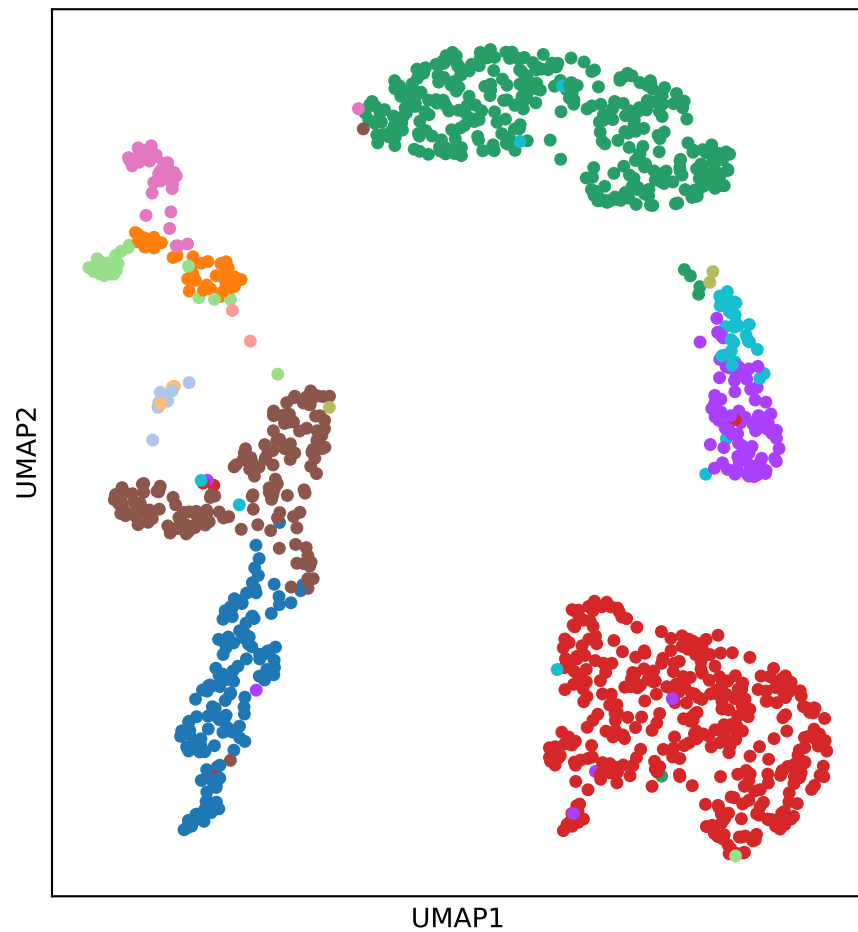
UMAP projection - Labels (val) (Res: 0.08, Iter: 19)



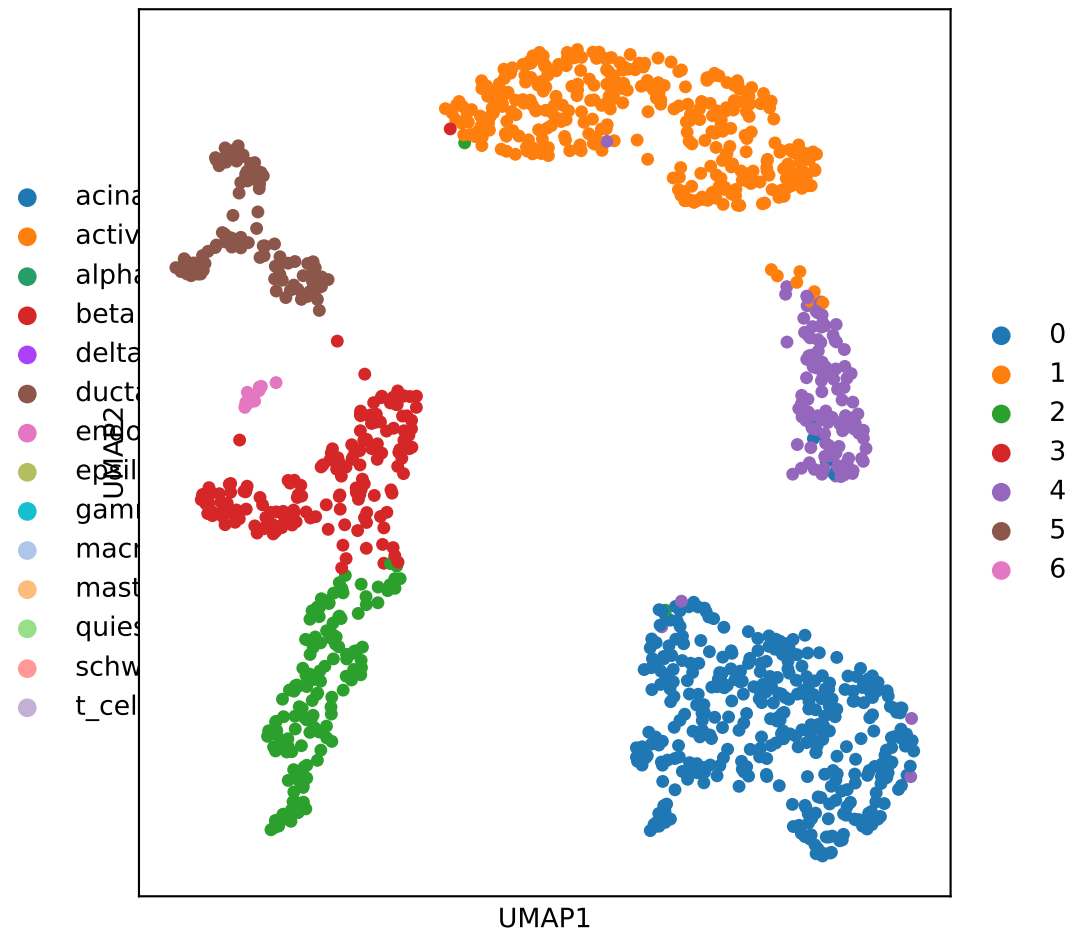
UMAP projection - Predictions (val) (Res: 0.08, Iter: 19)



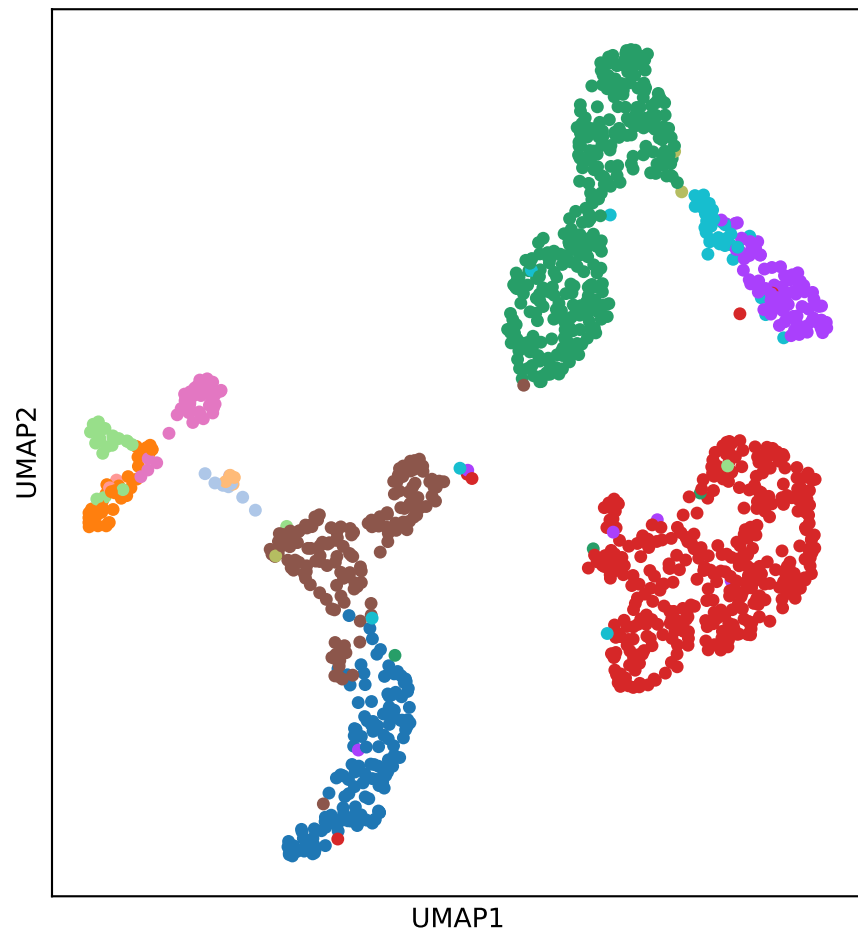
UMAP projection - Labels (val) (Res: 0.06, Iter: 20)



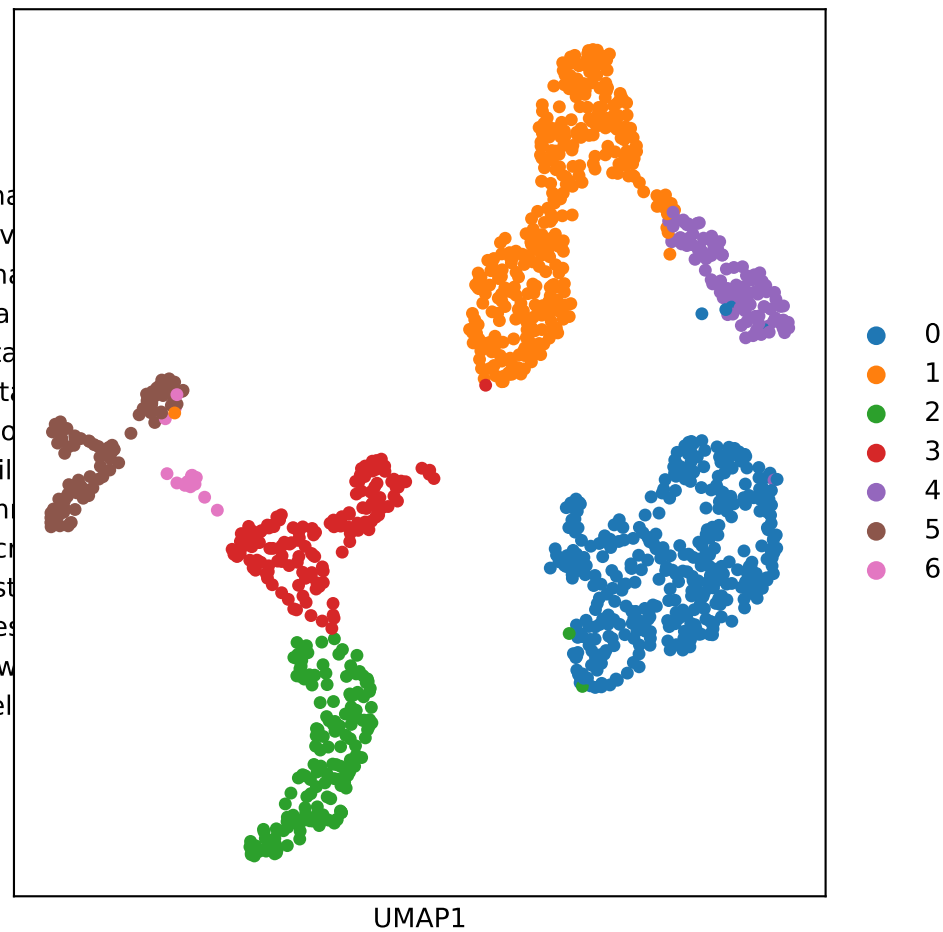
UMAP projection - Predictions (val) (Res: 0.06, Iter: 20)



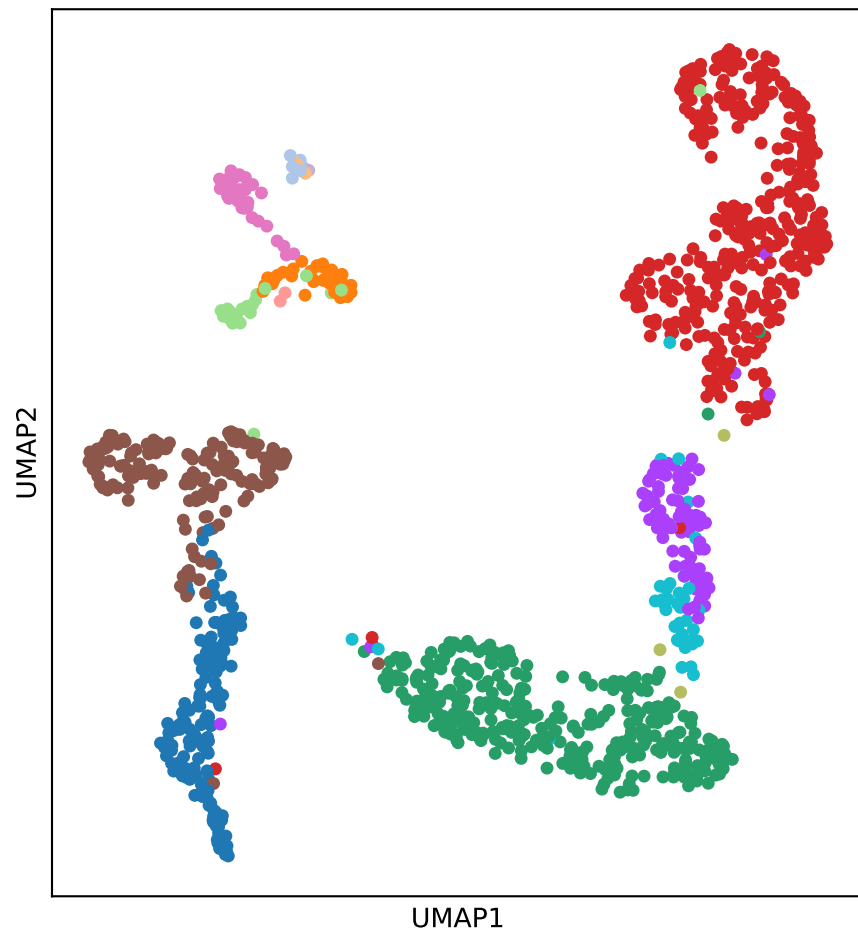
UMAP projection - Labels (val) (Res: 0.09, Iter: 21)



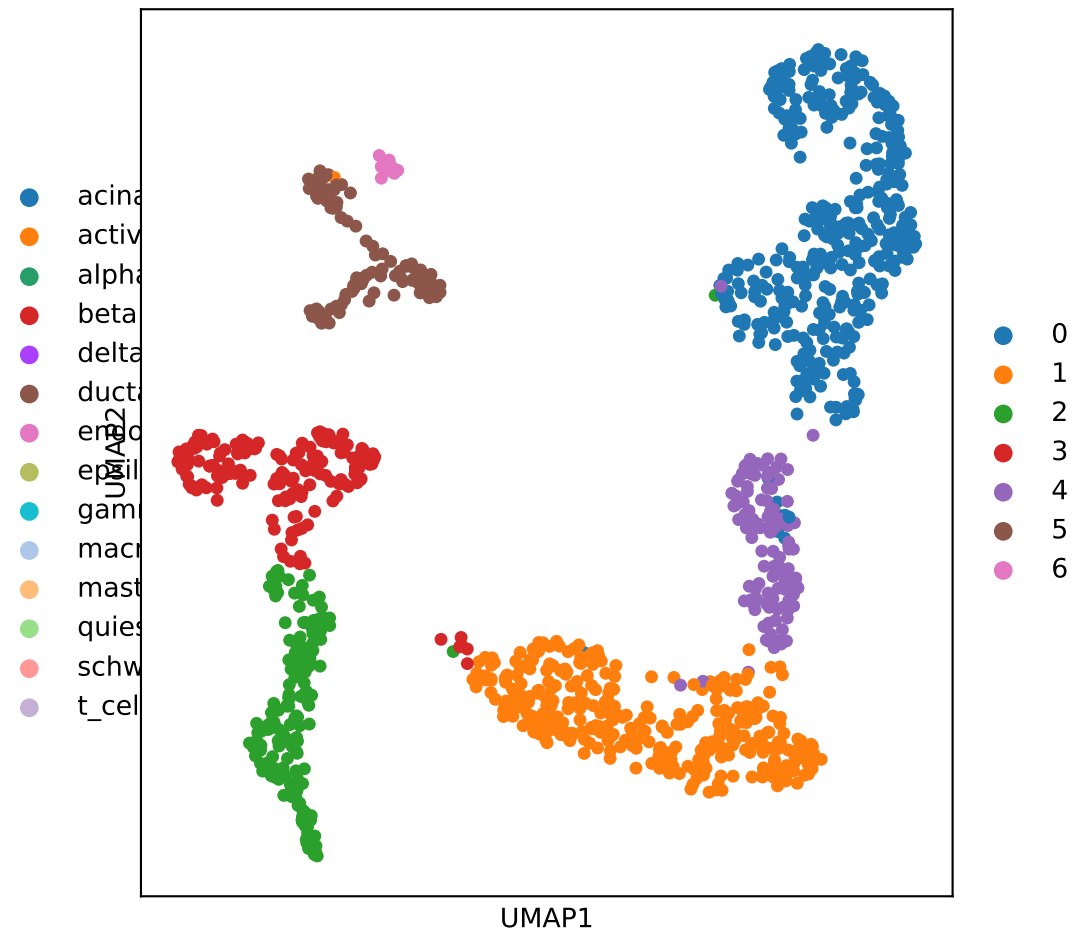
UMAP projection - Predictions (val) (Res: 0.09, Iter: 21)



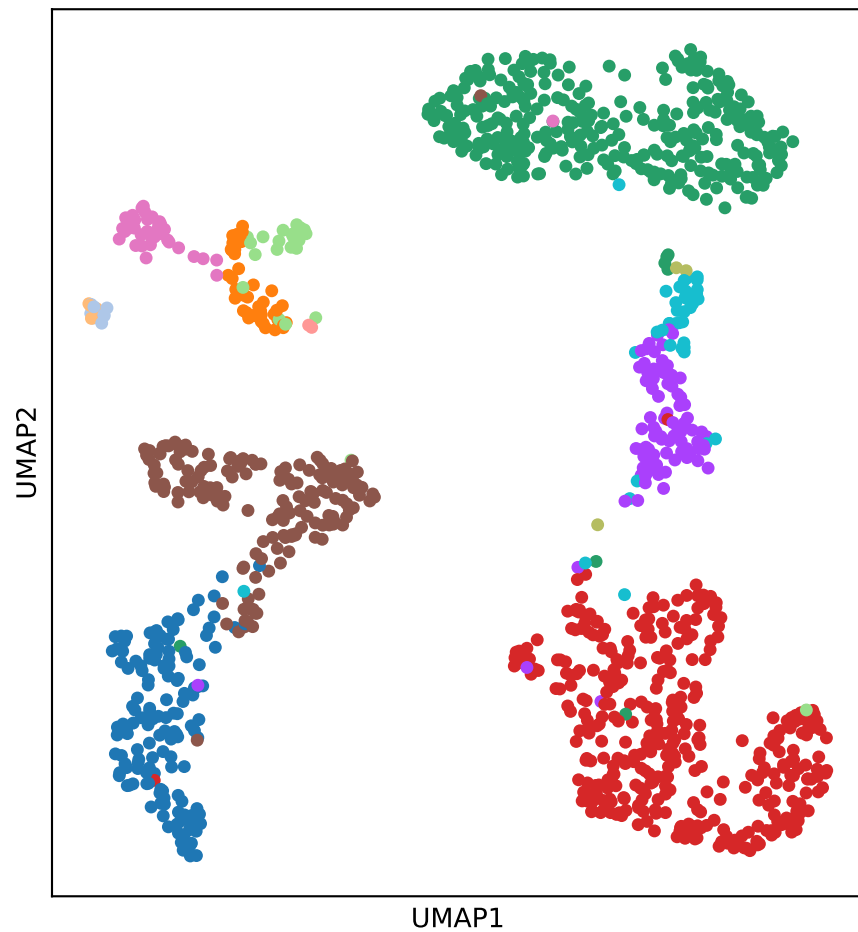
UMAP projection - Labels (val) (Res: 0.07, Iter: 22)



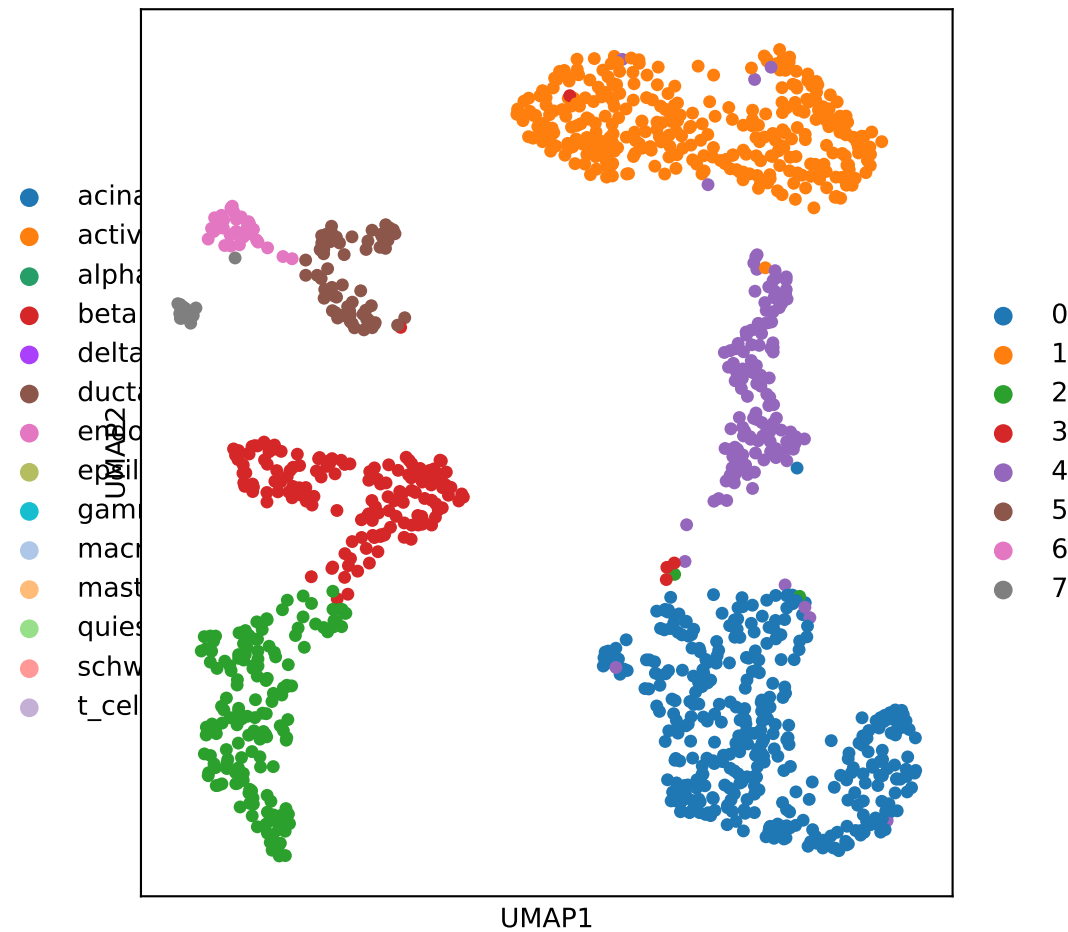
UMAP projection - Predictions (val) (Res: 0.07, Iter: 22)



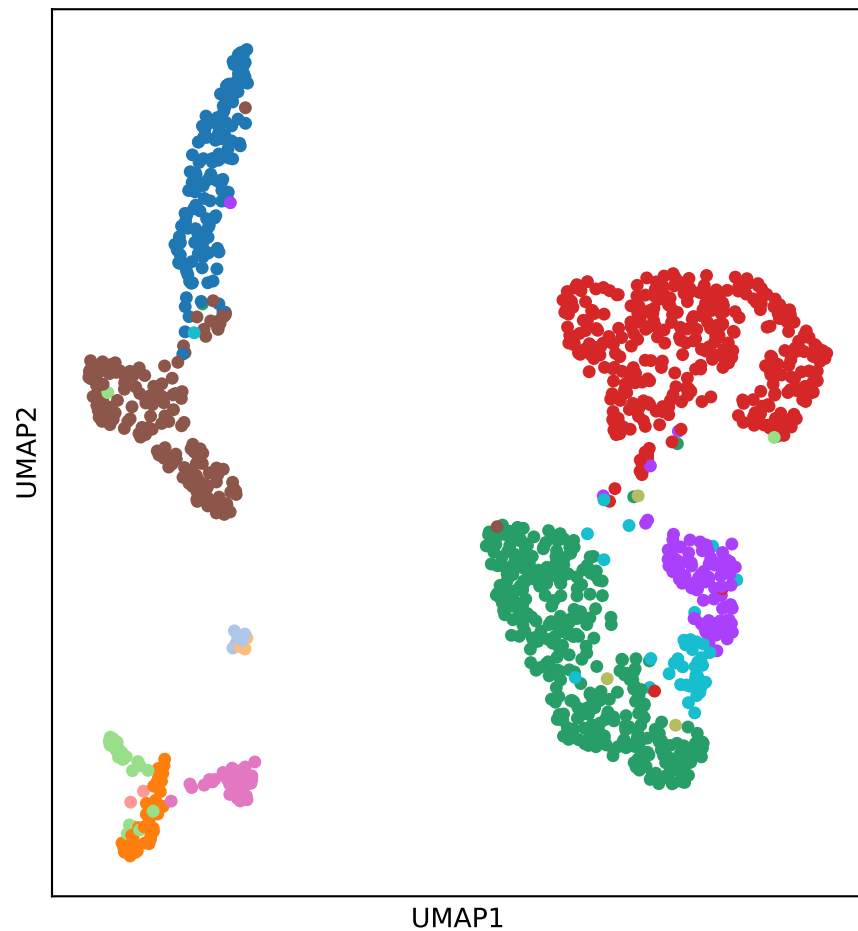
UMAP projection - Labels (val) (Res: 0.08, Iter: 23)



