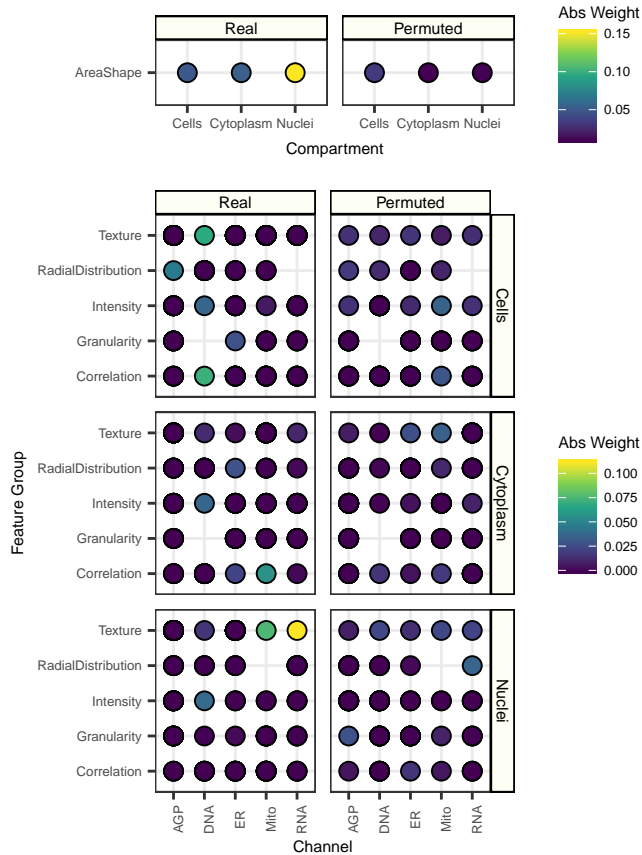
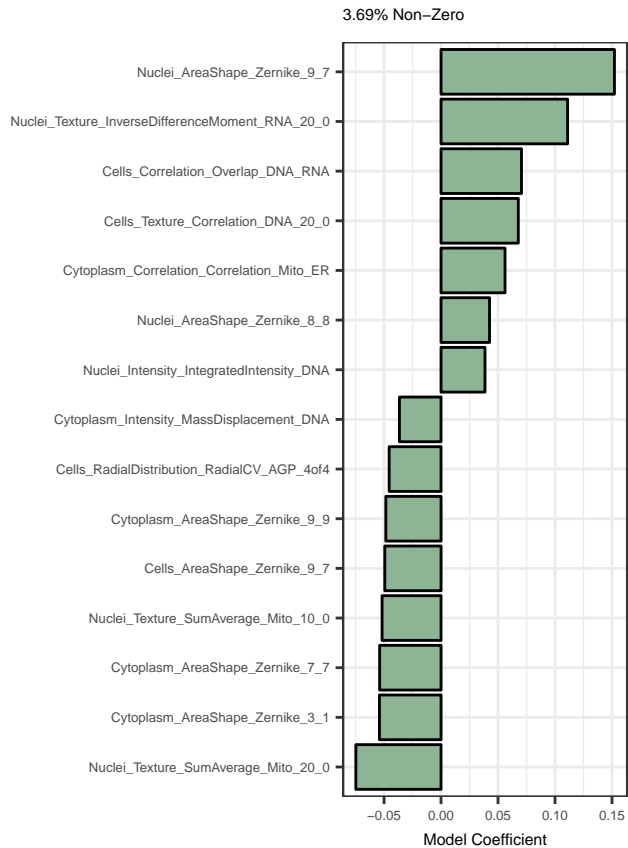
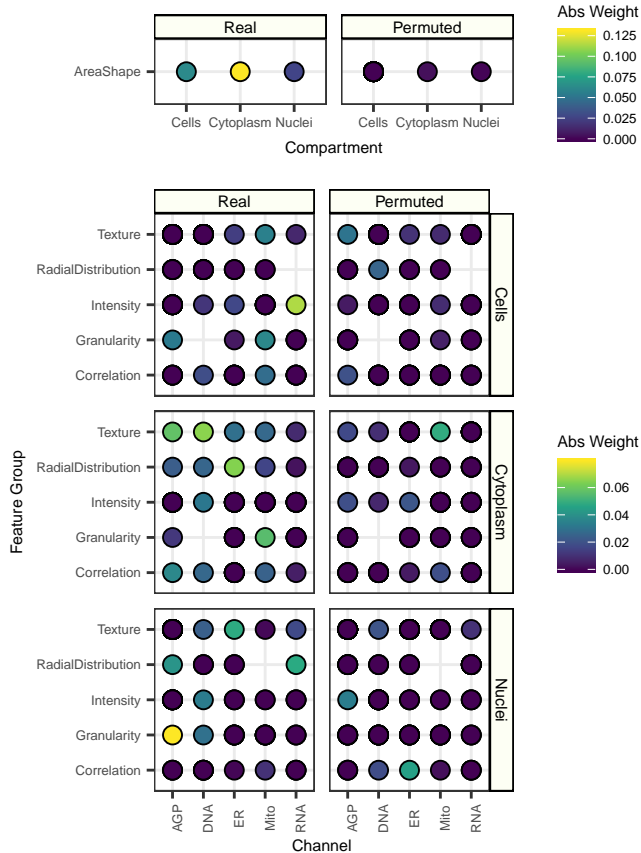
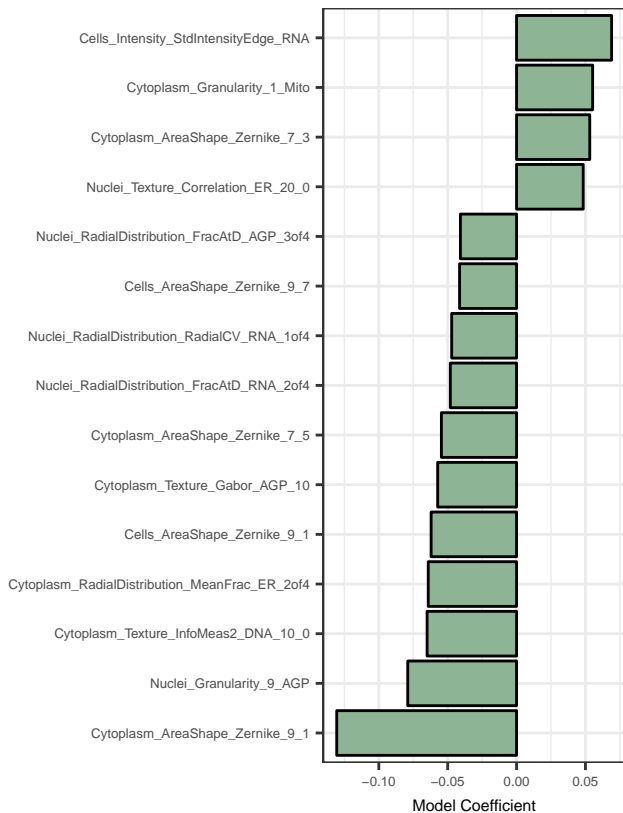


cc_all_high_h2ax



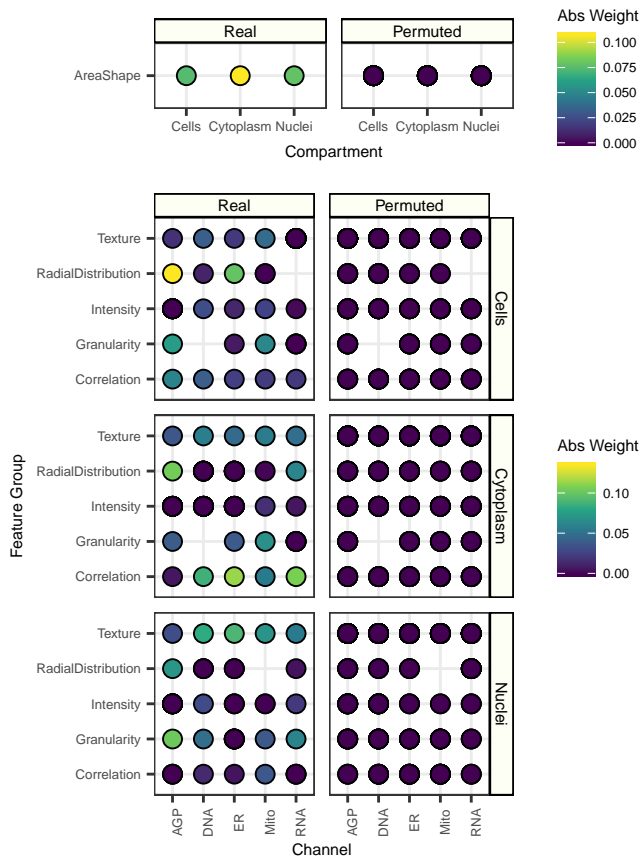
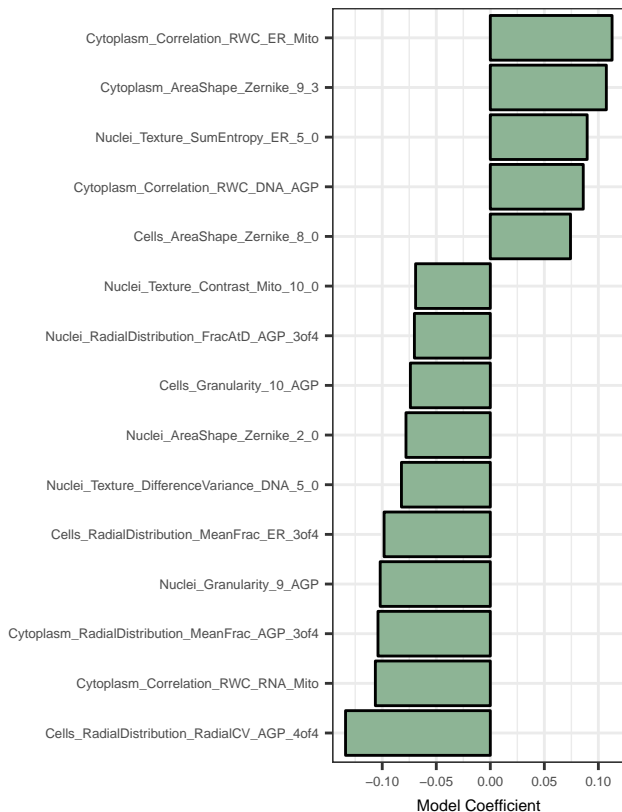
cc_all_large_notround_polynuclear_mean

14.54% Non-Zero

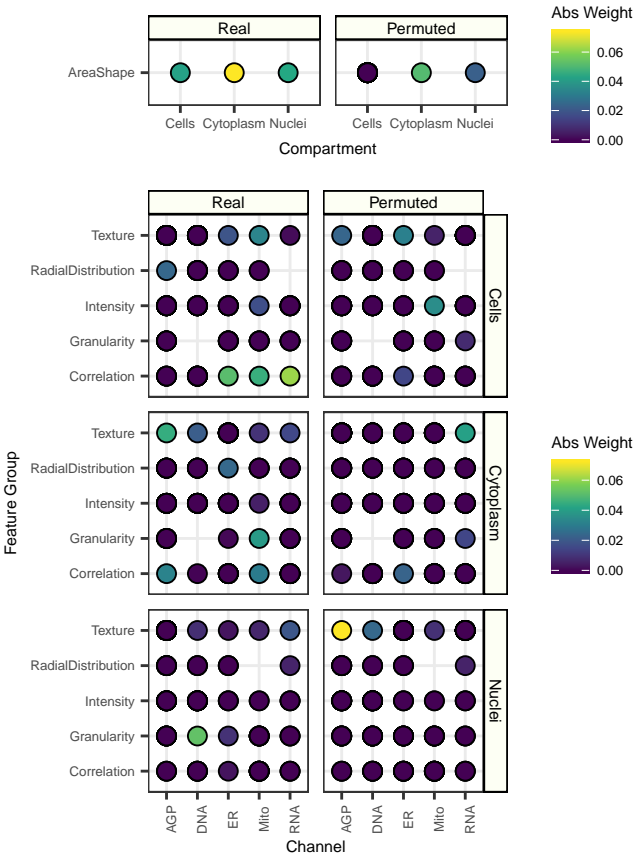
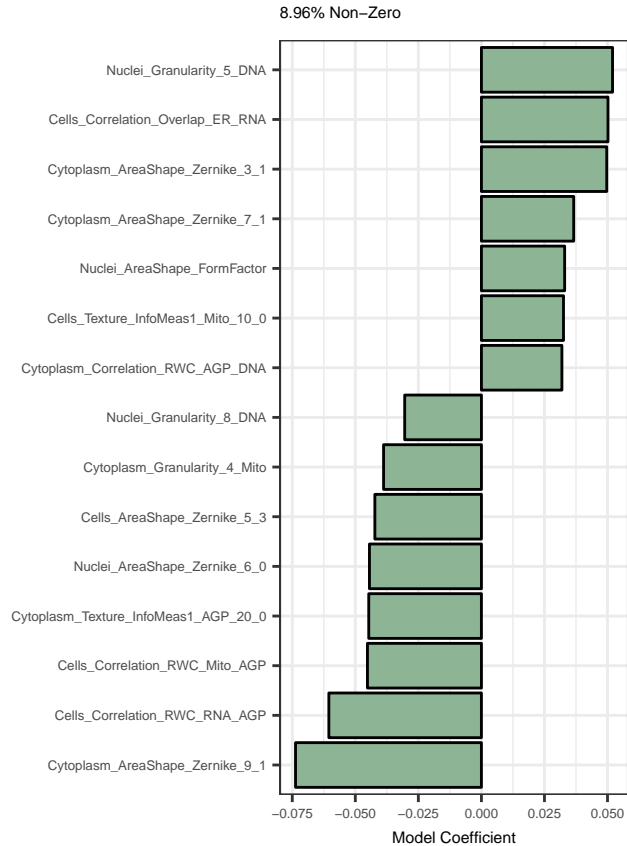


