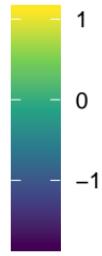


cell_health_modz_target_cc_late_mitosis_n_spots_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

10

cell_health_modz_target_cc_all_nucleus_roundness_m

UMAP 2

5

0

-5

-10

-10

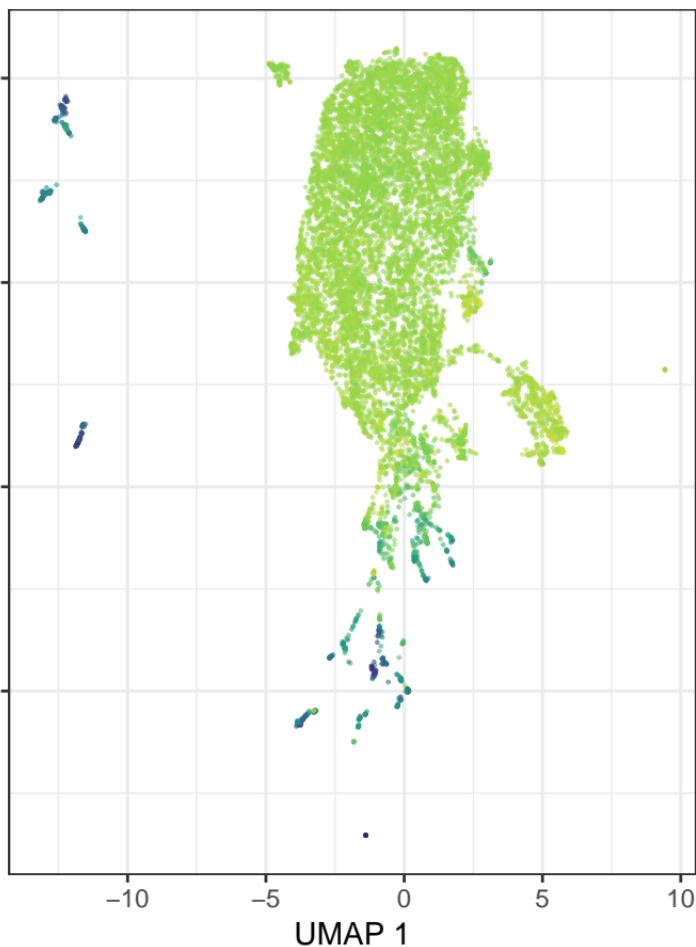
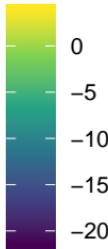
-5

5

10

UMAP 1

Prediction:

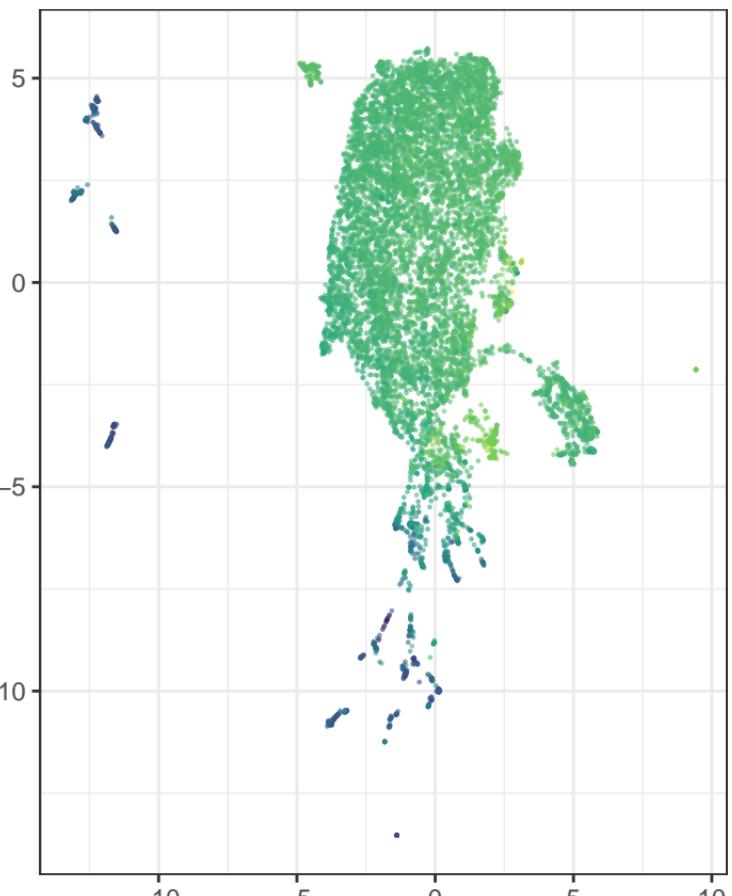
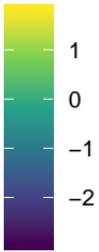


cell_health_modz_target_cc_polyplloid_n_objects

UMAP 2

UMAP 1

Prediction:

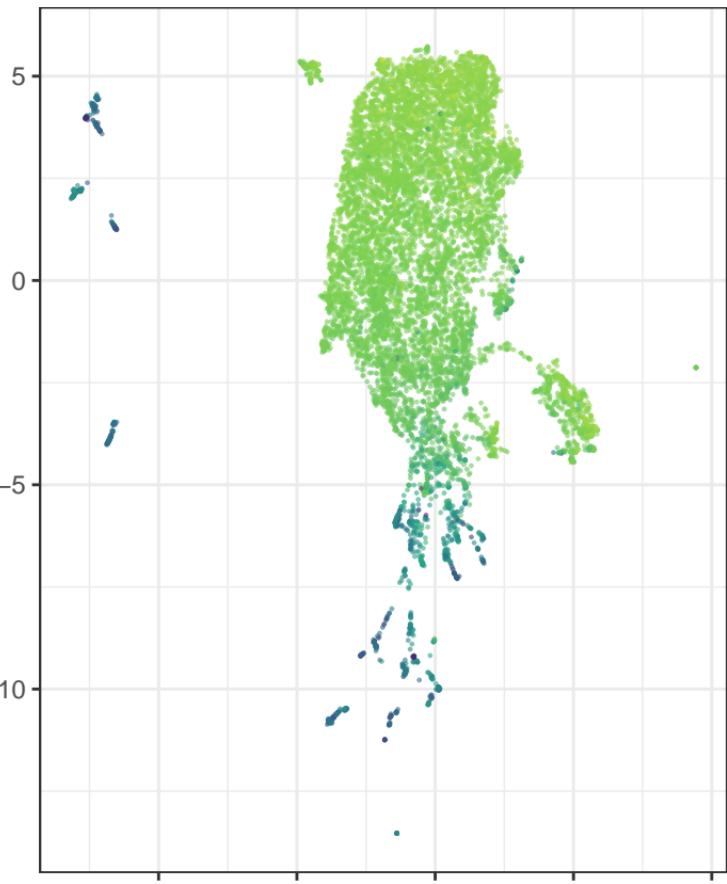
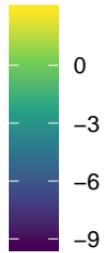


cell_health_modz_target_cc_g1_n_objects

UMAP 2

UMAP 1

Prediction:

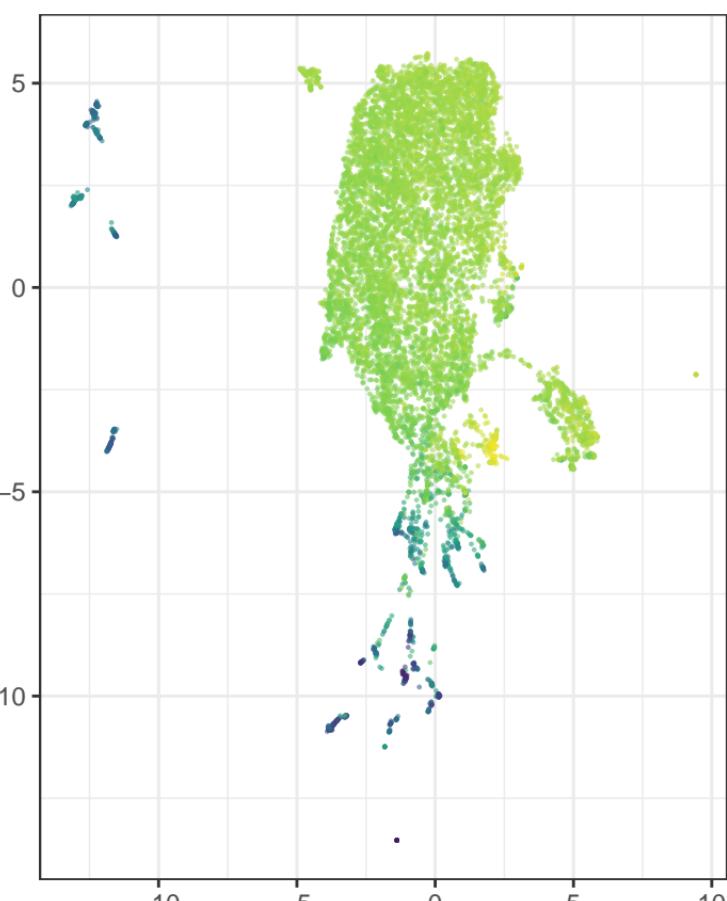
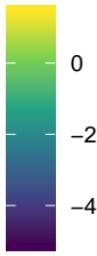


cell_health_modz_target_cc_mitosis_n_objects

UMAP 2

UMAP 1

Prediction:

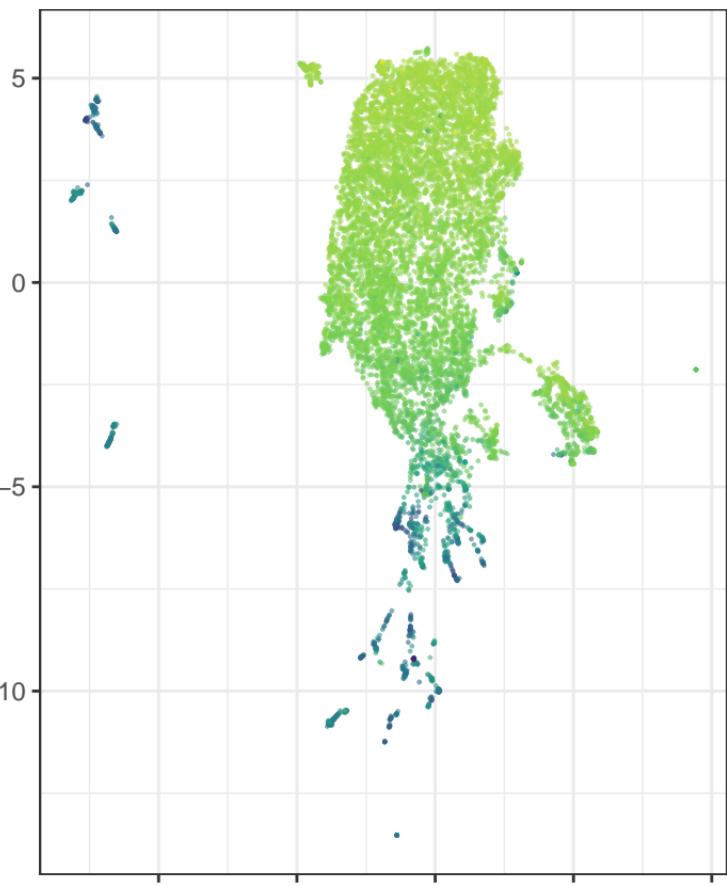
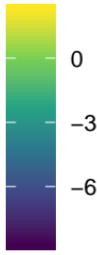


cell_health_modz_target_vb_num_live_cells

UMAP 2

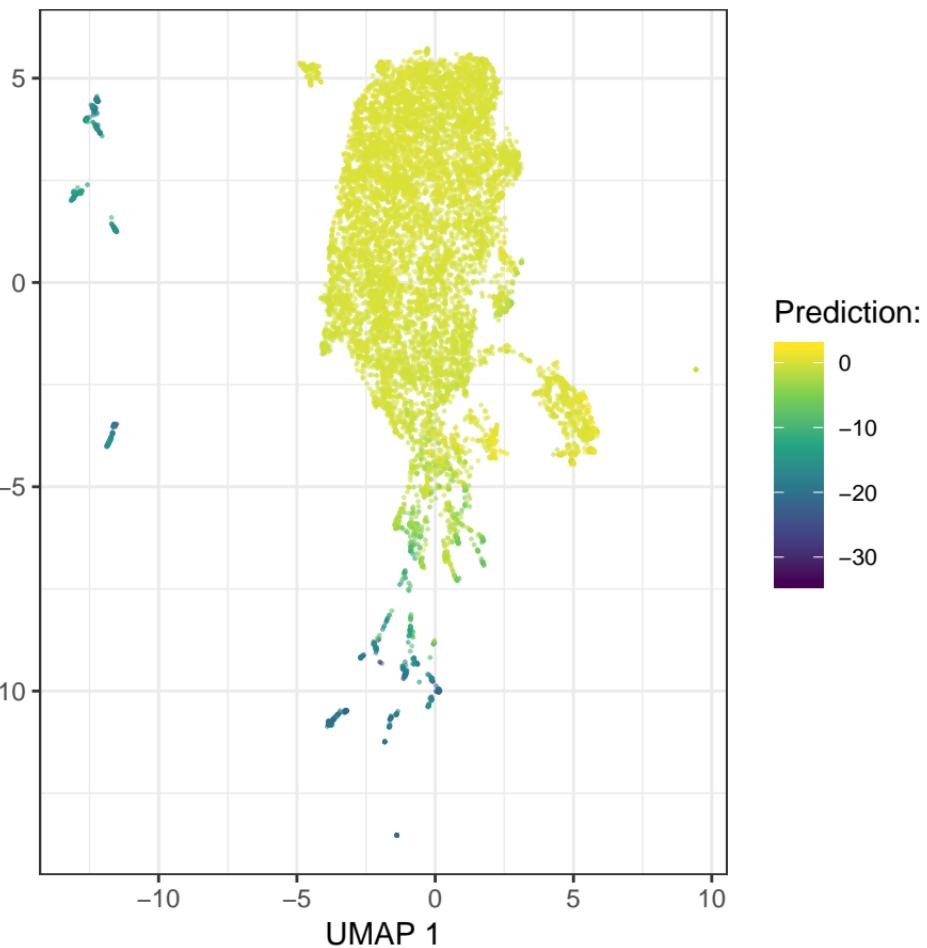
UMAP 1

Prediction:



cell_health_modz_target_vb_percent_live

UMAP 2



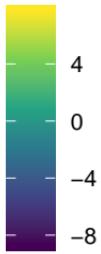
UMAP 1

cell_health_modz_target_vb_live_cell_roundness

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

5

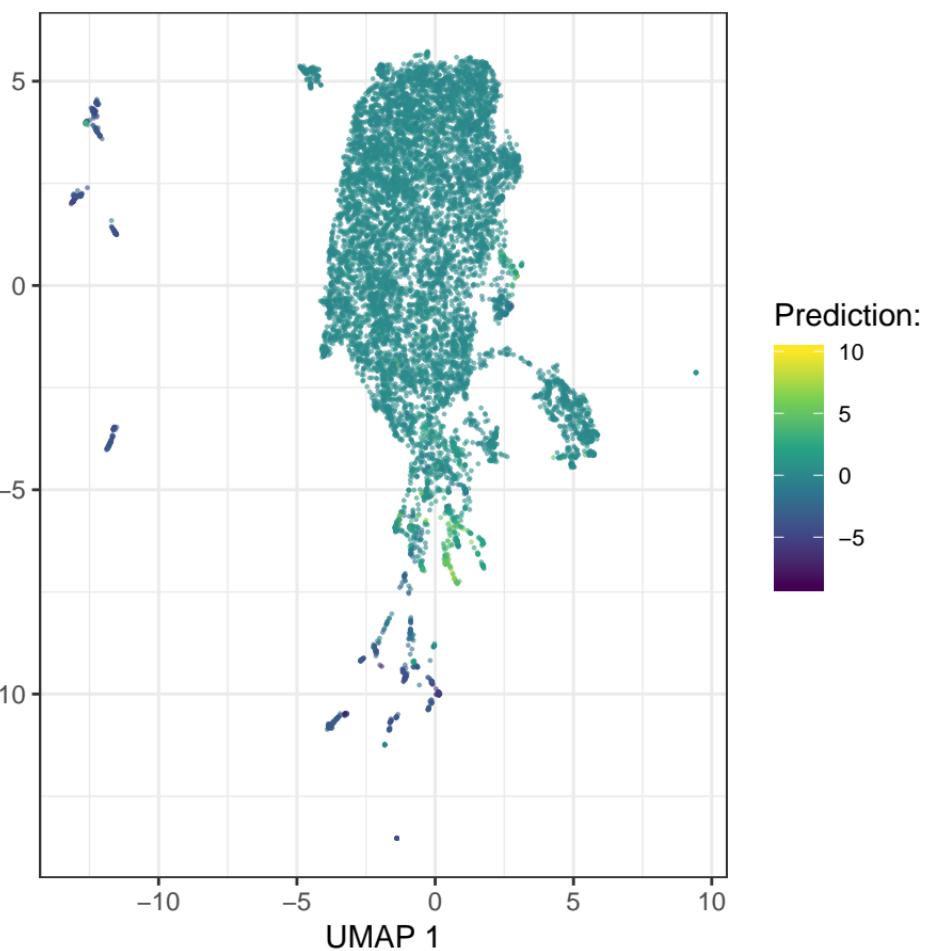
10

-10

-5

cell_health_modz_target_cc_all_large_notround_polynu

UMAP 2

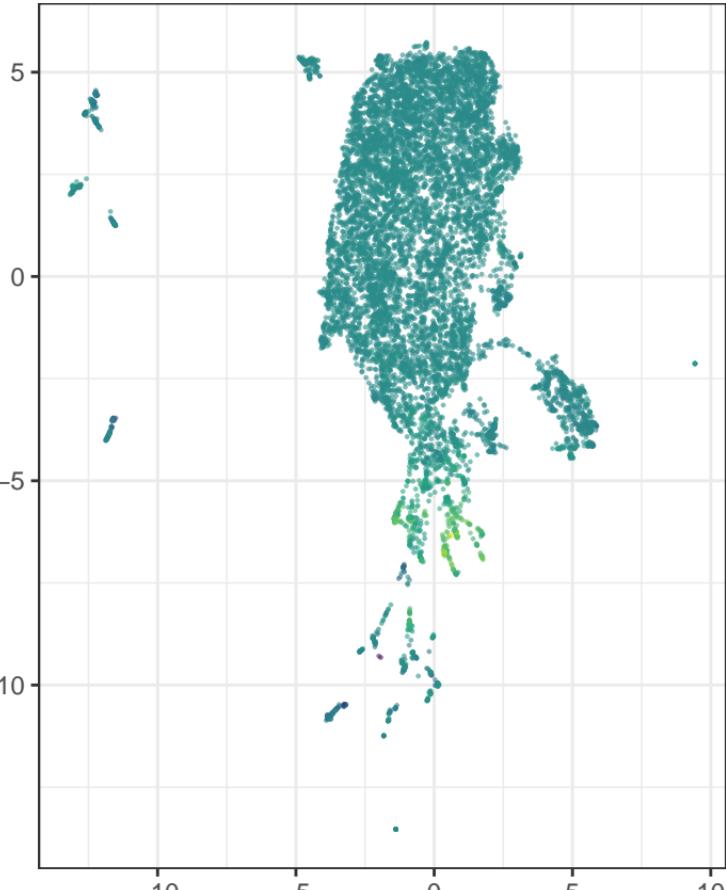
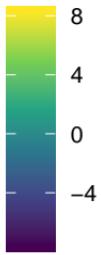


cell_health_modz_target_cc_all_high_h2ax

UMAP 2

UMAP 1

Prediction:

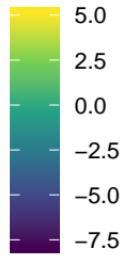


cell_health_modz_target_cc_early_mitosis_high_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

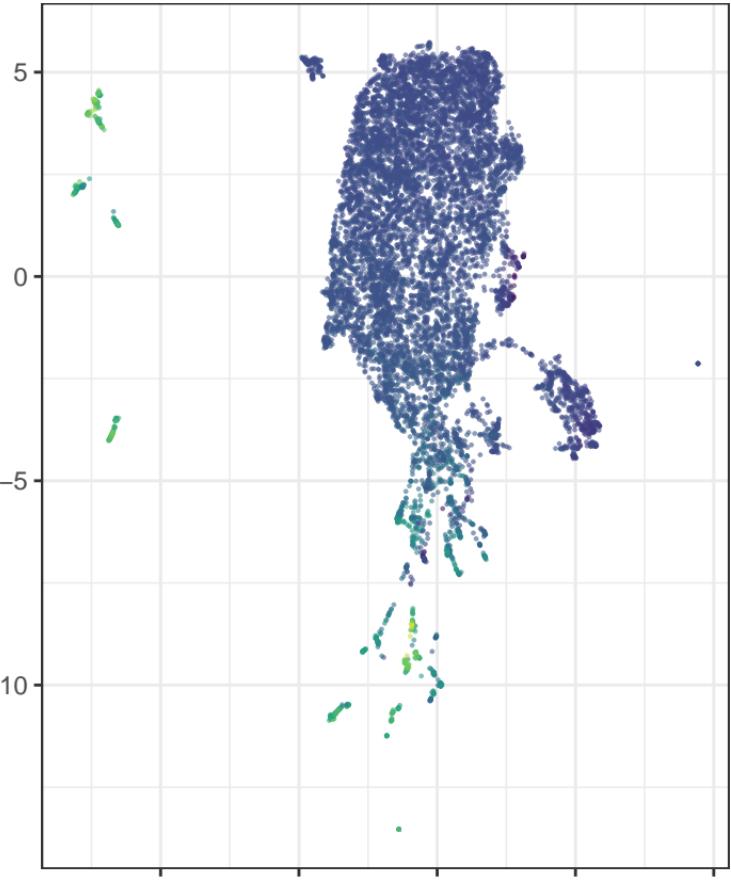
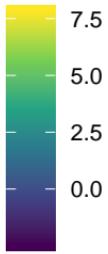
10

cell_health_modz_target_cc_g2_n_spots_h2ax_mean

UMAP 2

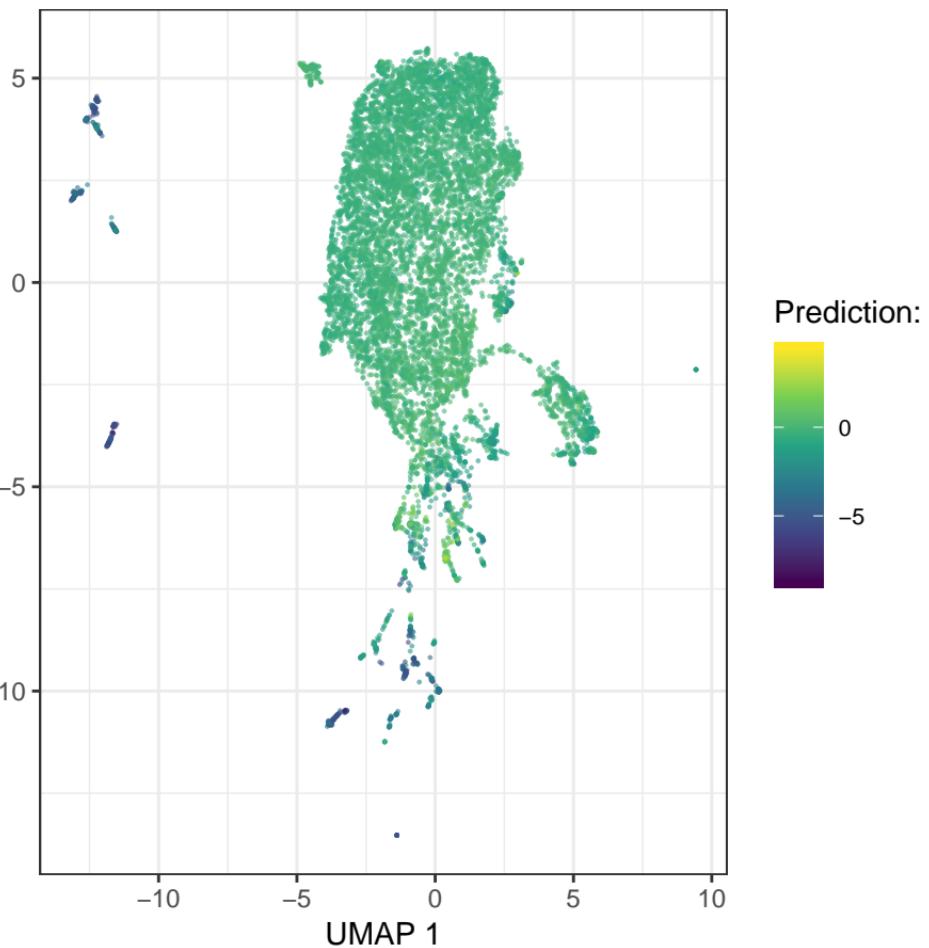
UMAP 1

Prediction:



cell_health_modz_target_cc_g2_plus_all_m_count

UMAP 2

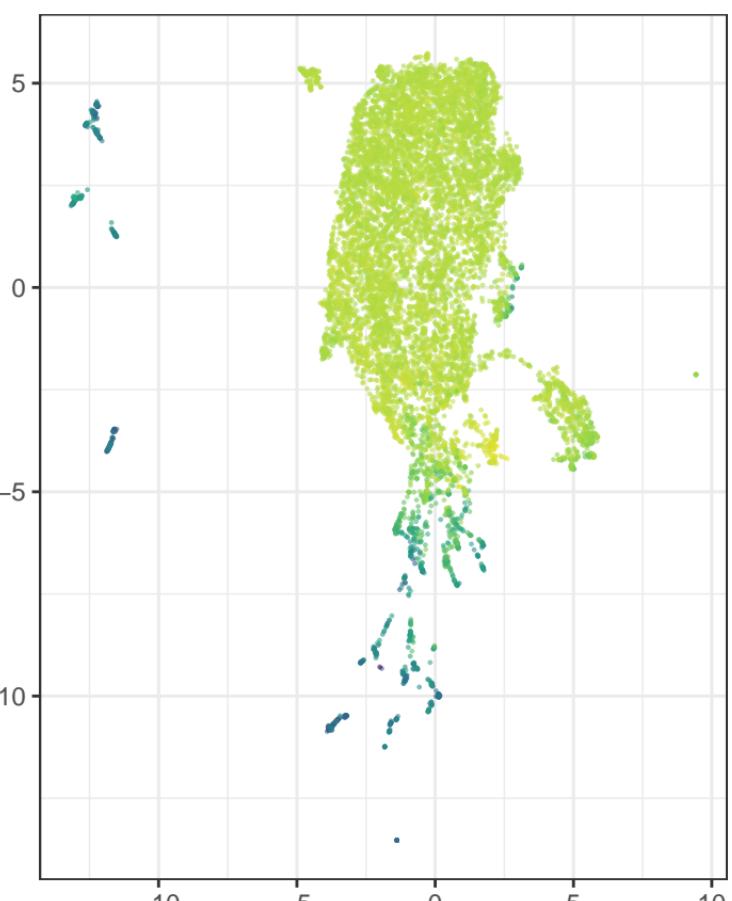
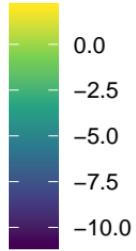


cell_health_modz_target_cc_early_mitosis_n_objects

UMAP 2

UMAP 1

Prediction:

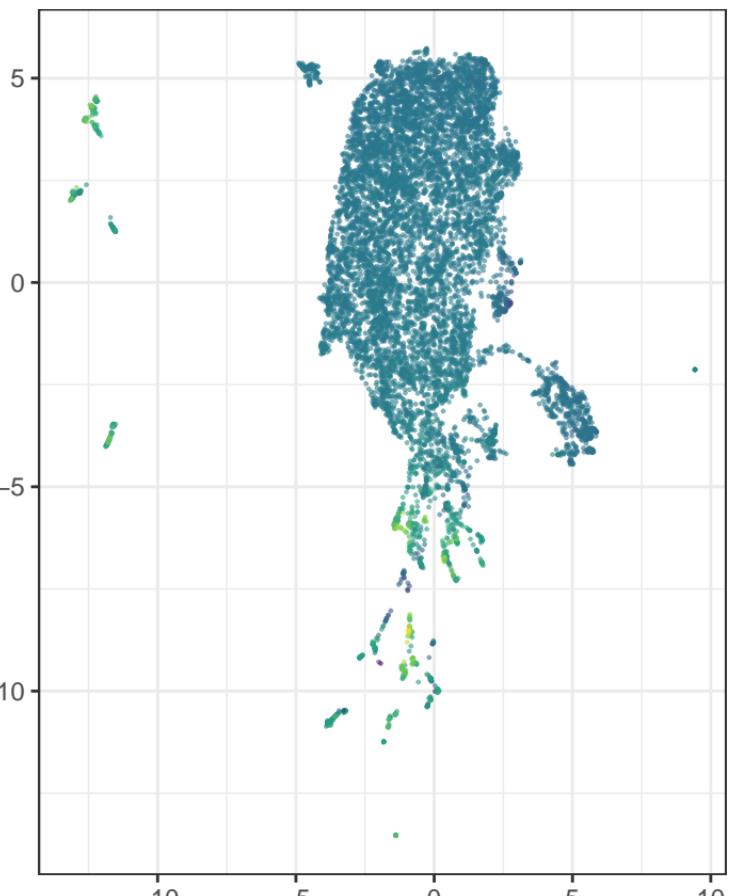
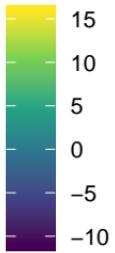


cell_health_modz_target_cc_all_nucleus_area_mean

UMAP 2

UMAP 1

Prediction:

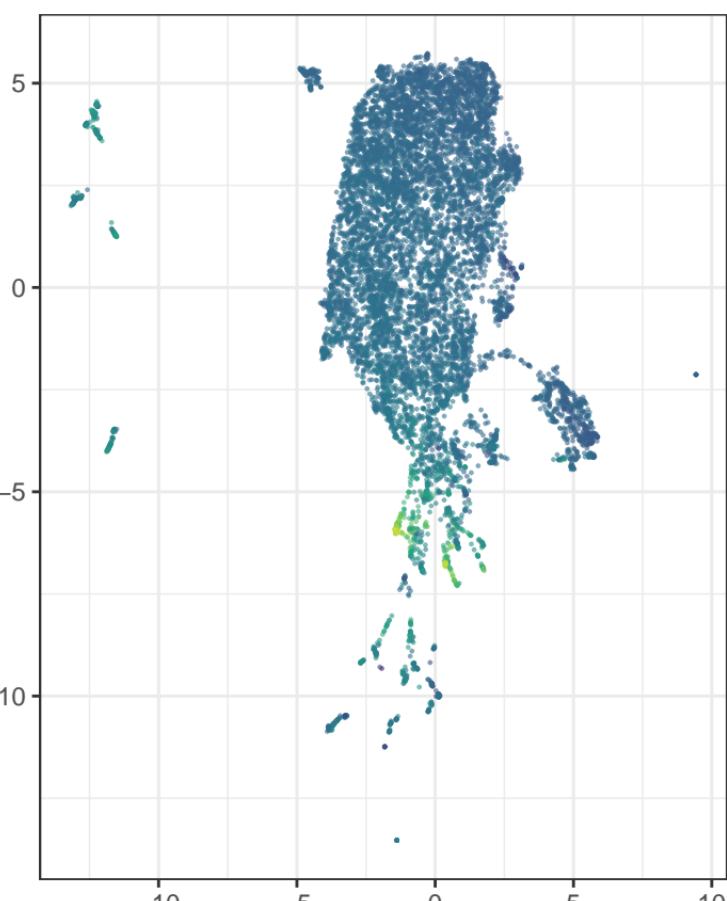
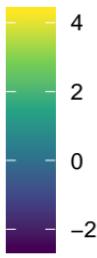


cell_health_modz_target_cc_g1_n_spots_h2ax_per_nuc

UMAP 2

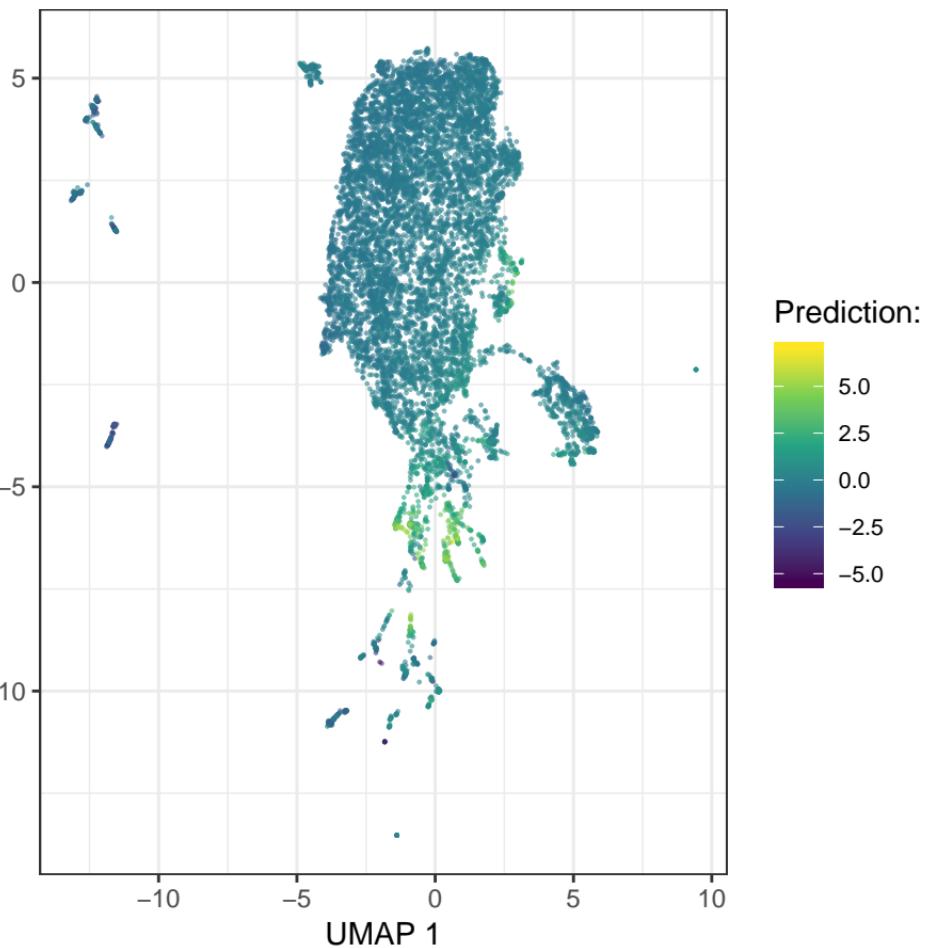
UMAP 1

Prediction:



cell_health_modz_target_vb_live_cell_area

UMAP 2

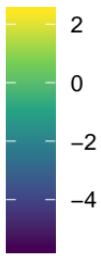


cell_health_modz_target_cc_polyplloid_n_spots_h2ax_m

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

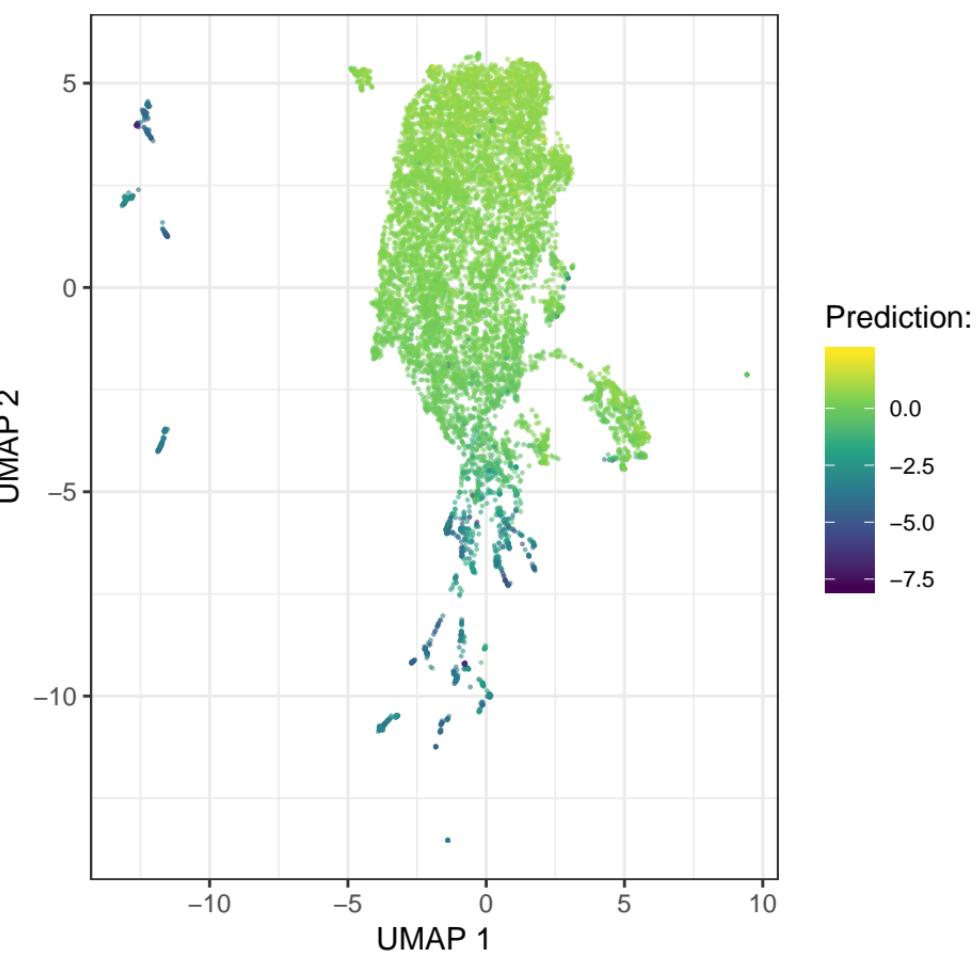
-5

5

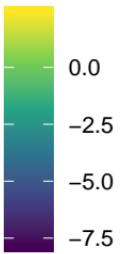
10

cell_health_modz_target_cc_all_n_objects

UMAP 2

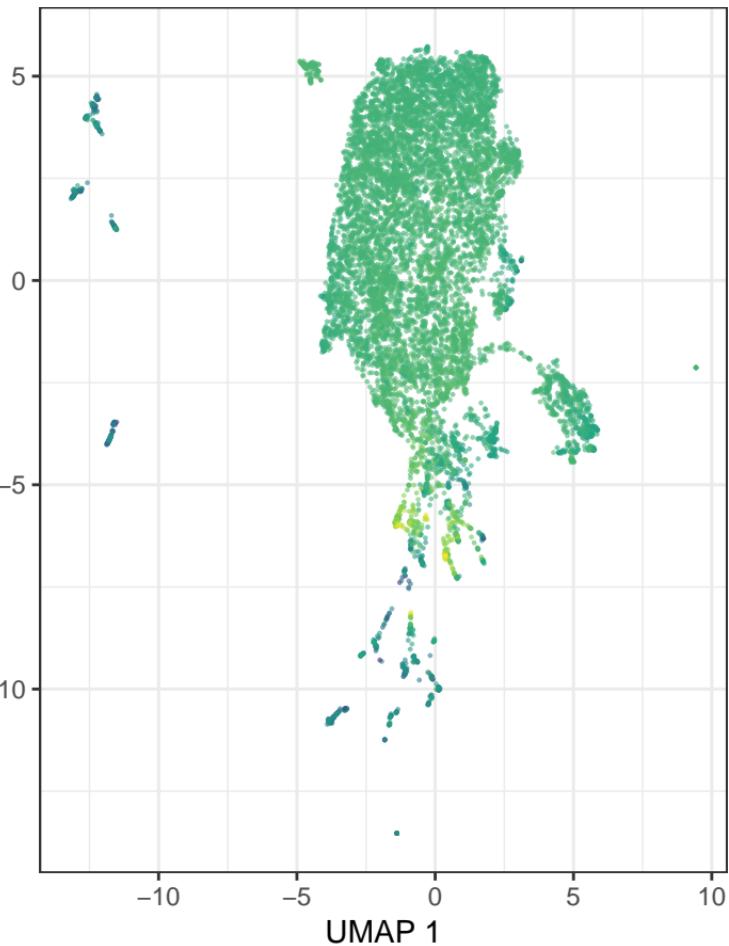


Prediction:

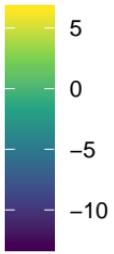


cell_health_modz_target_cc_cc_g2

UMAP 2



Prediction:

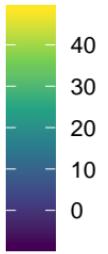


cell_health_modz_target_vb_ros_back_mean

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

10

5

-5

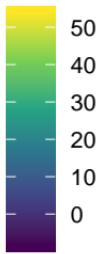
-10

cell_health_modz_target_vb_ros_mean

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

5

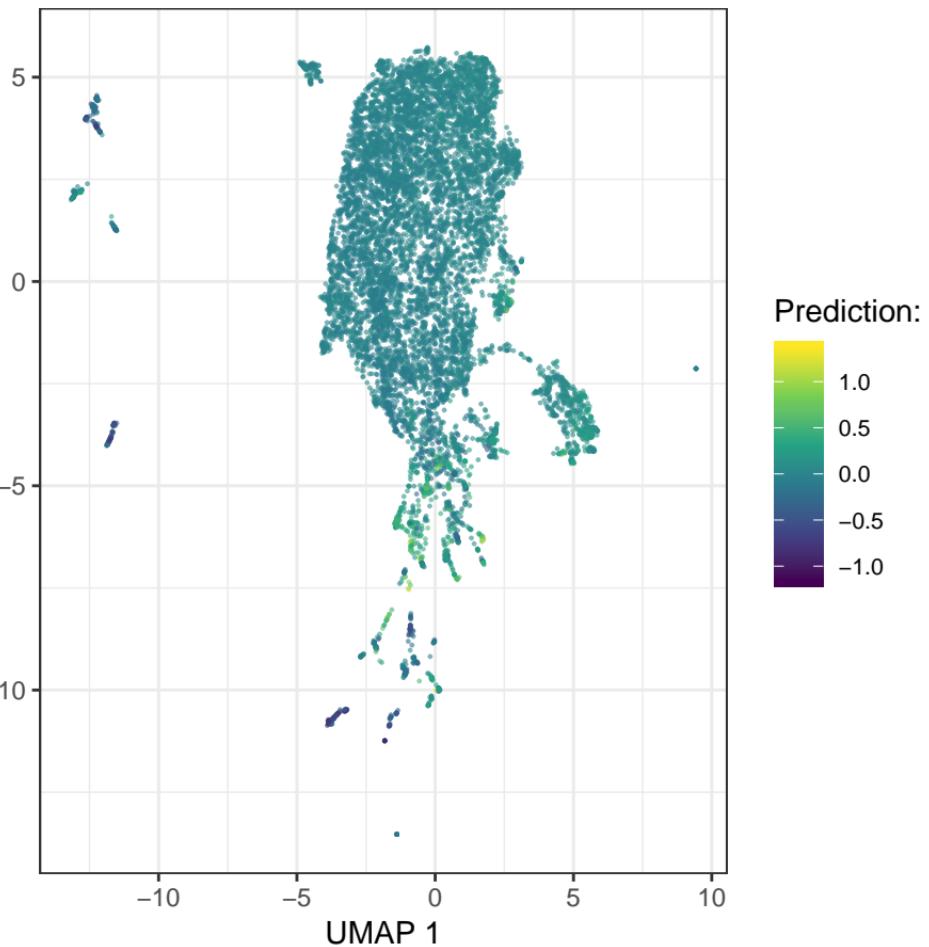
10

-5

-10

cell_health_modz_target_cc_infection_percentage

UMAP 2

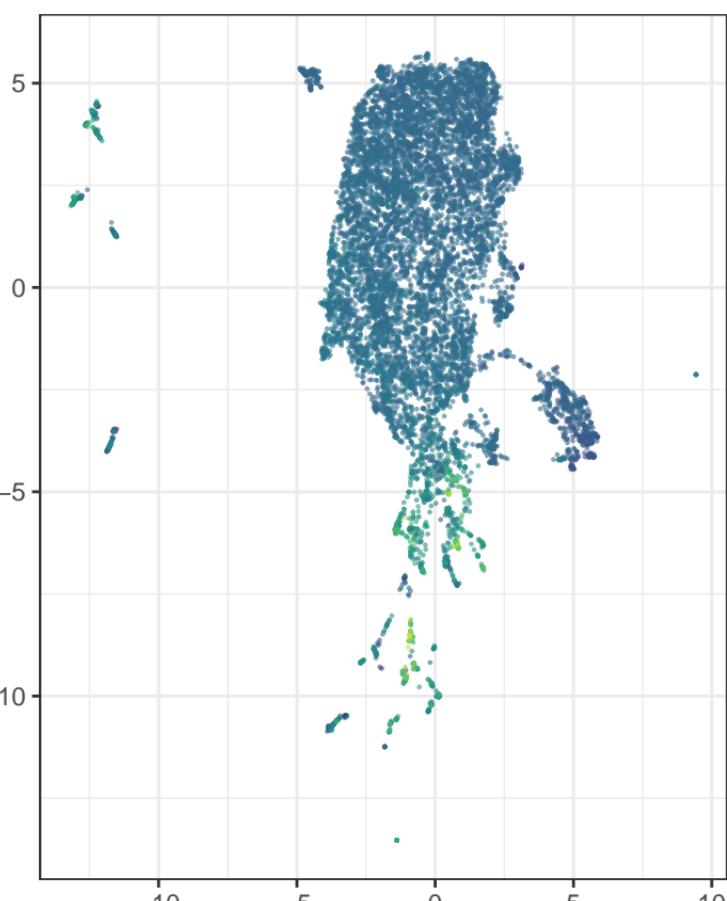
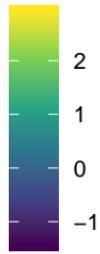


cell_health_modz_target_cc_early_mitosis_n_spots_h2

UMAP 2

UMAP 1

Prediction:

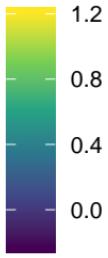


cell_health_modz_target_vb_percent_early_apoptosis

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

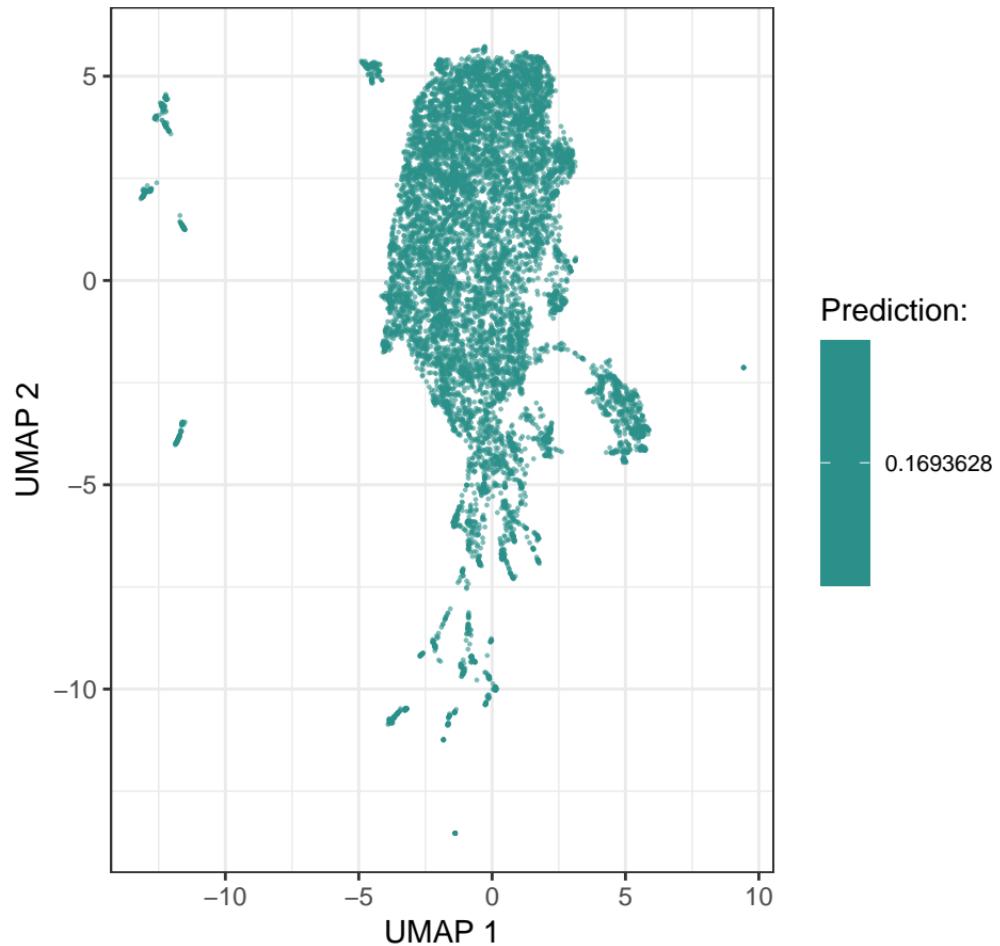
-10

-5

5

10

cell_health_modz_target_cc_mitosis_high_h2ax

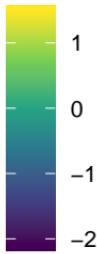


cell_health_modz_target_cc_polynuclear_n_spots_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

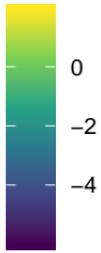
10

cell_health_modz_target_cc_polyplloid_high_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

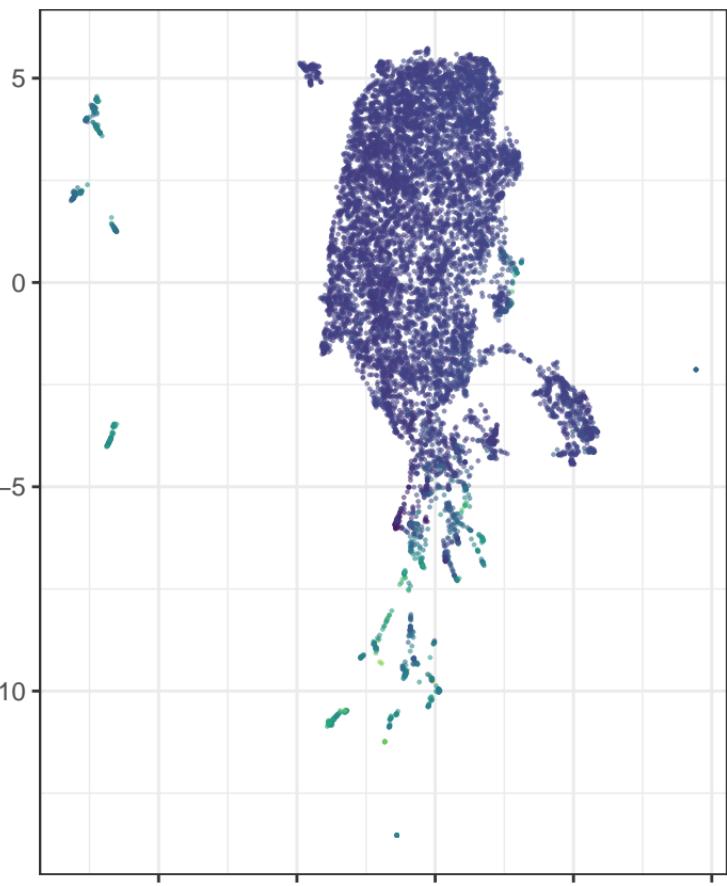
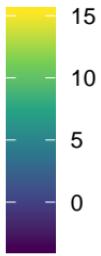
10

cell_health_modz_target_cc_cc_late_mitosis

UMAP 2

UMAP 1

Prediction:

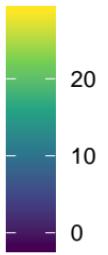


cell_health_modz_target_vb_percent_all_apoptosis

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

5 10

-5 -10

0

5

10

15

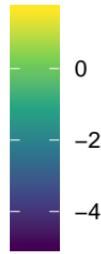
20

cell_health_modz_target_cc_polynuclear_n_spots_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

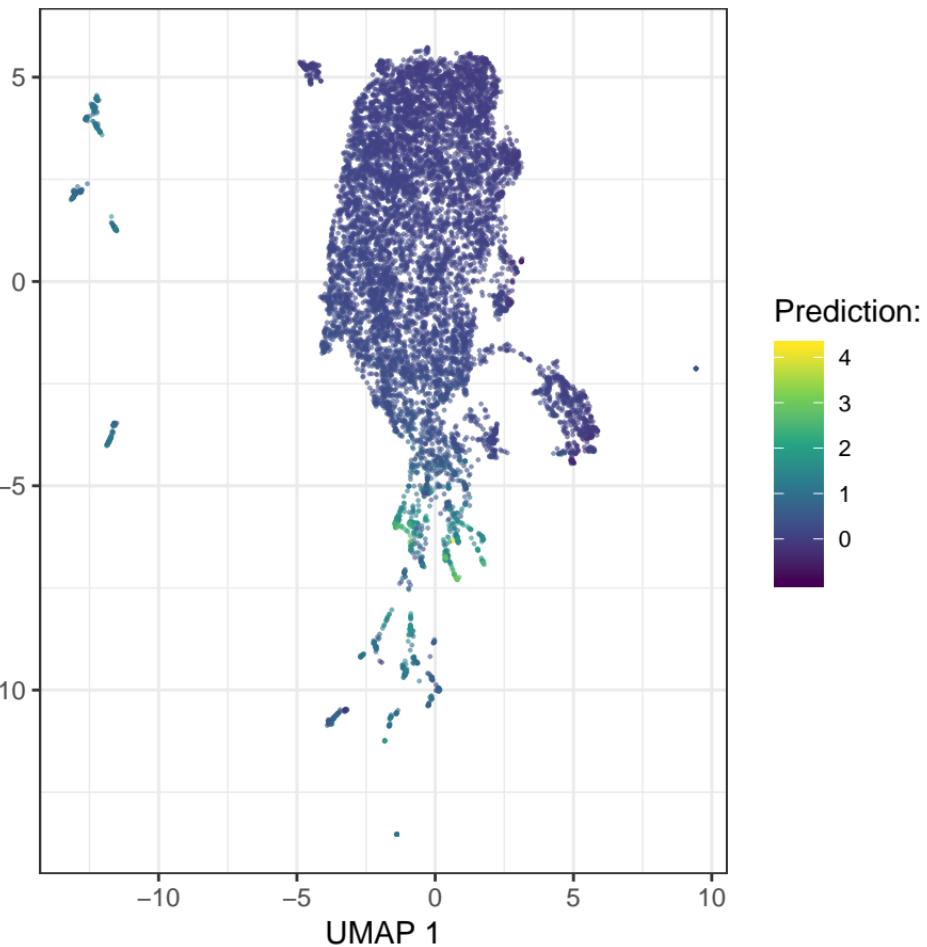
-5

5

10

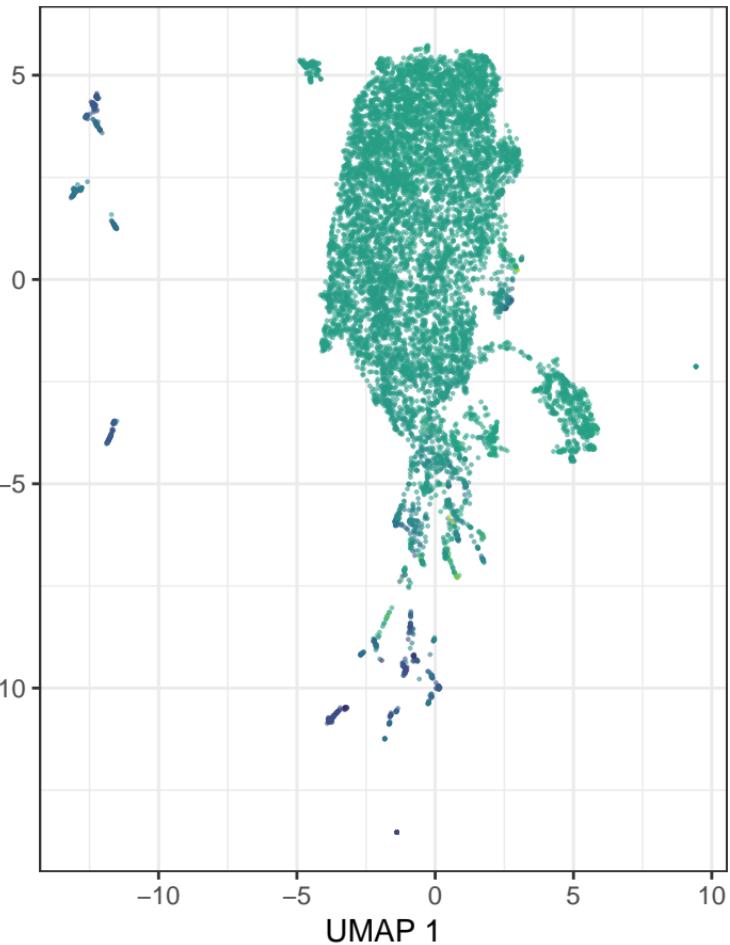
cell_health_modz_target_cc_g1_high_h2ax

UMAP 2

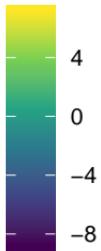


cell_health_modz_target_cc_cc_mitosis

UMAP 2



Prediction:

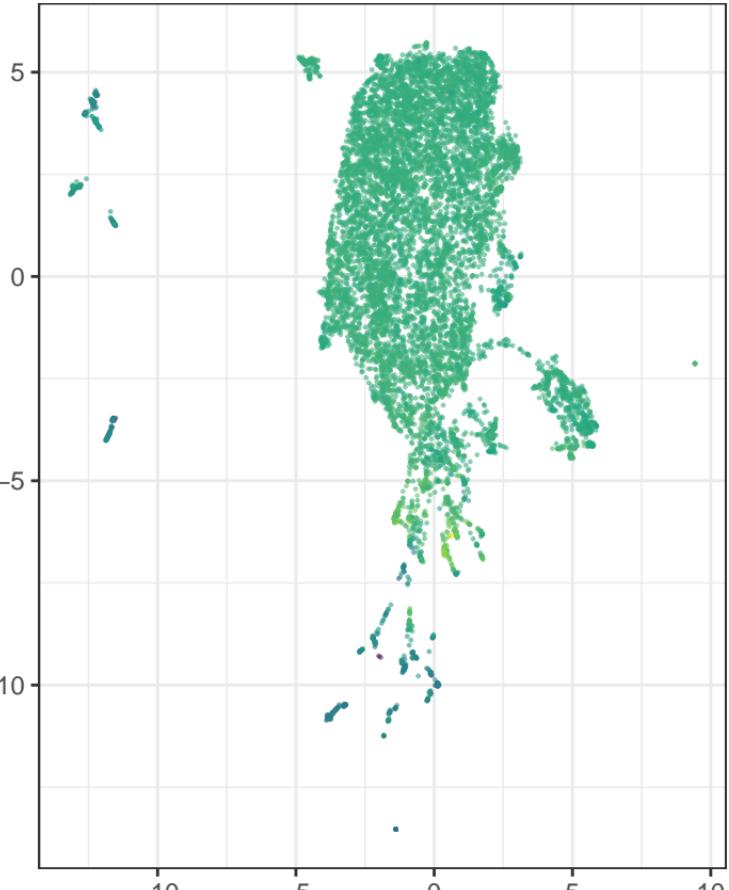
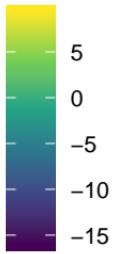


cell_health_modz_target_cc_all_n_spots_h2ax_mean

UMAP 2

UMAP 1

Prediction:

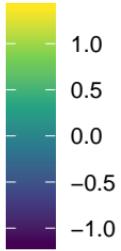


cell_health_modz_target_vb_infection_percentage

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

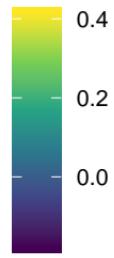
10

cell_health_modz_target_cc_mitosis_n_spots_h2ax_per

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

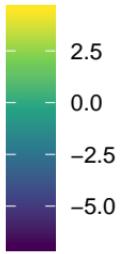
10

cell_health_modz_target_cc_s_n_spots_h2ax_mean

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

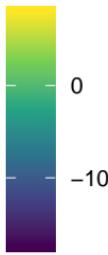
10

cell_health_modz_target_cc_all_n_spots_h2ax_per_nuc

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

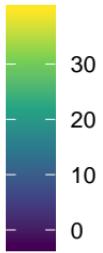
10

cell_health_modz_target_vb_percent_late_apoptosis

UMAP 2

UMAP 1

Prediction:



5

0

-5

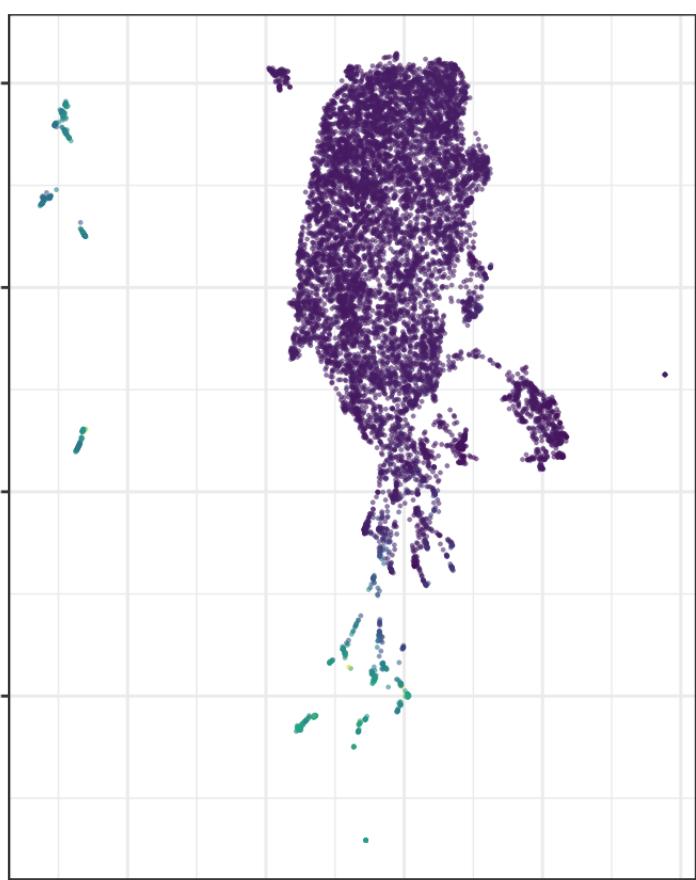
-10

10

5

-5

0

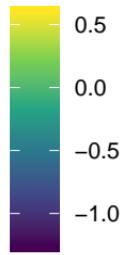


cell_health_modz_target_cc_mitosis_n_spots_h2ax_me

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

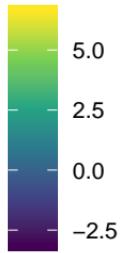
10

cell_health_modz_target_cc_g1_n_spots_h2ax_mean

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

10

cell_health_modz_target_cc_all_large_round_polyloid

UMAP 2

5

0

-5

-10

-10

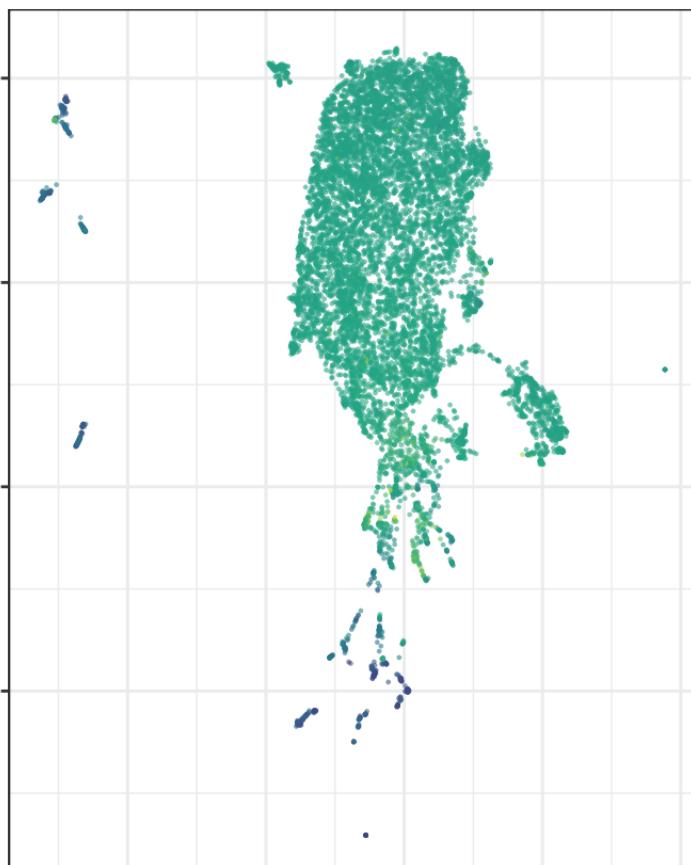
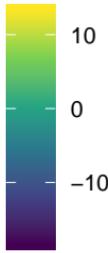
-5

5

10

UMAP 1

Prediction:

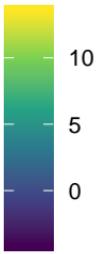


cell_health_modz_target_cc_cc_g1

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

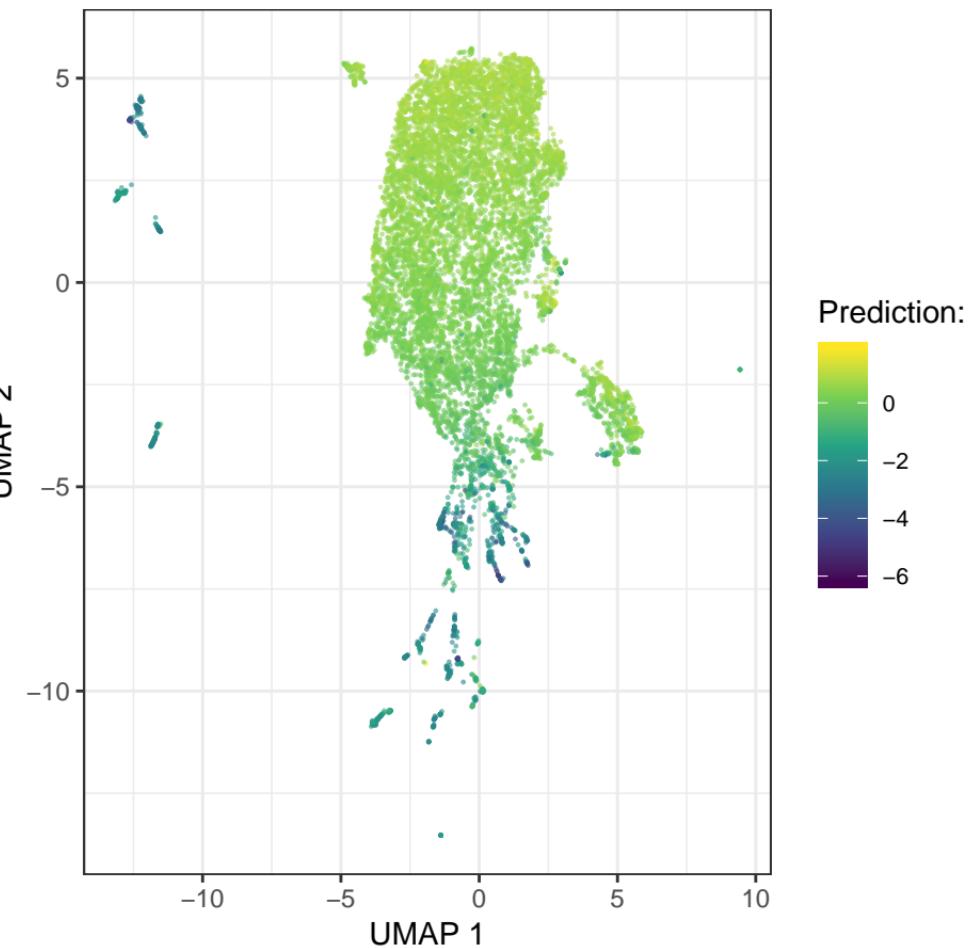
-5

5

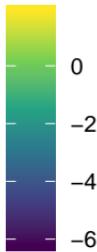
10

cell_health_modz_target_cc_s_intensity_nucleus_area

UMAP 2



Prediction:

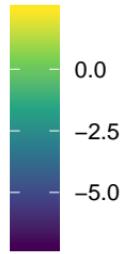


cell_health_modz_target_cc_polyplloid_n_spots_h2ax_p

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

10

cell_health_modz_target_cc_s_n_objects

UMAP 2

-10 -5 0 5 10

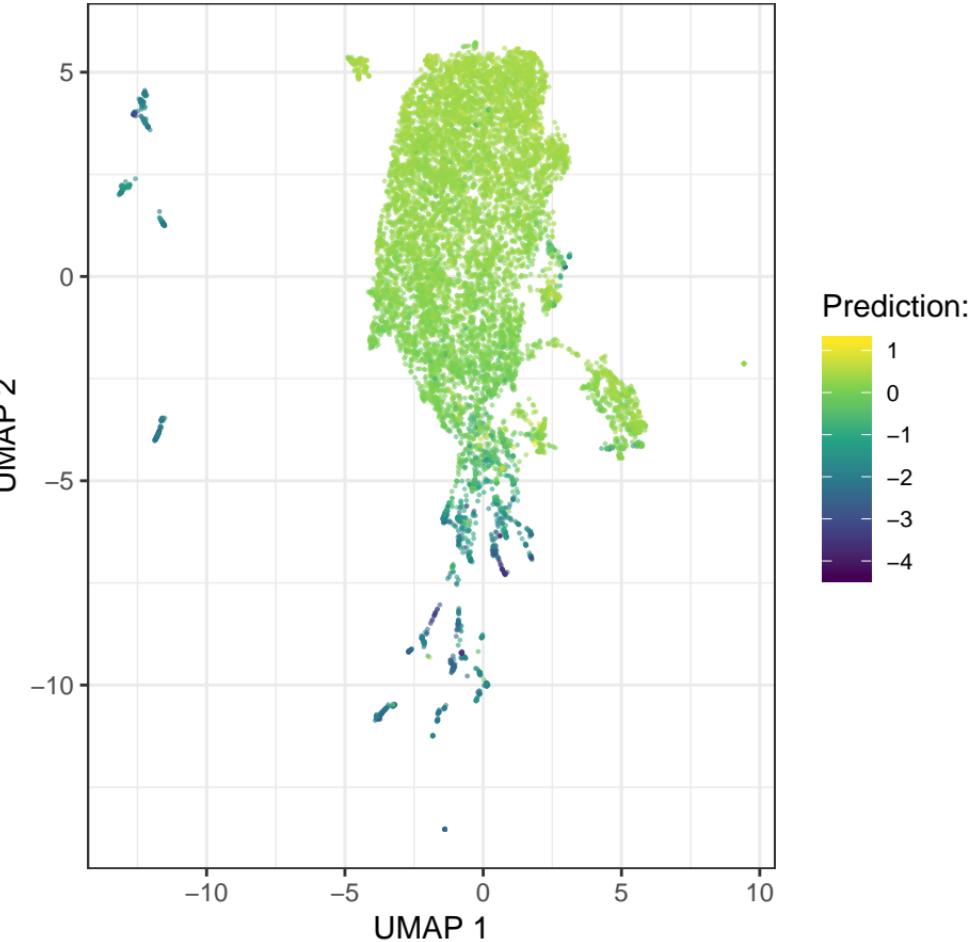
UMAP 1

Prediction:



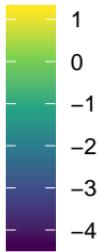
cell_health_modz_target_cc_g2_n_objects

UMAP 2



UMAP 1

Prediction:

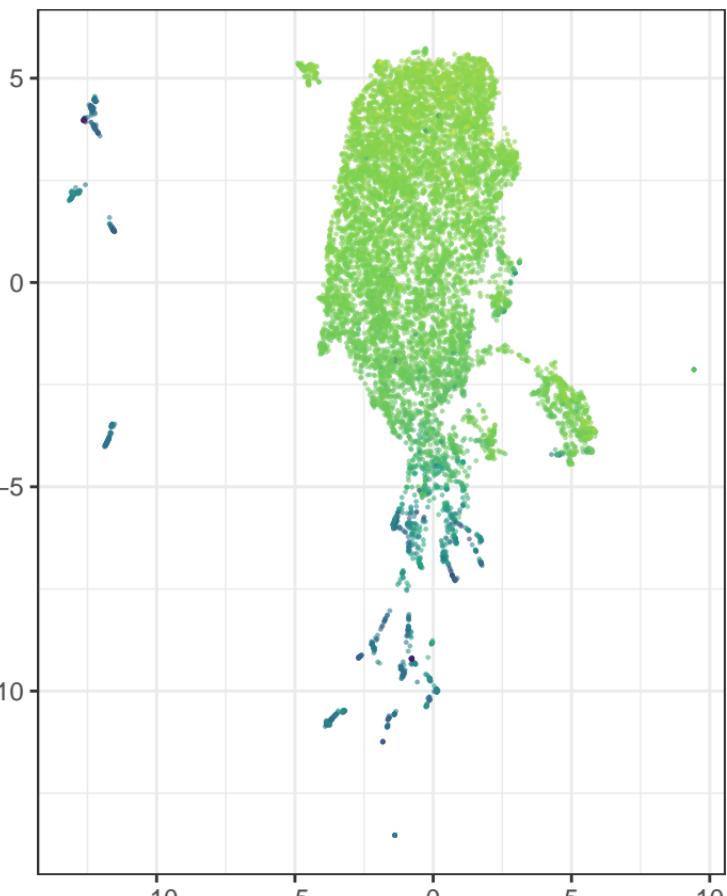
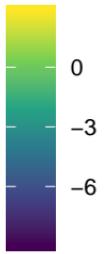


cell_health_modz_target_cc_cc_n_objects

UMAP 2

UMAP 1

Prediction:

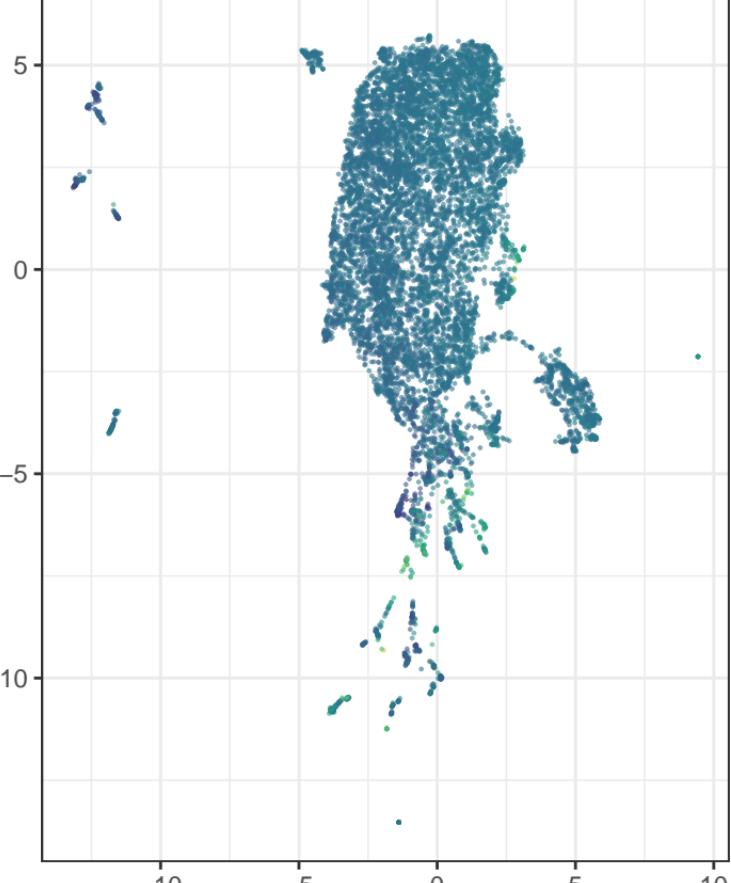
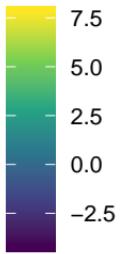


cell_health_modz_target_cc_late_mitosis_n_objects

UMAP 2

UMAP 1

Prediction:

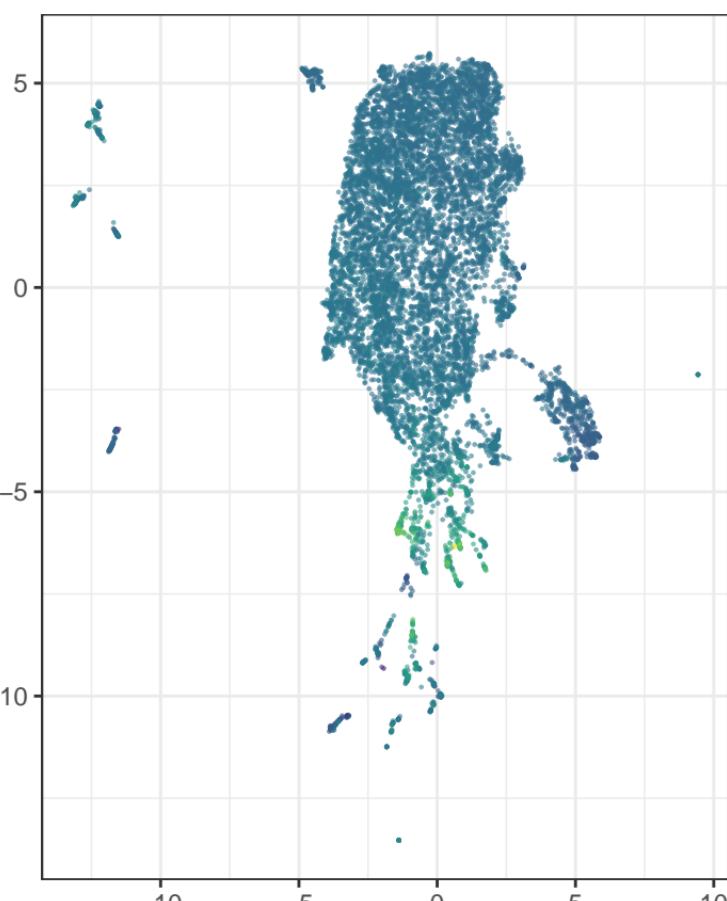
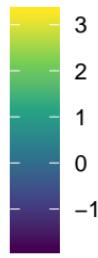


cell_health_modz_target_cc_early_mitosis_n_spots_h2

UMAP 2

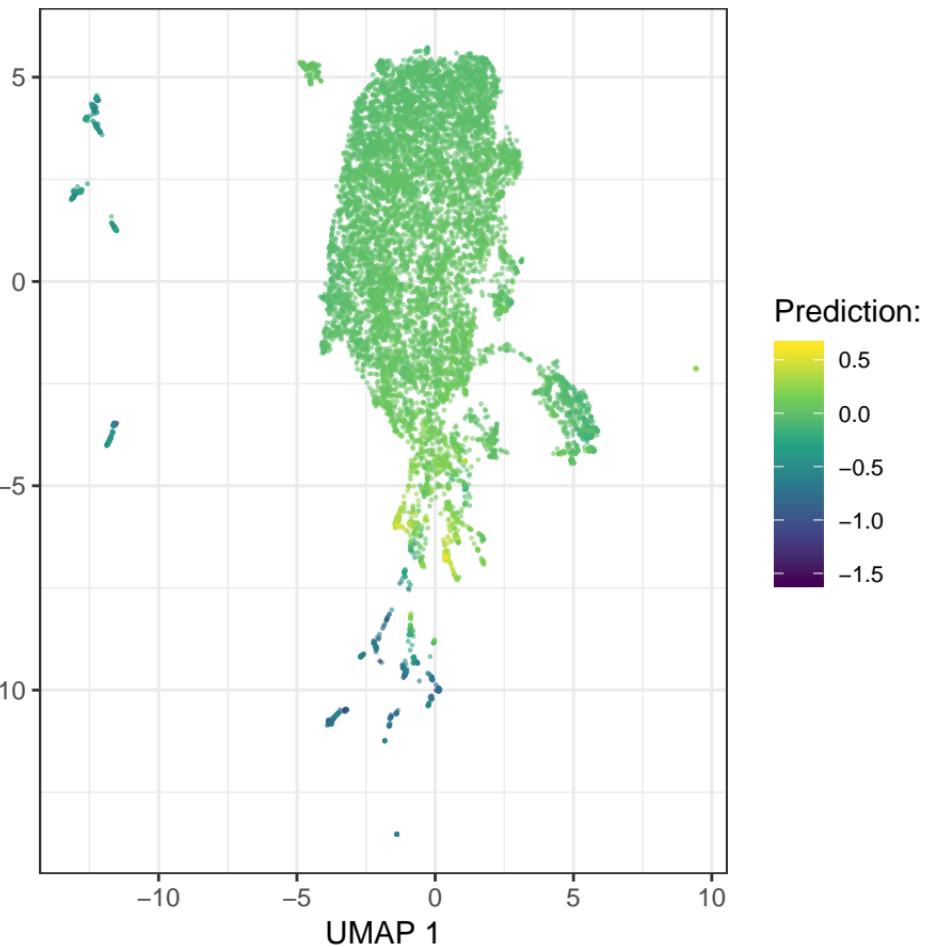
UMAP 1

Prediction:



cell_health_modz_target_cc_g2_g1_count

UMAP 2

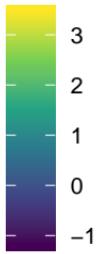


cell_health_modz_target_vb_percent_caspase_dead_on

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

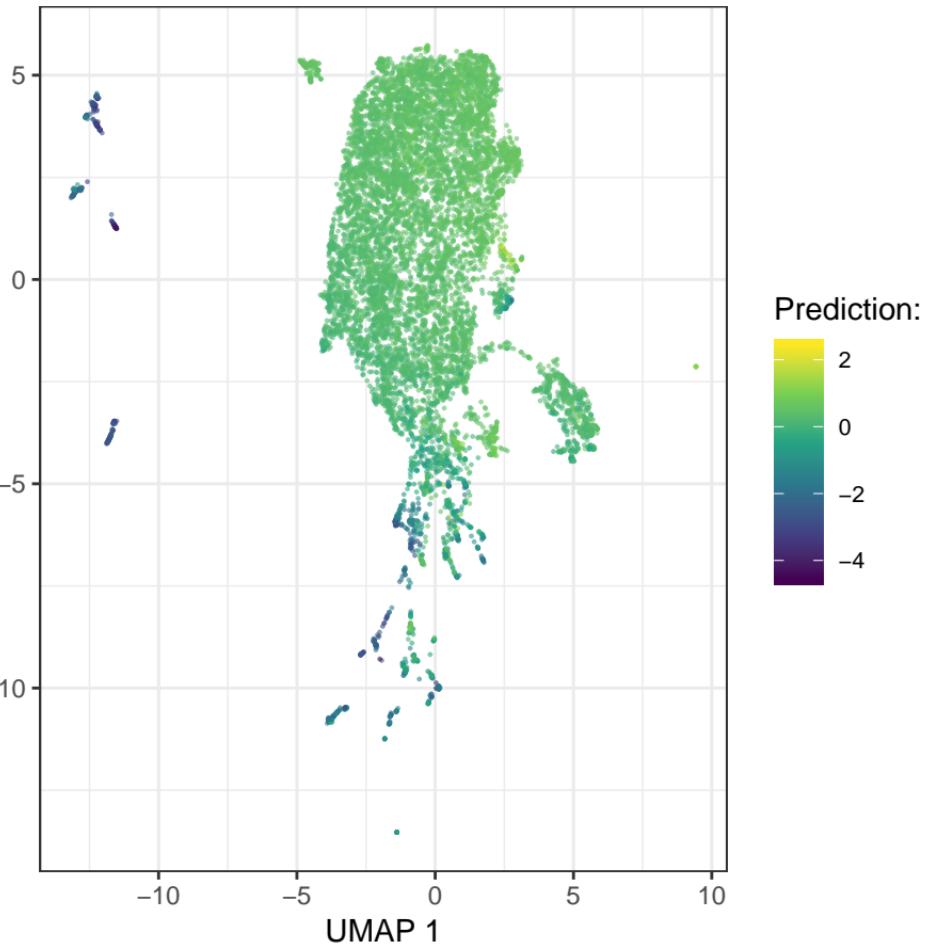
-5

5

10

cell_health_modz_target_cc_polynuclear_n_objects

UMAP 2

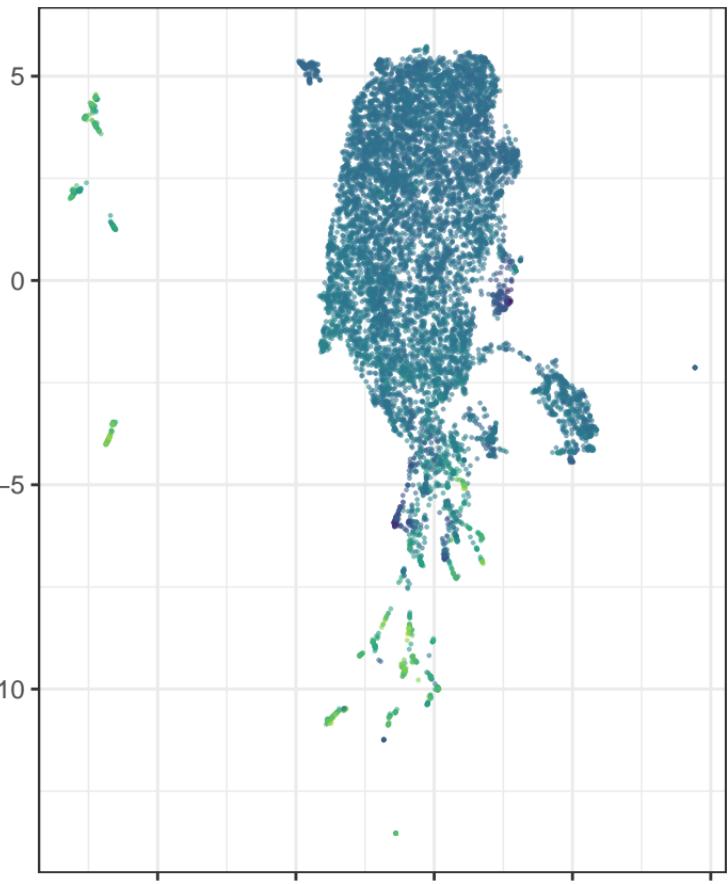
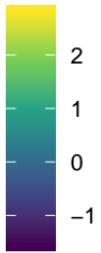


cell_health_modz_target_cc_cc_early_mitosis

UMAP 2

UMAP 1

Prediction:

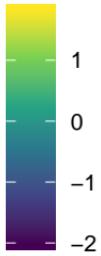


cell_health_modz_target_cc_polynuclear_high_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

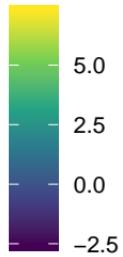
10

cell_health_modz_target_cc_g2_n_spots_h2ax_per_nuc

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

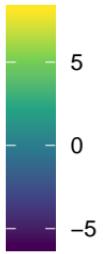
10

cell_health_modz_target_cc_cc_n_spots_h2ax_mean

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

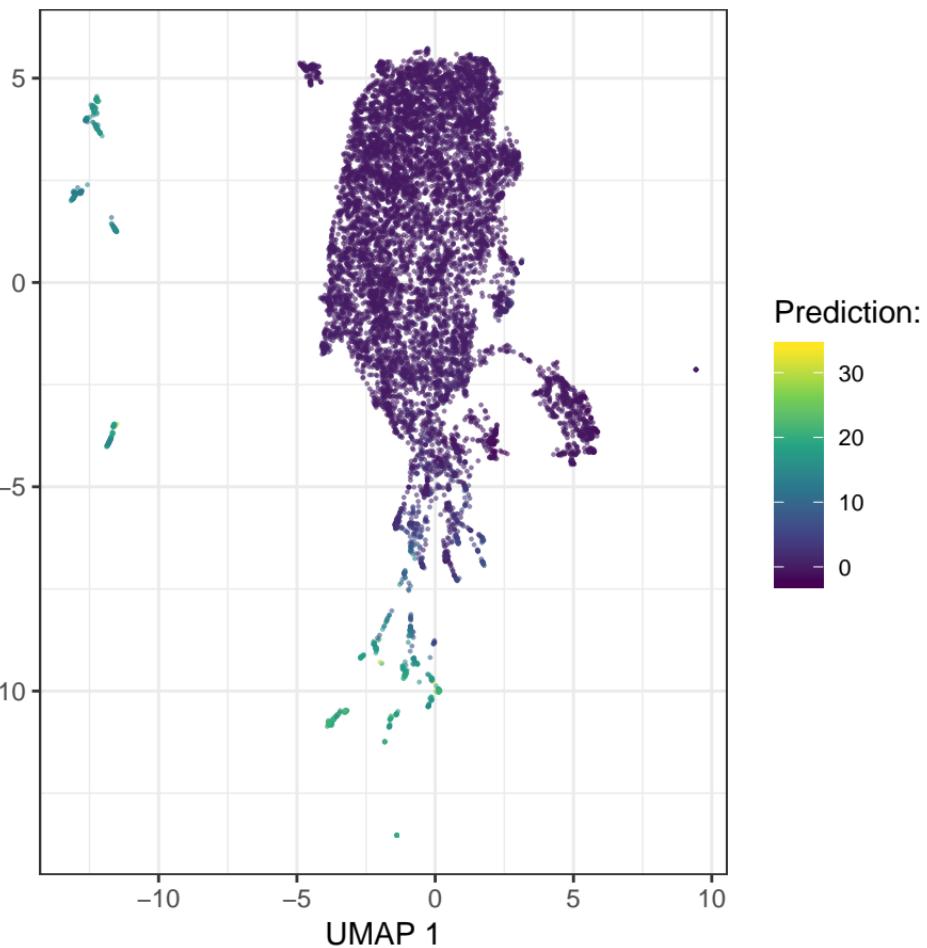
-5

5

10

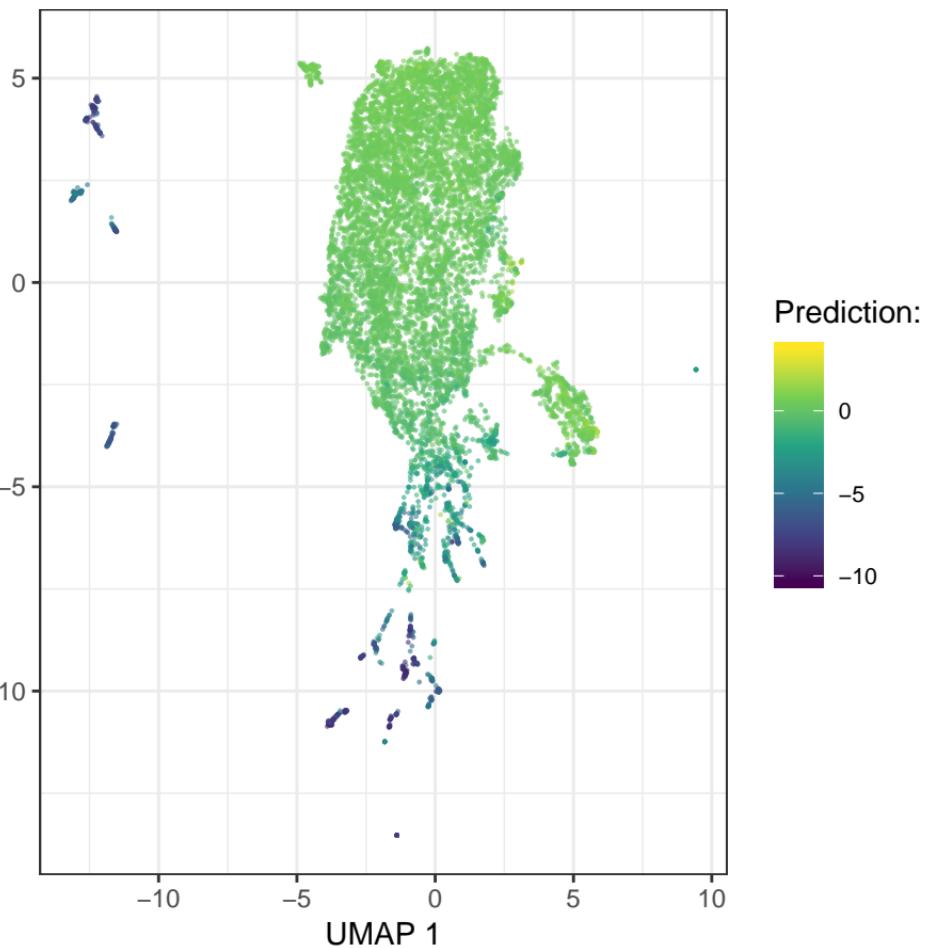
cell_health_modz_target_vb_percent_dead

UMAP 2



cell_health_modz_target_cc_cc_s

UMAP 2



cell_health_modz_target_cc_late_mitosis_high_h2ax

UMAP 2

5

0

-5

-10

-10

-5

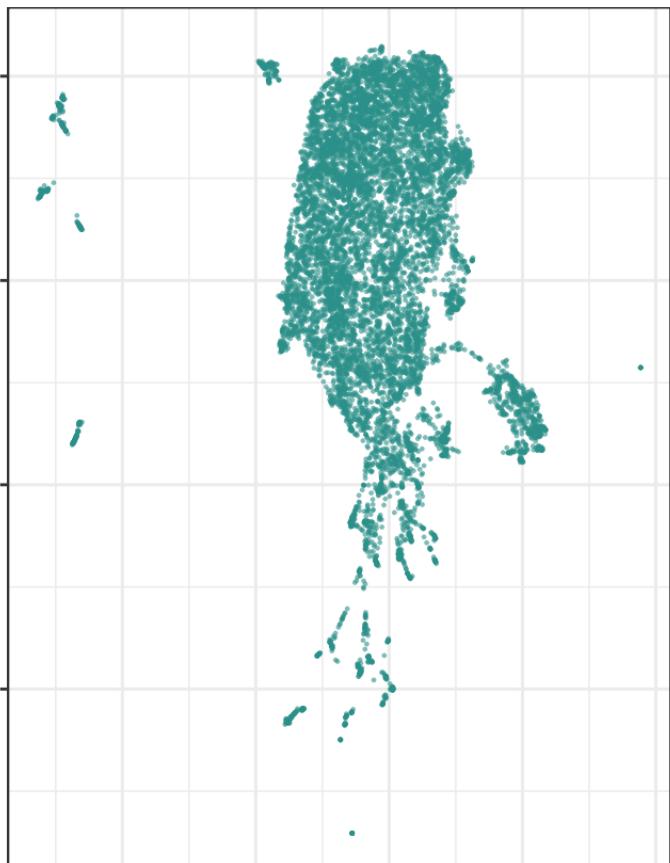
5

10

UMAP 1

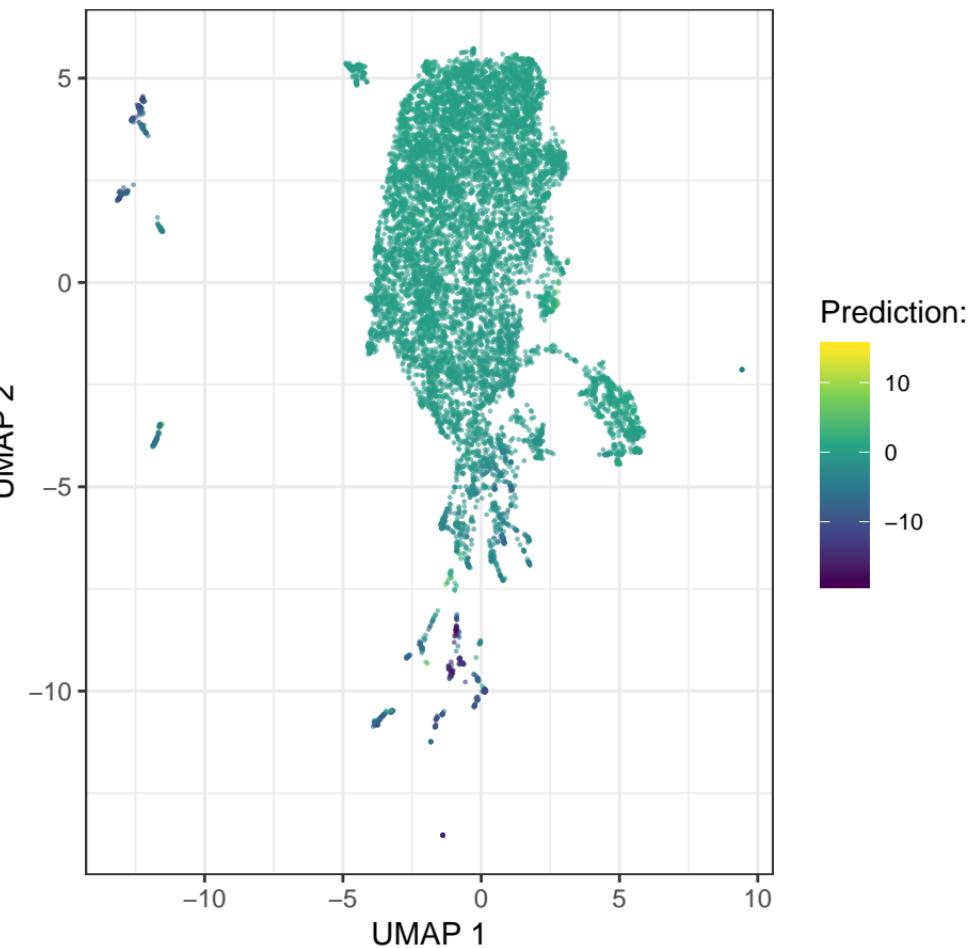
Prediction:

0.2245825



cell_health_modz_target_cc_s_intensity_nucleus_area

UMAP 2

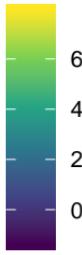


cell_health_modz_target_cc_g2_high_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