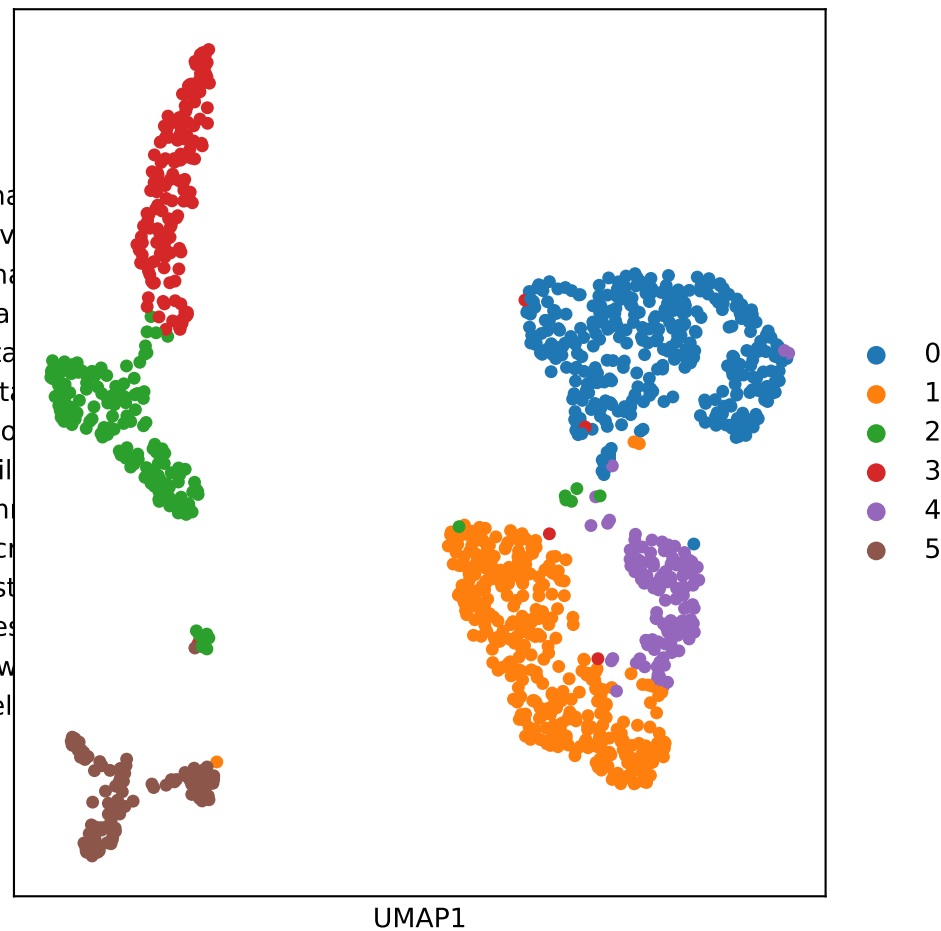
UMAP projection - Predictions (val) (Res: 0.08, Iter: 23)



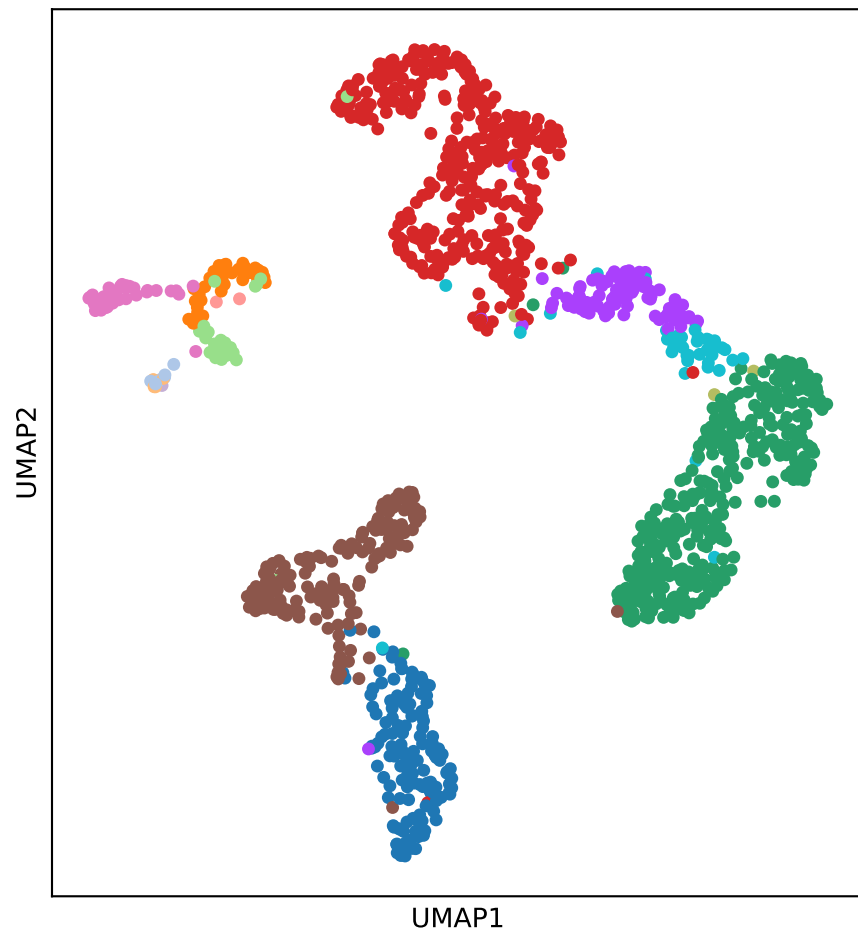
UMAP projection - Labels (val) (Res: 0.04, Iter: 24)



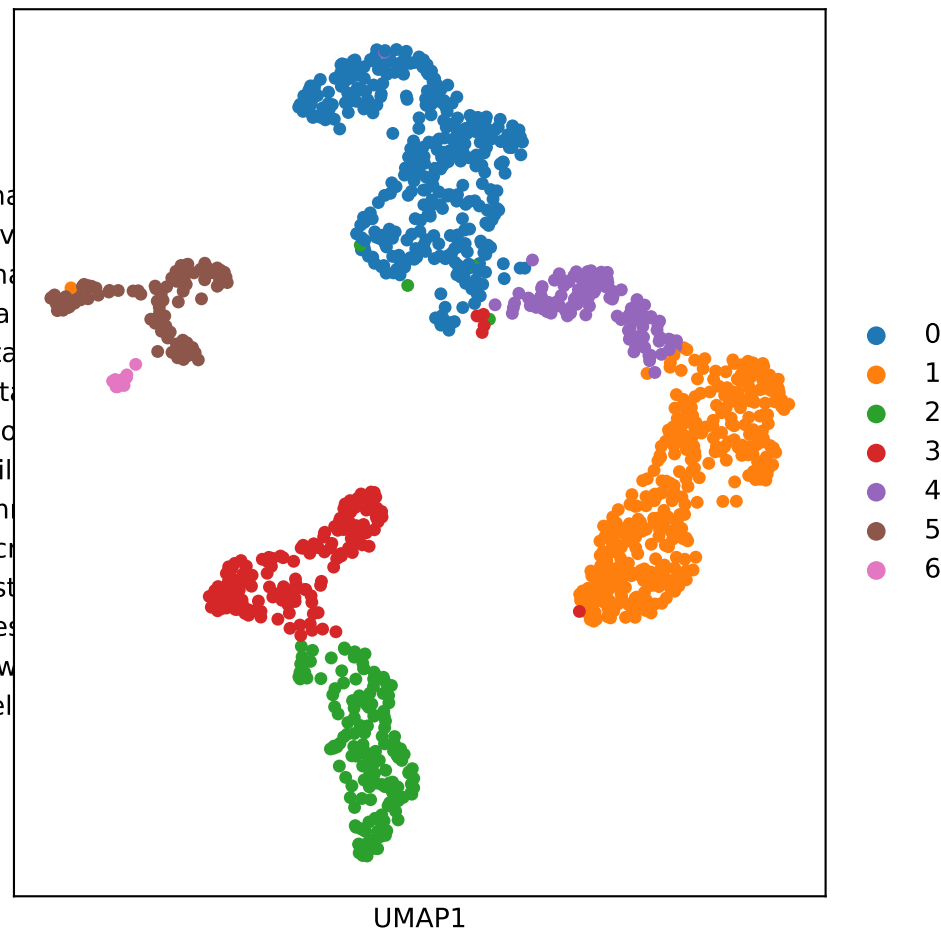
UMAP projection - Predictions (val) (Res: 0.04, Iter: 24)



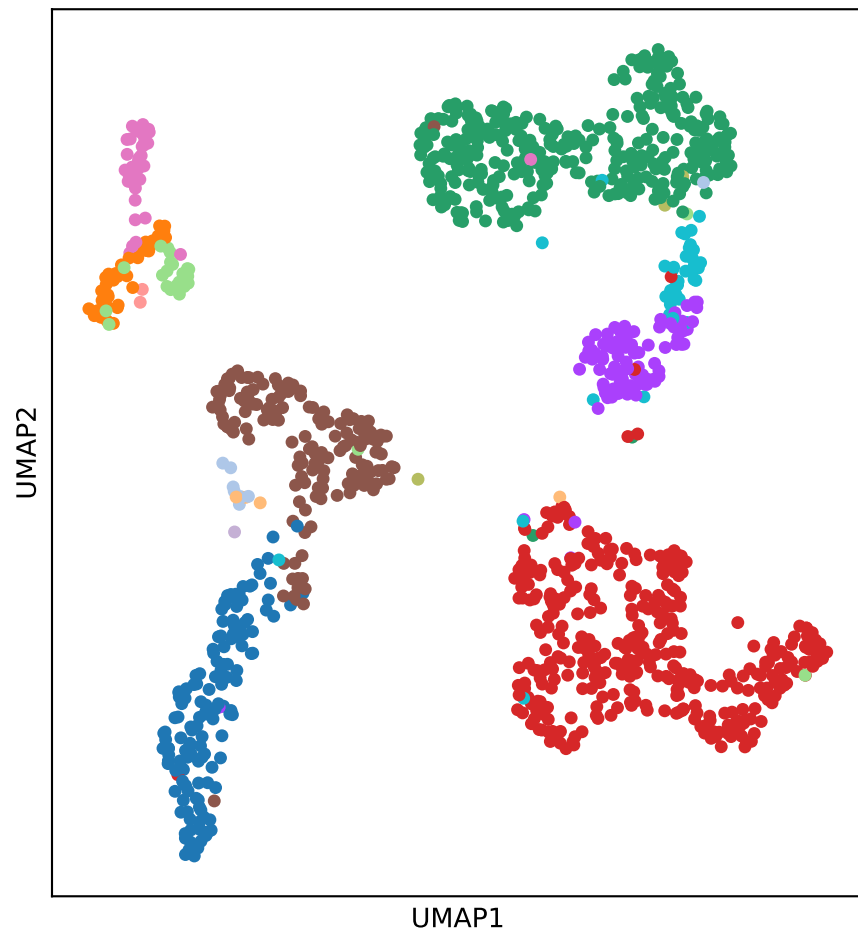
UMAP projection - Labels (val) (Res: 0.06, Iter: 25)



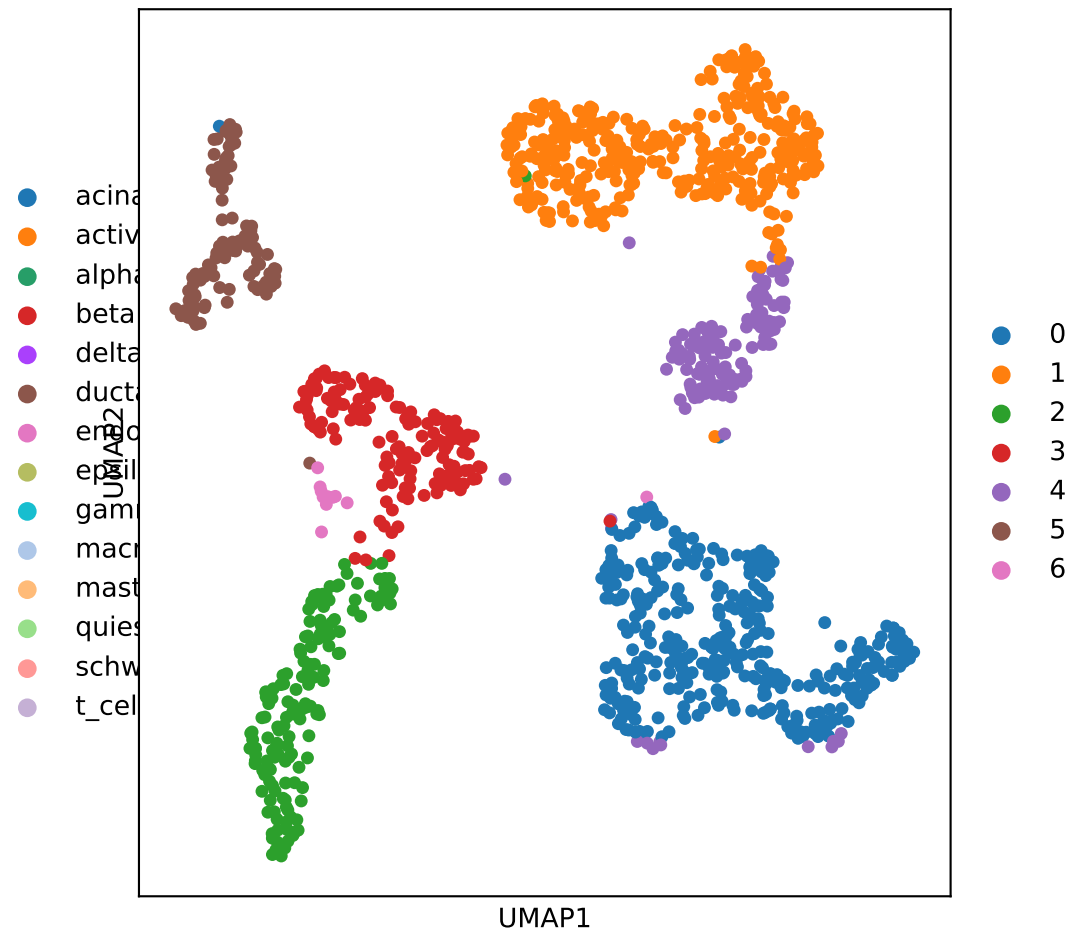
UMAP projection - Predictions (val) (Res: 0.06, Iter: 25)



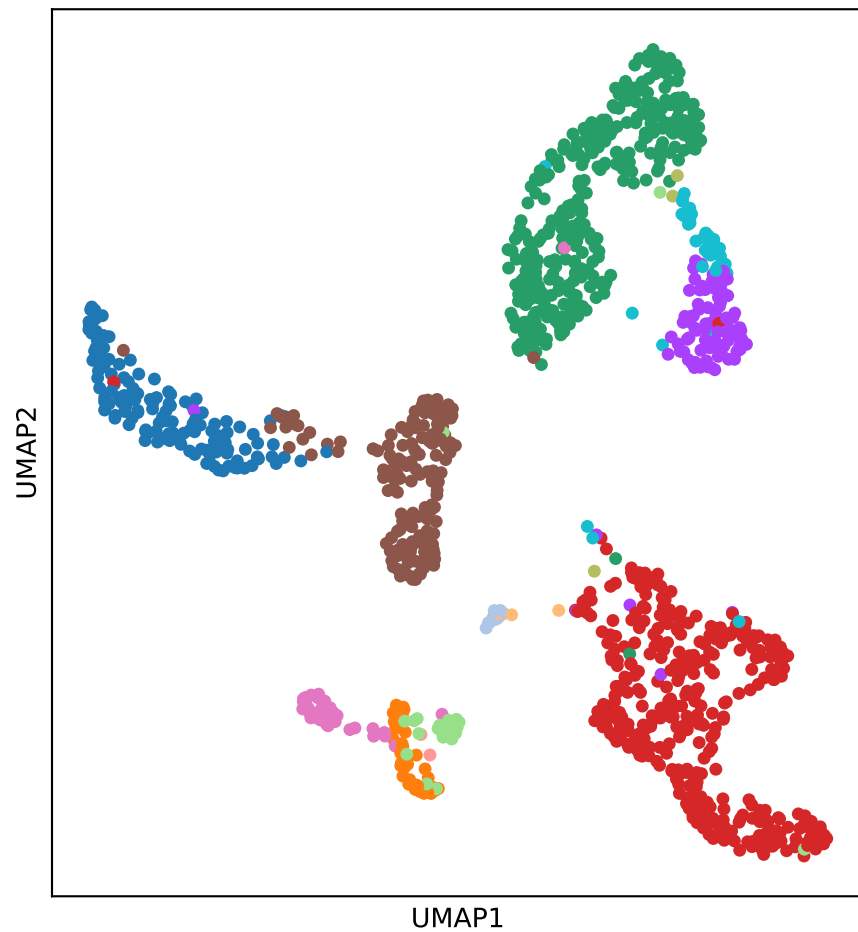
UMAP projection - Labels (val) (Res: 0.08, Iter: 26)



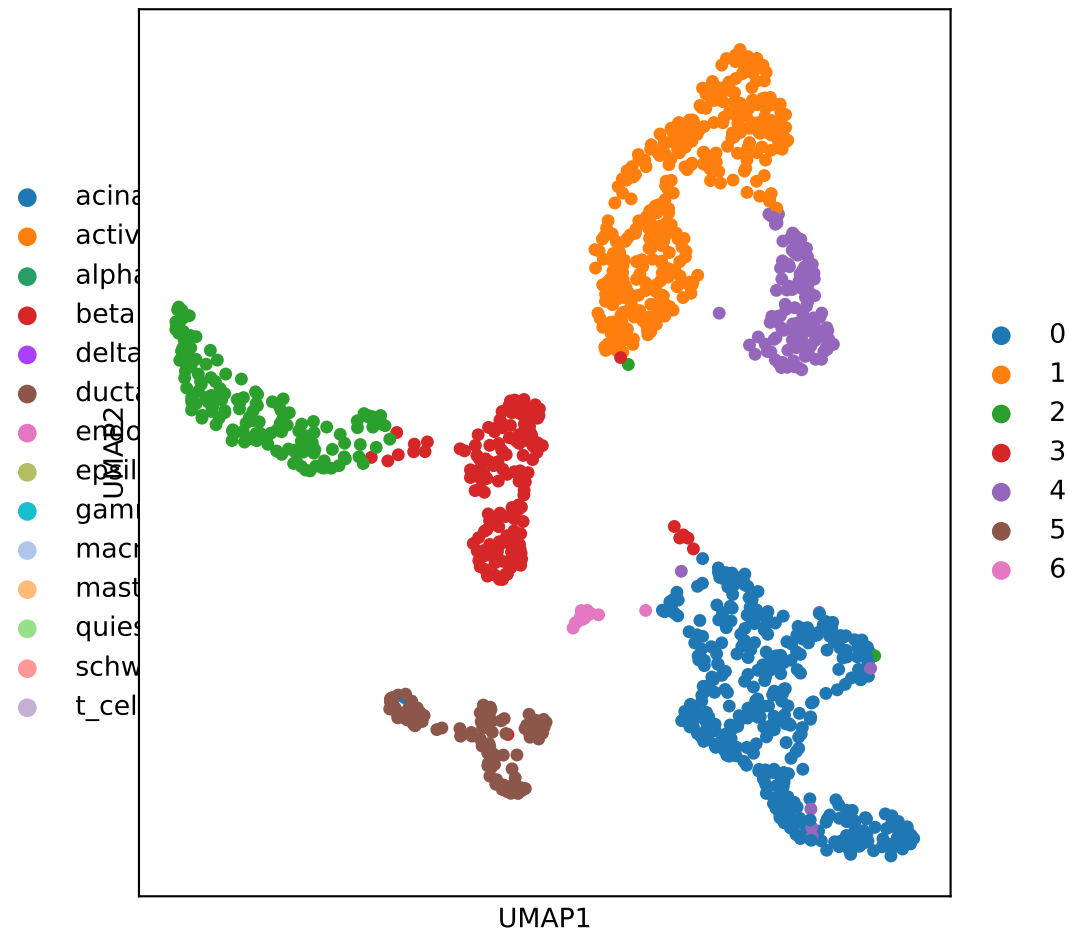
UMAP projection - Predictions (val) (Res: 0.08, Iter: 26)



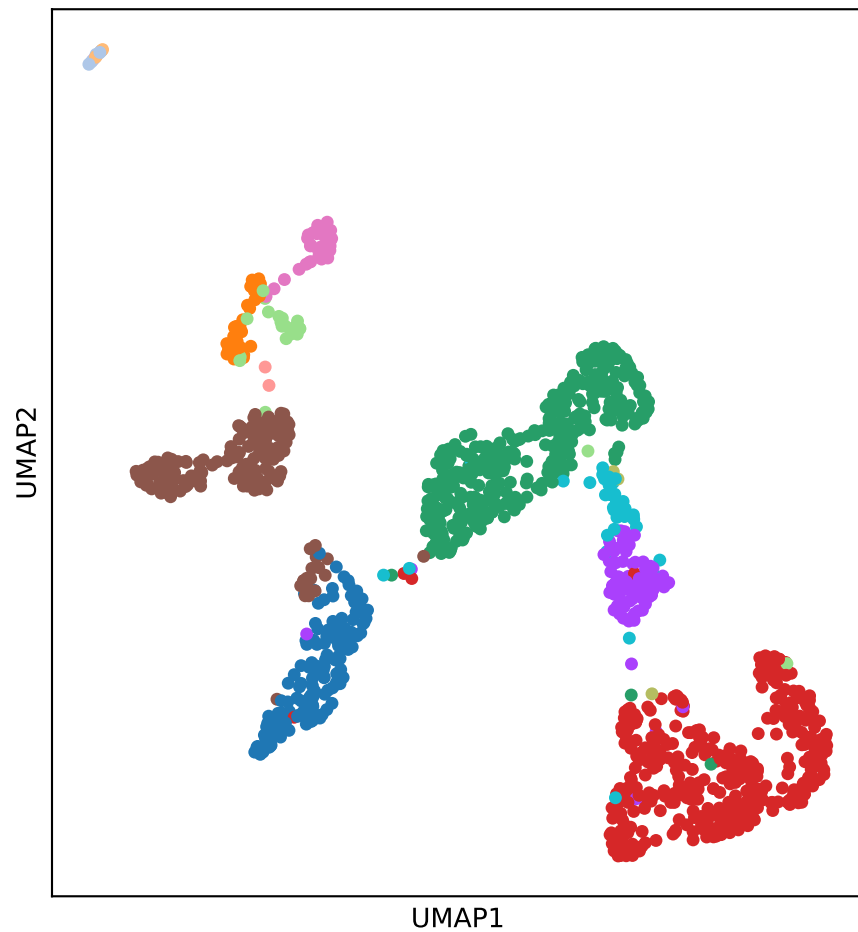
UMAP projection - Labels (val) (Res: 0.09, Iter: 27)



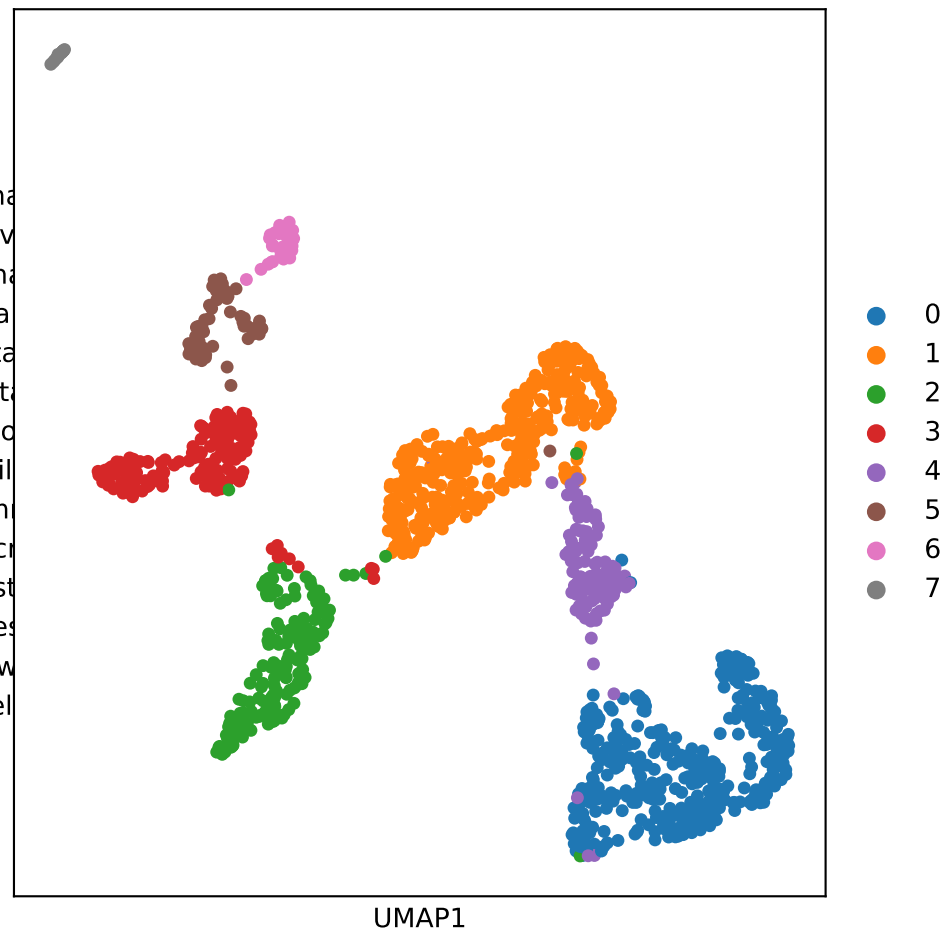
UMAP projection - Predictions (val) (Res: 0.09, Iter: 27)