cc_all_large_round_polyploid_mean

24.24% Non-Zero

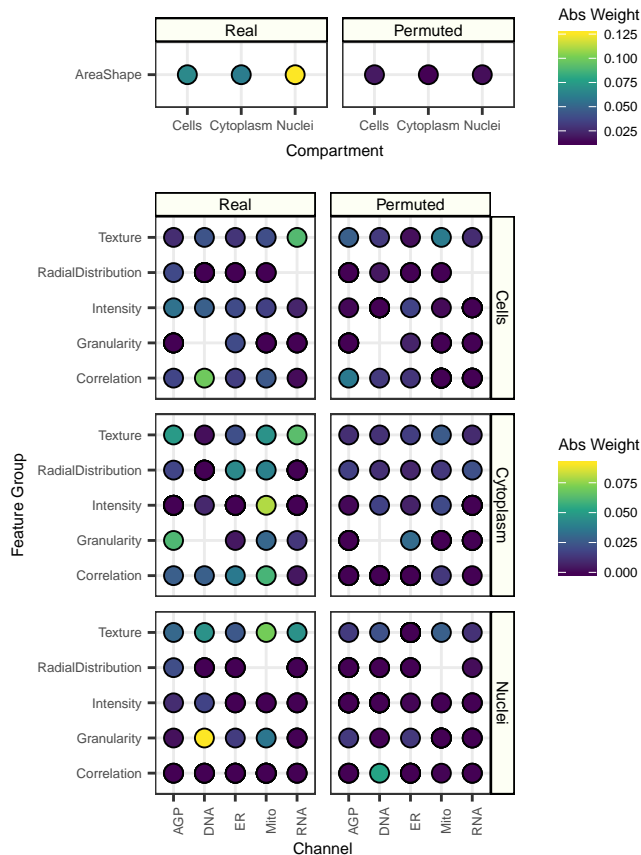
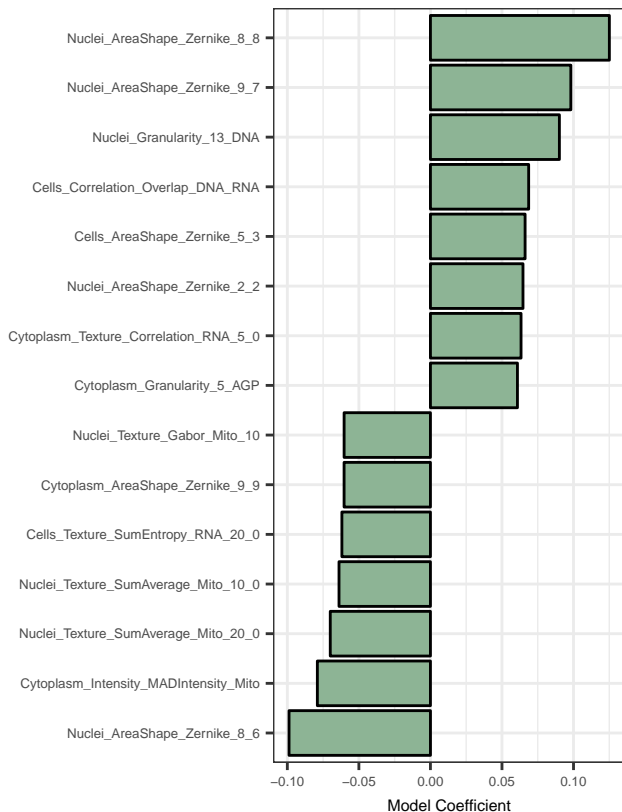


cc_all_n_objects



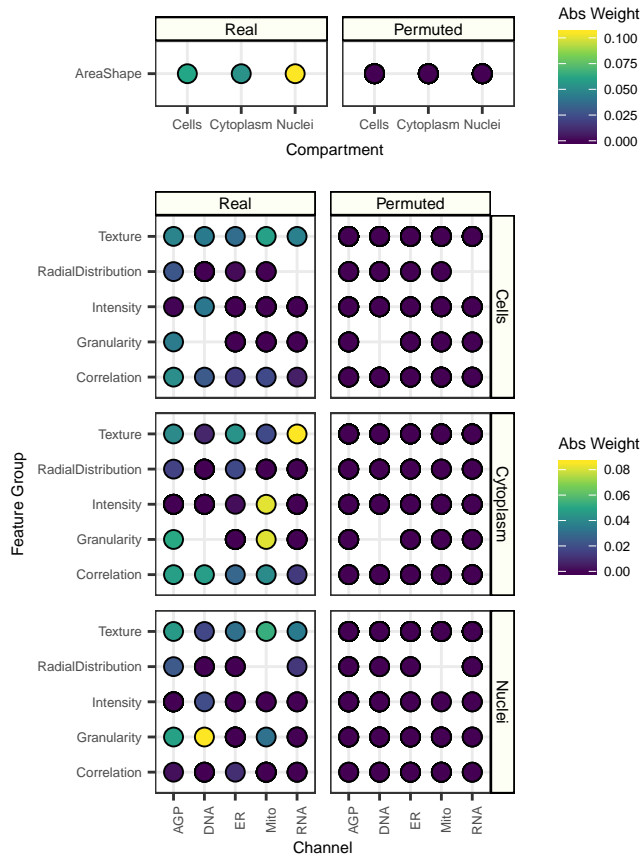
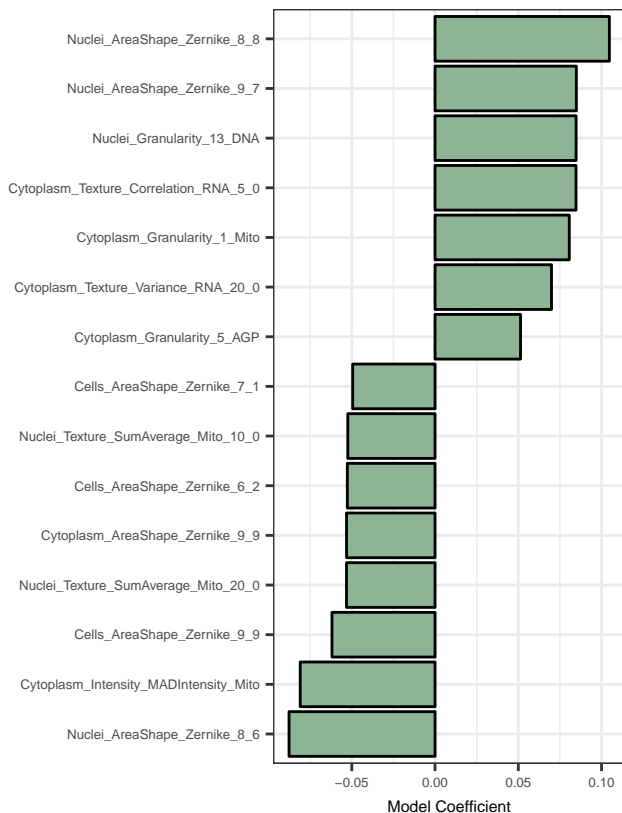
cc_all_n_spots_h2ax_mean

17.18% Non-Zero

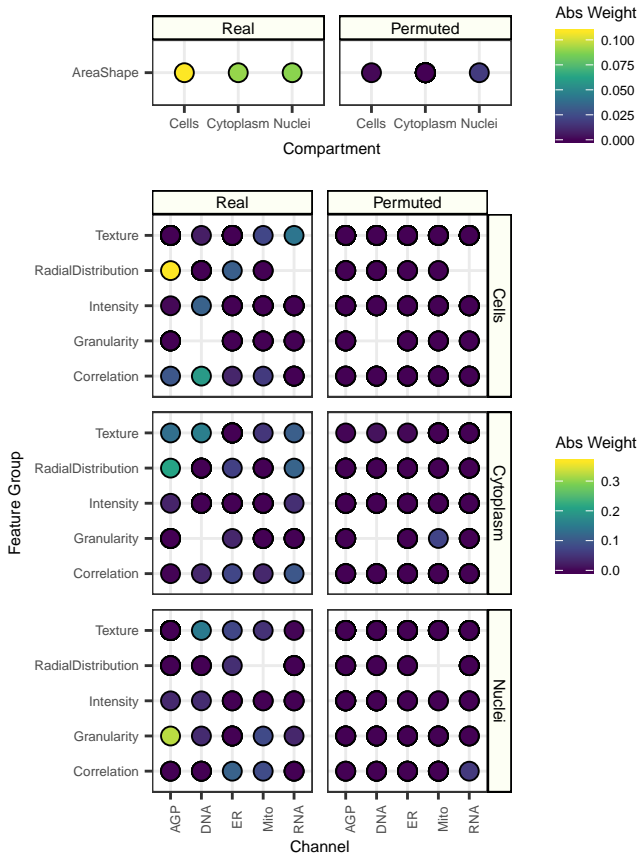
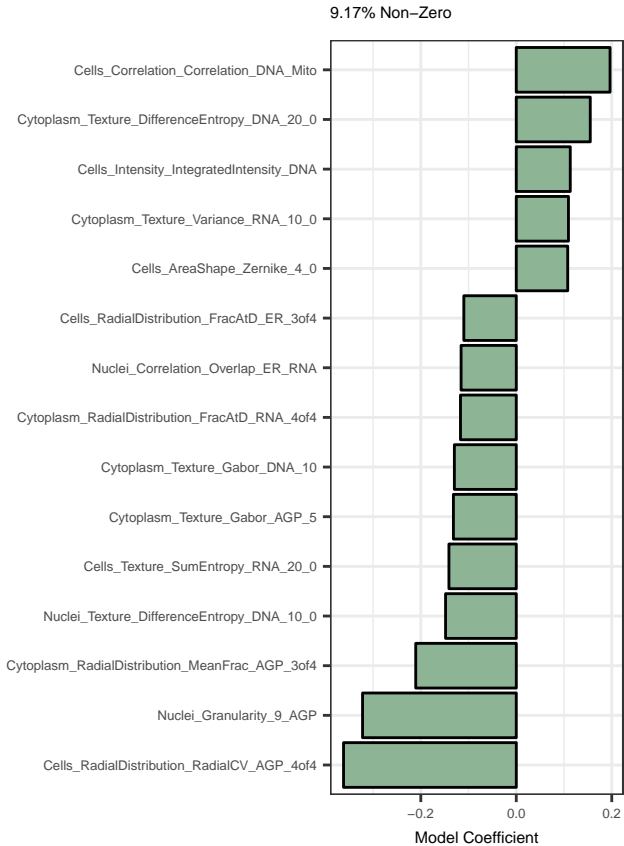


cc_all_n_spots_h2ax_per_nucleus_area_mean

17.39% Non-Zero

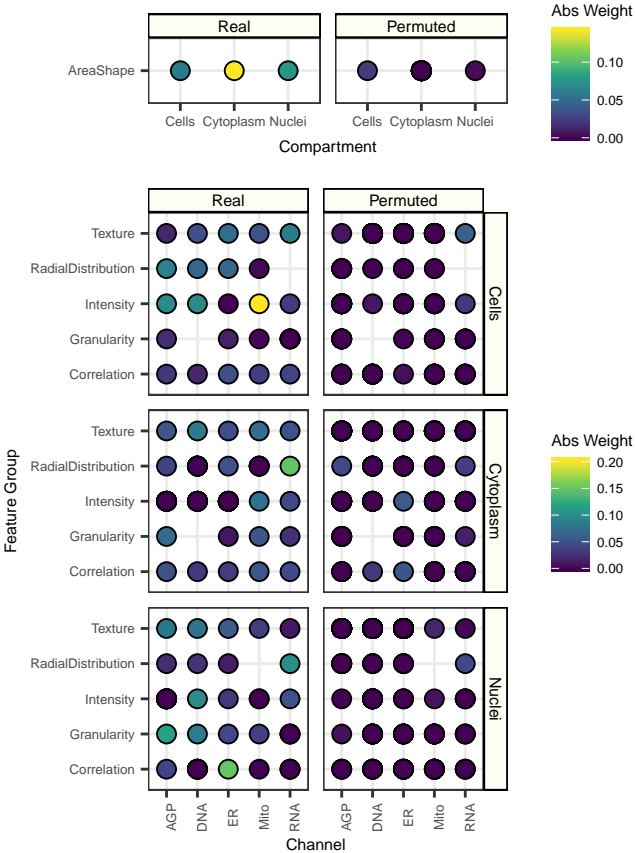
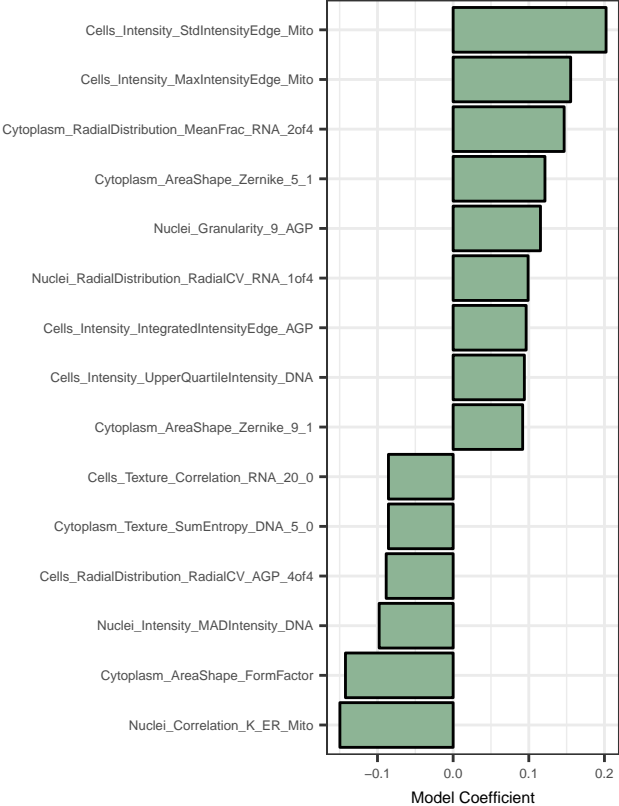


cc_all_nucleus_area_mean



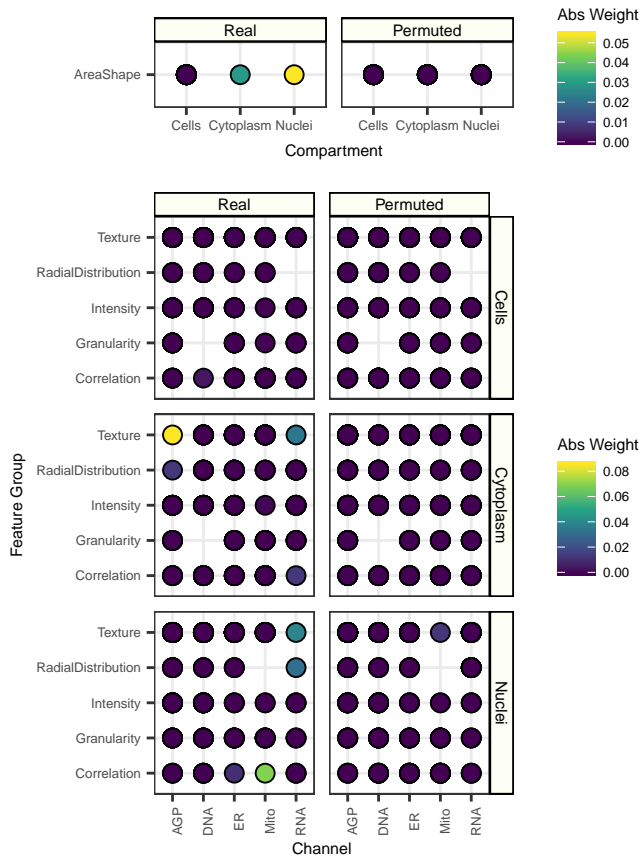
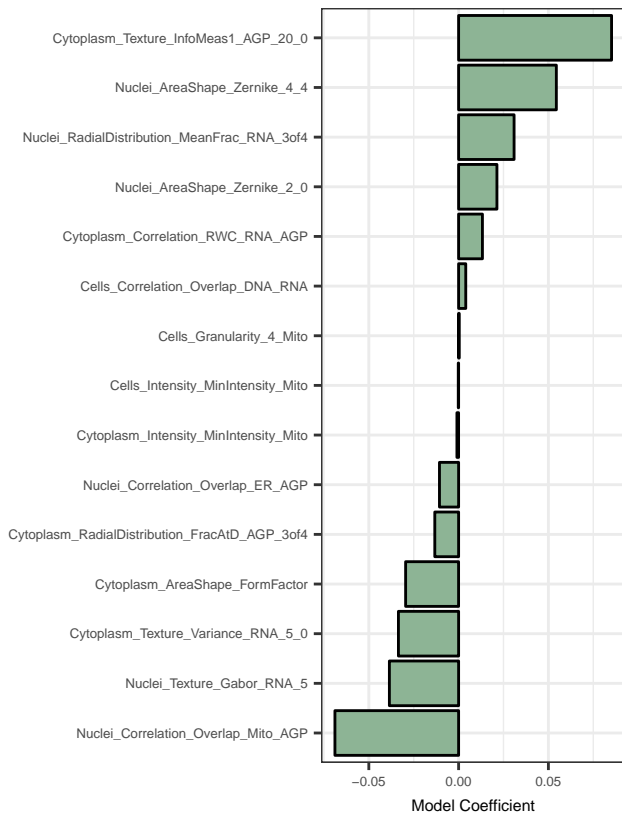
cc_all_nucleus_roundness_mean