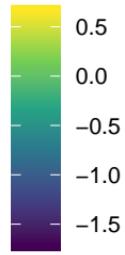
10

cell_health_modz_target_cc_late_mitosis_n_spots_h2ax

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

-5

5

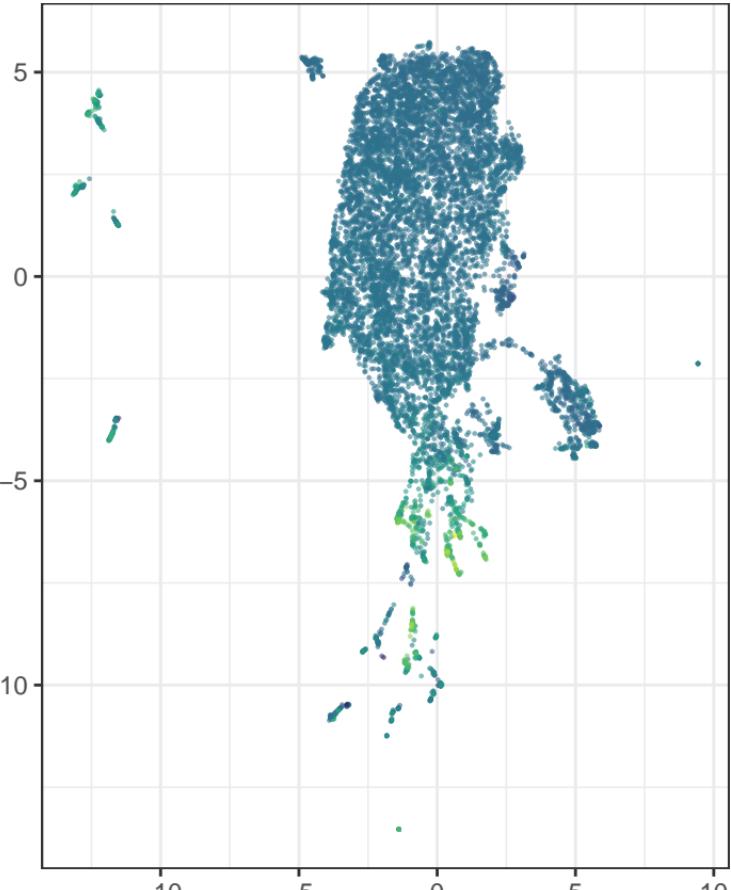
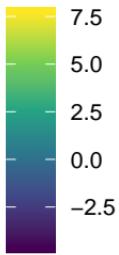
10

cell_health_modz_target_cc_cc_high_h2ax

UMAP 2

UMAP 1

Prediction:

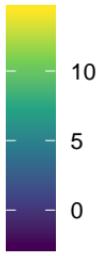


cell_health_modz_target_vb_percent_dead_only

UMAP 2

UMAP 1

Prediction:



5

0

-5

-10

-10

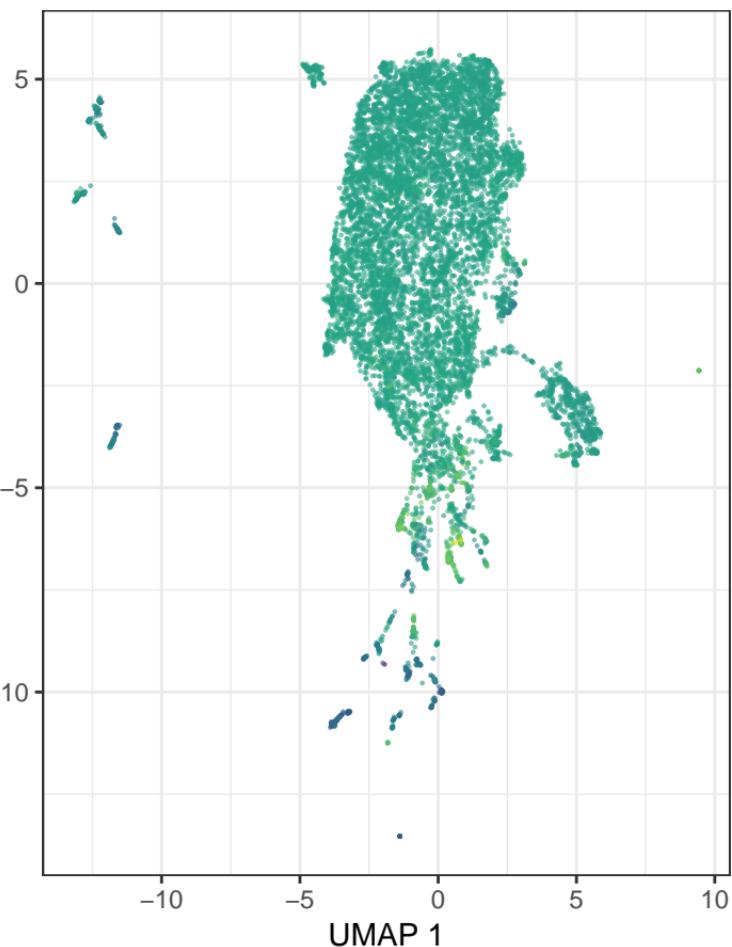
-5

5

10

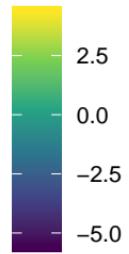
cell_health_modz_target_cc_s_high_h2ax

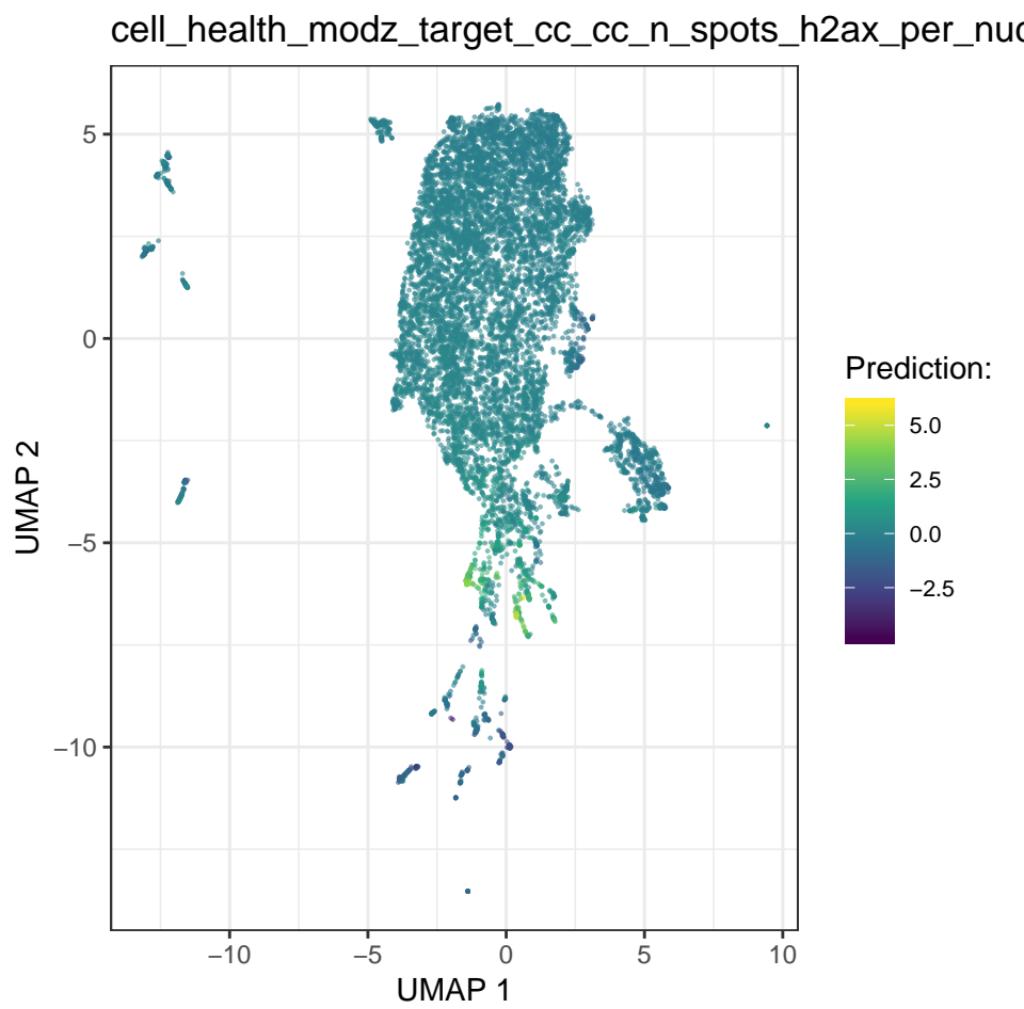
UMAP 2



UMAP 1

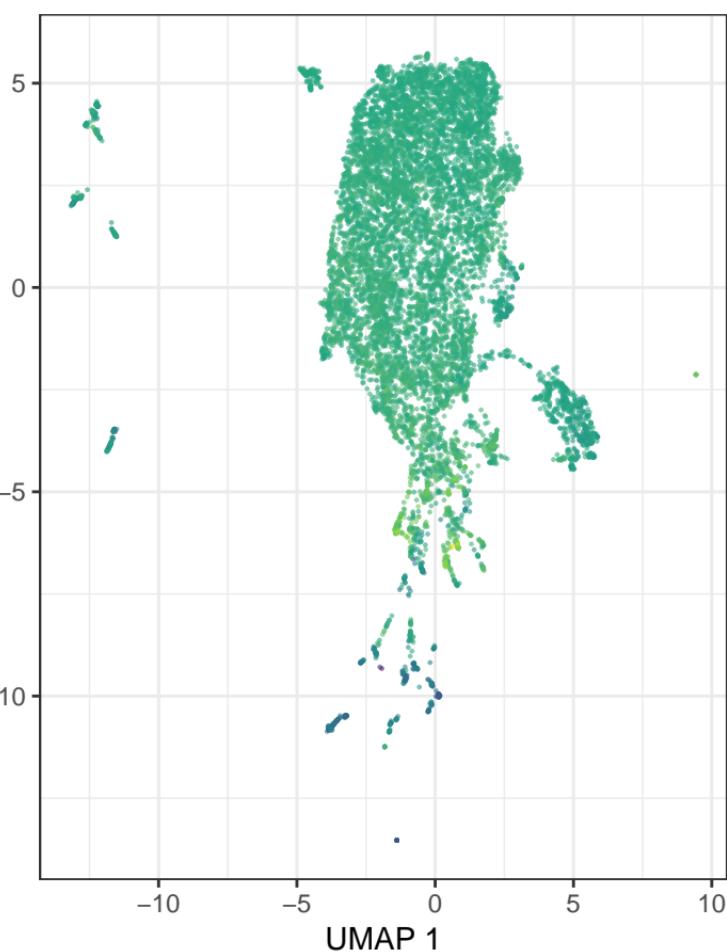
Prediction:



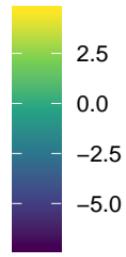


cell_health_modz_target_cc_s_n_spots_h2ax_per_nucleus

UMAP 2



Prediction:



cell_health_modz_target_cc_g1_plus_g2_count

UMAP 2

5

0

-5

-10

-10

-5

5

10

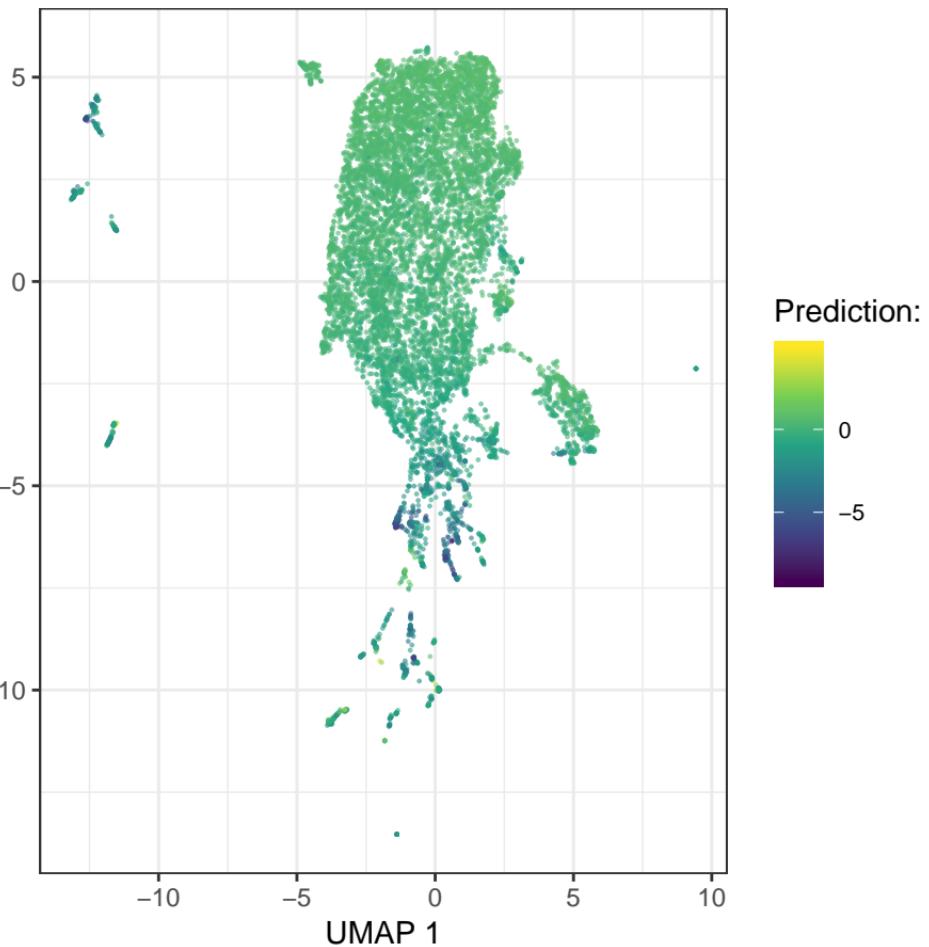
UMAP 1

Prediction:



cell_health_modz_target_vb_live_cell_width_length

UMAP 2



UMAP 1

Prediction:

0

-5