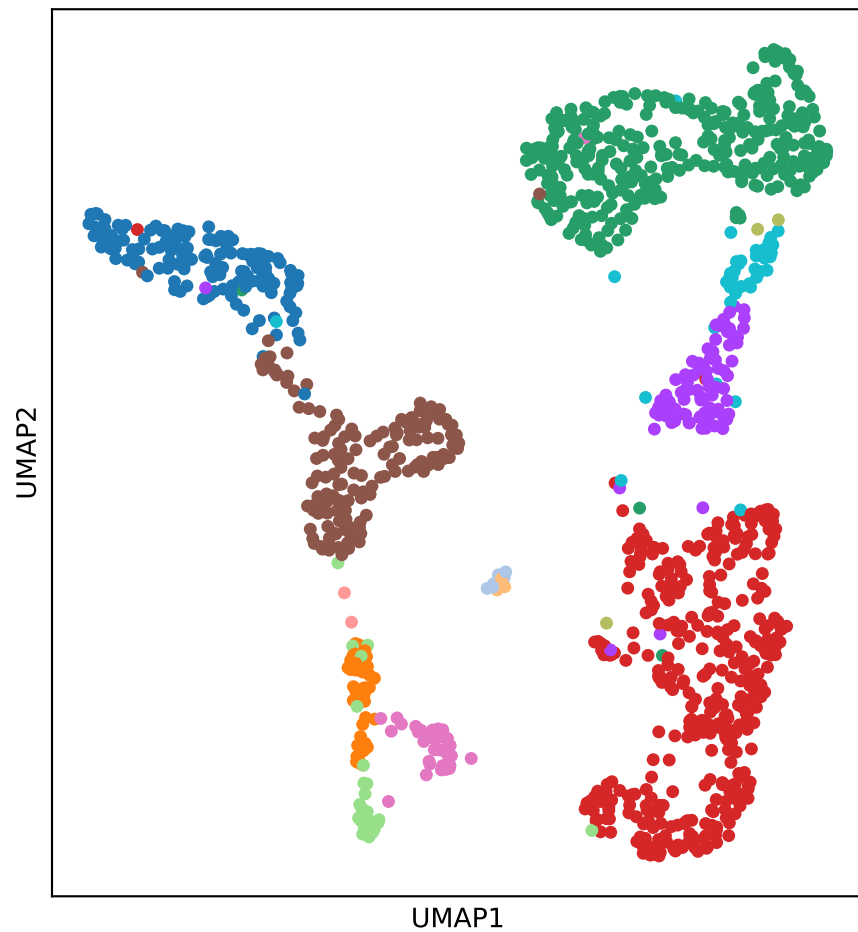
UMAP projection - Labels (val) (Res: 0.1, Iter: 28)



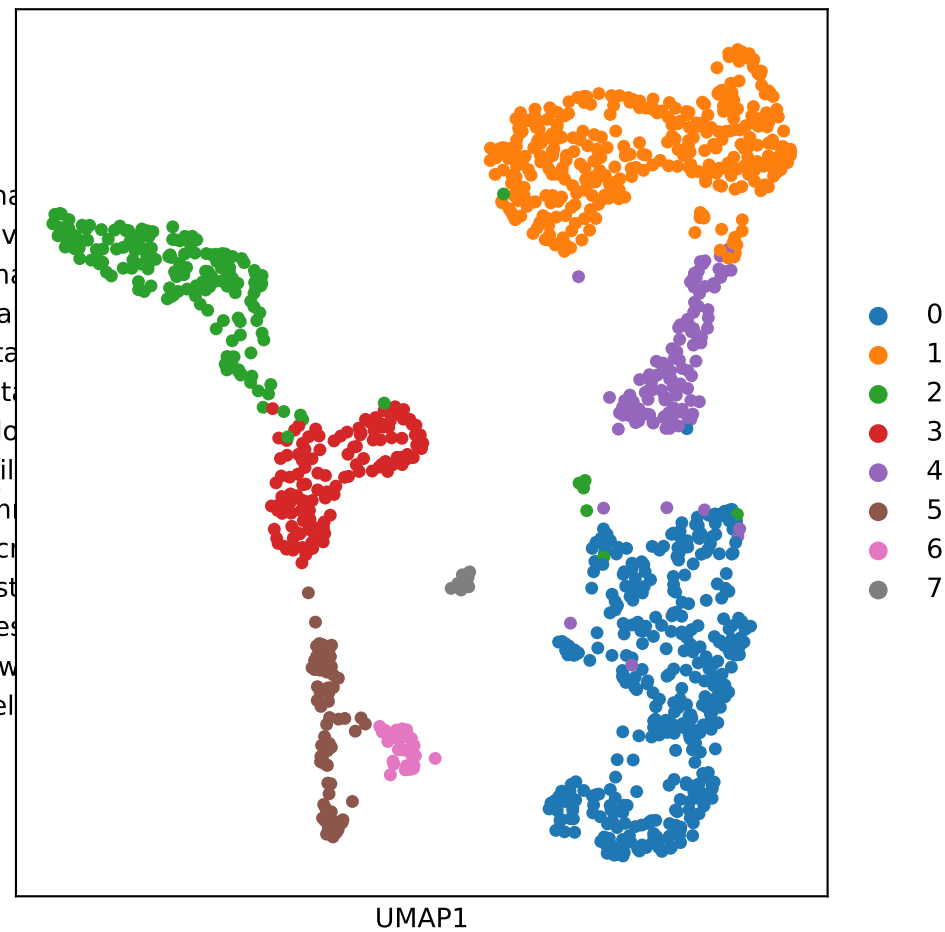
UMAP projection - Predictions (val) (Res: 0.1, Iter: 28)



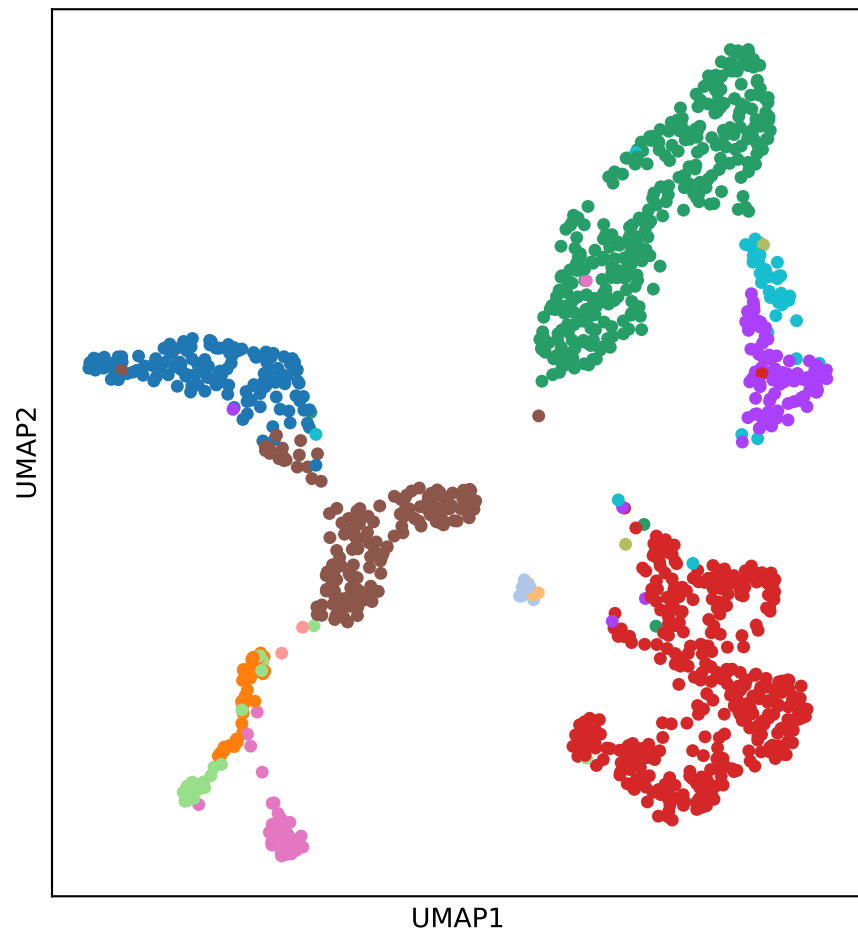
UMAP projection - Labels (val) (Res: 0.08, Iter: 29)



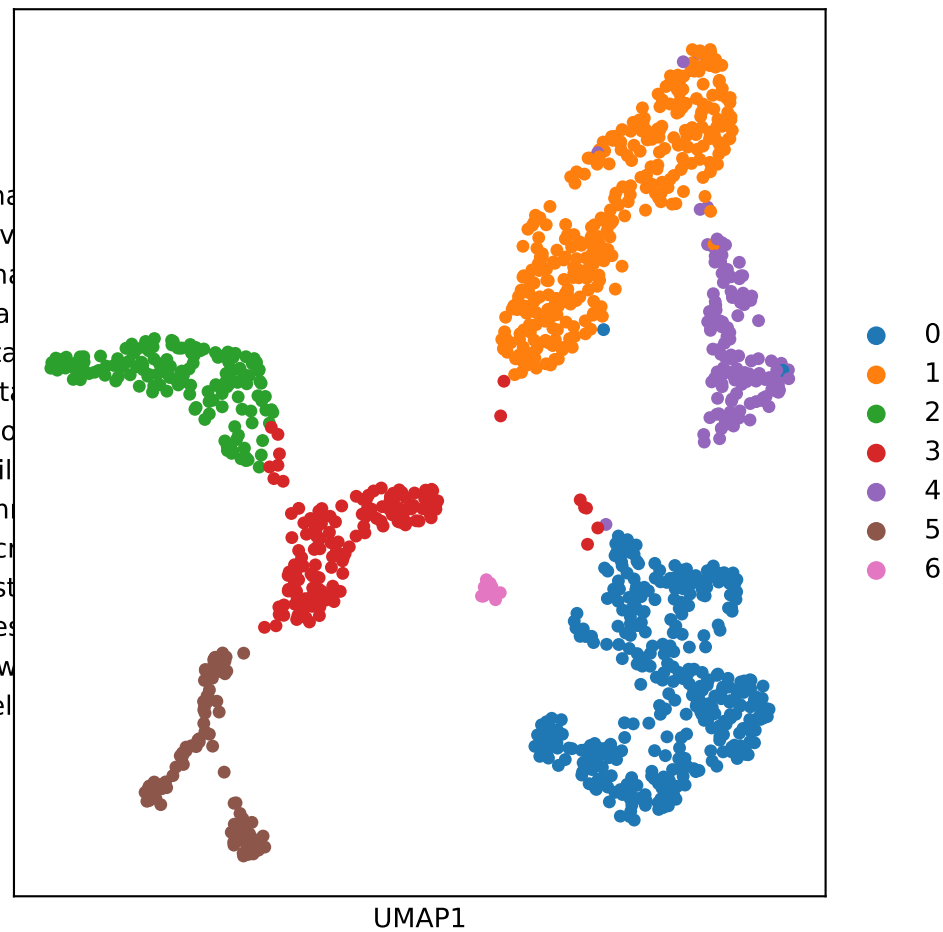
UMAP projection - Predictions (val) (Res: 0.08, Iter: 29)



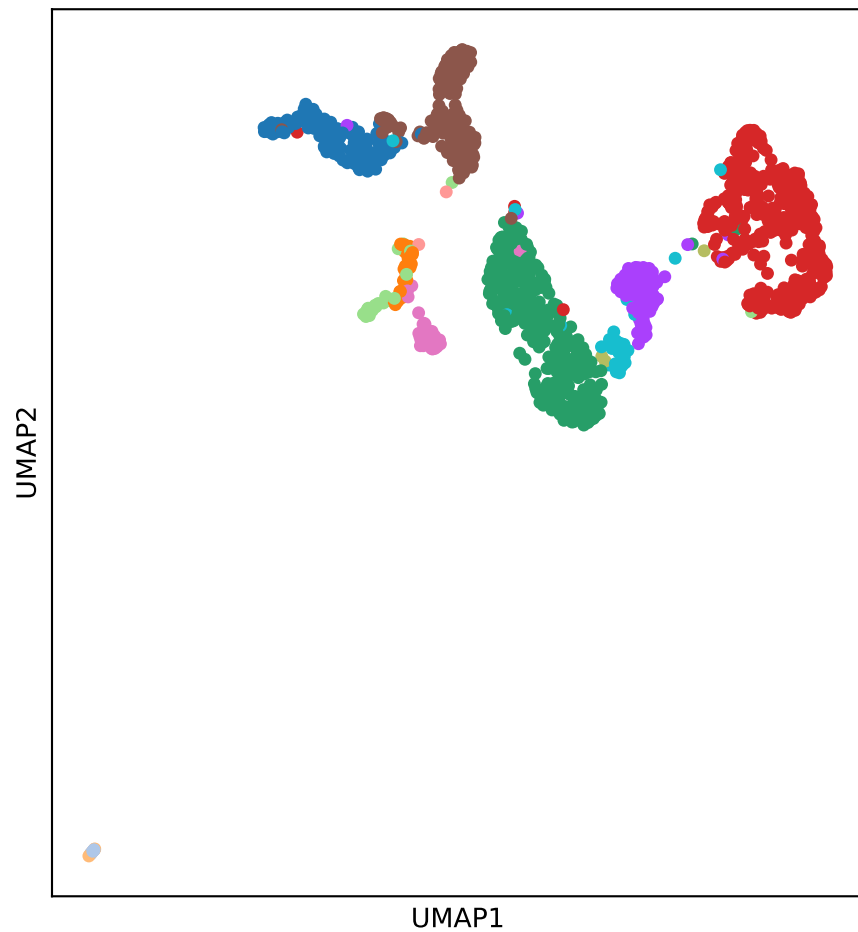
UMAP projection - Labels (val) (Res: 0.04, Iter: 30)



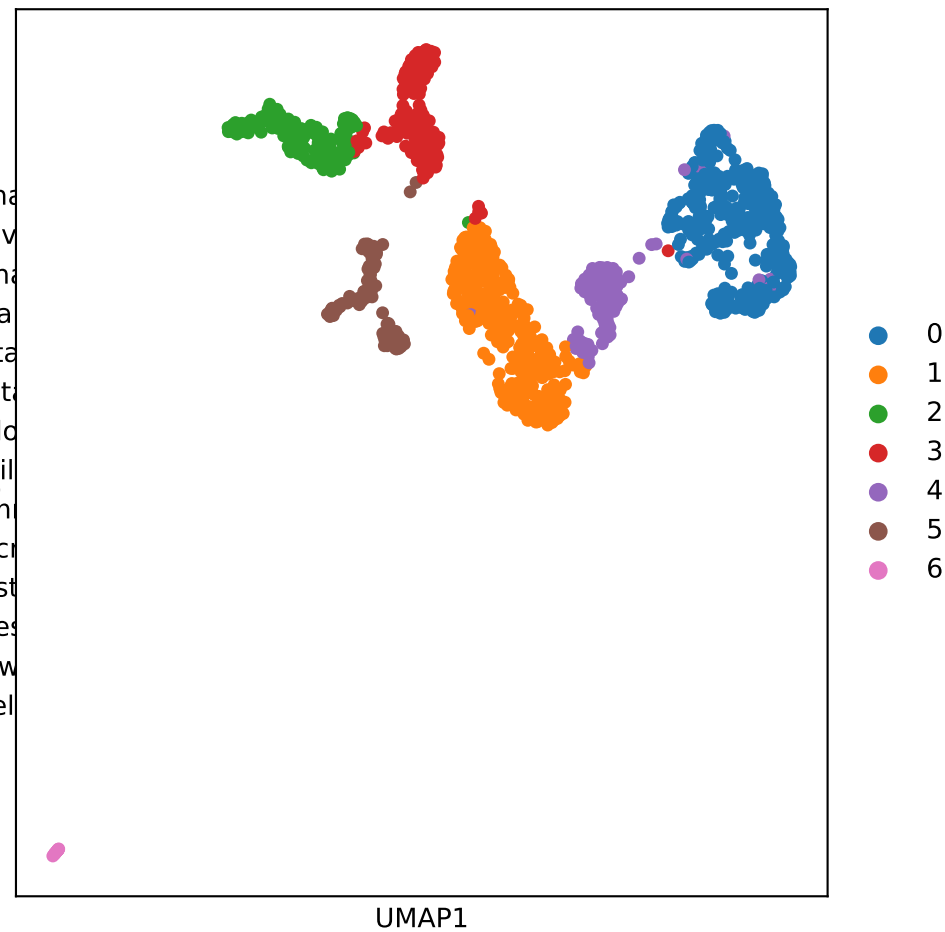
UMAP projection - Predictions (val) (Res: 0.04, Iter: 30)



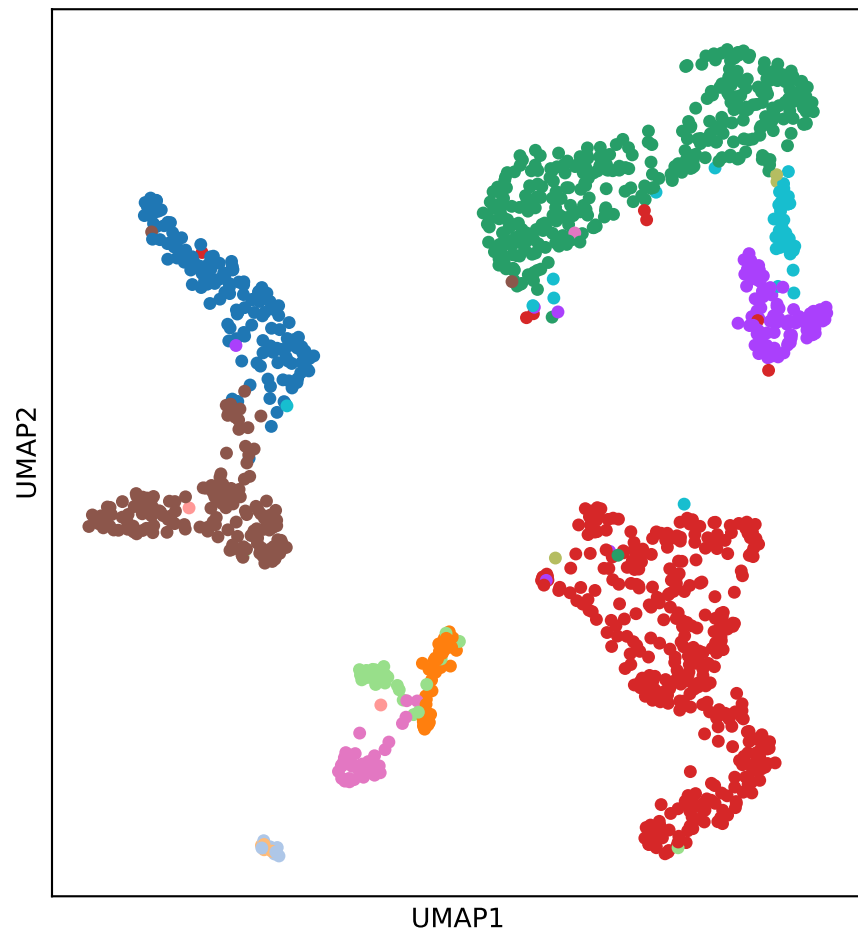
UMAP projection - Labels (val) (Res: 0.05, Iter: 31)



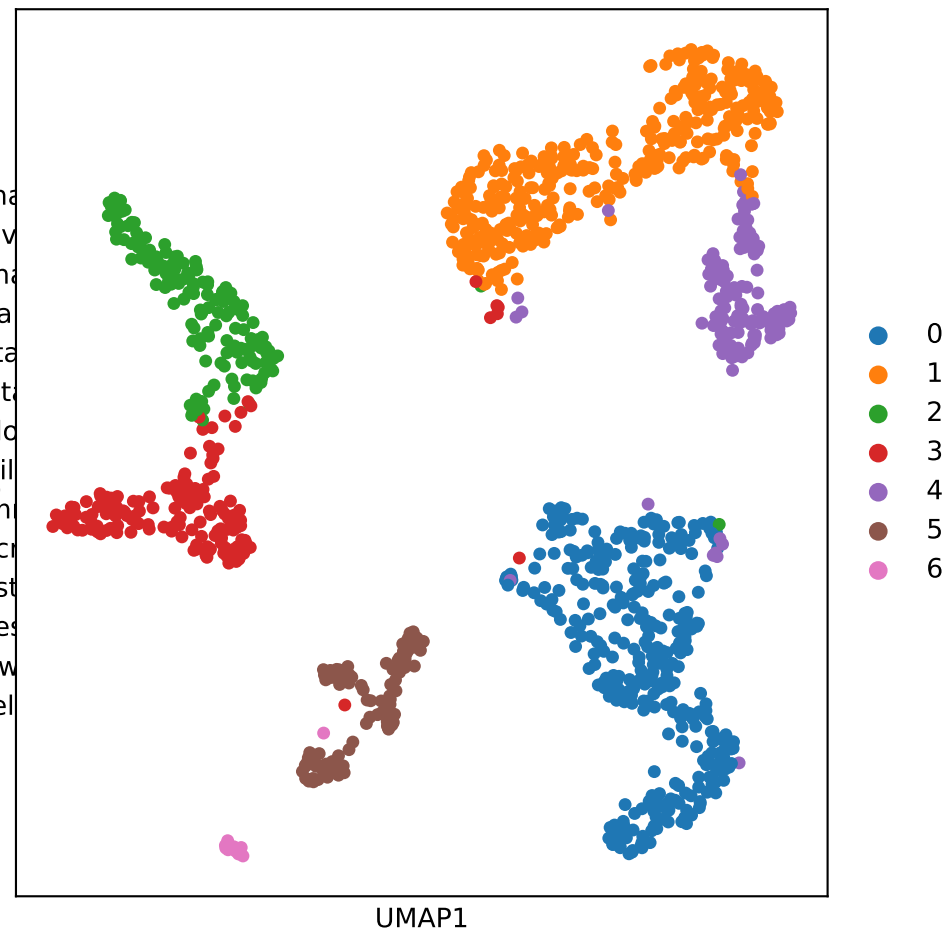
UMAP projection - Predictions (val) (Res: 0.05, Iter: 31)



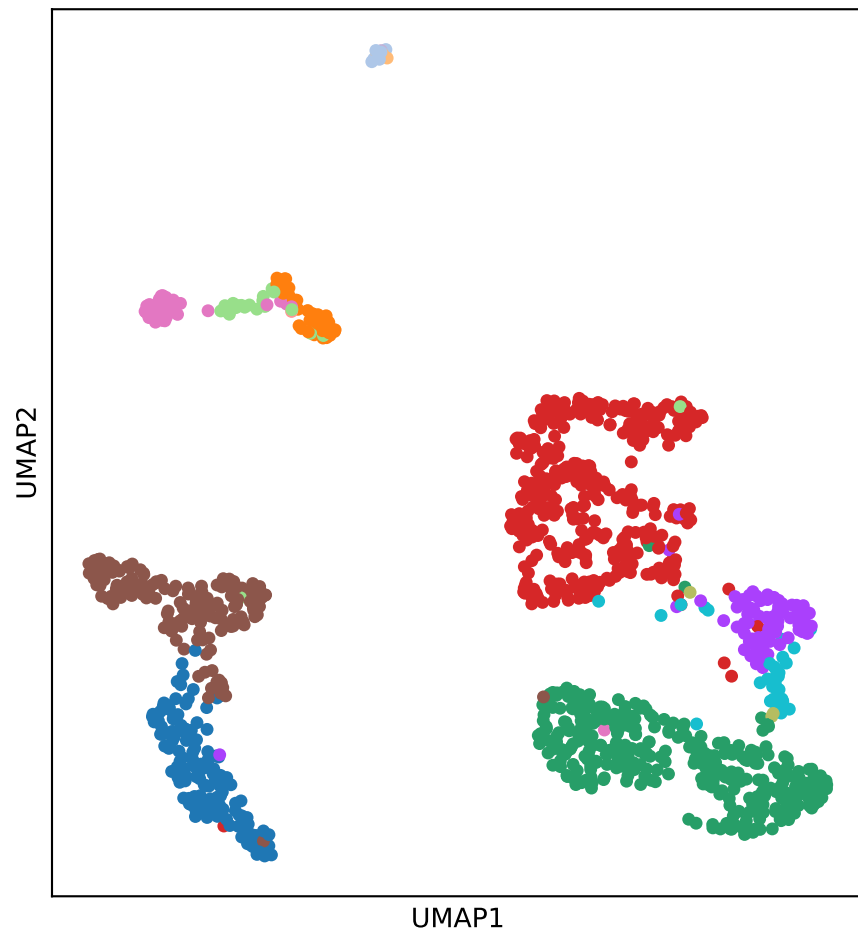
UMAP projection - Labels (val) (Res: 0.06, Iter: 32)



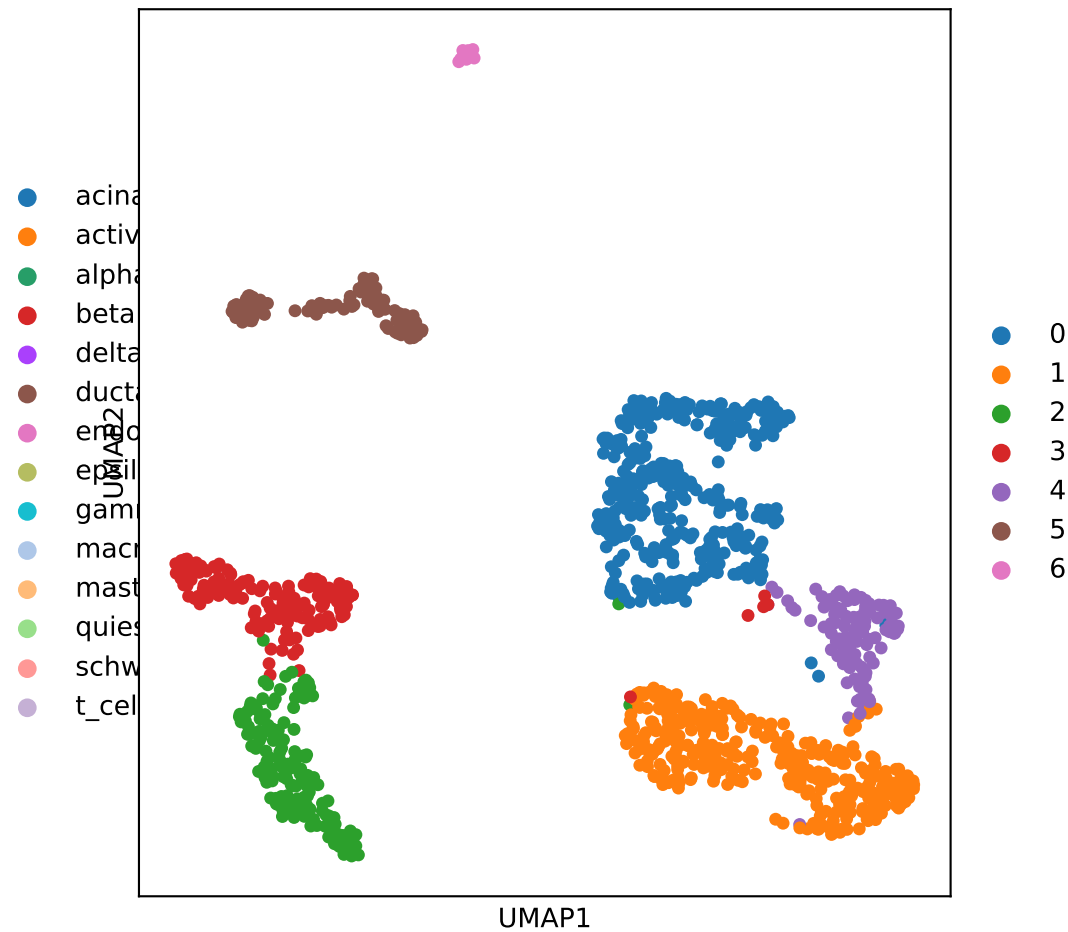
UMAP projection - Predictions (val) (Res: 0.06, Iter: 32)