29.4% Non-Zero

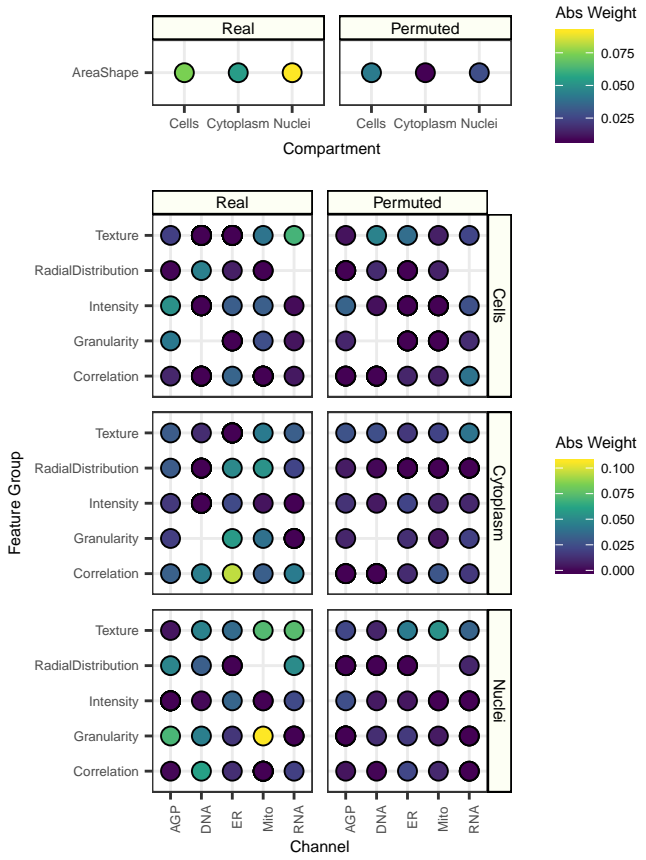
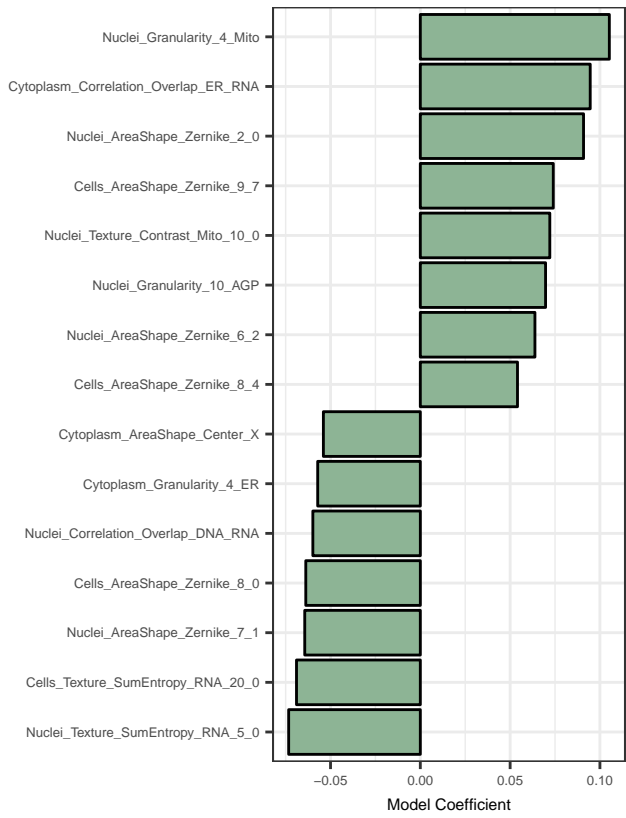


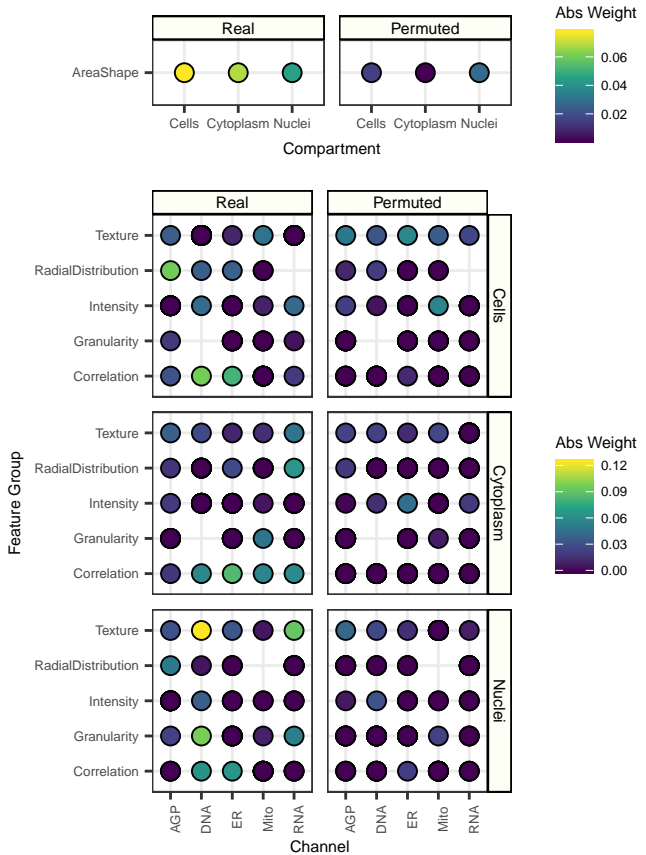
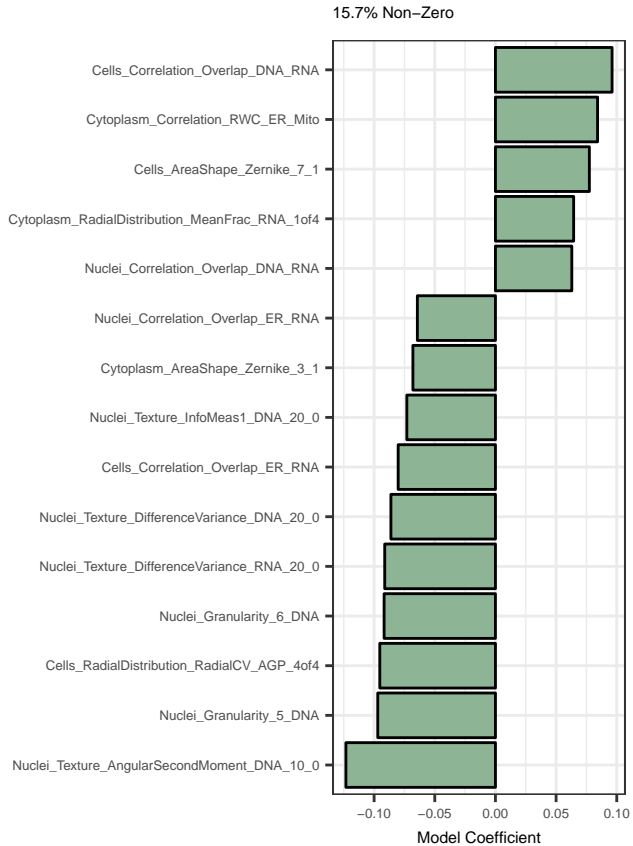
cc_cc_early_mitosis

1.58% Non-Zero

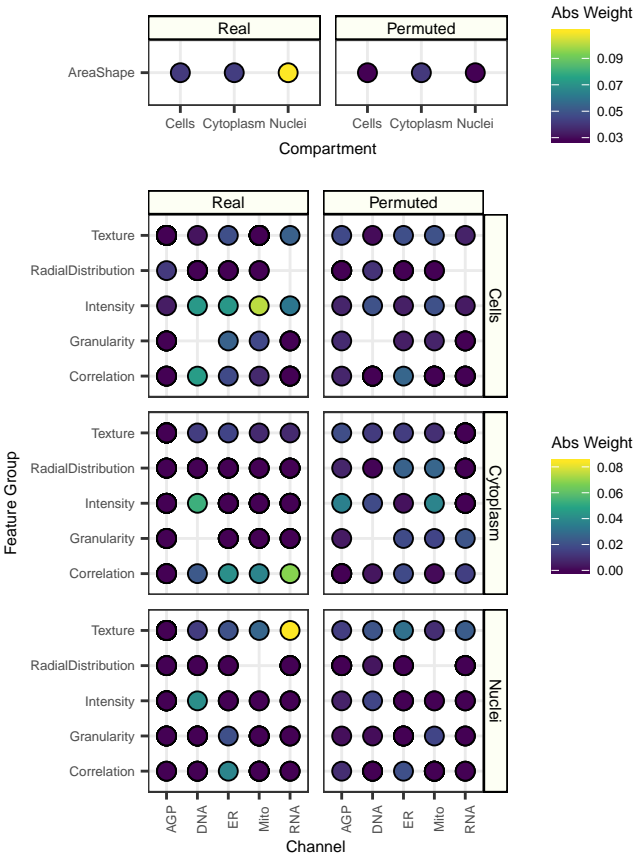
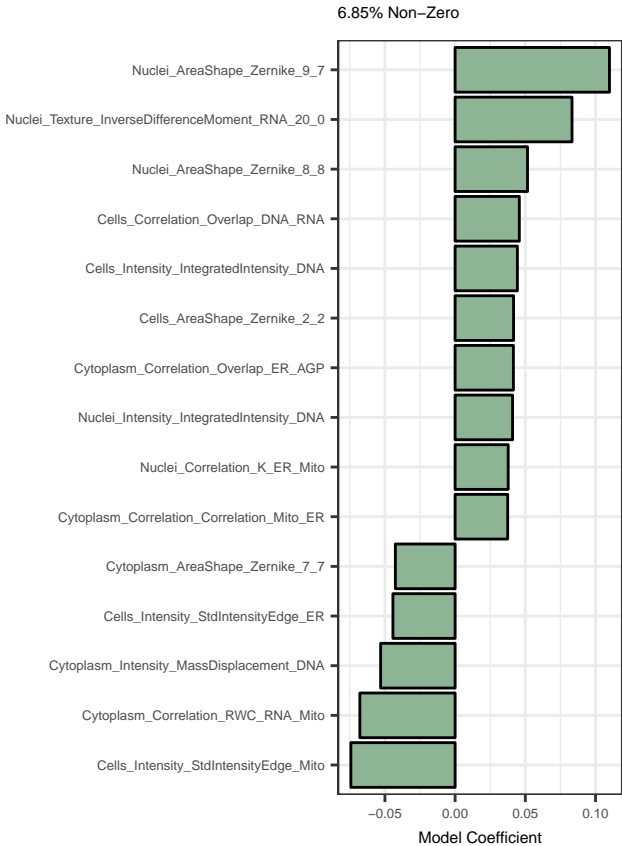


20.44% Non-Zero

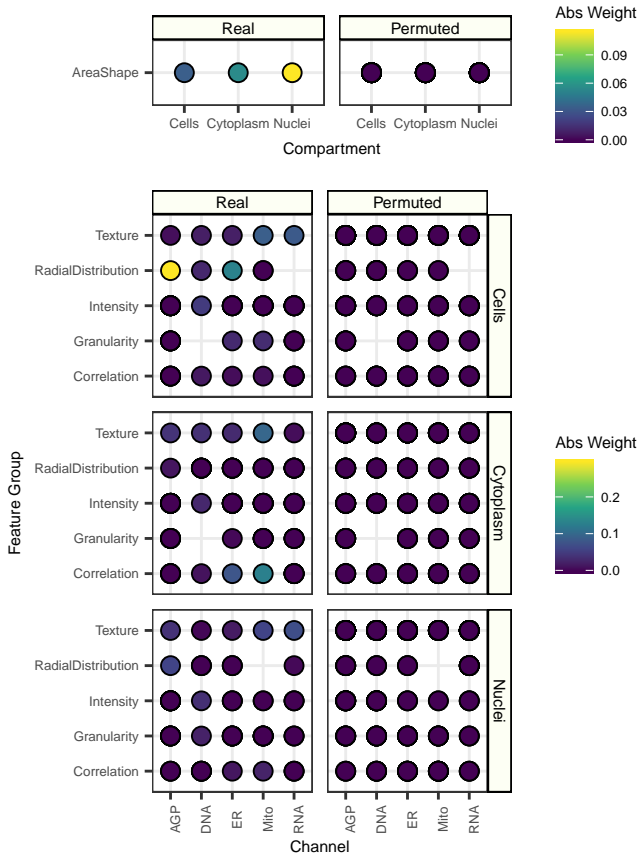
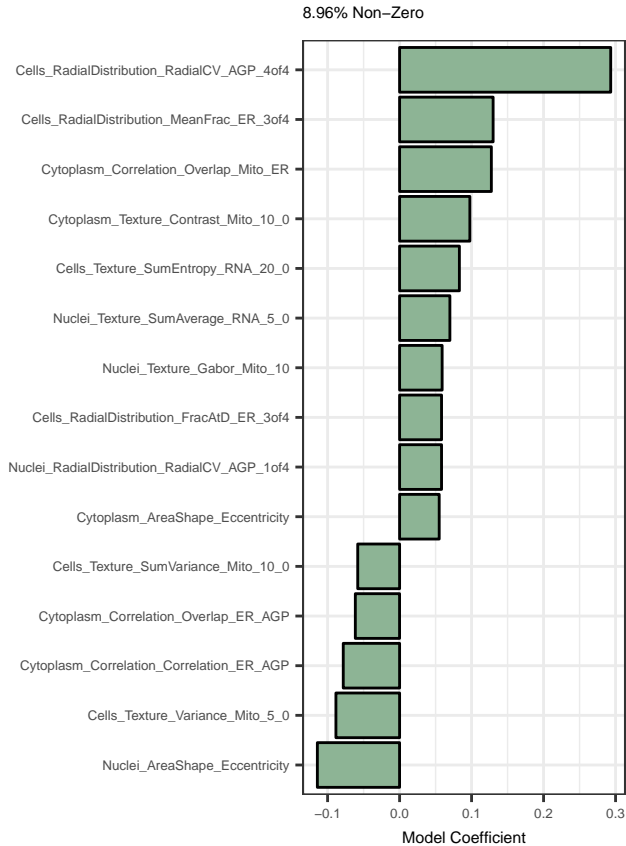




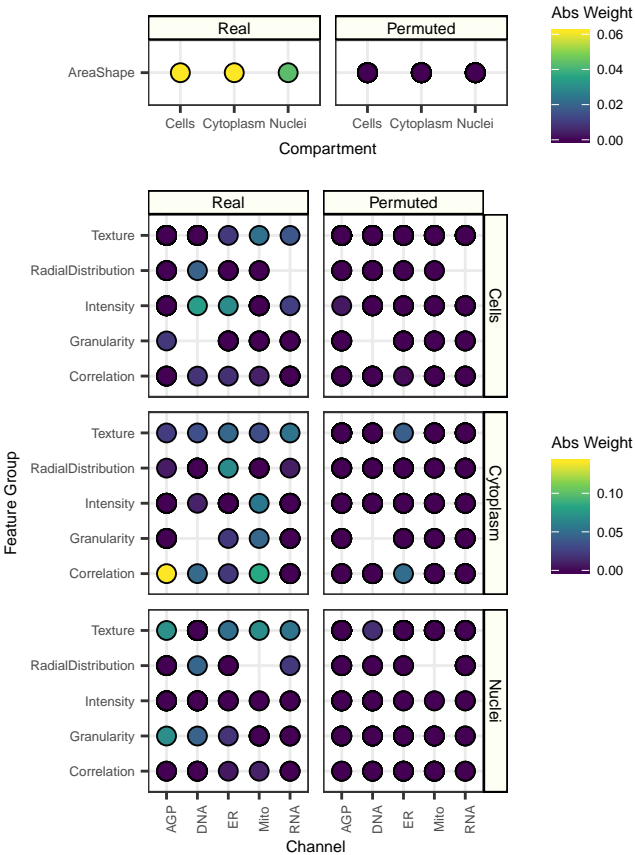
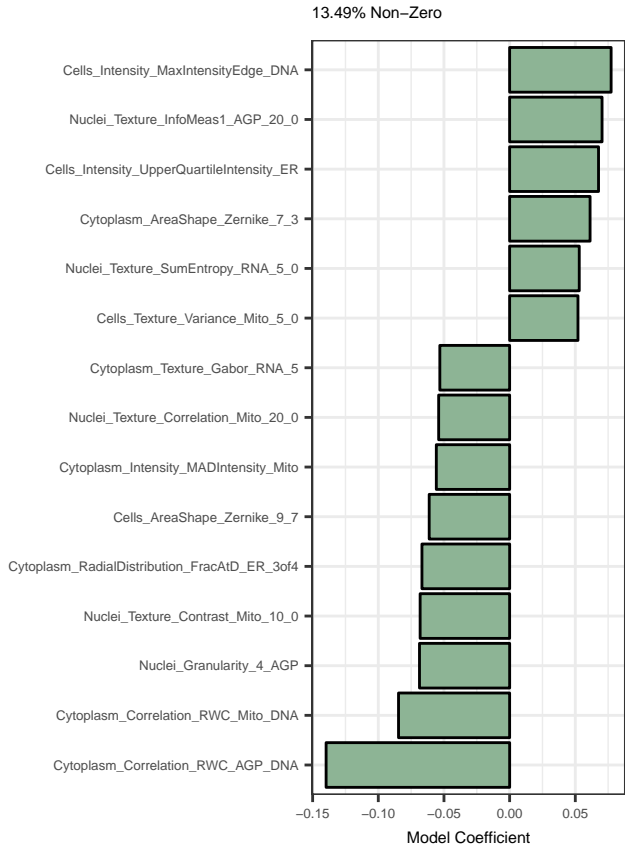
cc_cc_high_h2ax



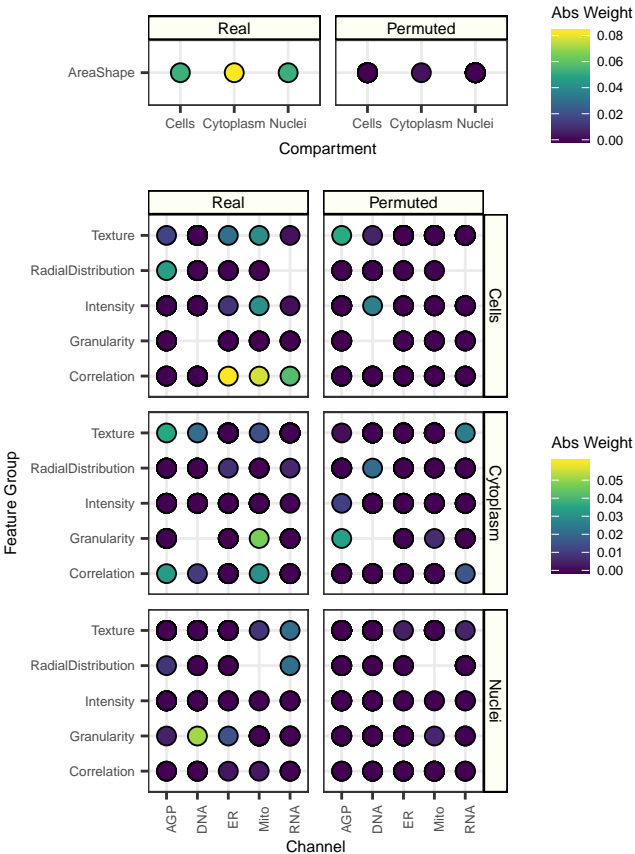
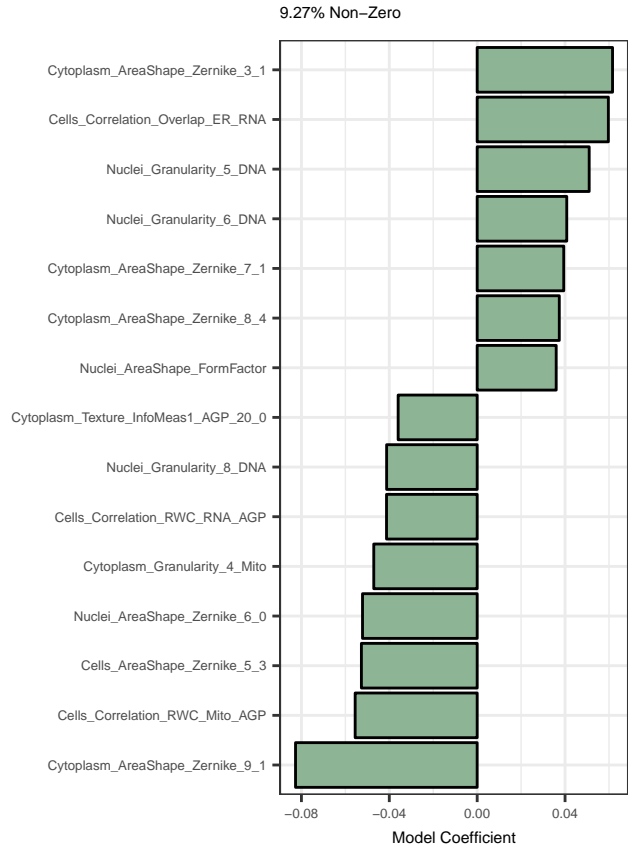
cc_cc_late_mitosis



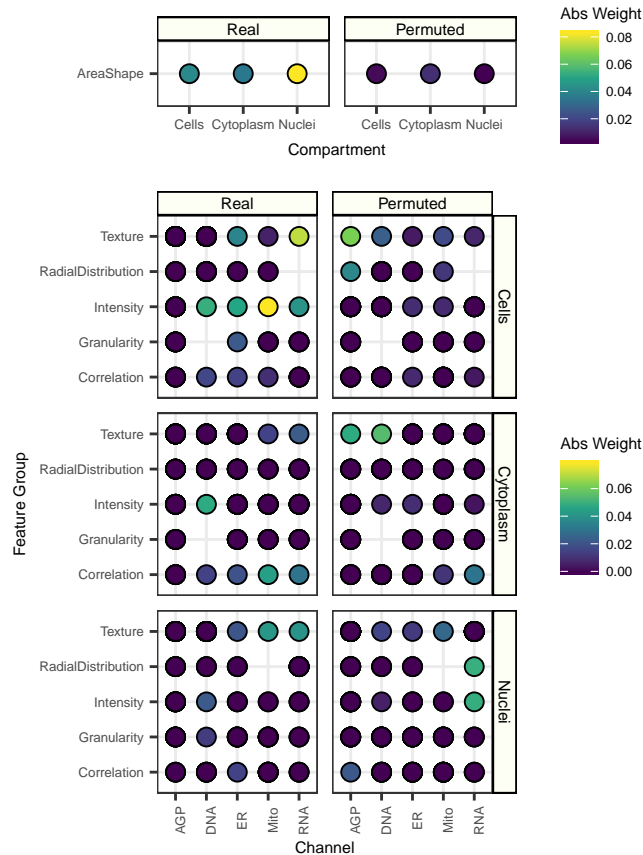
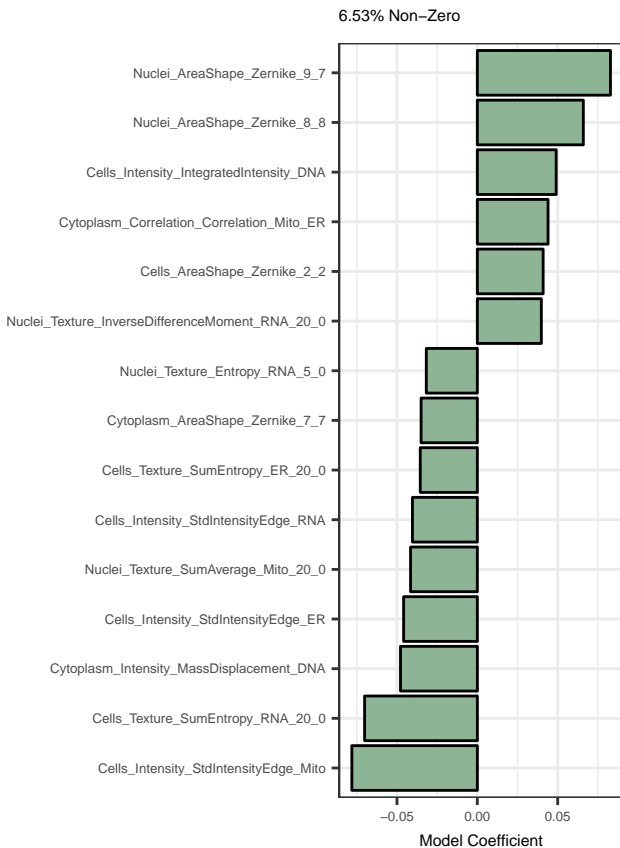
cc_cc_mitosis



cc_cc_n_objects

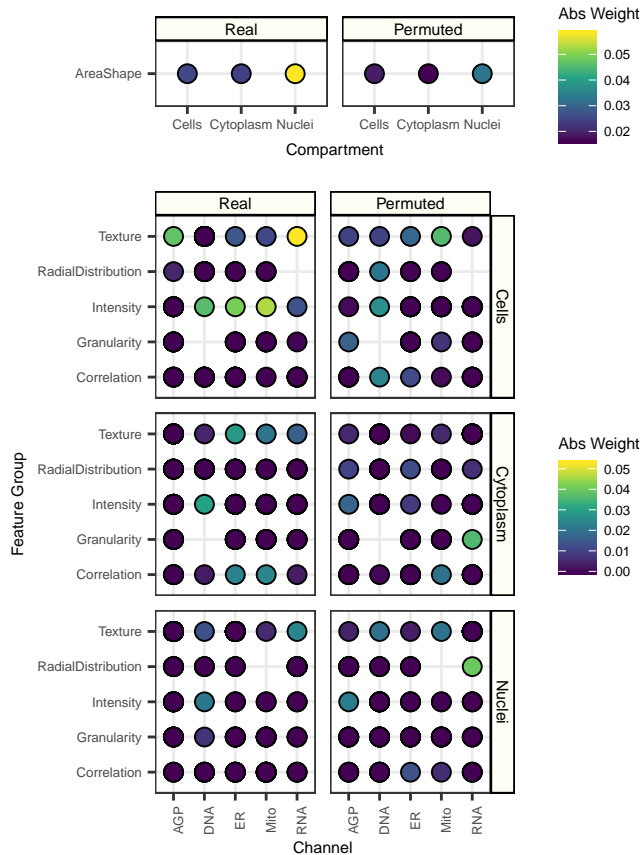
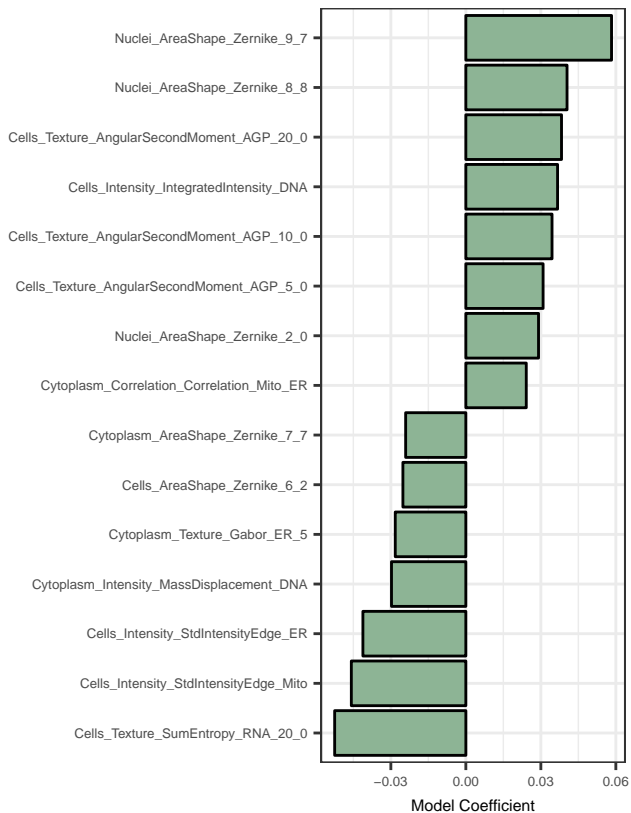


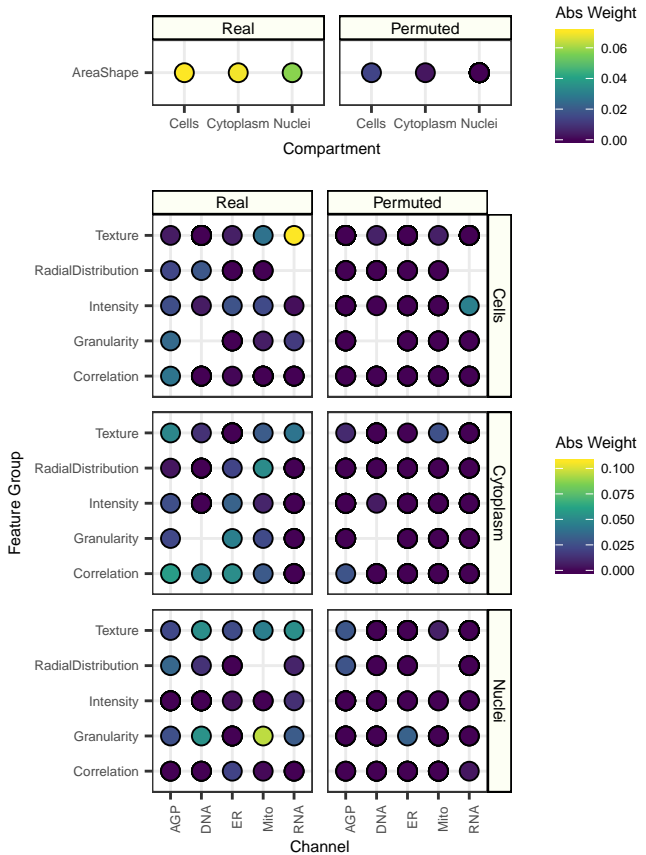
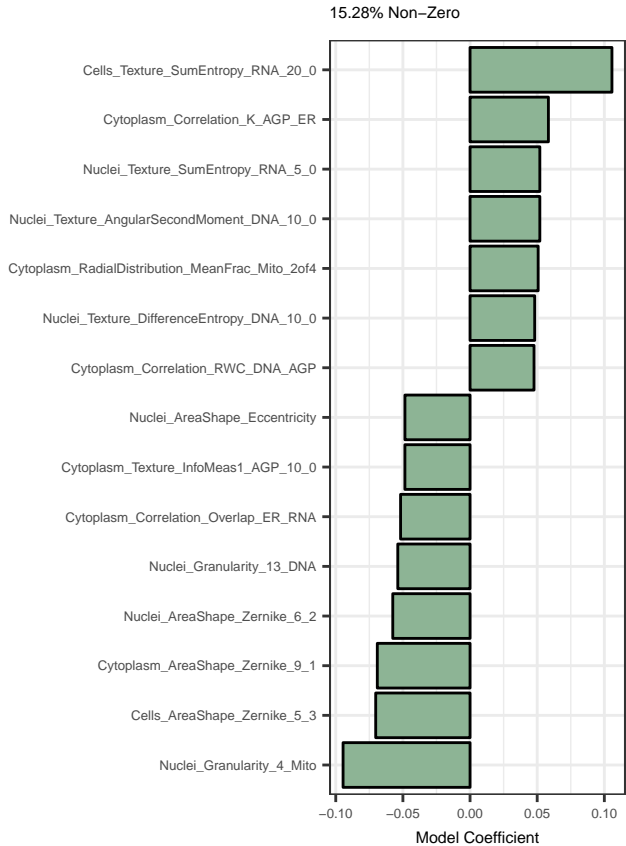
cc_cc_n_spots_h2ax_mean