UMAP projection - Labels (val) (Res: 0.05, Iter: 33)

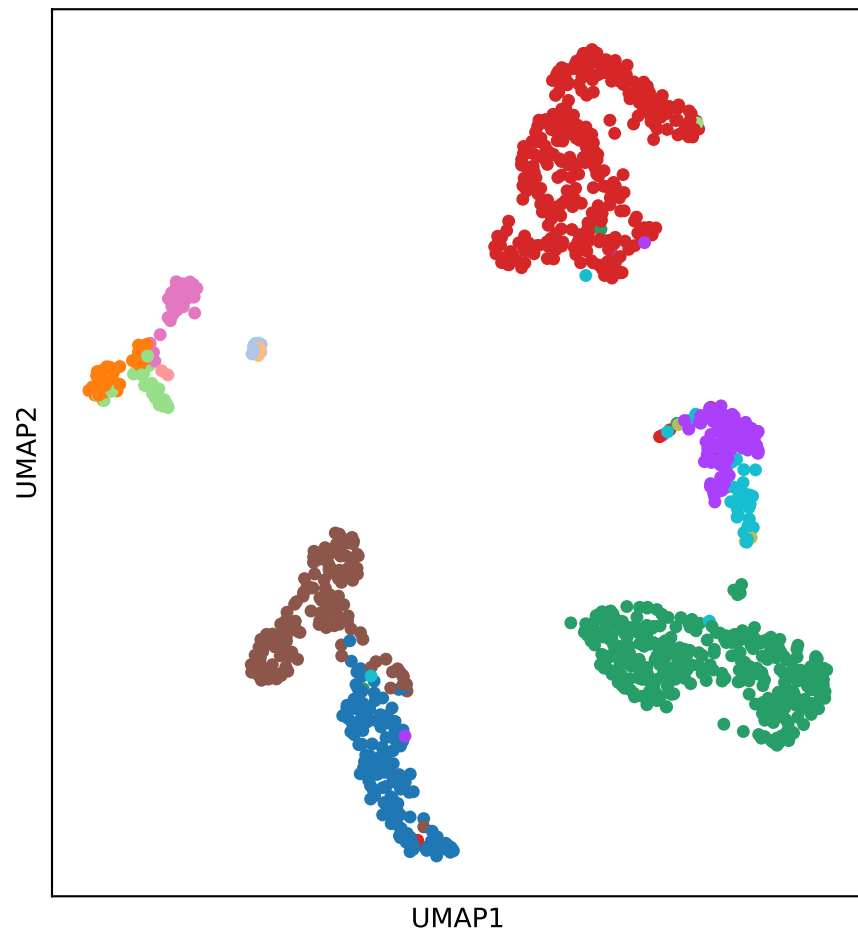


UMAP projection - Predictions (val) (Res: 0.05, Iter: 33)

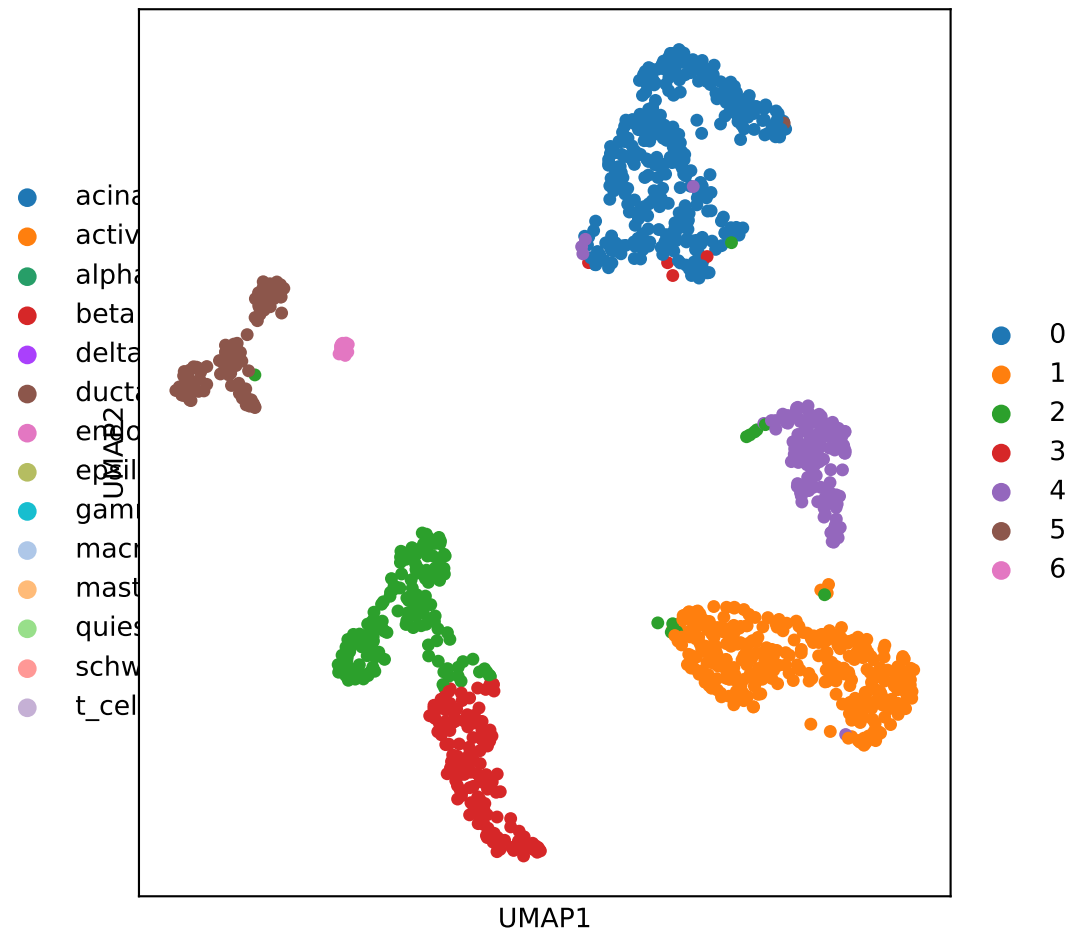


acina
 activ
 alpha
 beta
 delta
 duct
 endo
 epil
 gam
 macr
 mast
 quies
 schw
 t_cel

UMAP projection - Labels (val) (Res: 0.07, Iter: 34)

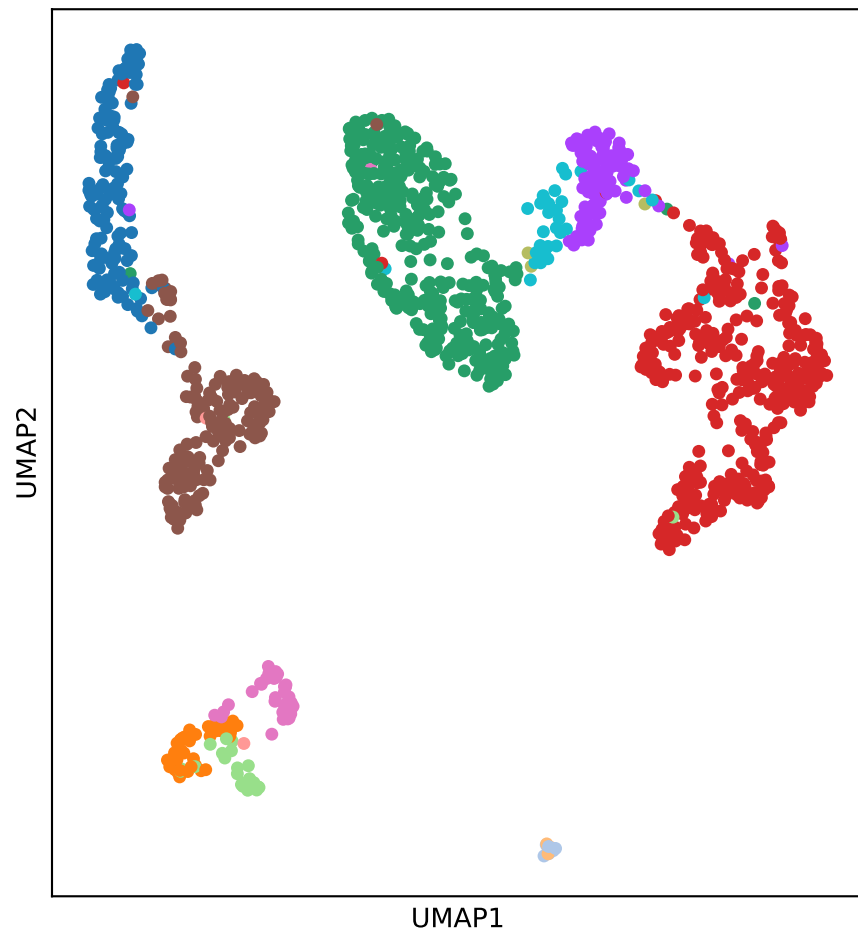


UMAP projection - Predictions (val) (Res: 0.07, Iter: 34)

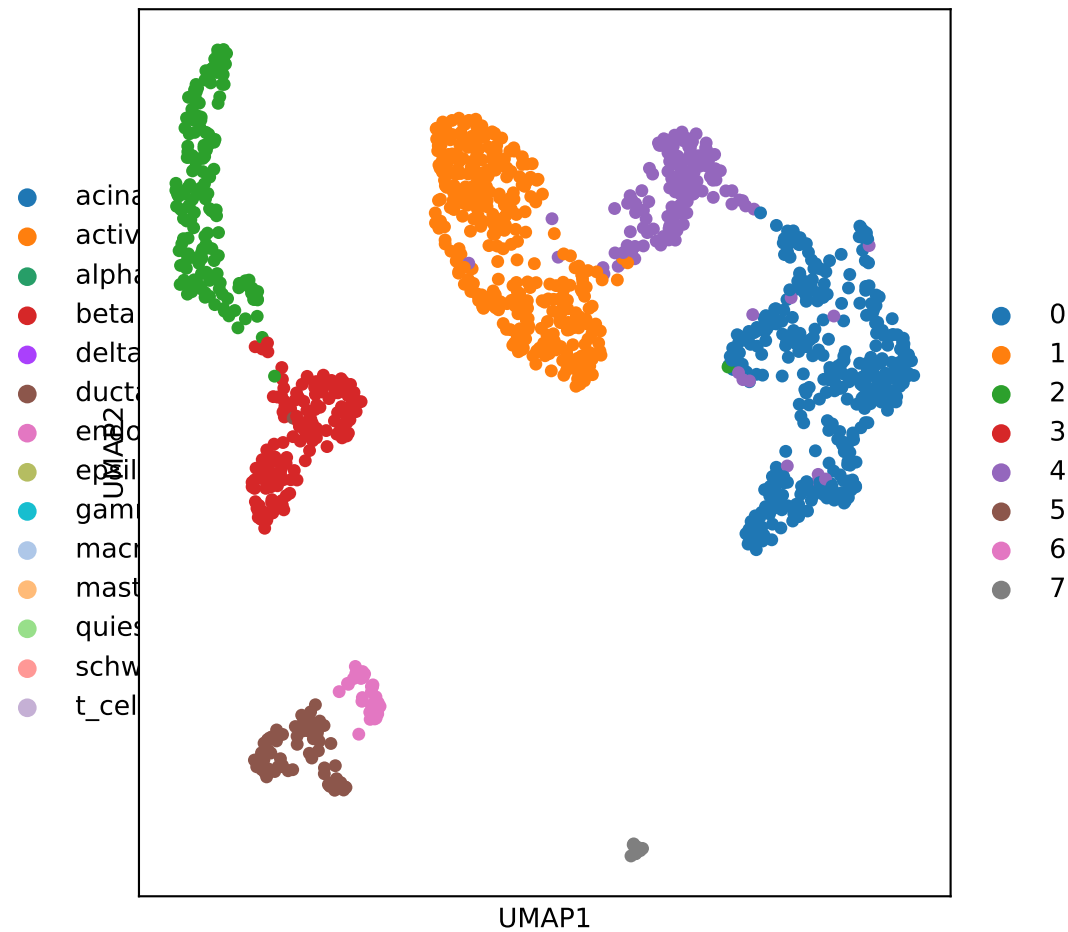


● acina
● activ
● alpha
● beta
● delta
● duct
● endo
● epil
● gam
● macr
● mast
● quies
● schw
● t_cel

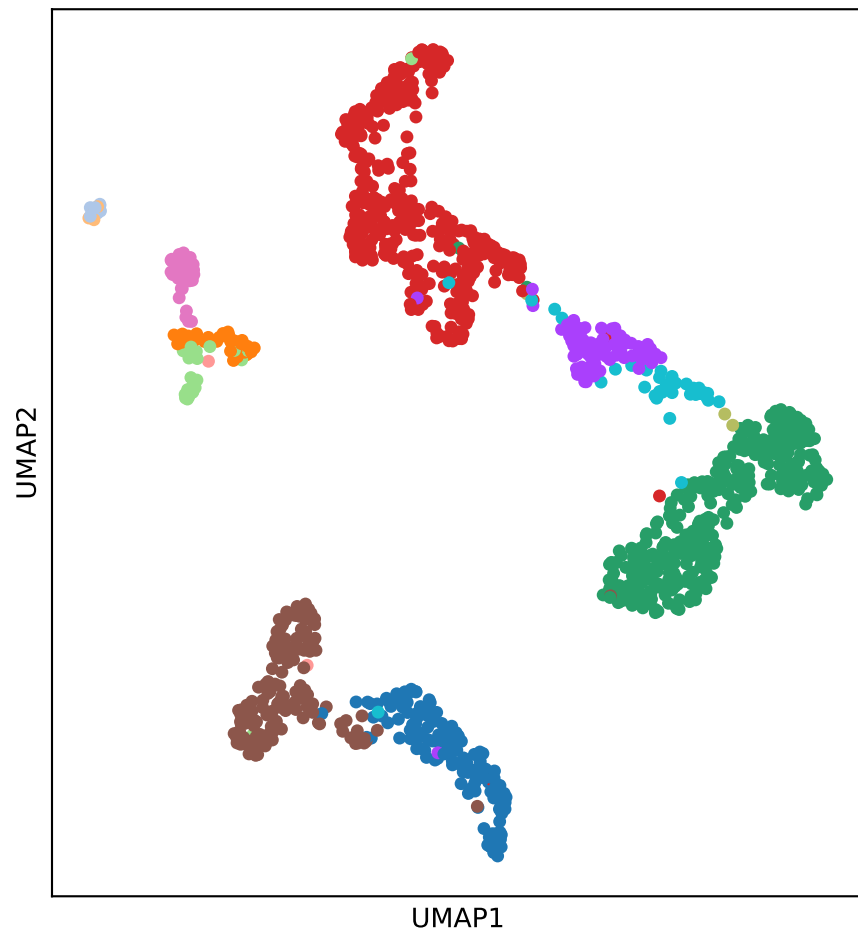
UMAP projection - Labels (val) (Res: 0.08, Iter: 35)



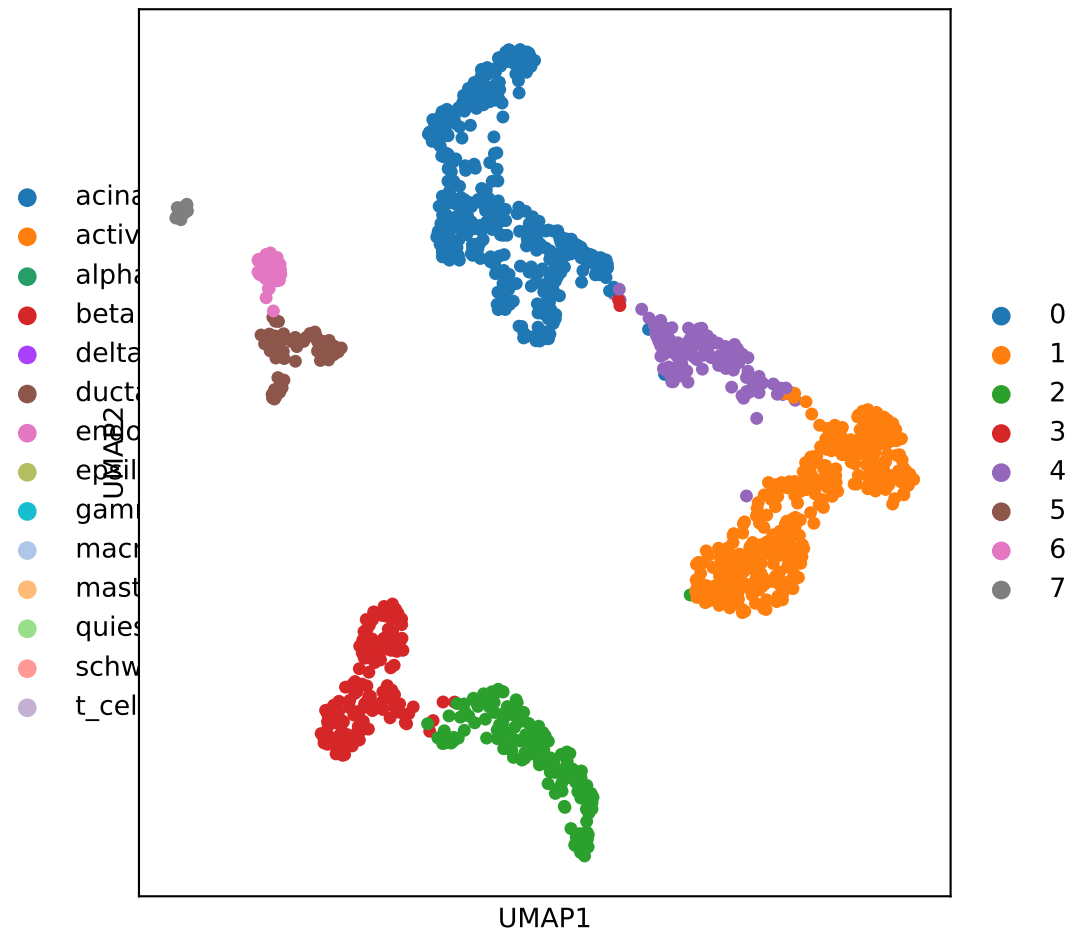
UMAP projection - Predictions (val) (Res: 0.08, Iter: 35)



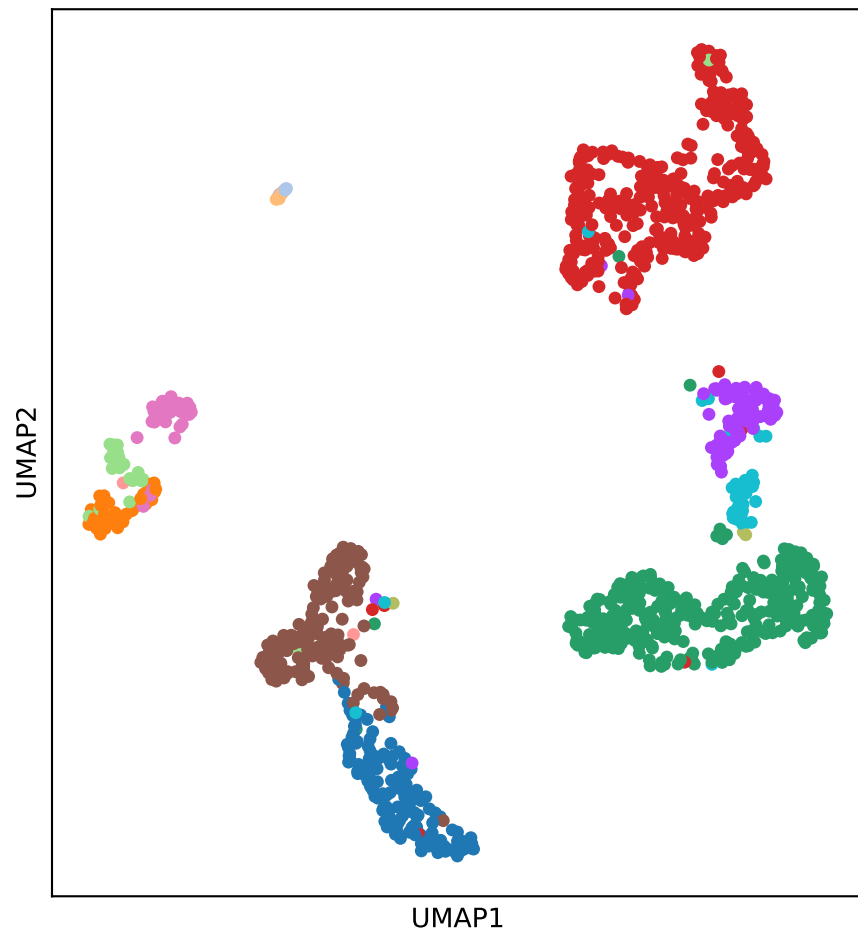
UMAP projection - Labels (val) (Res: 0.07, Iter: 36)



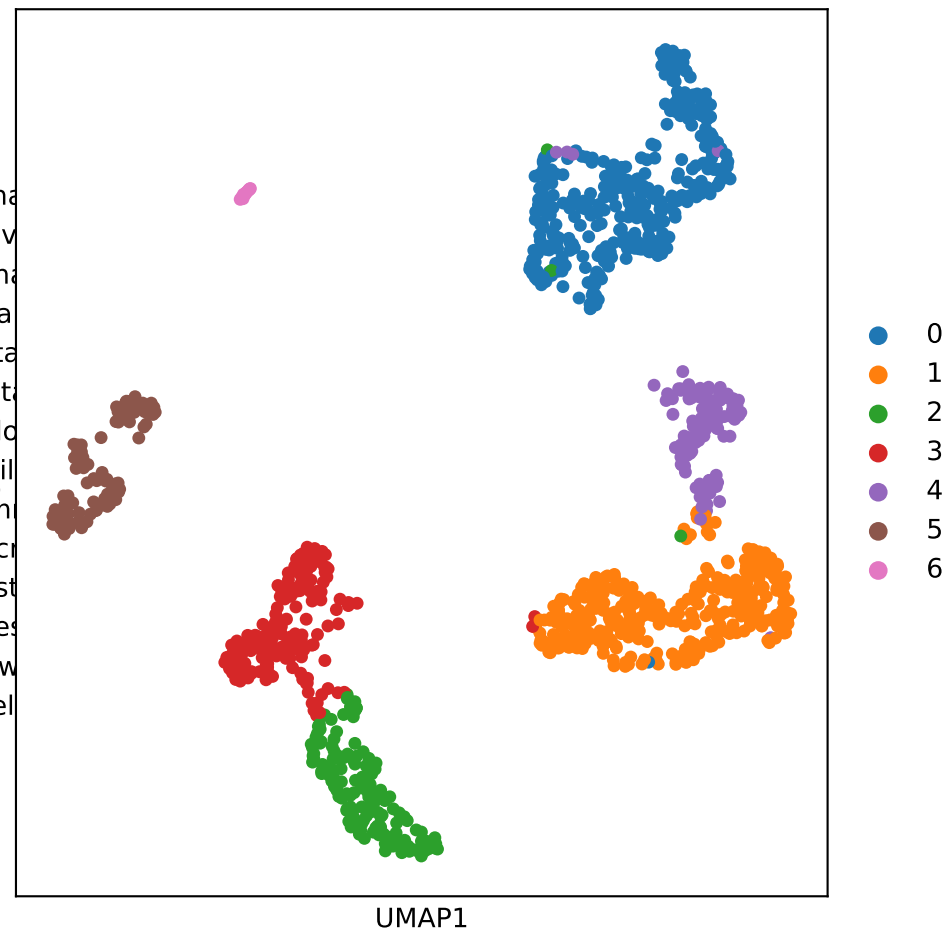
UMAP projection - Predictions (val) (Res: 0.07, Iter: 36)



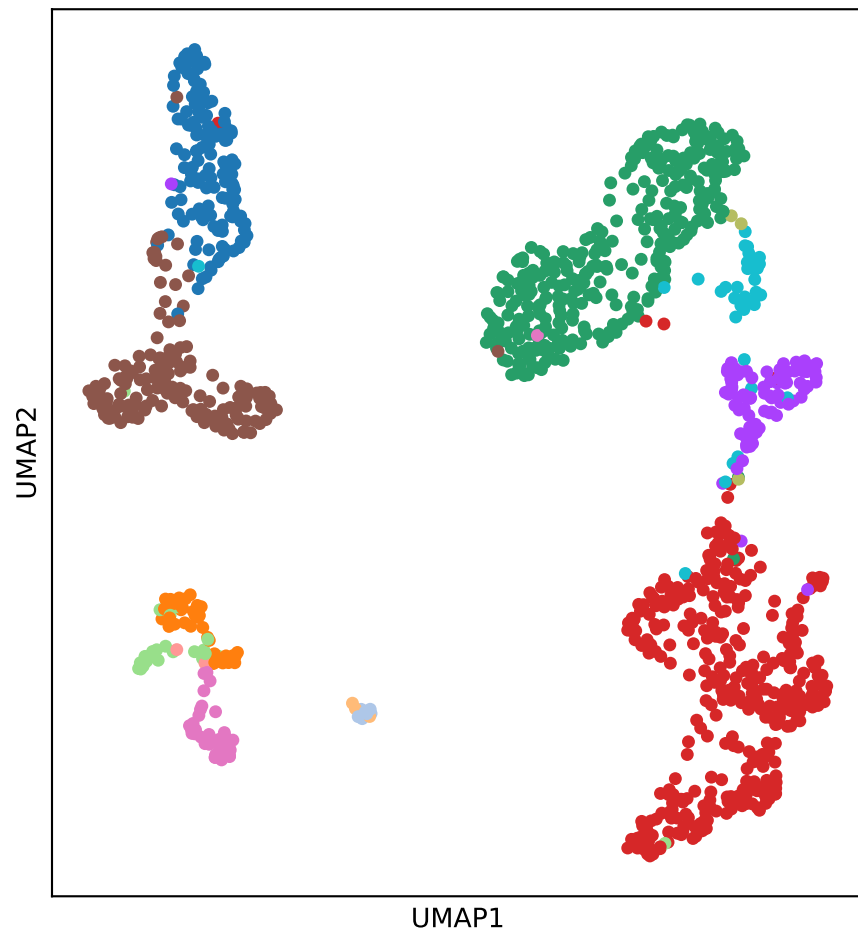
UMAP projection - Labels (val) (Res: 0.07, Iter: 37)