cc_cc_n_spots_h2ax_per_nucleus_area_mean

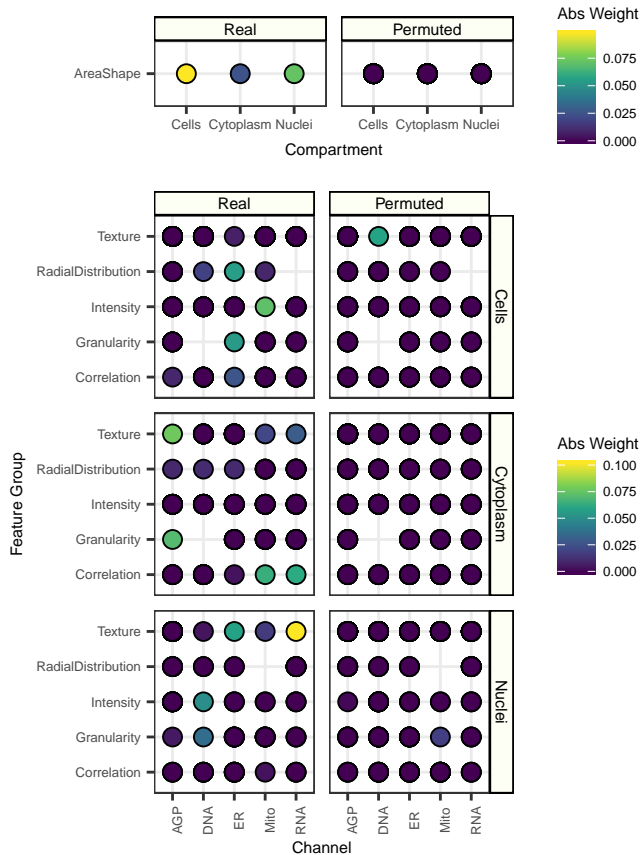
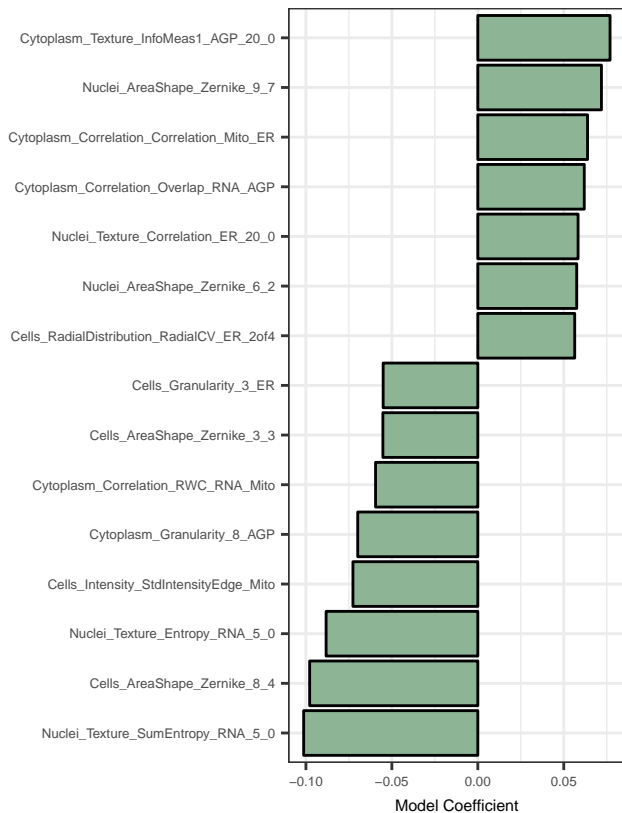
7.38% Non-Zero



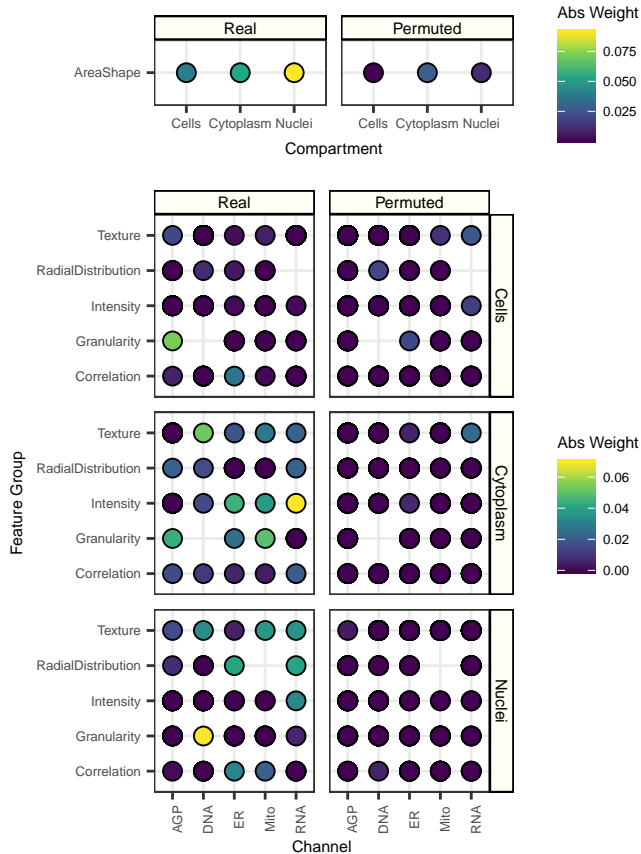
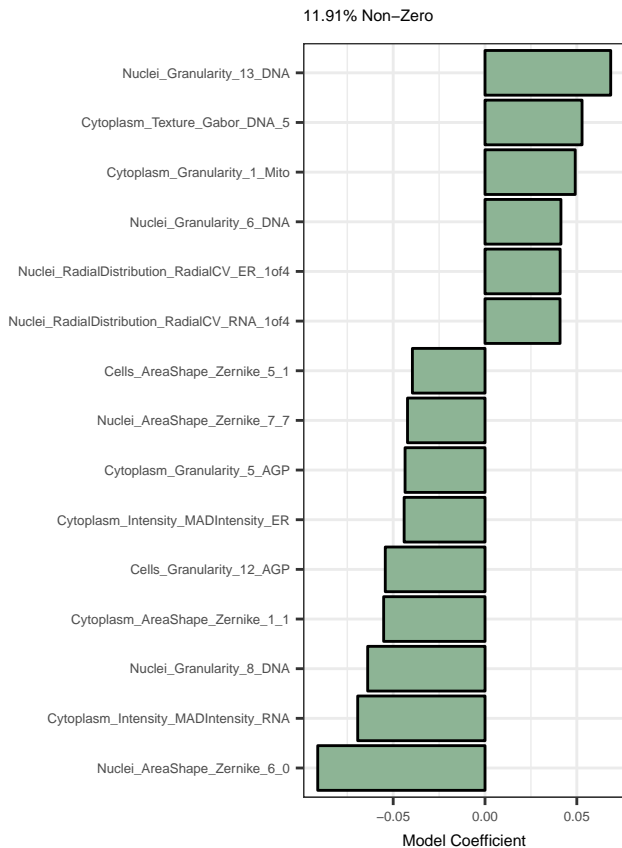


cc_early_mitosis_high_h2ax

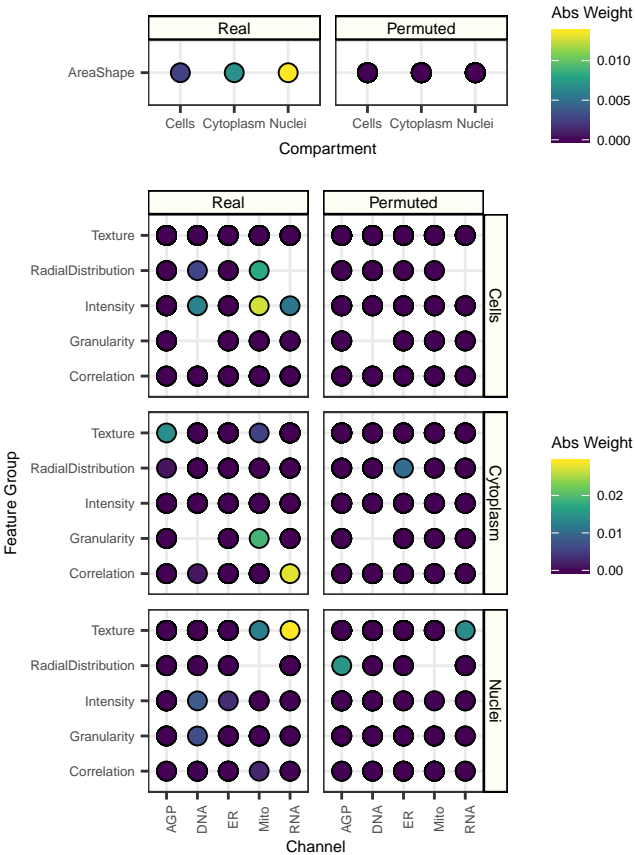
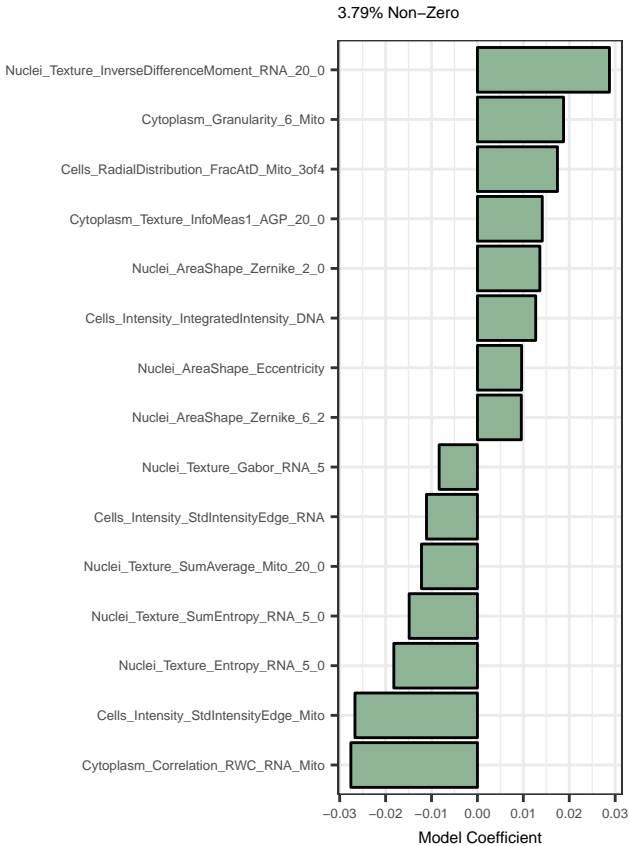
7.06% Non-Zero



cc_early_mitosis_n_objects

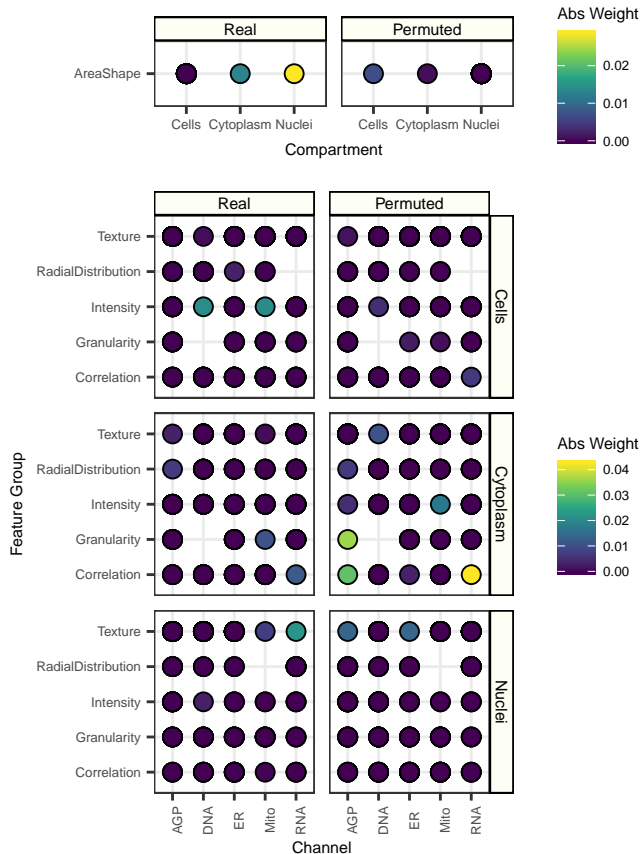
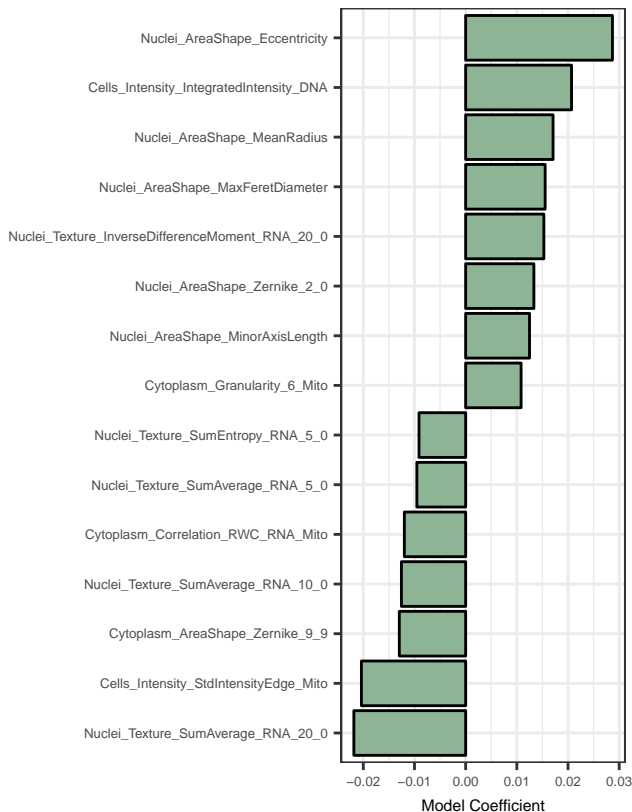


cc_early_mitosis_n_spots_h2ax_mean



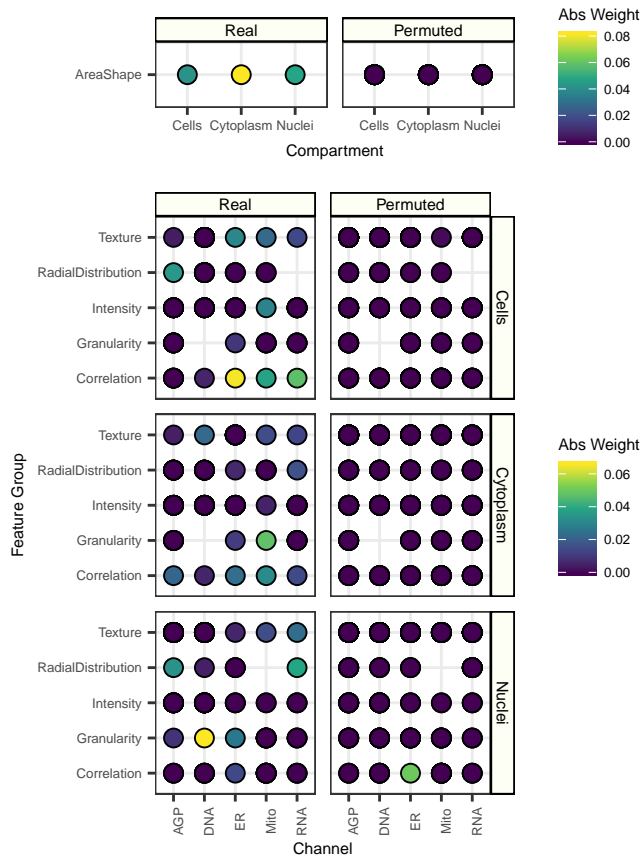
cc_early_mitosis_n_spots_h2ax_per_nucleus_area_mean