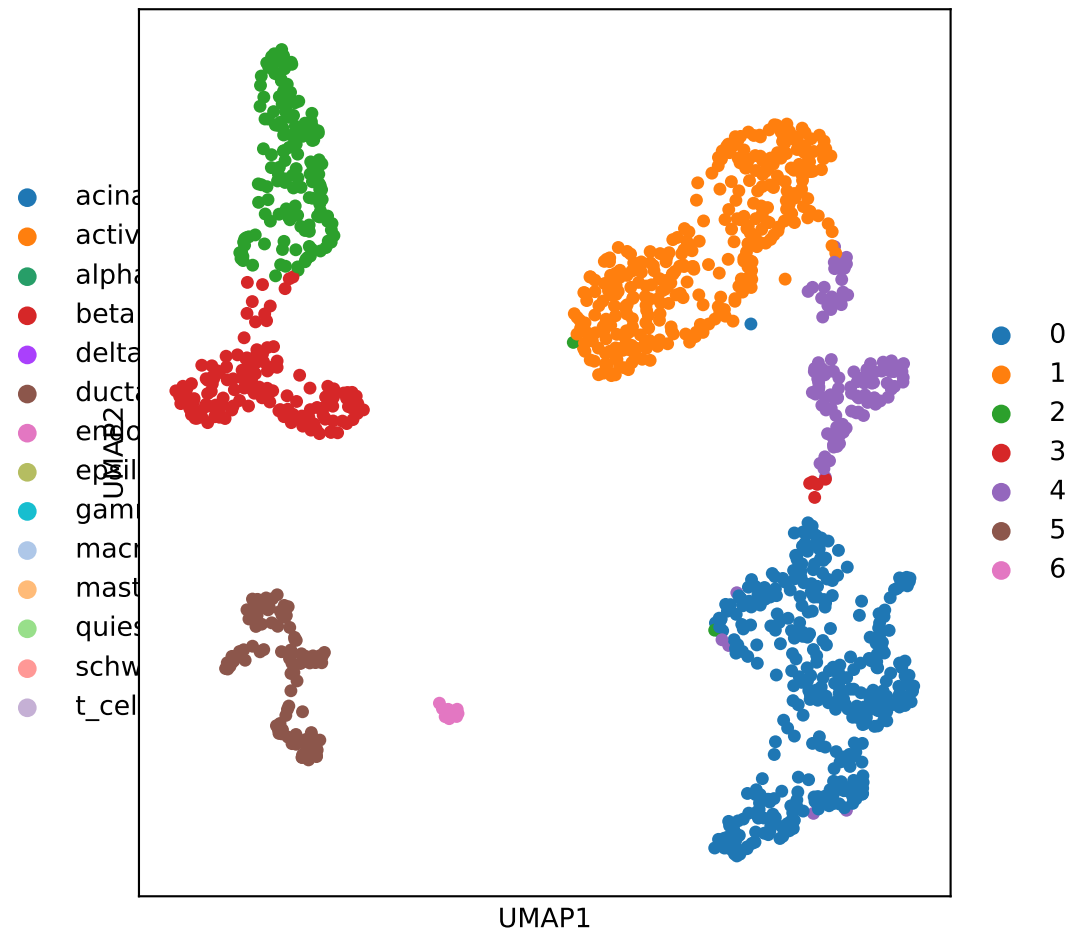
UMAP projection - Predictions (val) (Res: 0.07, Iter: 37)



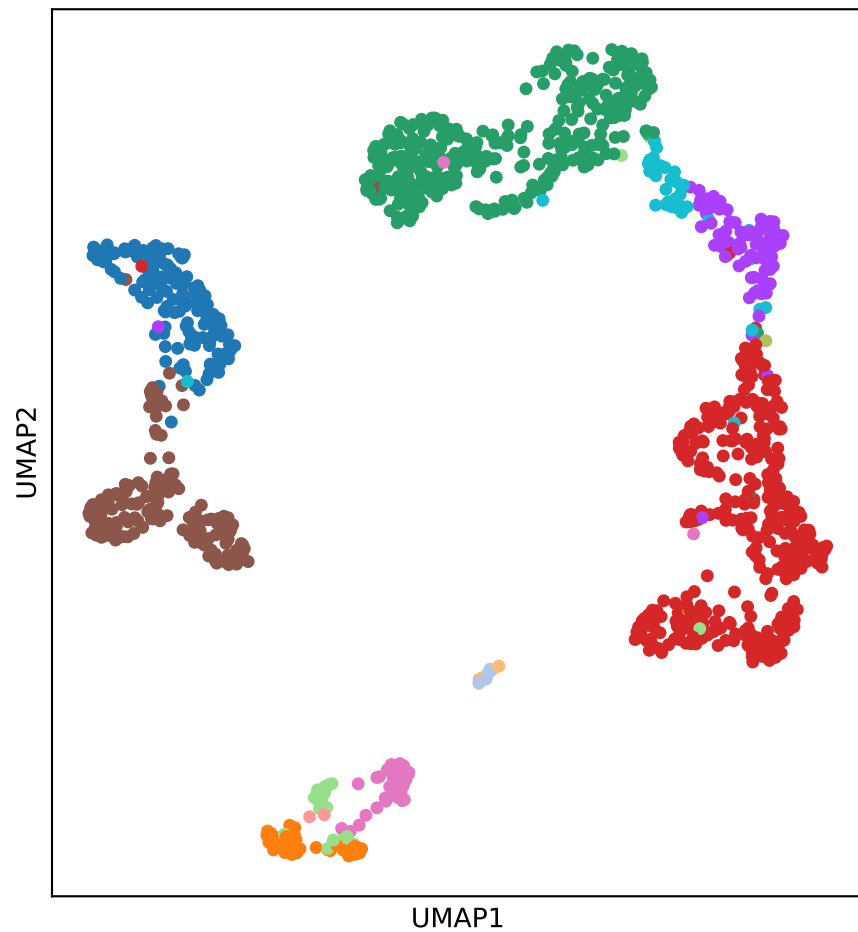
UMAP projection - Labels (val) (Res: 0.04, Iter: 38)



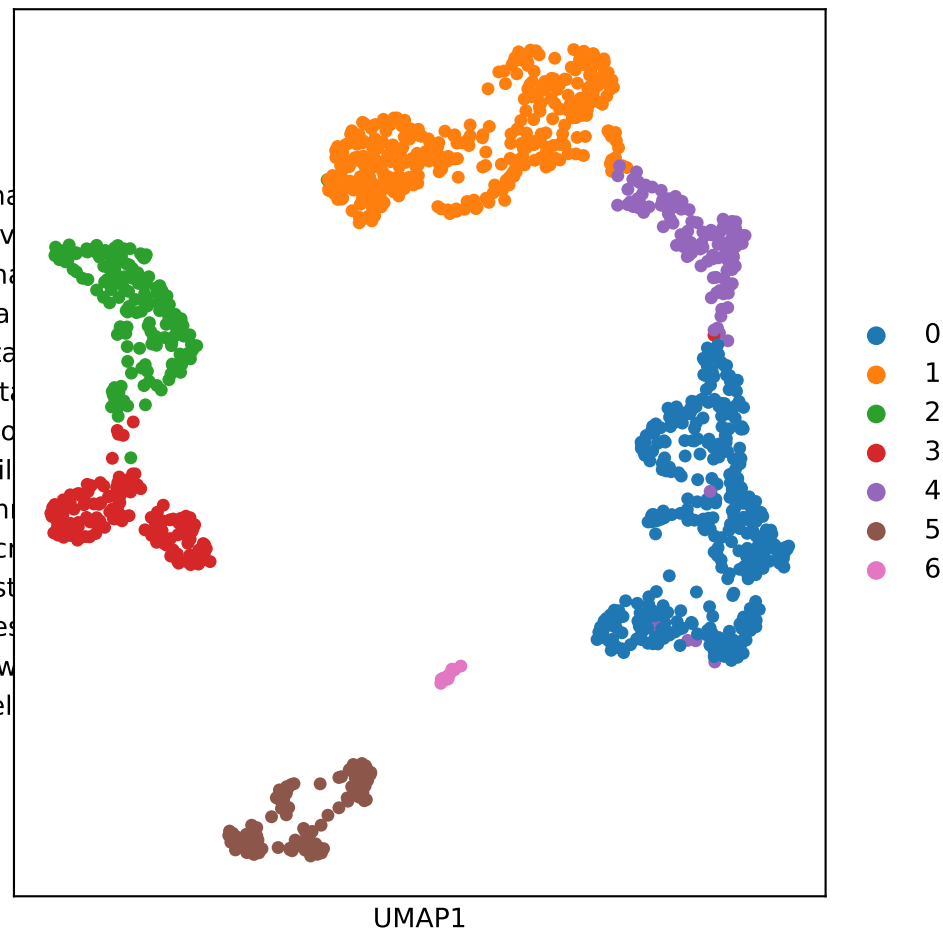
UMAP projection - Predictions (val) (Res: 0.04, Iter: 38)



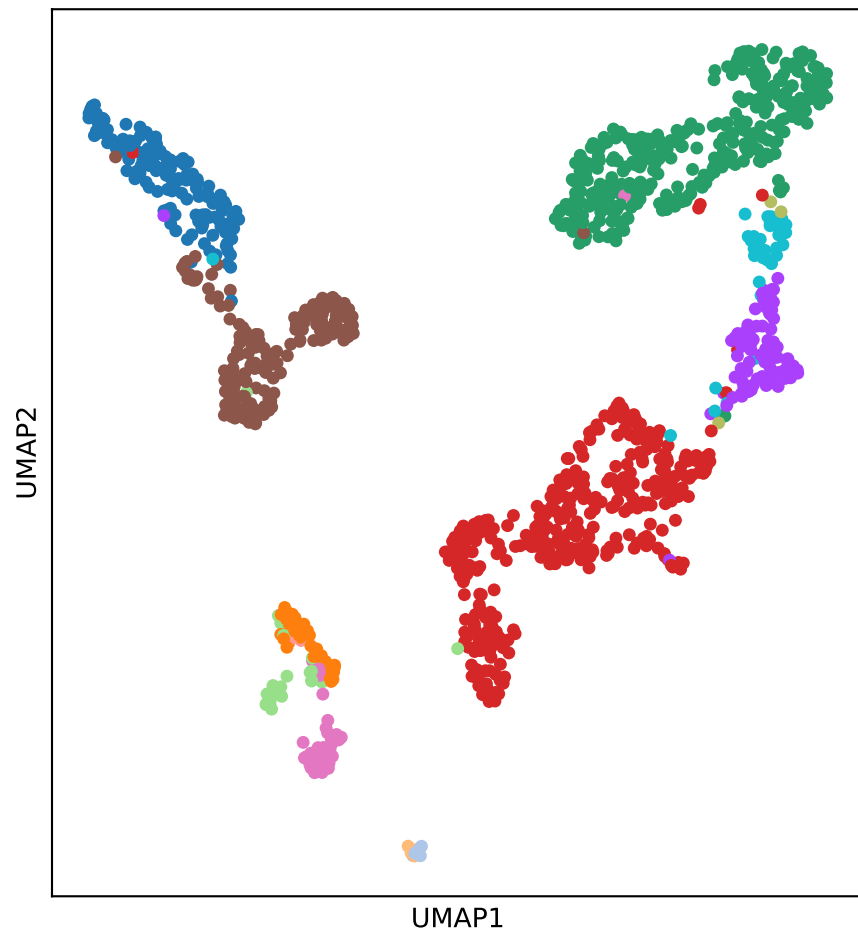
UMAP projection - Labels (val) (Res: 0.04, Iter: 39)



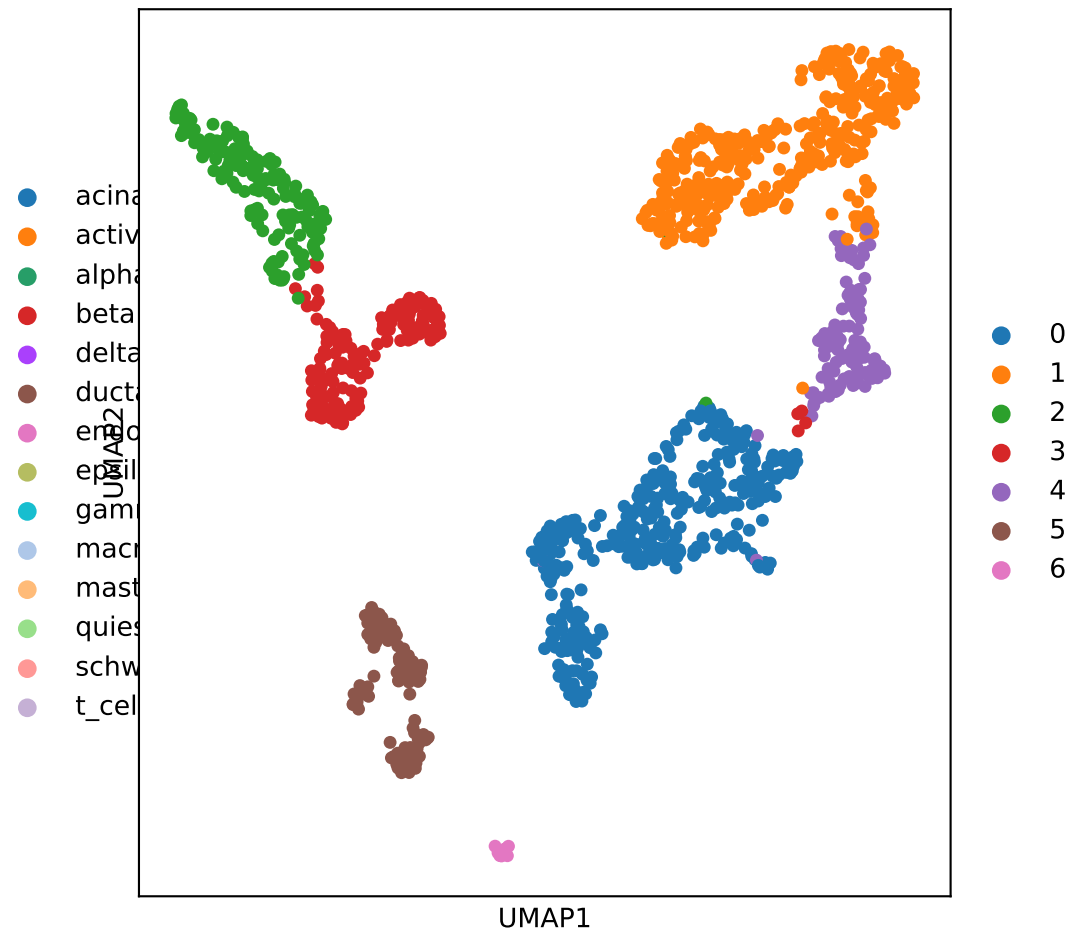
UMAP projection - Predictions (val) (Res: 0.04, Iter: 39)



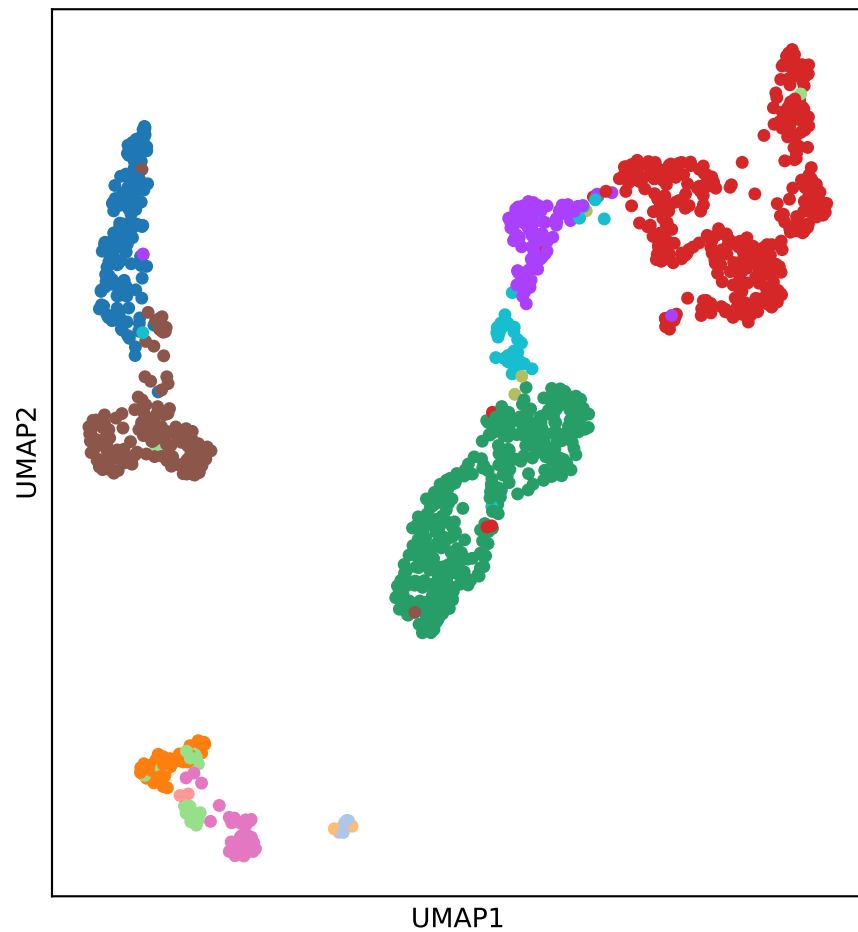
UMAP projection - Labels (val) (Res: 0.05, Iter: 40)



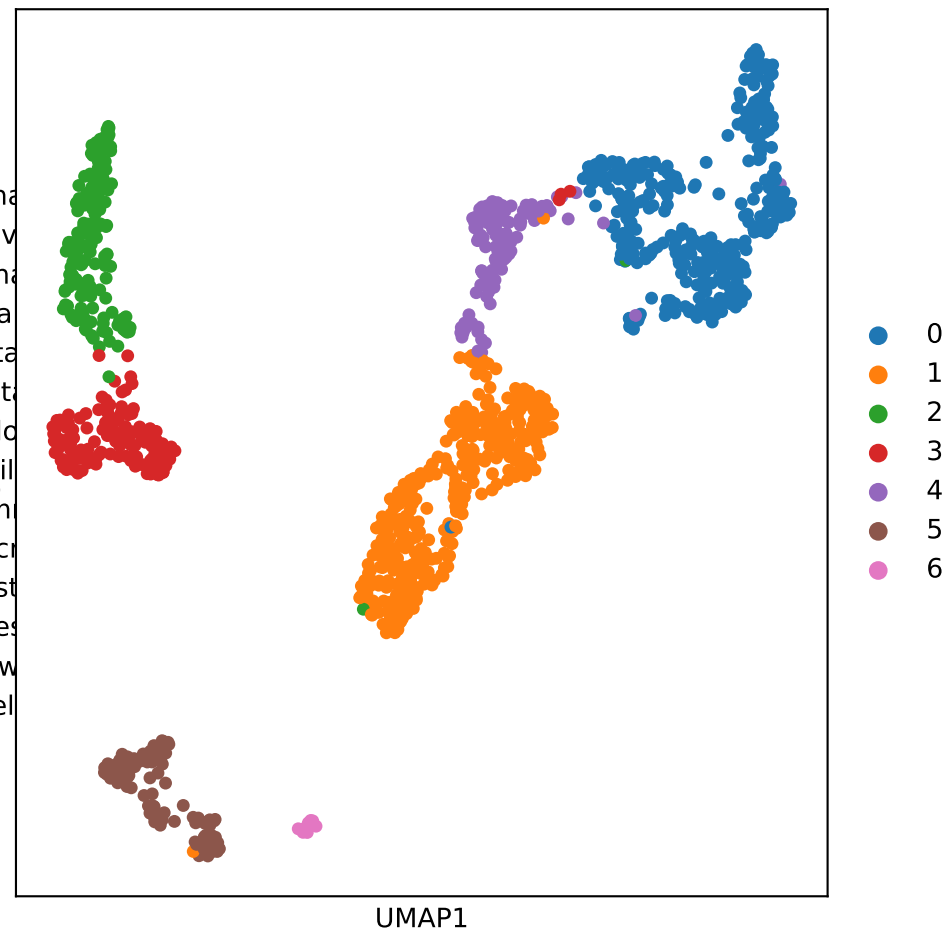
UMAP projection - Predictions (val) (Res: 0.05, Iter: 40)



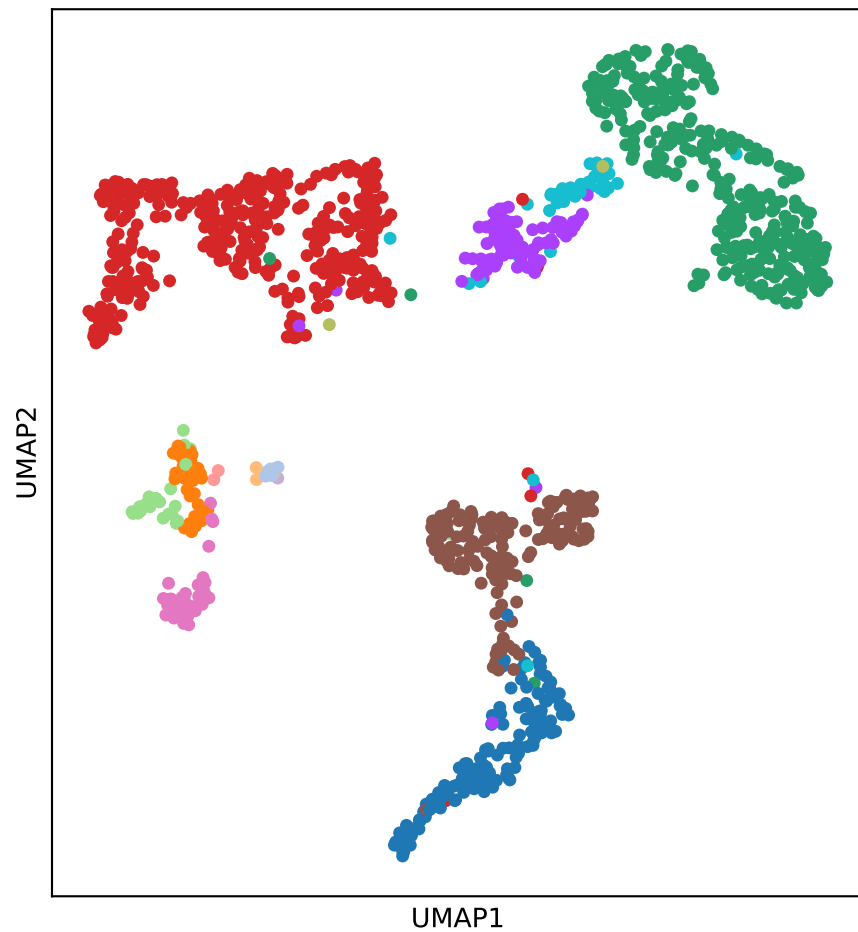
UMAP projection - Labels (val) (Res: 0.07, Iter: 41)



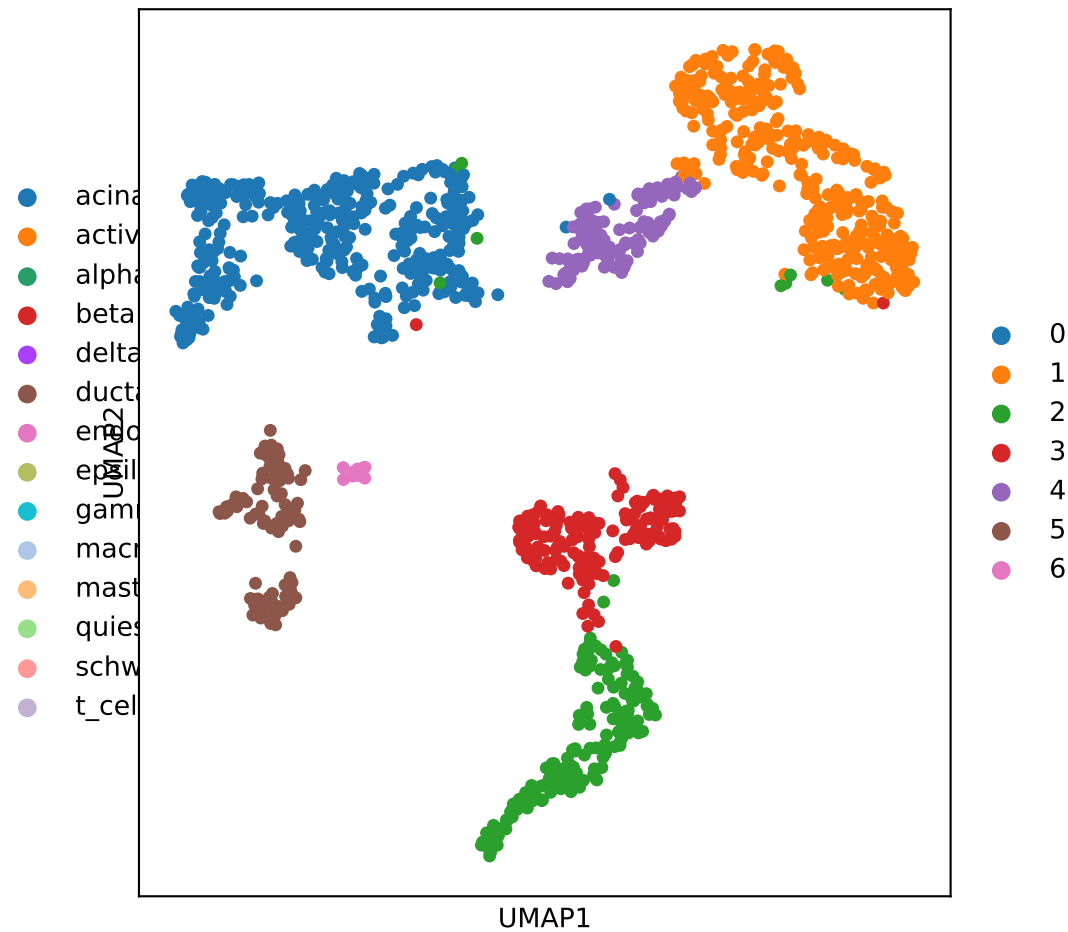
UMAP projection - Predictions (val) (Res: 0.07, Iter: 41)



UMAP projection - Labels (val) (Res: 0.08, Iter: 42)

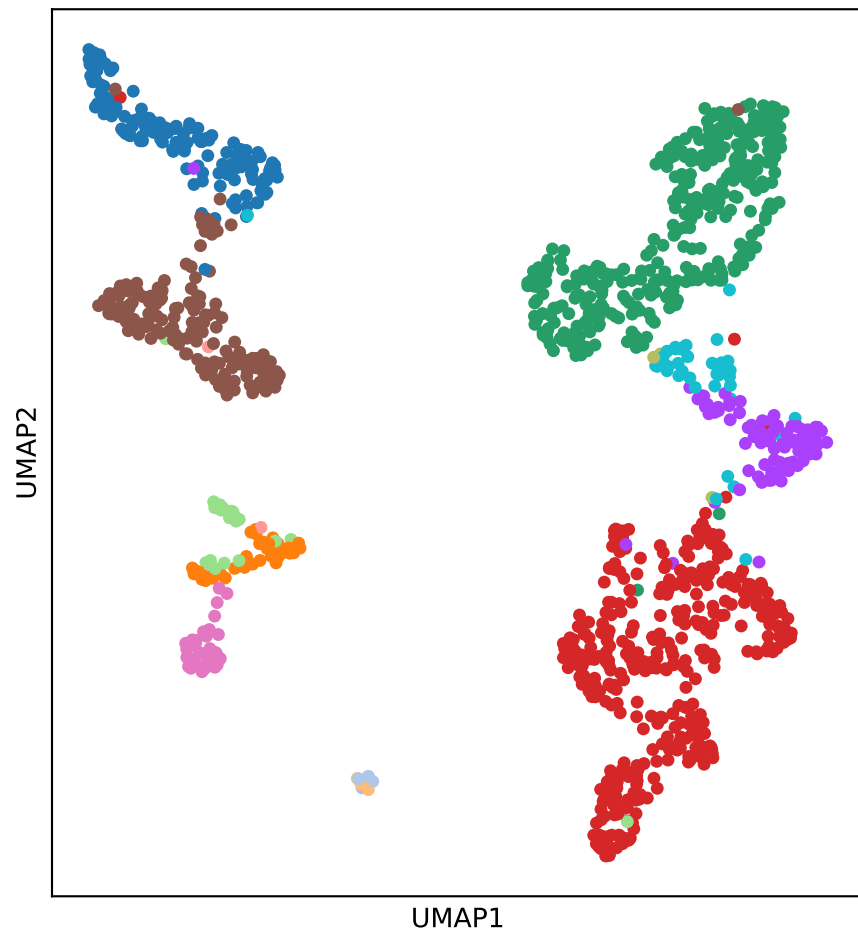


UMAP projection - Predictions (val) (Res: 0.08, Iter: 42)

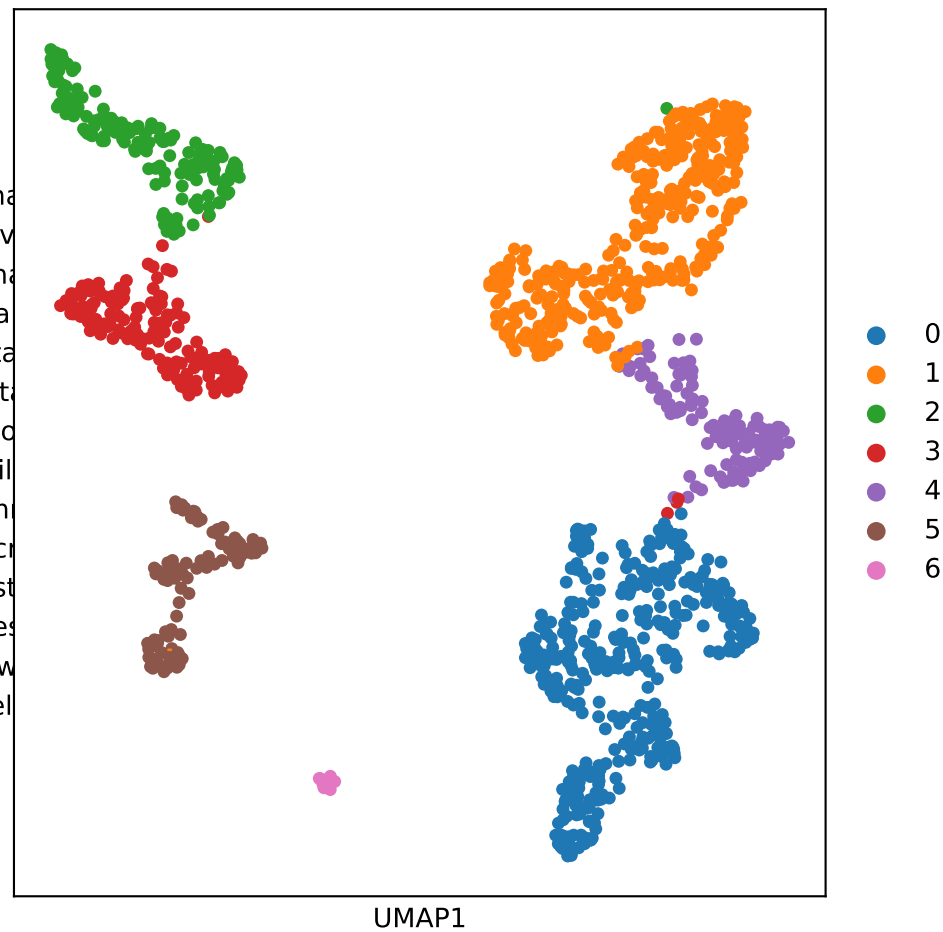


● acina
 ● activ
 ● alpha
 ● beta
 ● delta
 ● duct
 ● endo
 ● epil
 ● gam
 ● macr
 ● mast
 ● quies
 ● schw
 ● t_cel

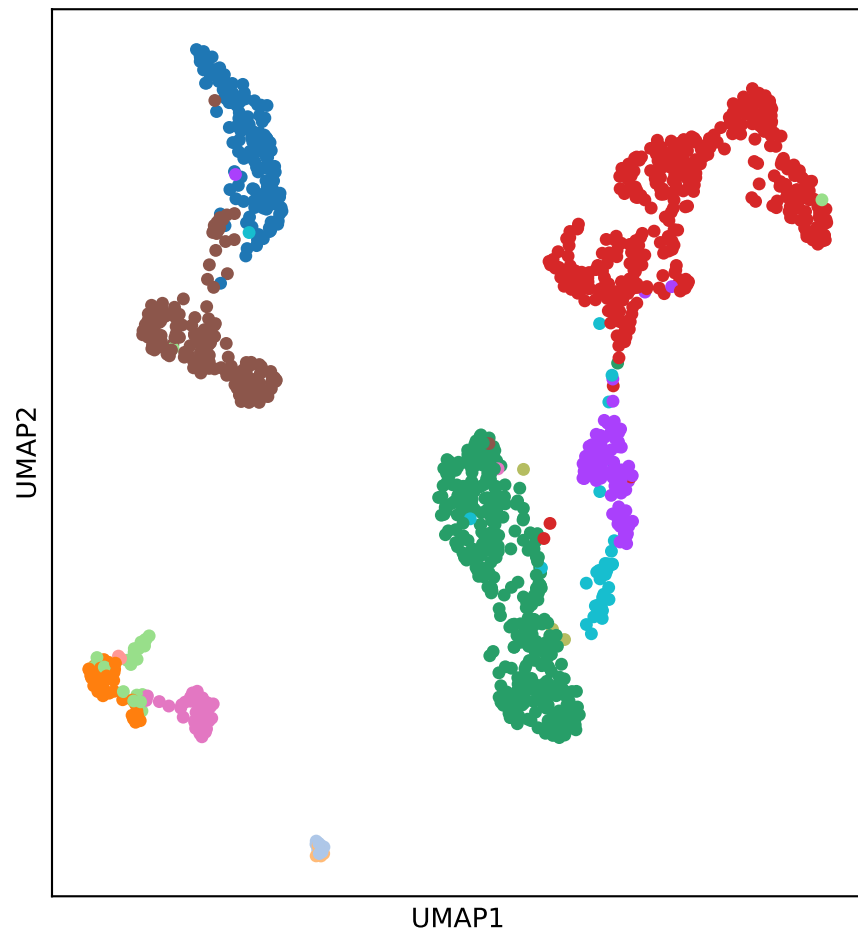
UMAP projection - Labels (val) (Res: 0.05, Iter: 43)



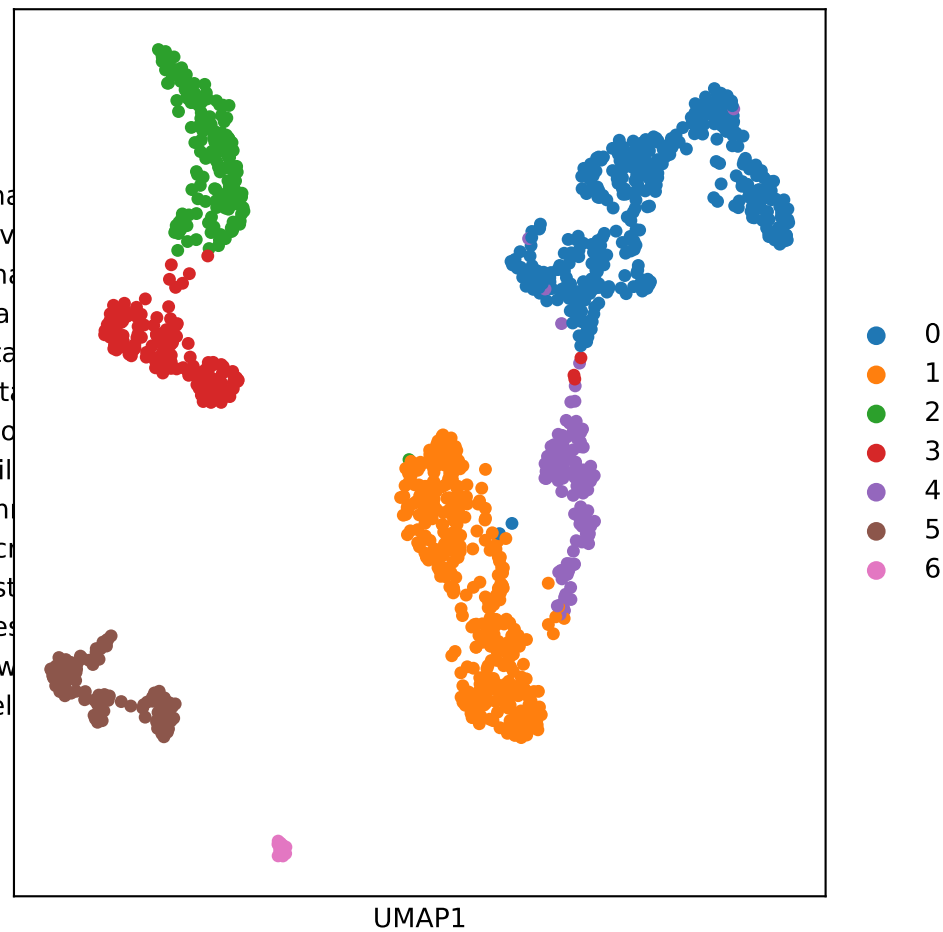
UMAP projection - Predictions (val) (Res: 0.05, Iter: 43)



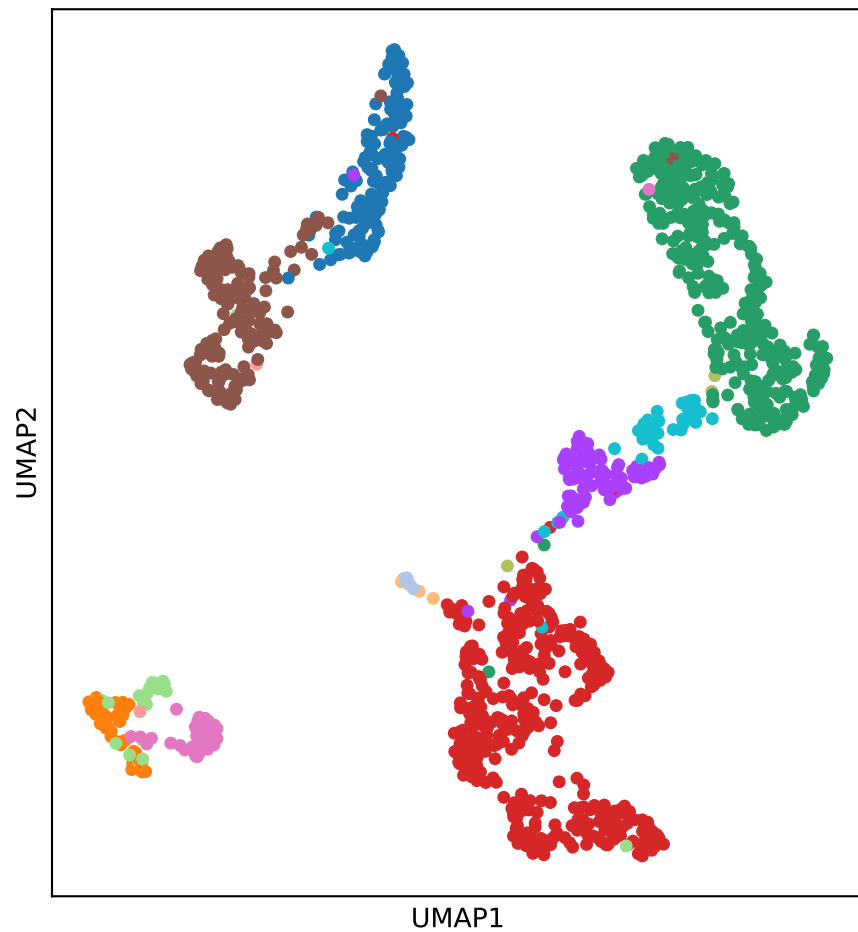
UMAP projection - Labels (val) (Res: 0.04, Iter: 44)



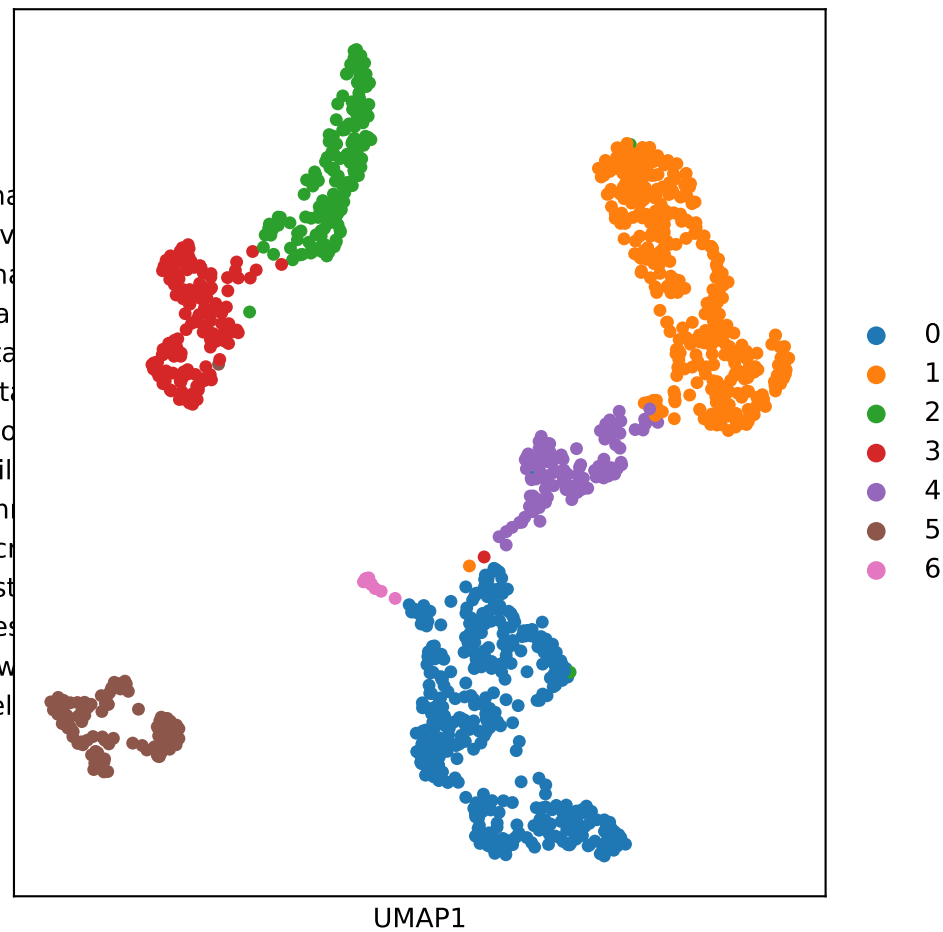
UMAP projection - Predictions (val) (Res: 0.04, Iter: 44)



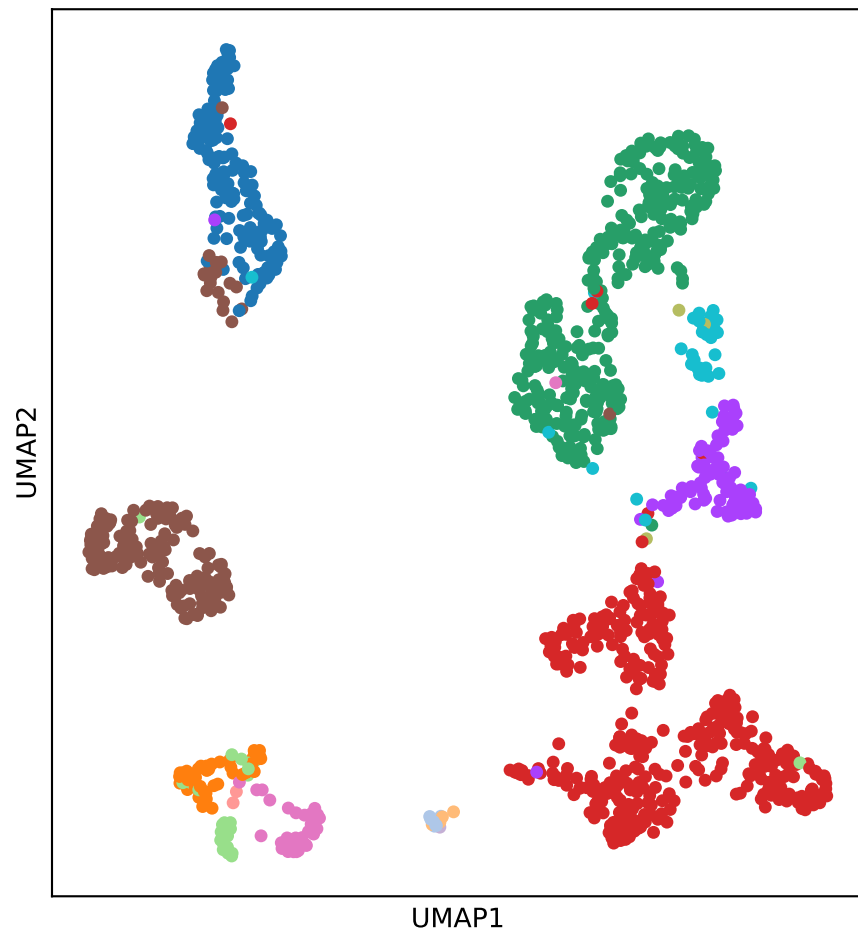
UMAP projection - Labels (val) (Res: 0.07, Iter: 45)



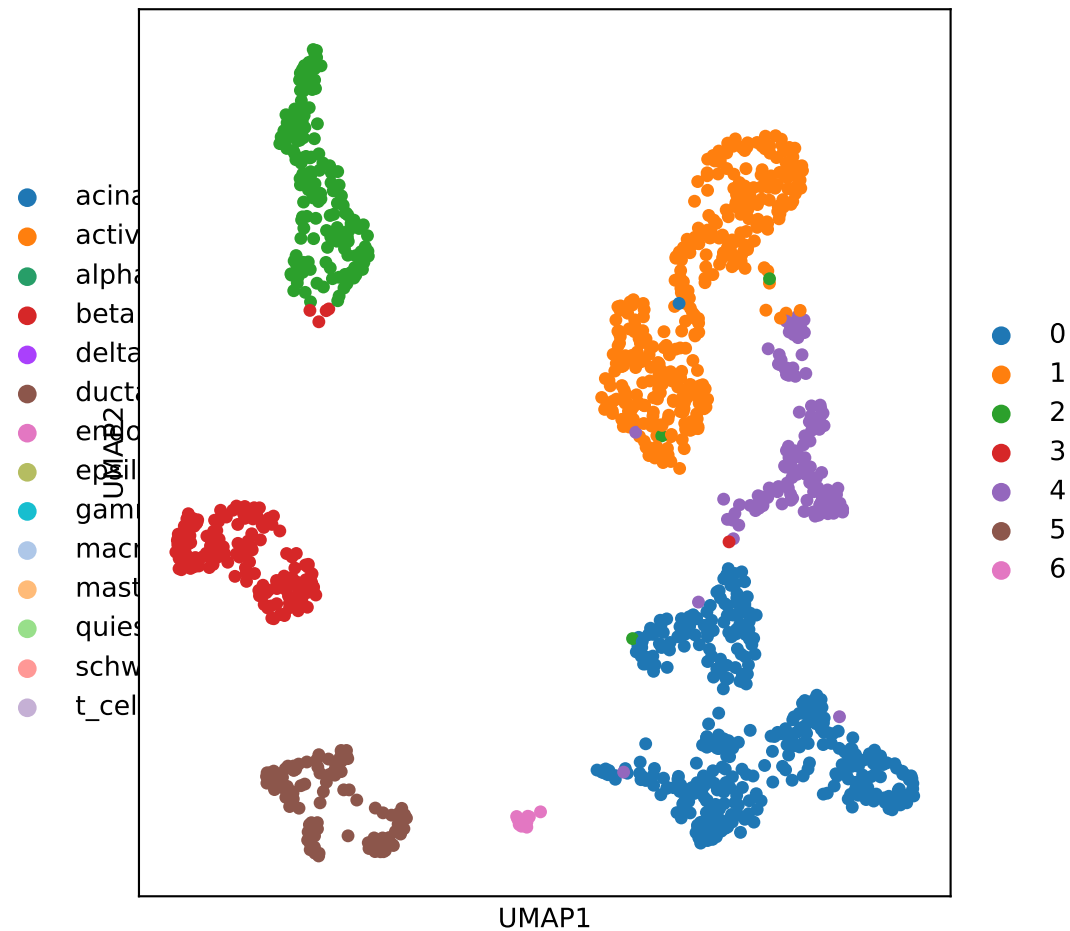
UMAP projection - Predictions (val) (Res: 0.07, Iter: 45)



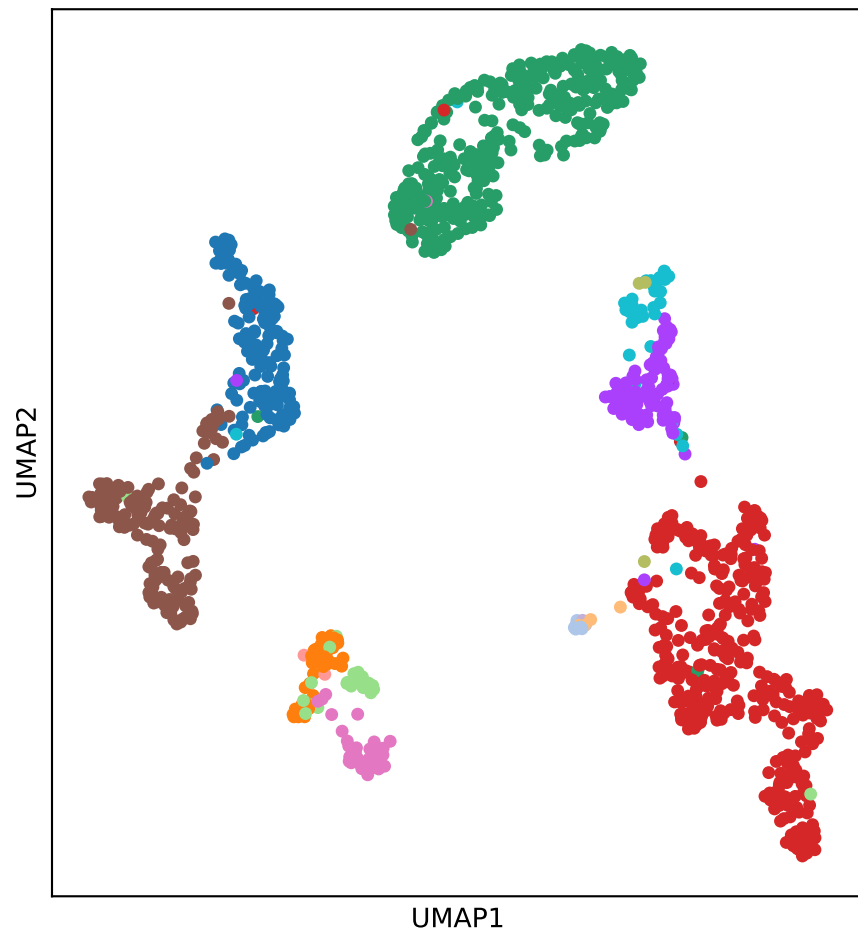
UMAP projection - Labels (val) (Res: 0.05, Iter: 46)