3.69% Non-Zero

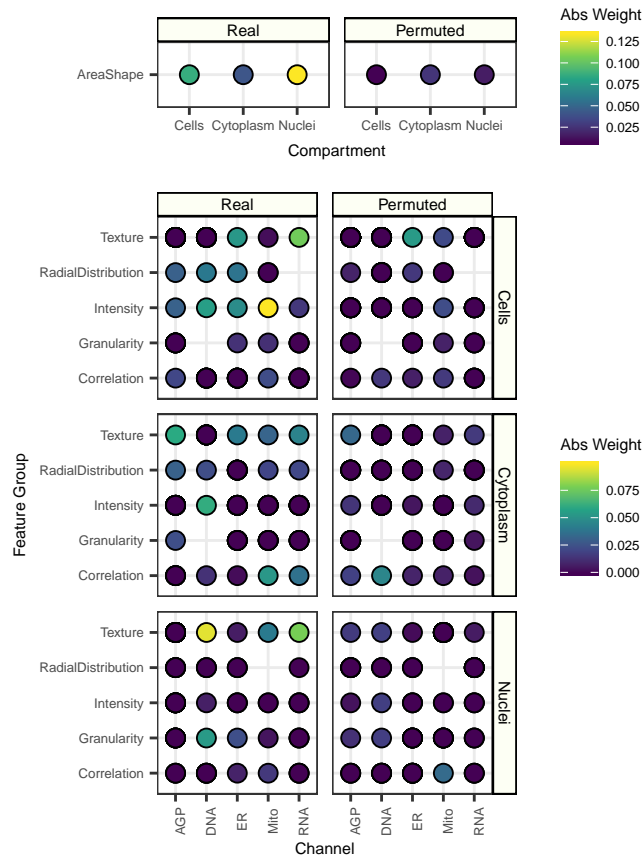


cc_g1_n_objects

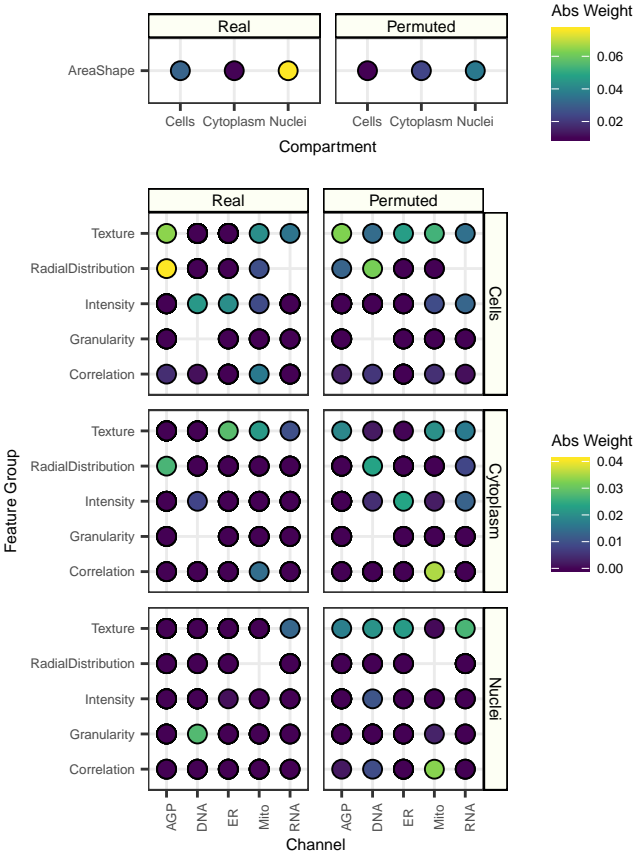
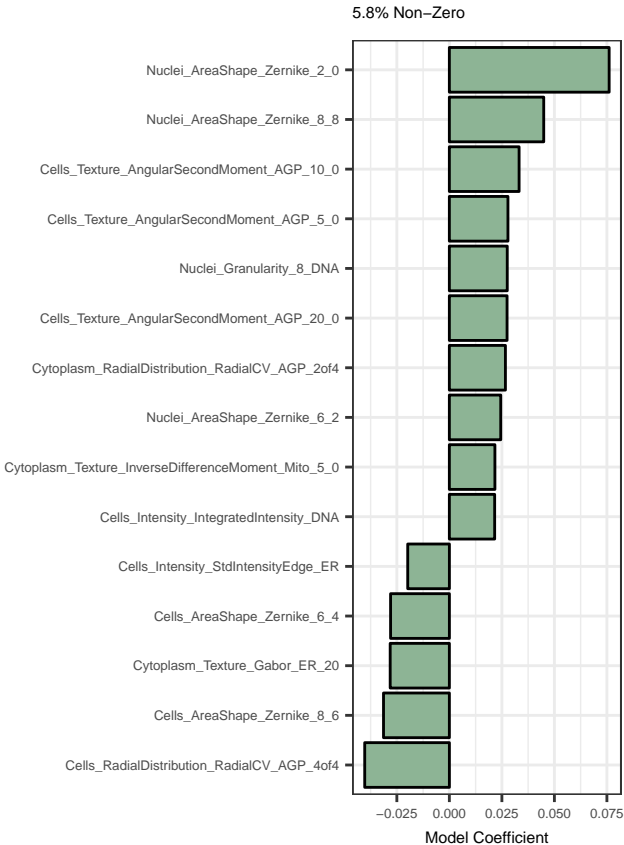
11.59% Non-Zero



cc_g1_n_spots_h2ax_mean

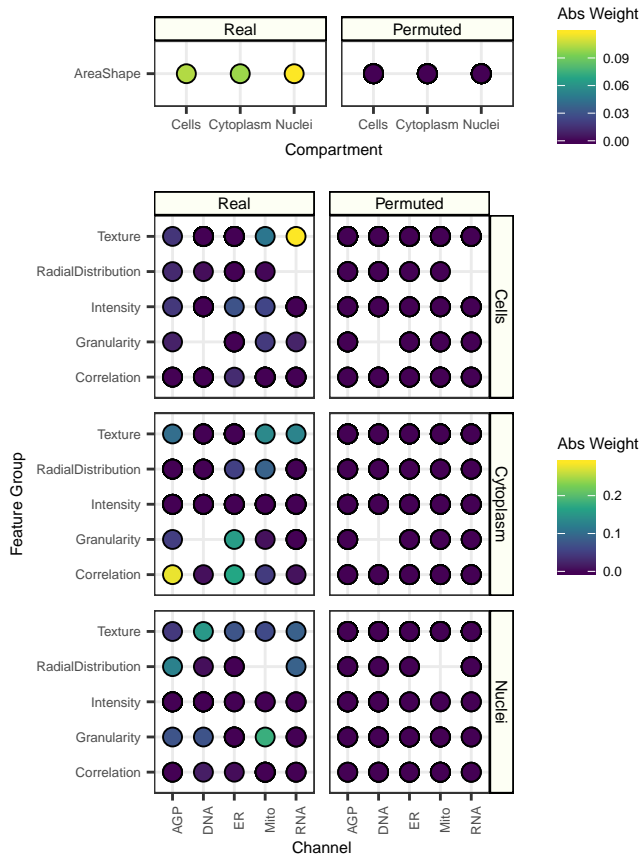
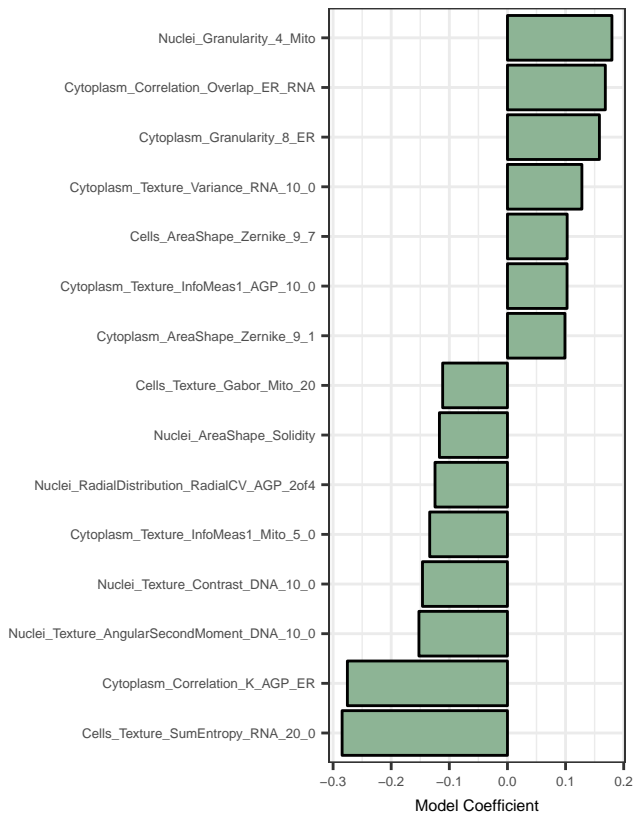


cc_g1_n_spots_h2ax_per_nucleus_area_mean



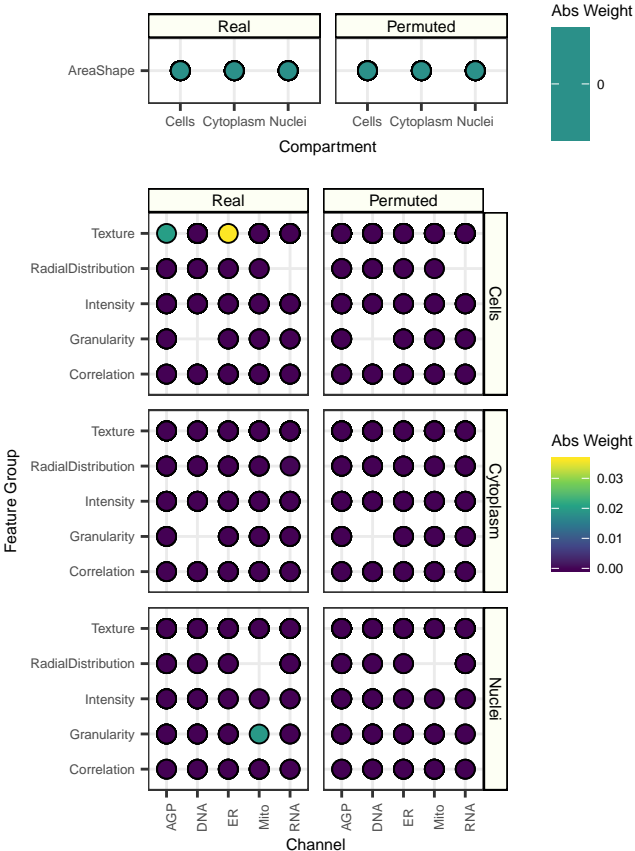
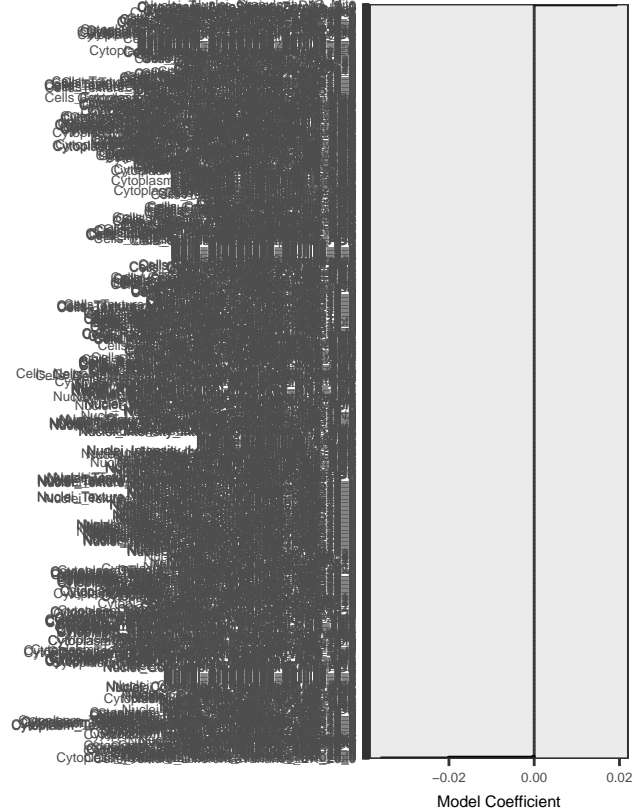
cc_g1_plus_g2_count

9.91% Non-Zero

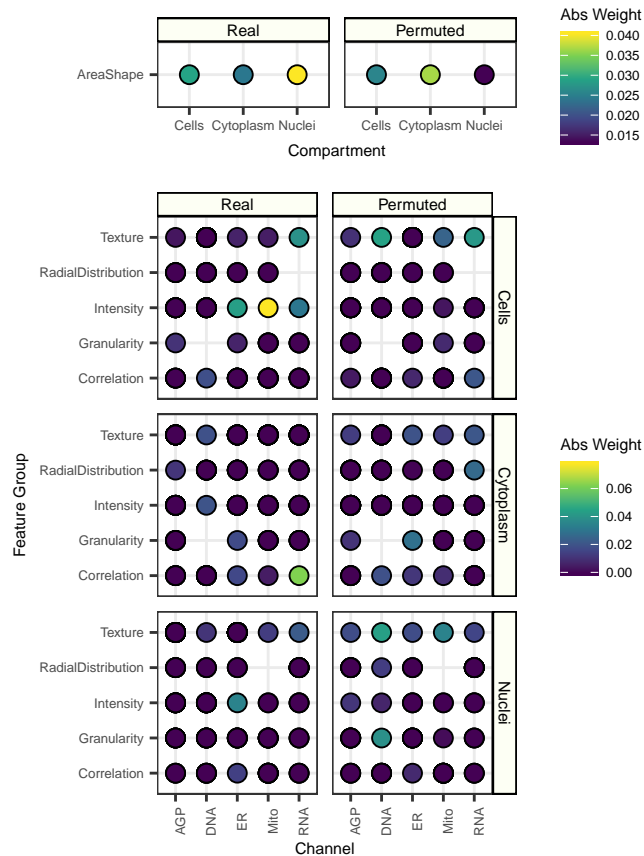
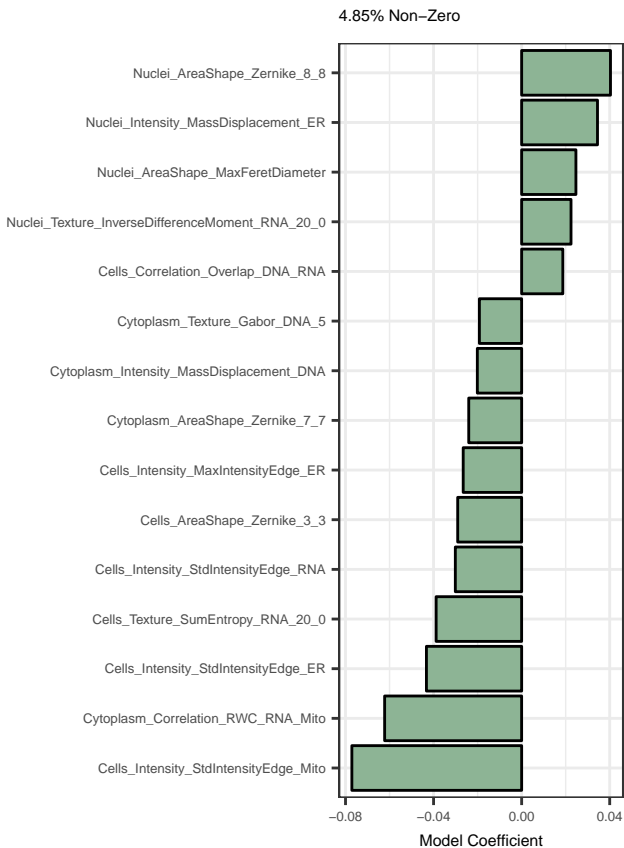


cc_g2_g1_count

0.42% Non-Zero

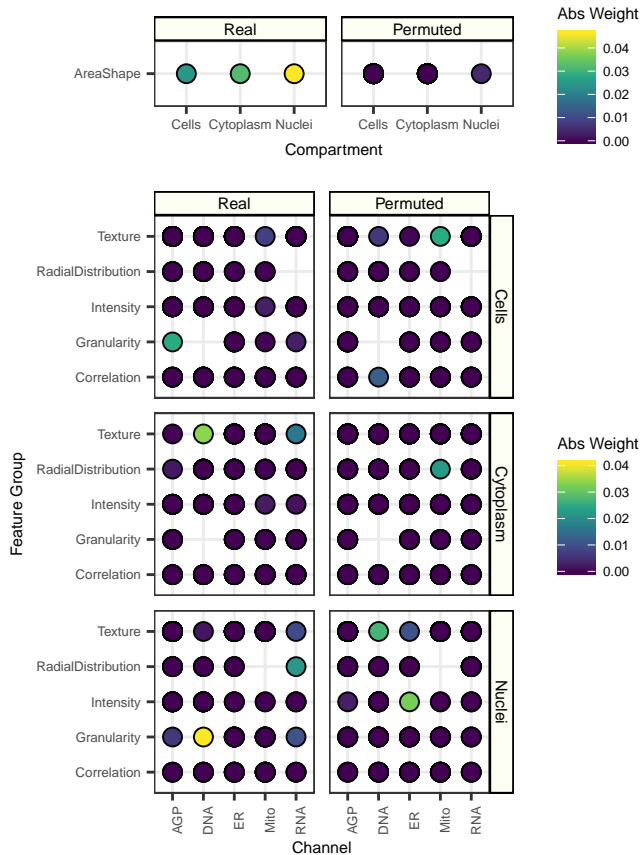
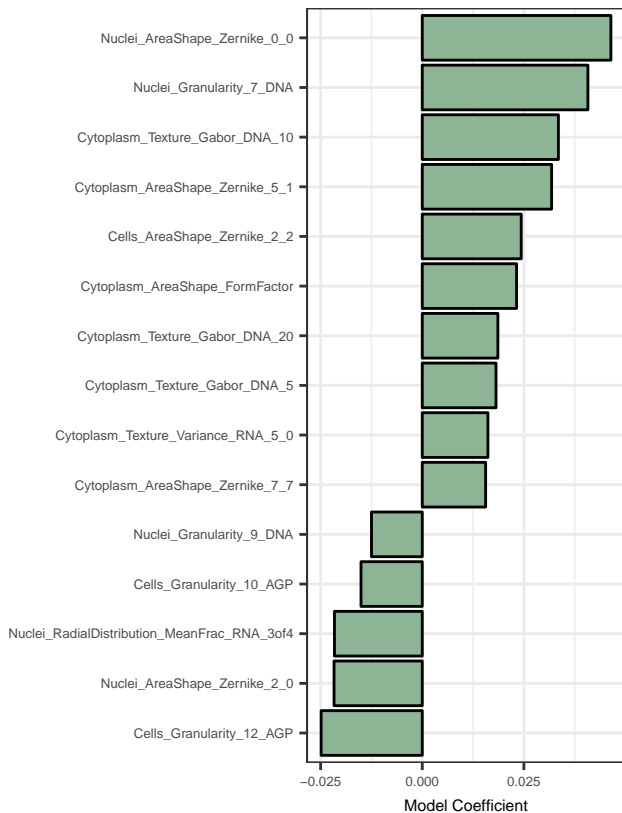


cc_g2_high_h2ax



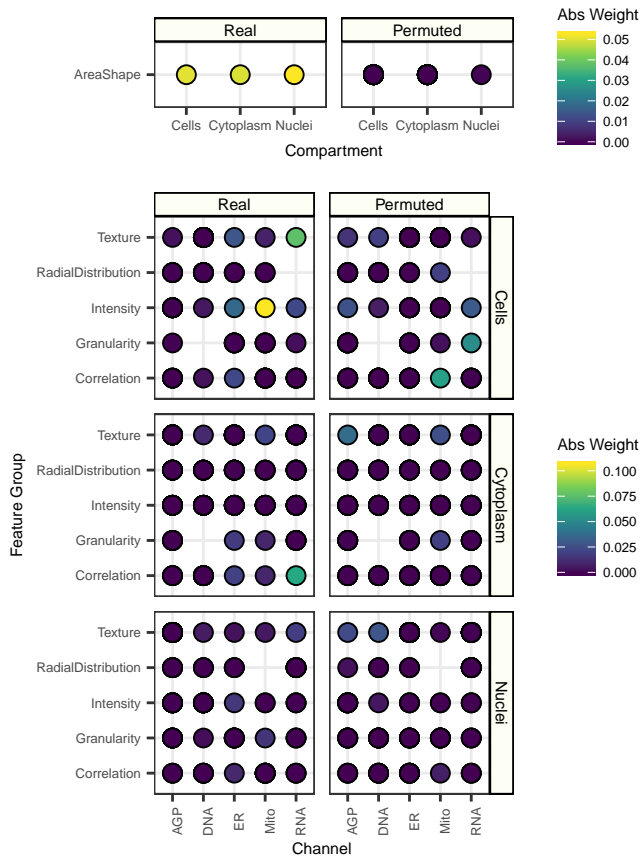
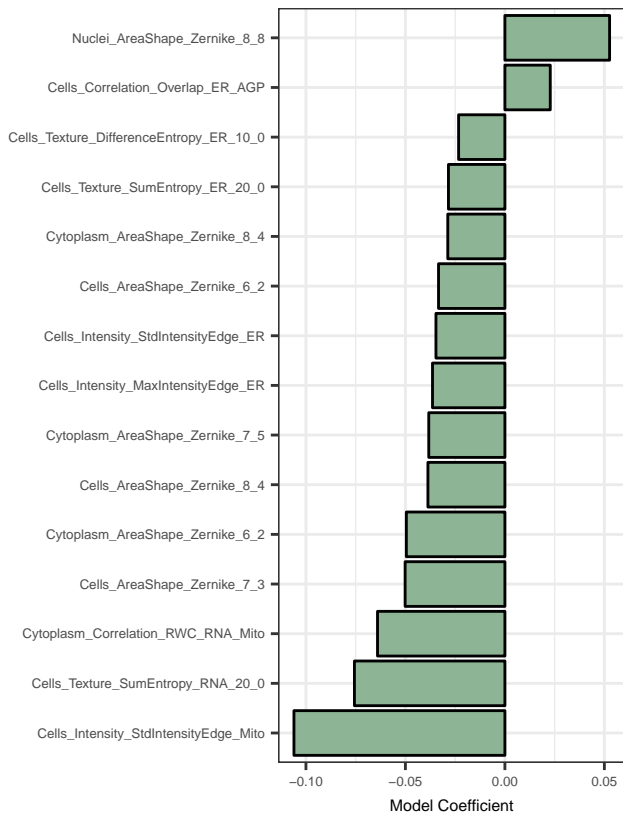
cc_g2_n_objects

6.01% Non-Zero



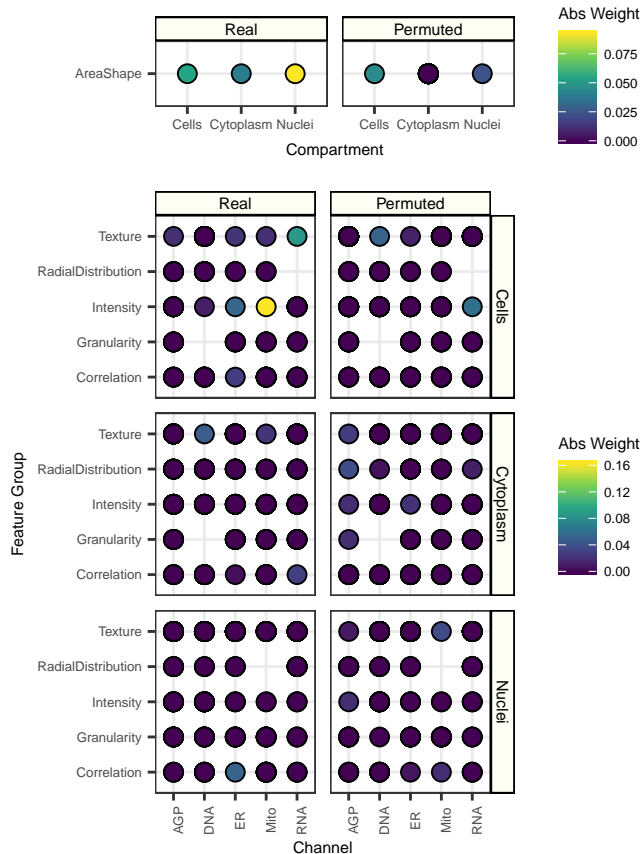
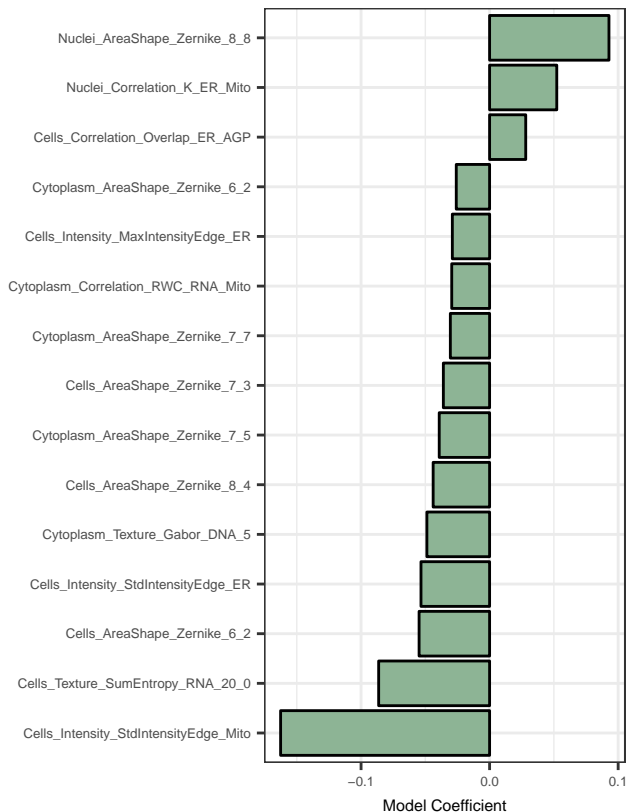
cc_g2_n_spots_h2ax_mean

6.85% Non-Zero

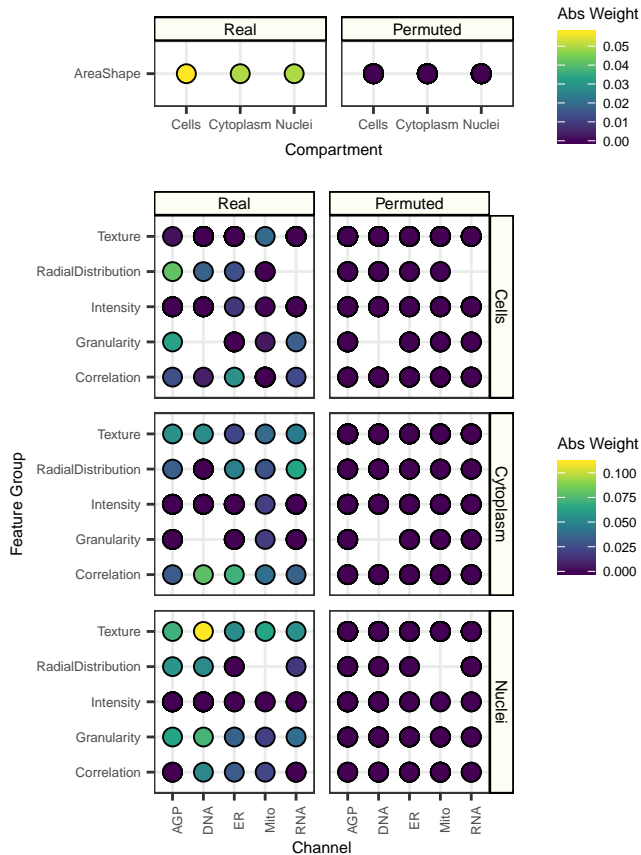


cc_g2_n_spots_h2ax_per_nucleus_area_mean

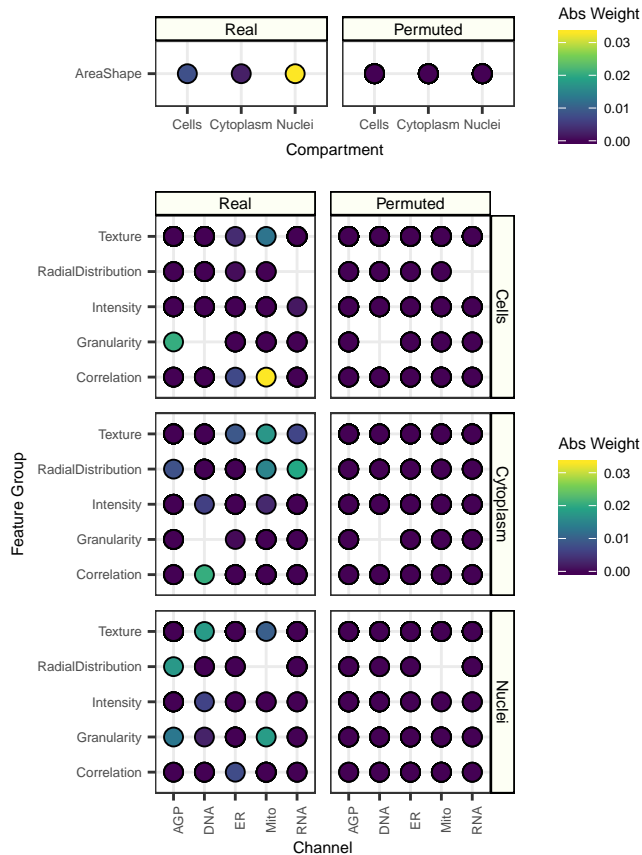
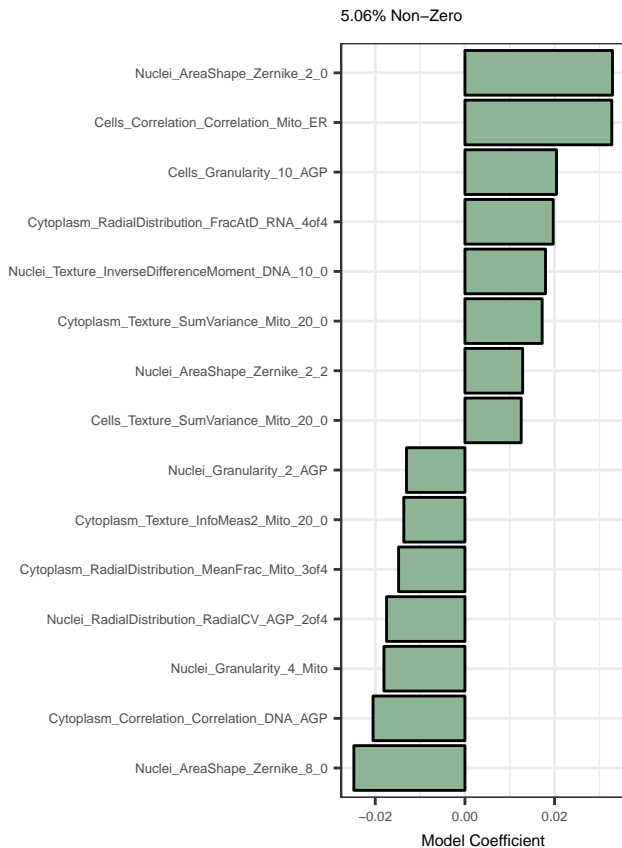
2.63% Non-Zero