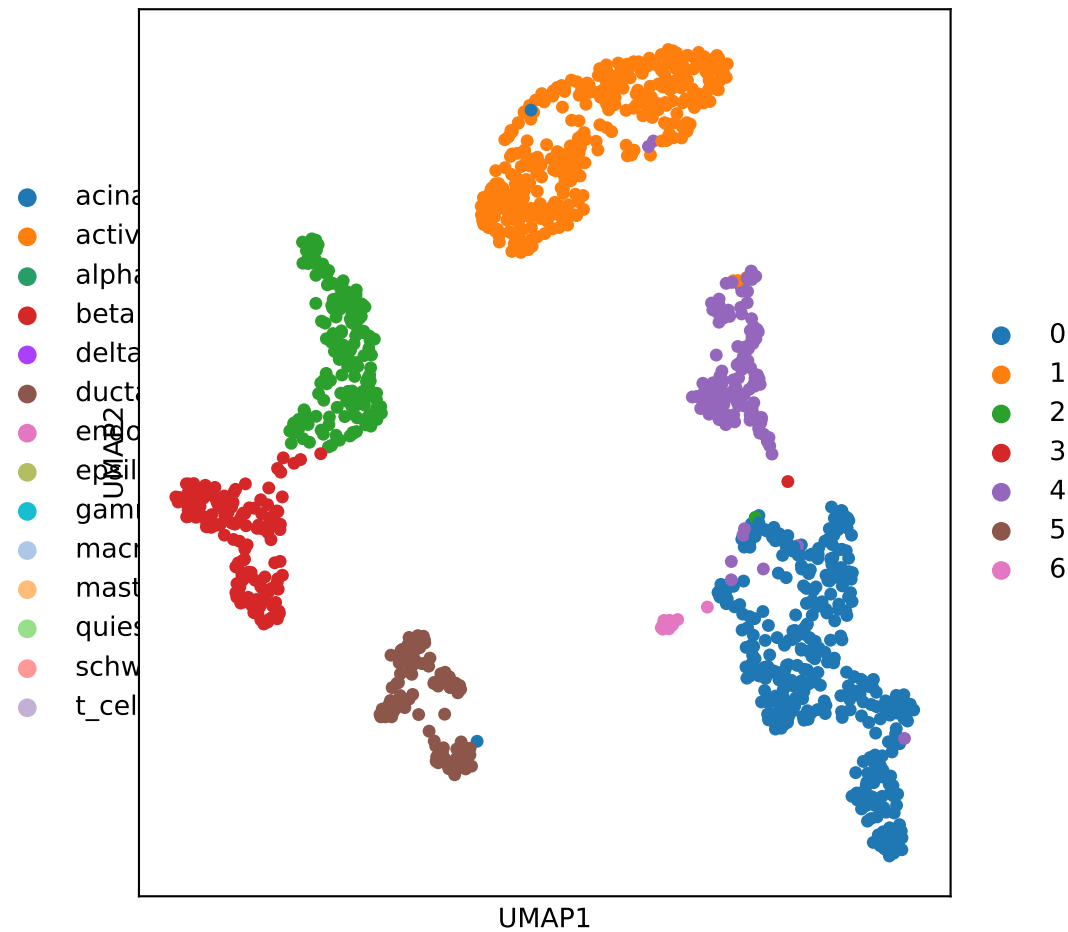
UMAP projection - Predictions (val) (Res: 0.05, Iter: 46)



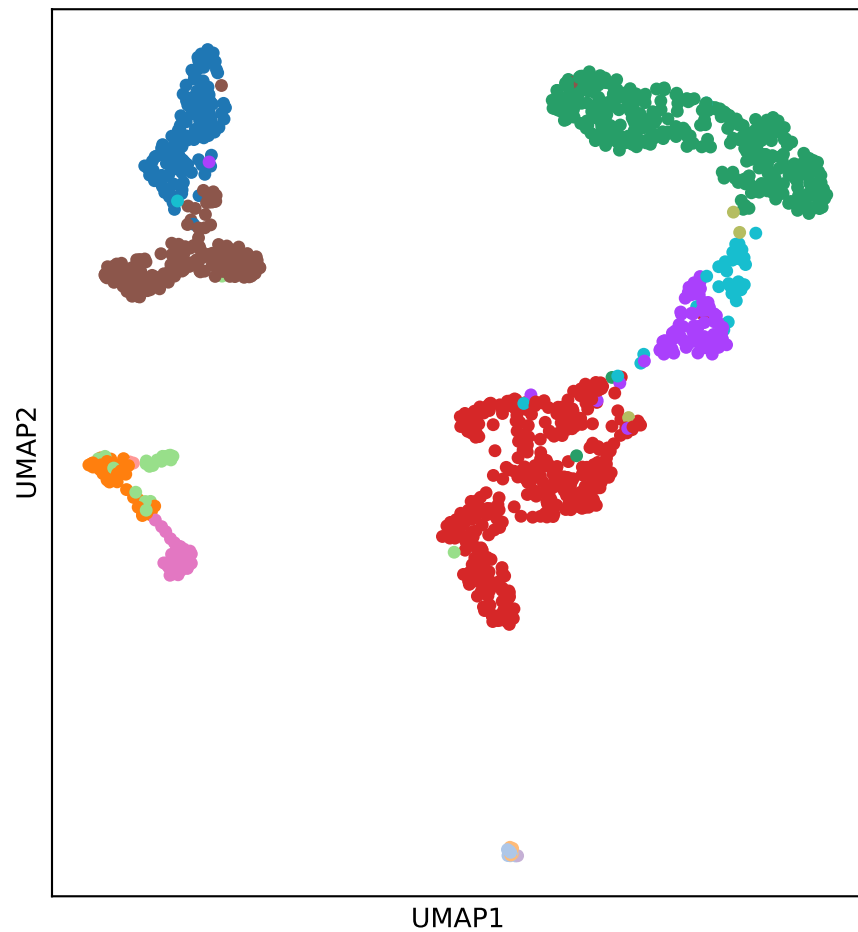
UMAP projection - Labels (val) (Res: 0.06, Iter: 47)



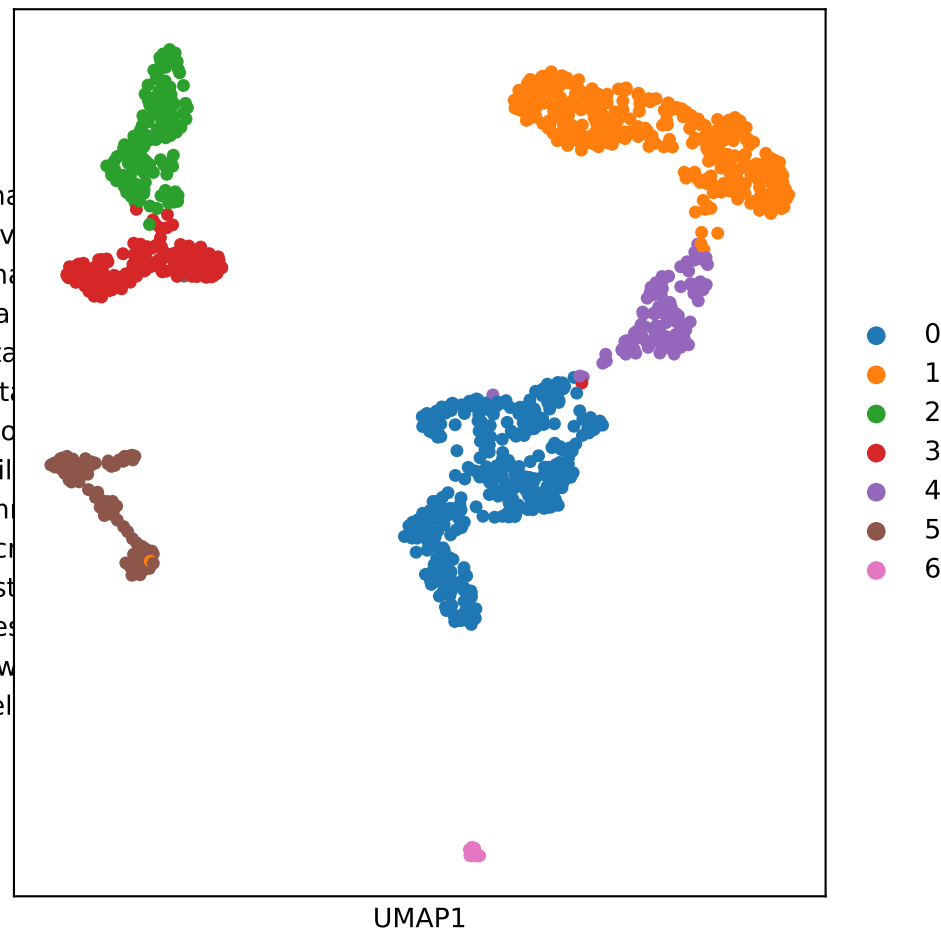
UMAP projection - Predictions (val) (Res: 0.06, Iter: 47)



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



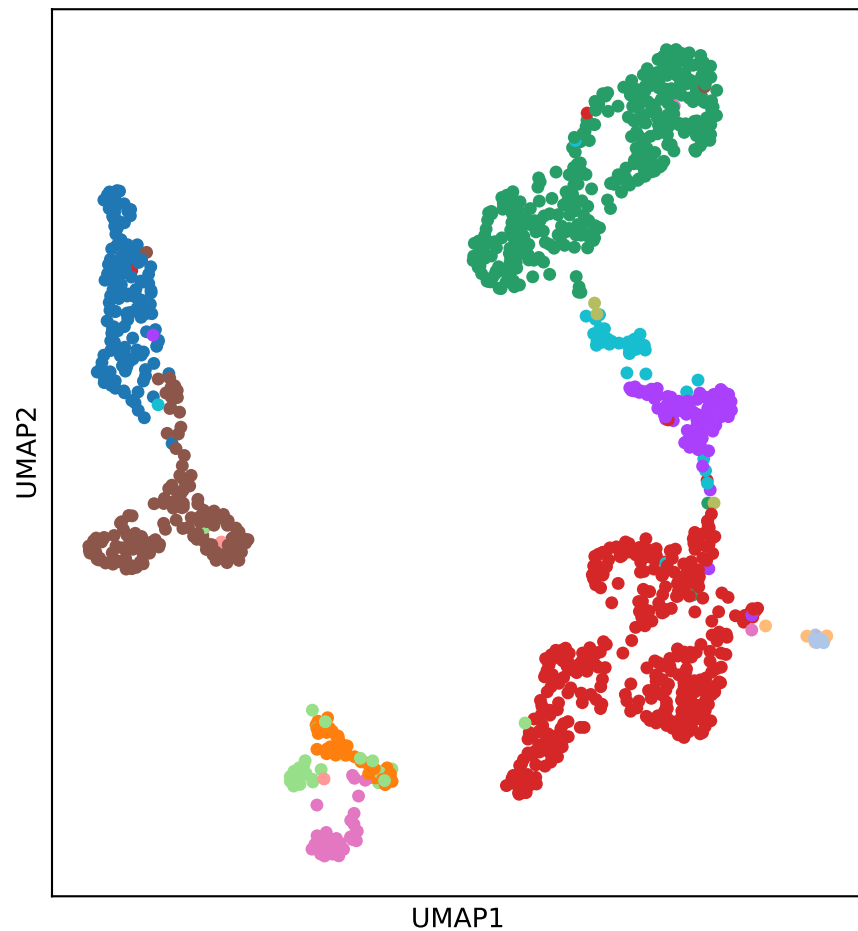
UMAP projection - Predictions (val) (Res: 0.07, Iter: 48)



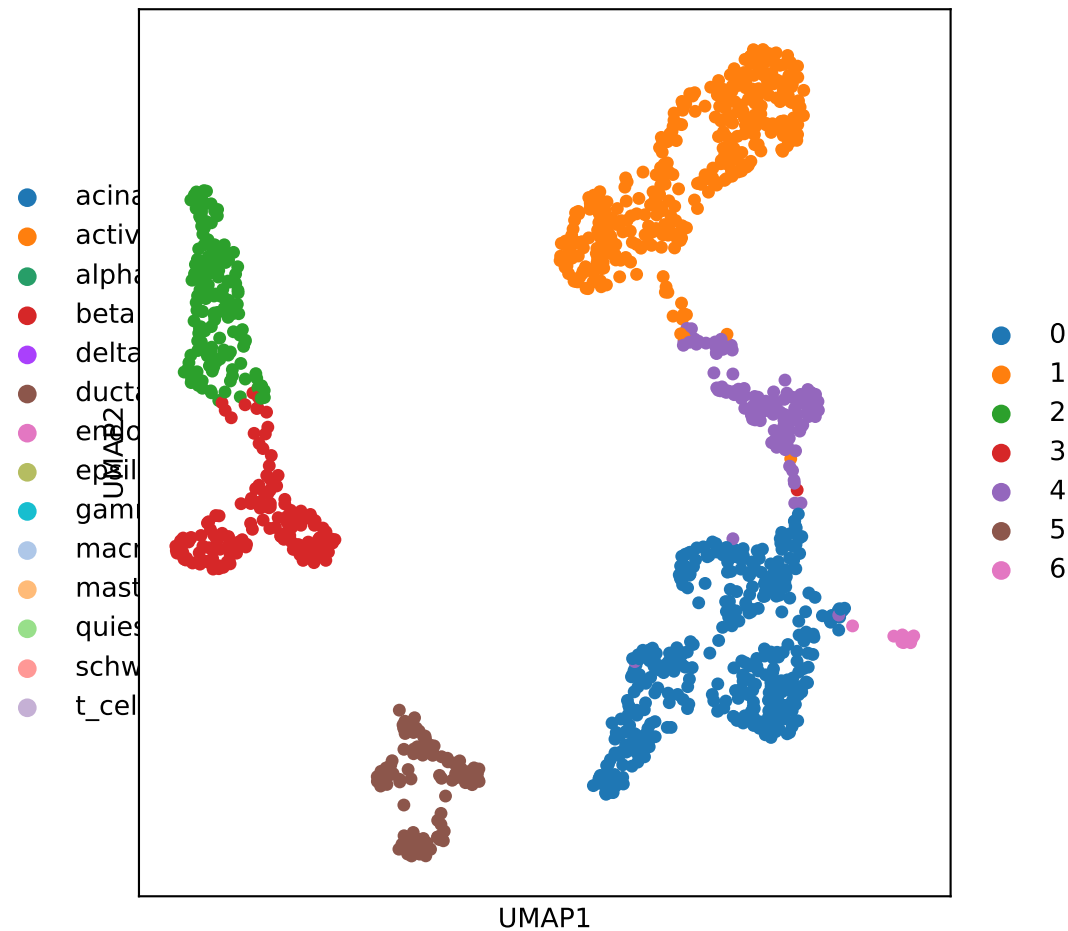
- acina
- activ
- alpha
- beta
- delta
- duct
- endo
- epil
- gam
- macr
- mast
- quies
- schw
- t_cel

- 0
- 1
- 2
- 3
- 4
- 5
- 6

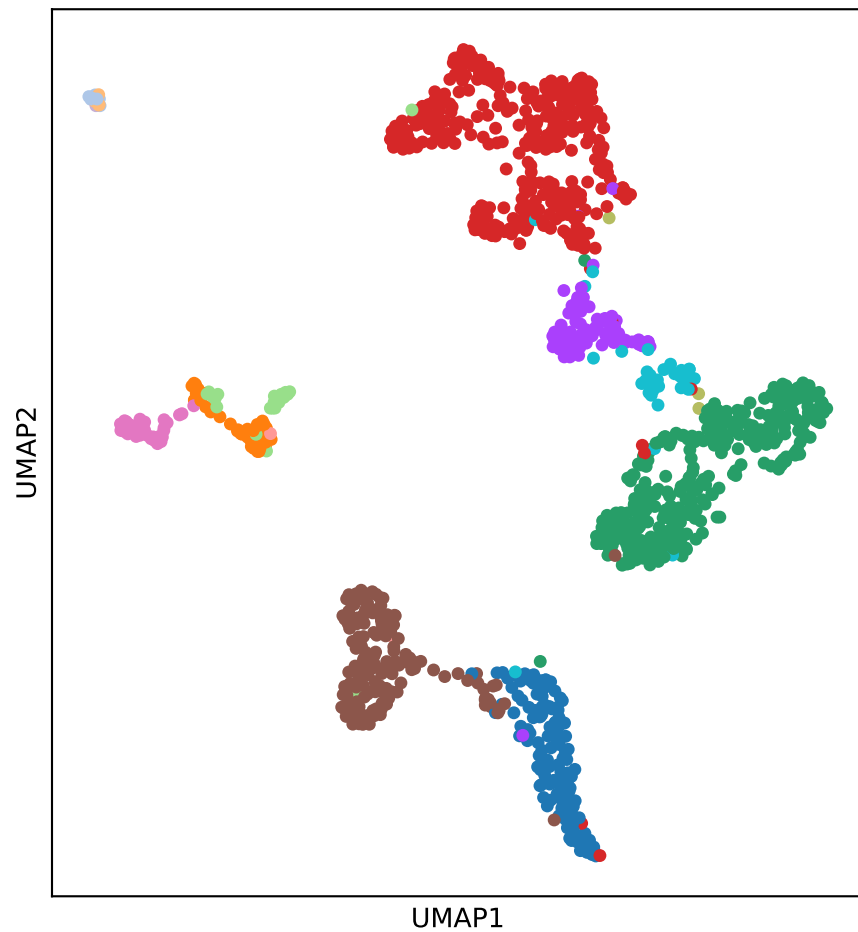
UMAP projection - Labels (val) (Res: 0.05, Iter: 49)



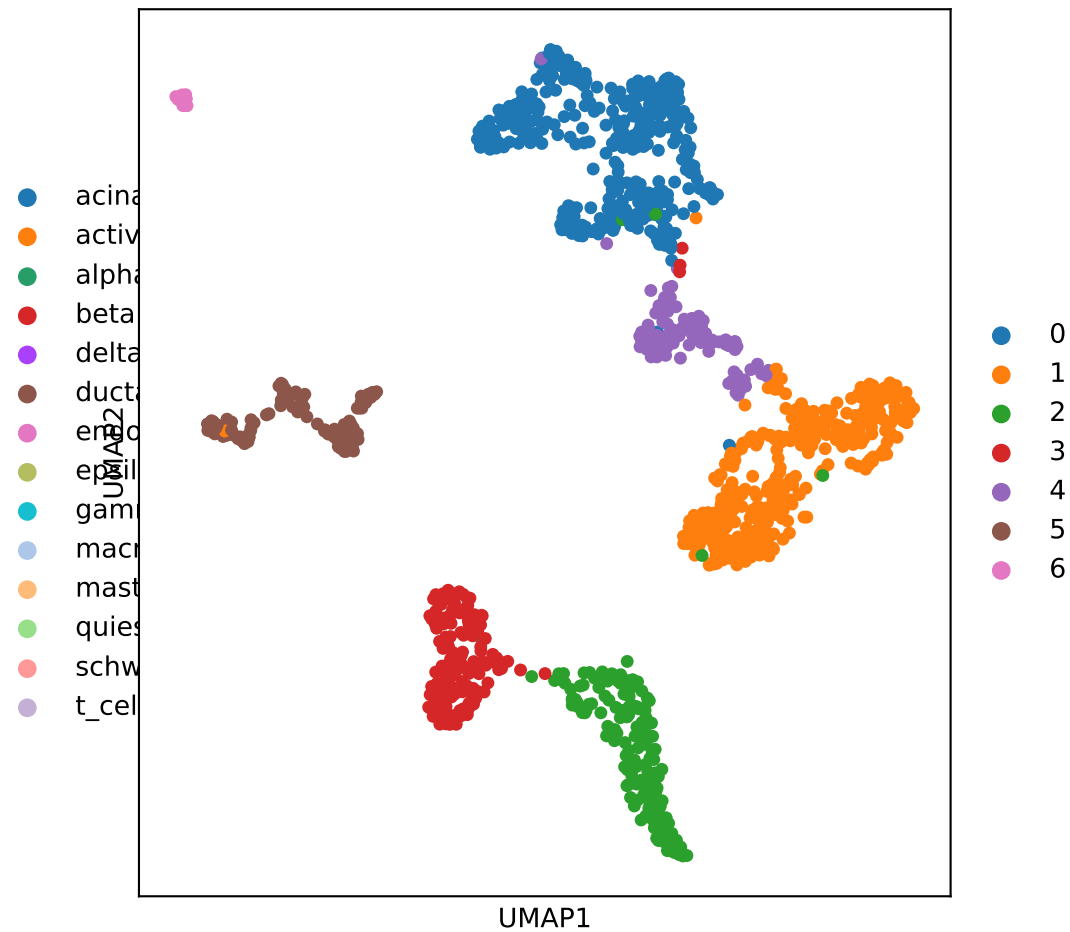
UMAP projection - Predictions (val) (Res: 0.05, Iter: 49)



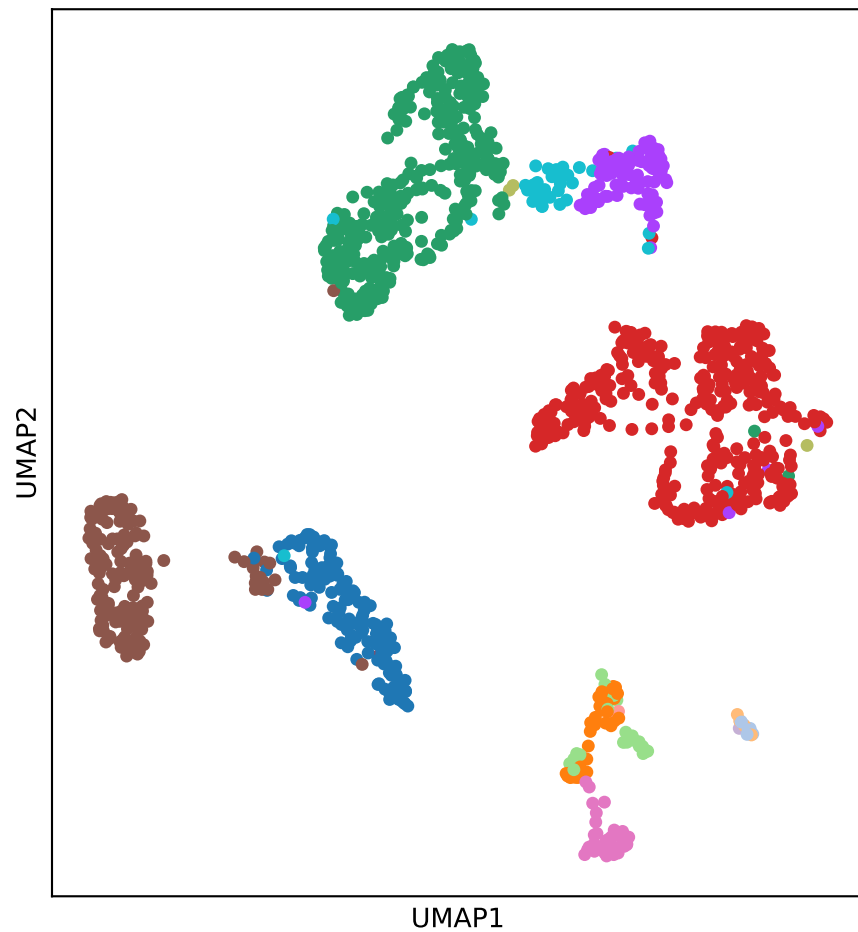
UMAP projection - Labels (val) (Res: 0.06, Iter: 50)



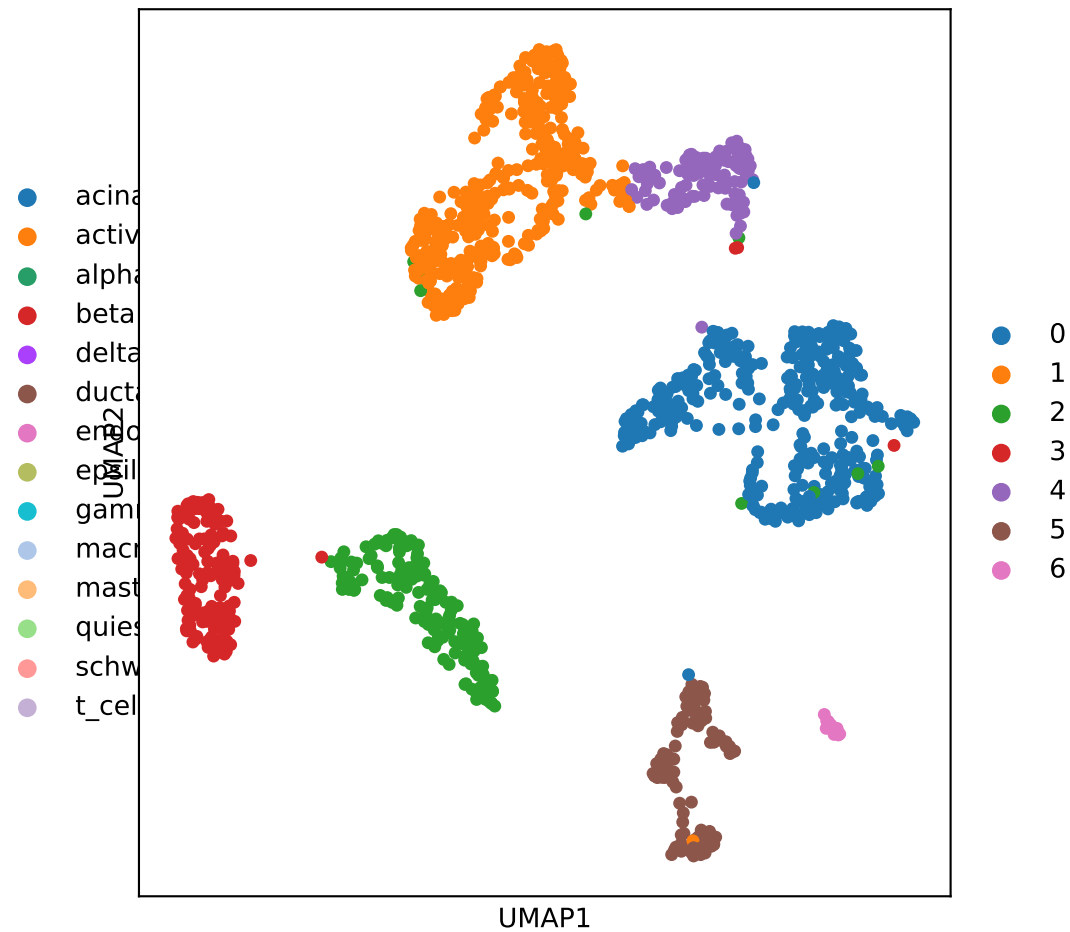
UMAP projection - Predictions (val) (Res: 0.06, Iter: 50)



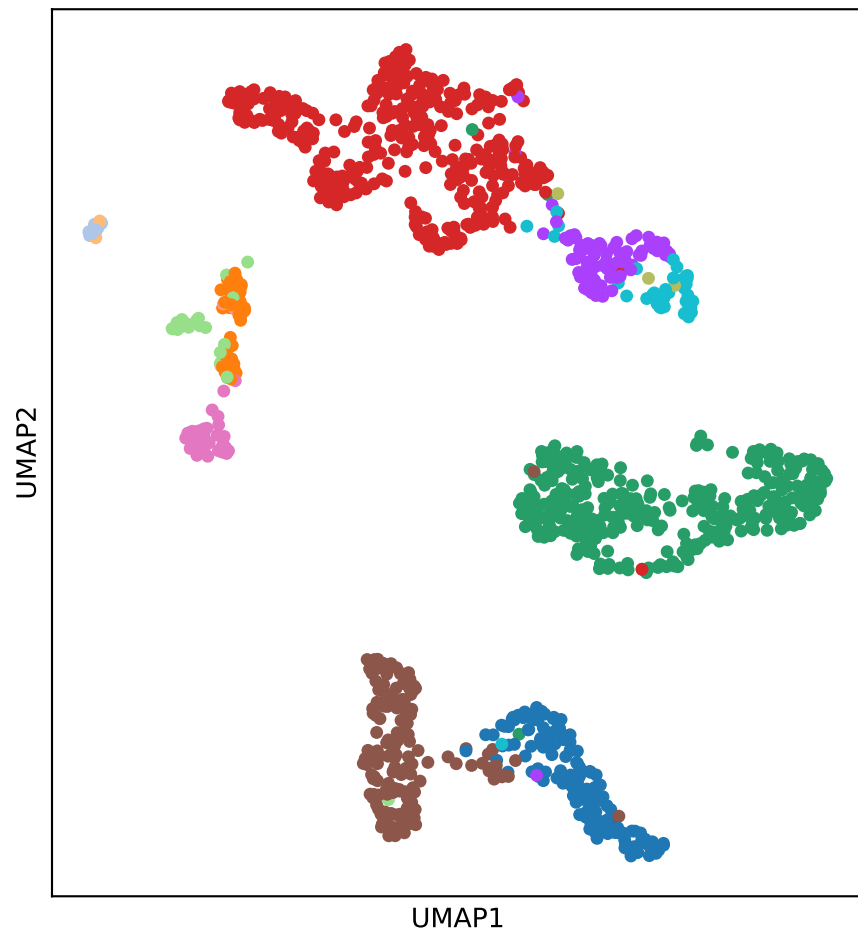
UMAP projection - Labels (val) (Res: 0.05, Iter: 51)



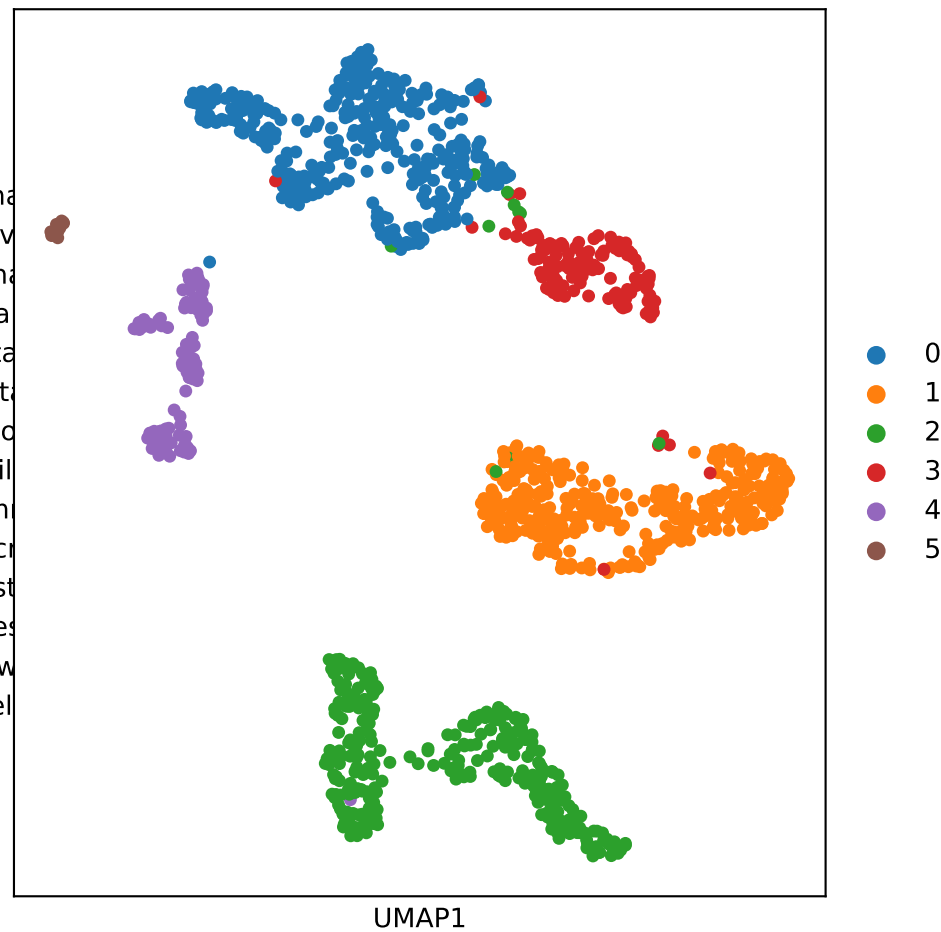
UMAP projection - Predictions (val) (Res: 0.05, Iter: 51)



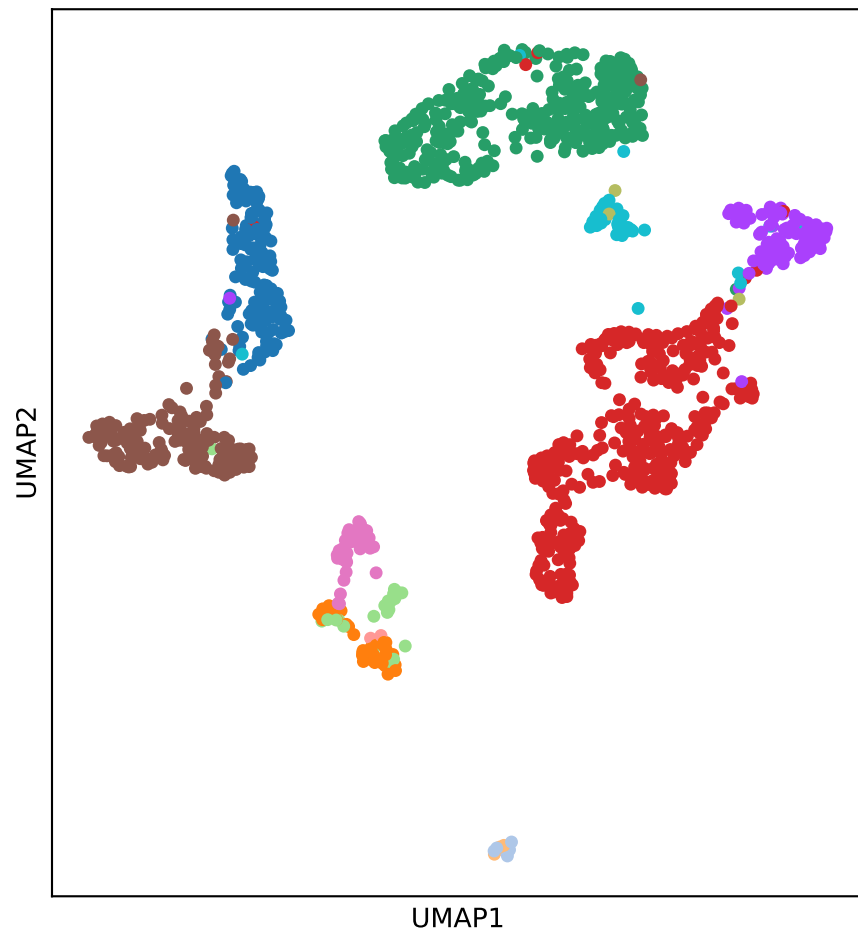
UMAP projection - Labels (val) (Res: 0.03, Iter: 52)



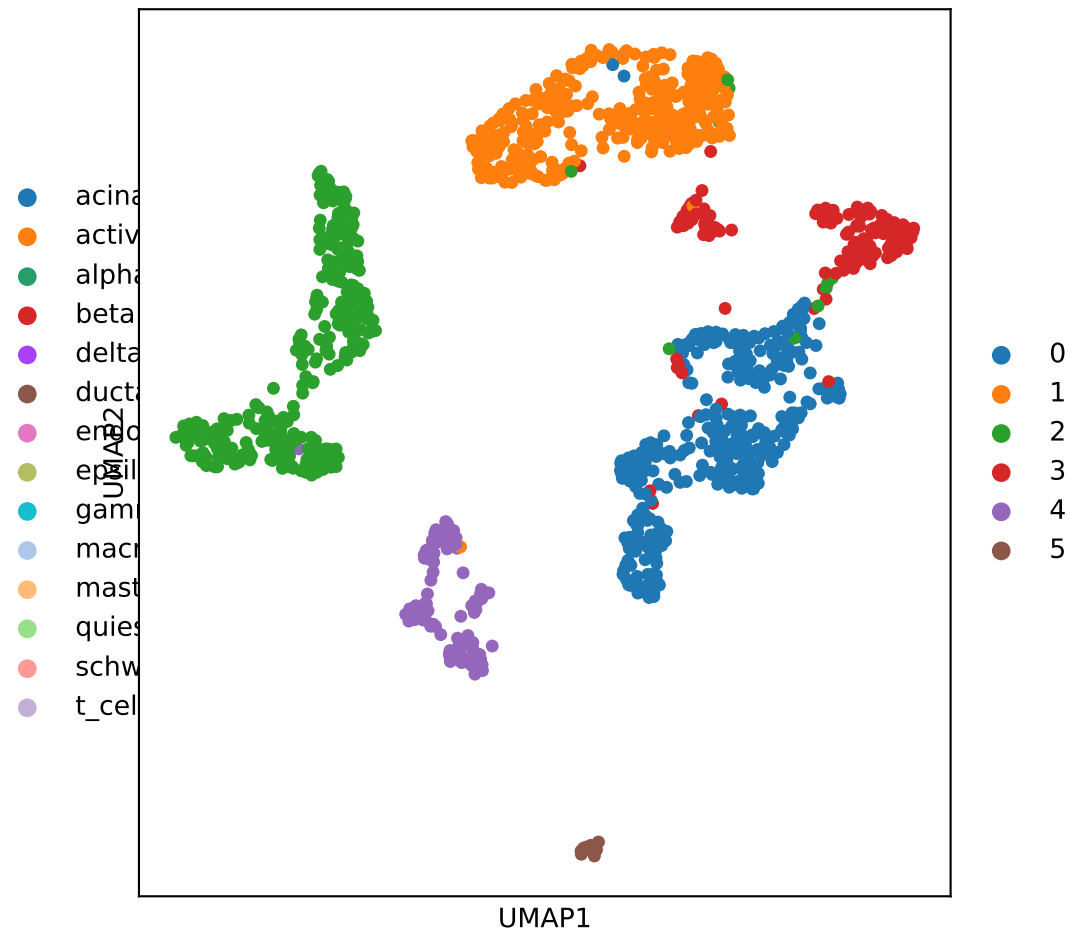
UMAP projection - Predictions (val) (Res: 0.03, Iter: 52)



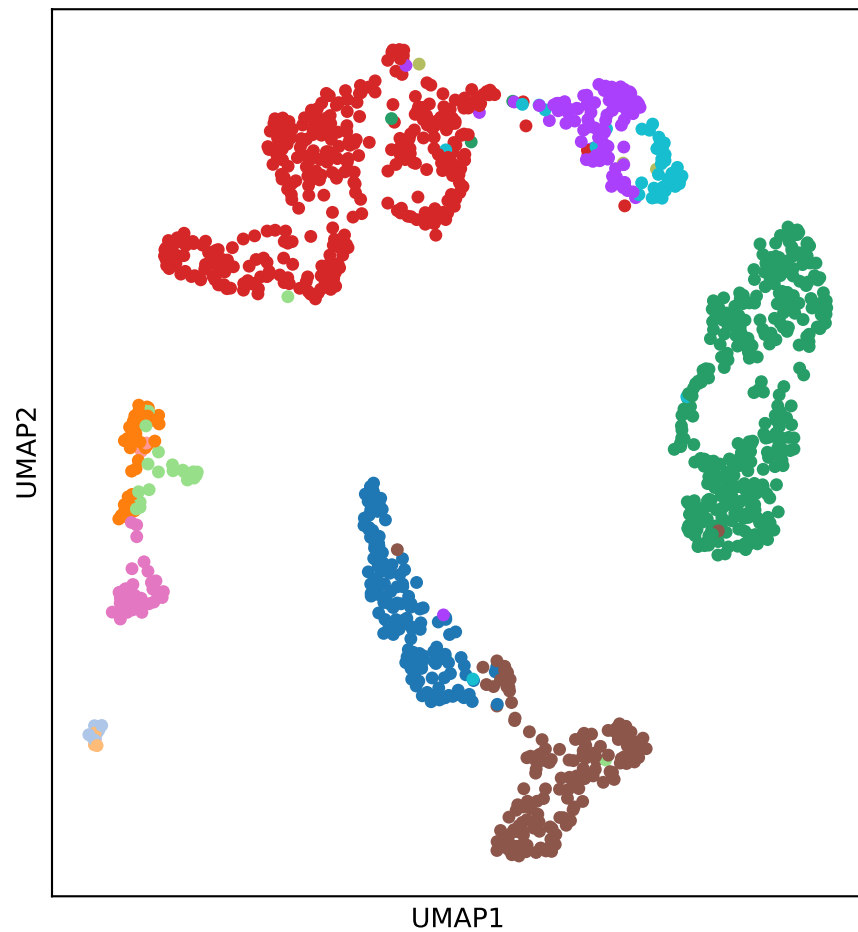
UMAP projection - Labels (val) (Res: 0.04, Iter: 53)



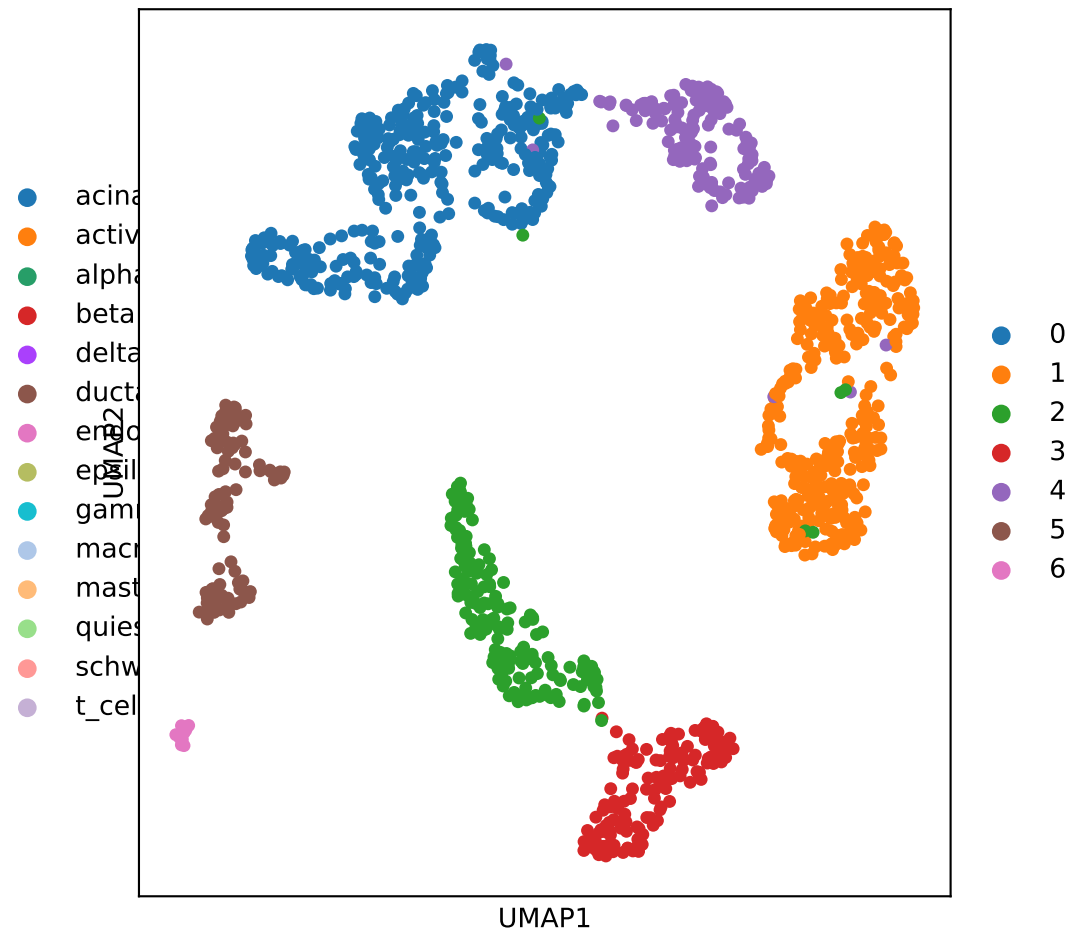
UMAP projection - Predictions (val) (Res: 0.04, Iter: 53)



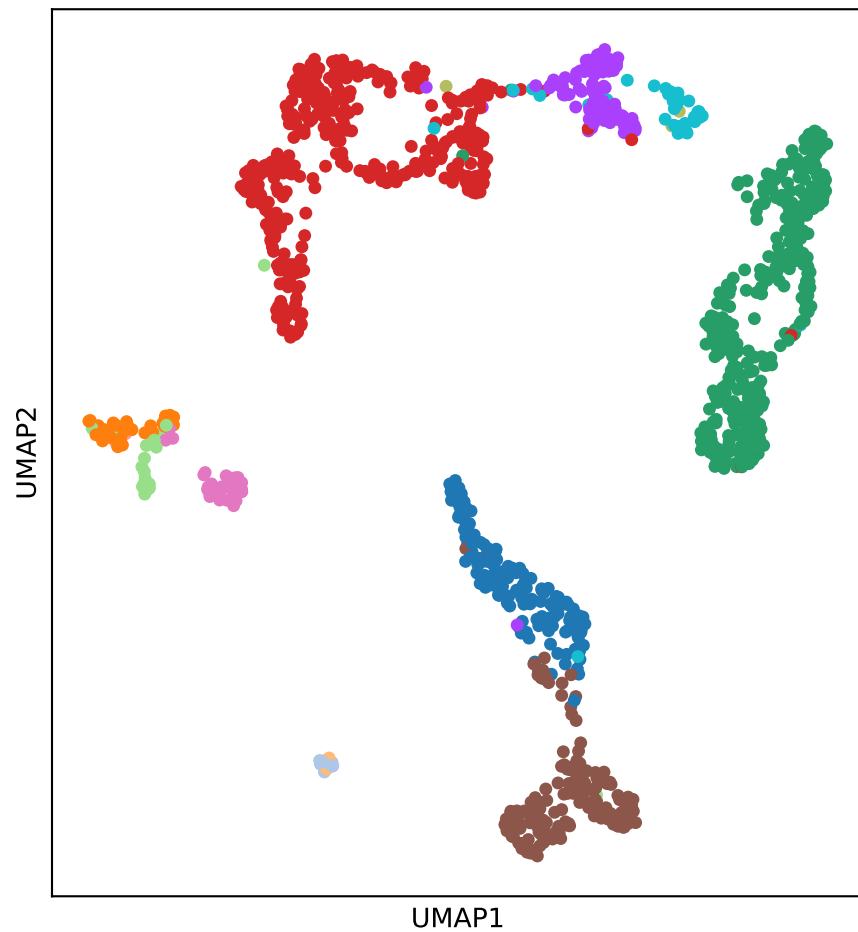
UMAP projection - Labels (val) (Res: 0.06, Iter: 54)



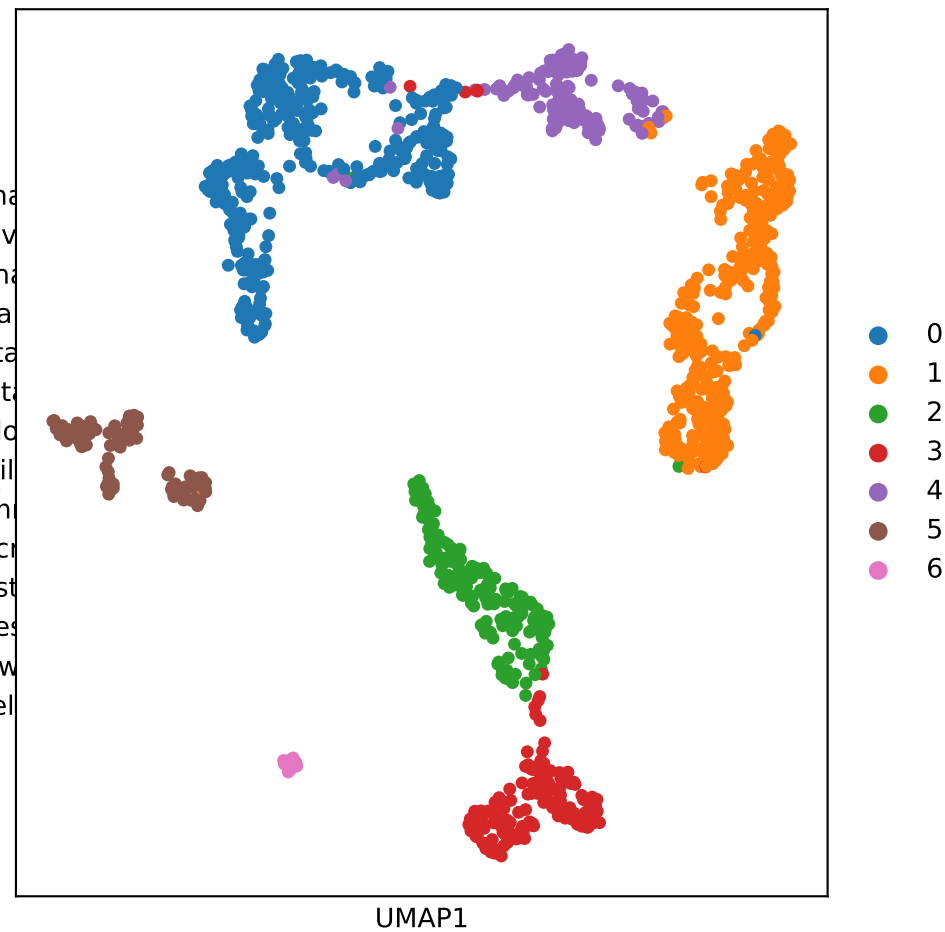
UMAP projection - Predictions (val) (Res: 0.06, Iter: 54)



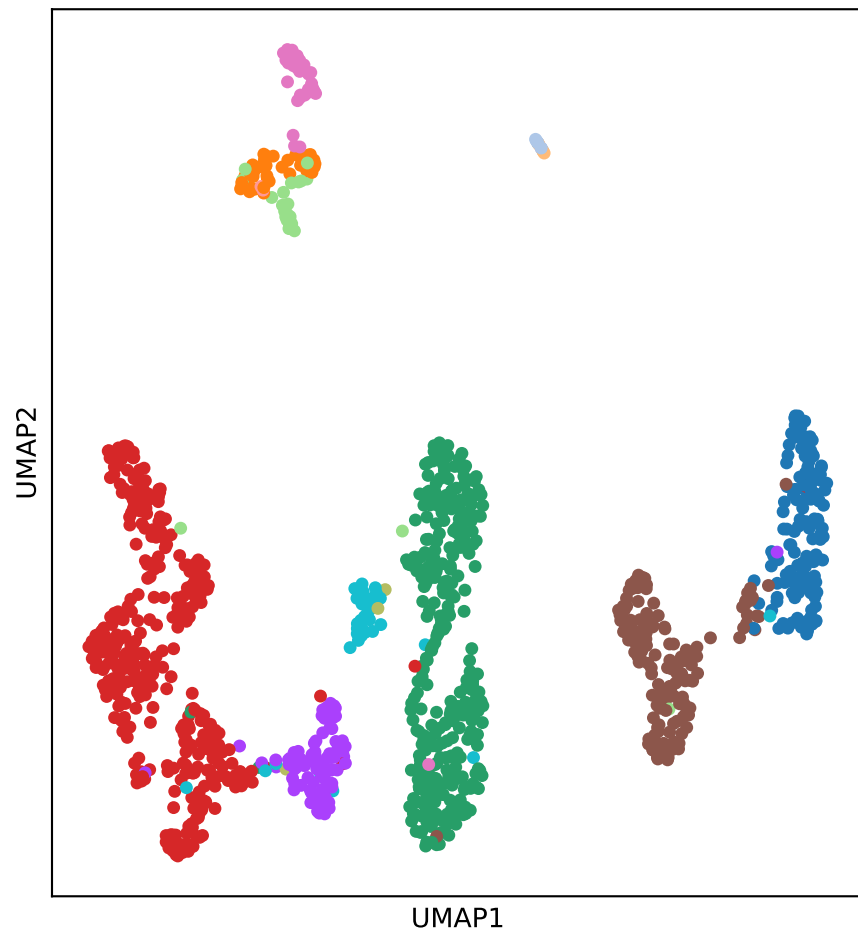
UMAP projection - Labels (val) (Res: 0.05, Iter: 55)



UMAP projection - Predictions (val) (Res: 0.05, Iter: 55)



UMAP projection - Labels (val) (Res: 0.06, Iter: 56)



UMAP projection - Predictions (val) (Res: 0.06, Iter: 56)