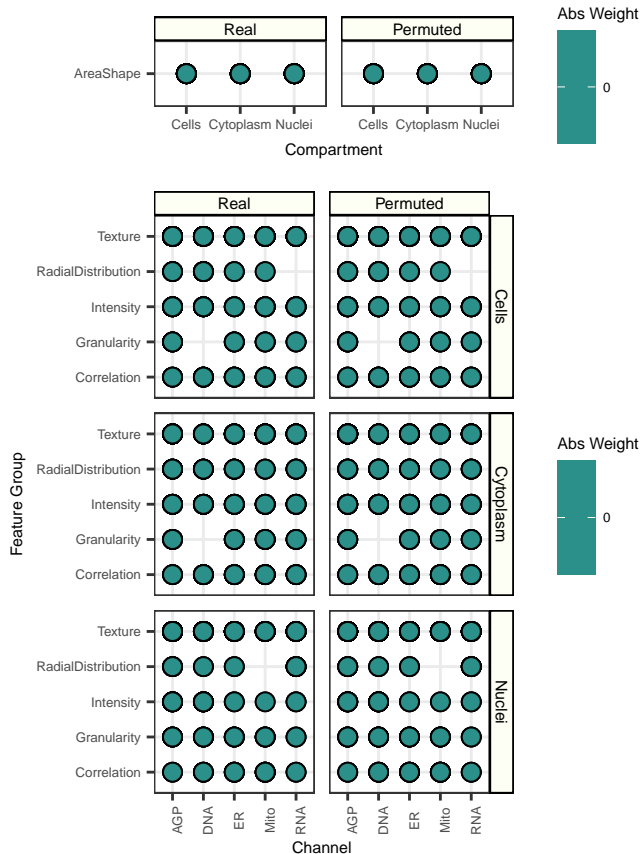
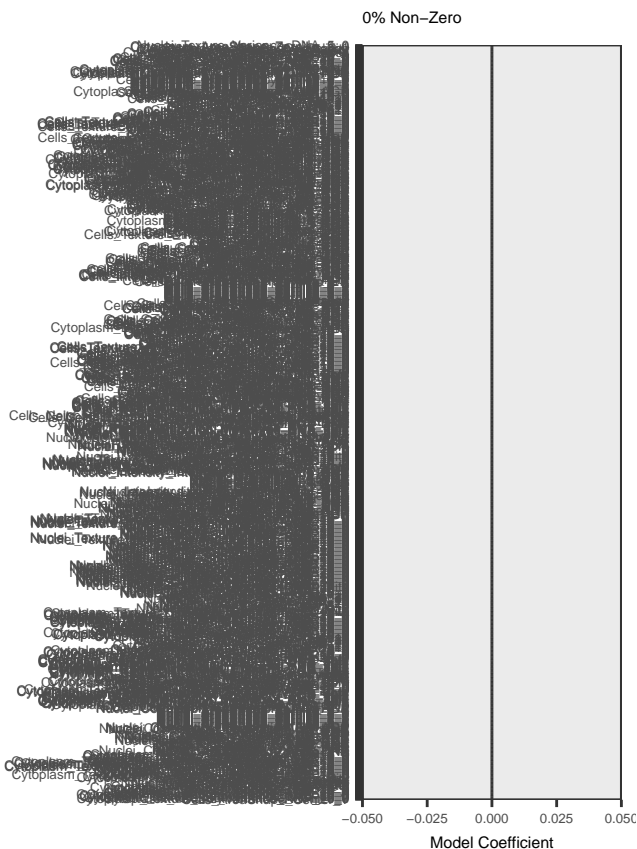
cc_g2_plus_all_m_count



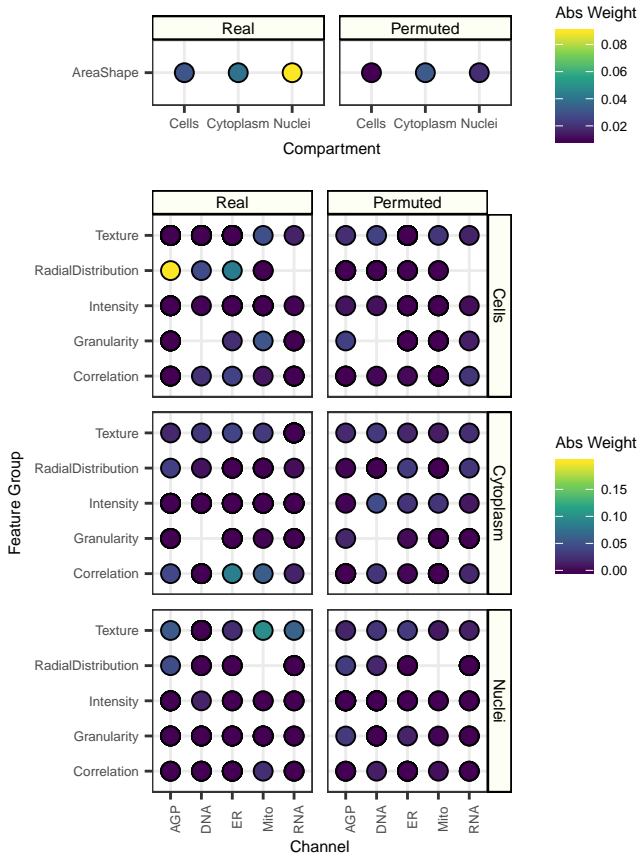
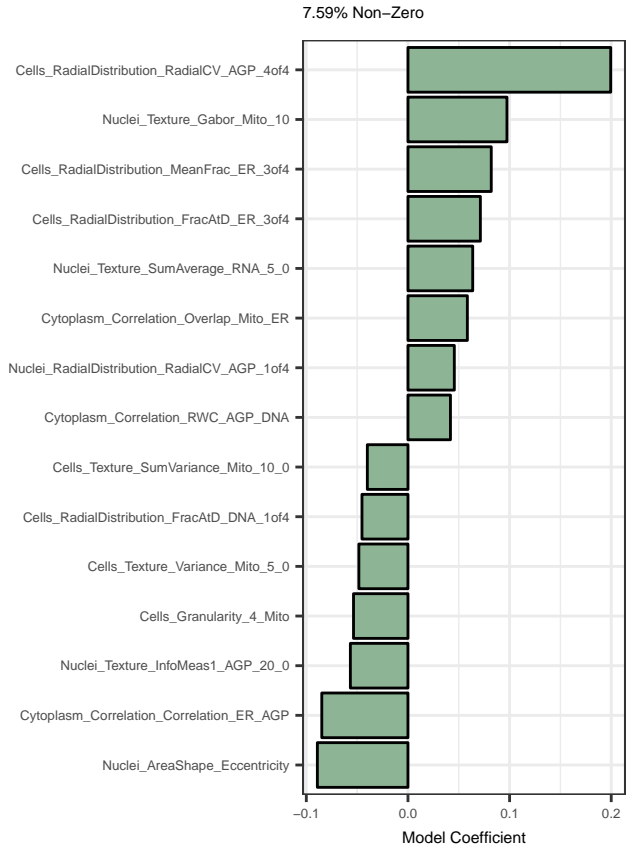
cc_infection_percentage



cc_late_mitosis_high_h2ax

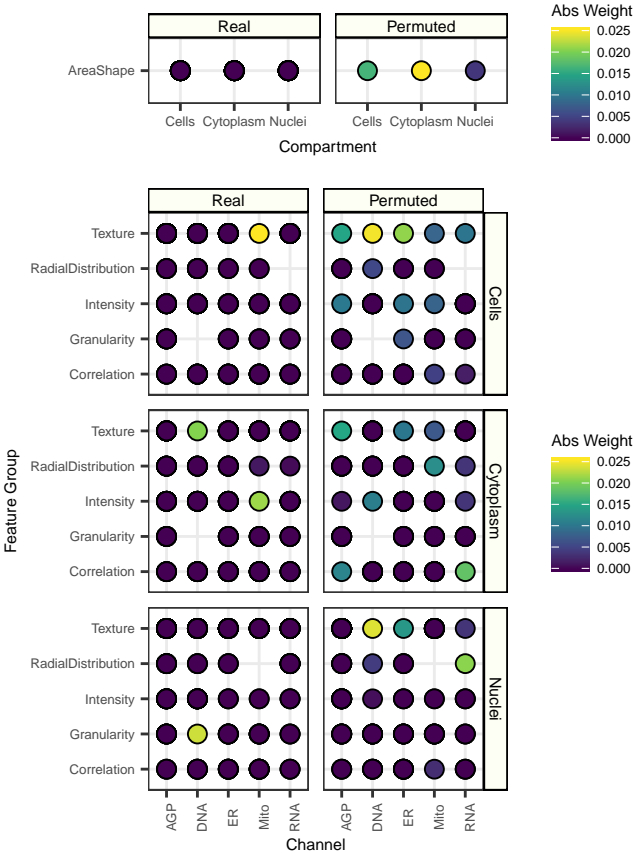
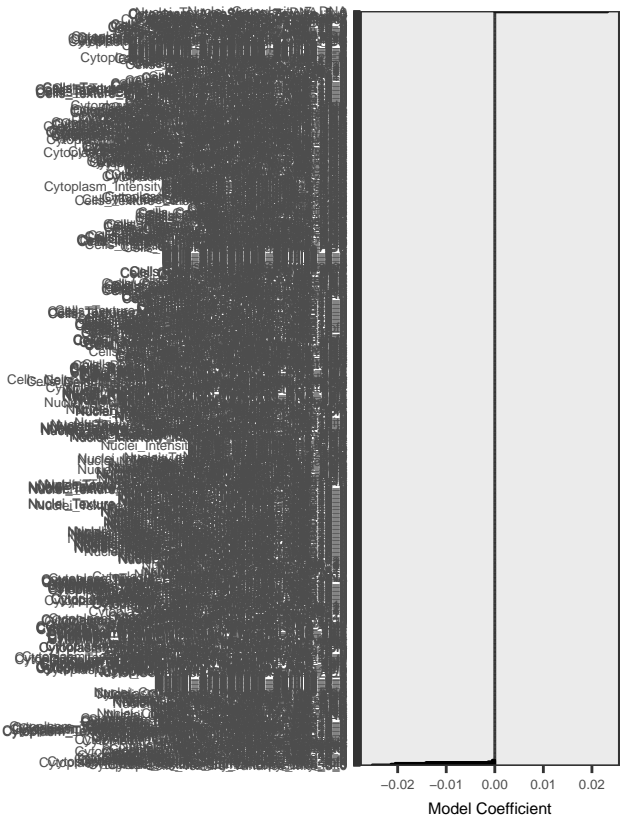


cc_late_mitosis_n_objects

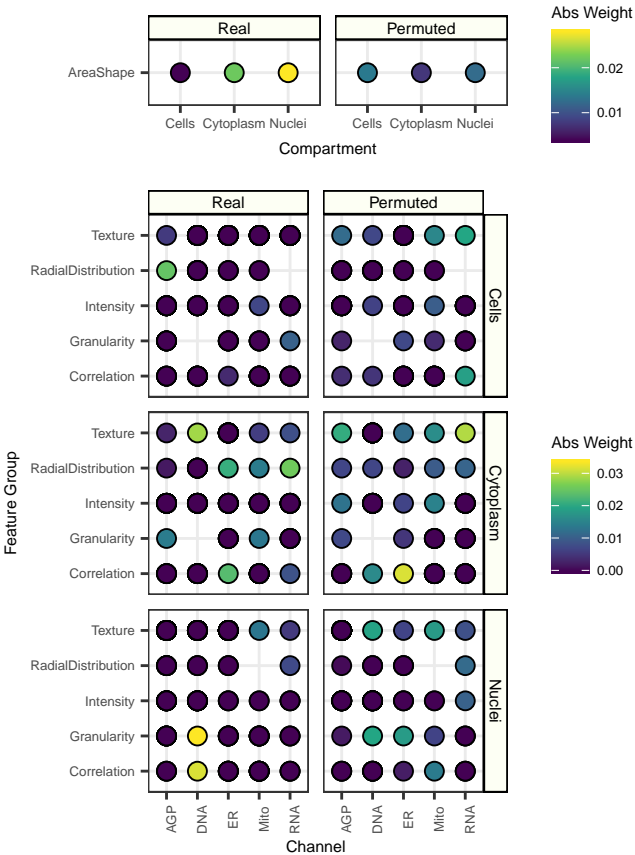
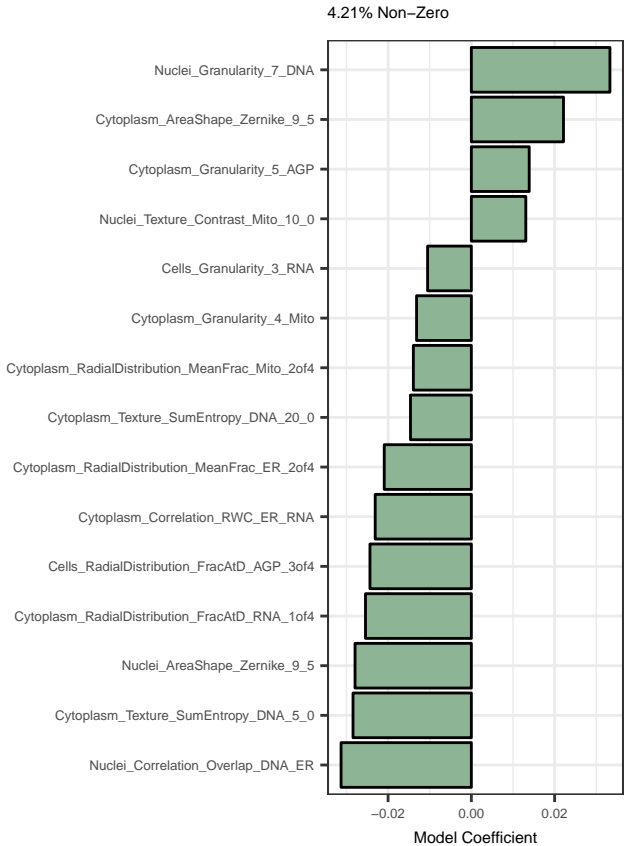


cc_late_mitosis_n_spots_h2ax_mean

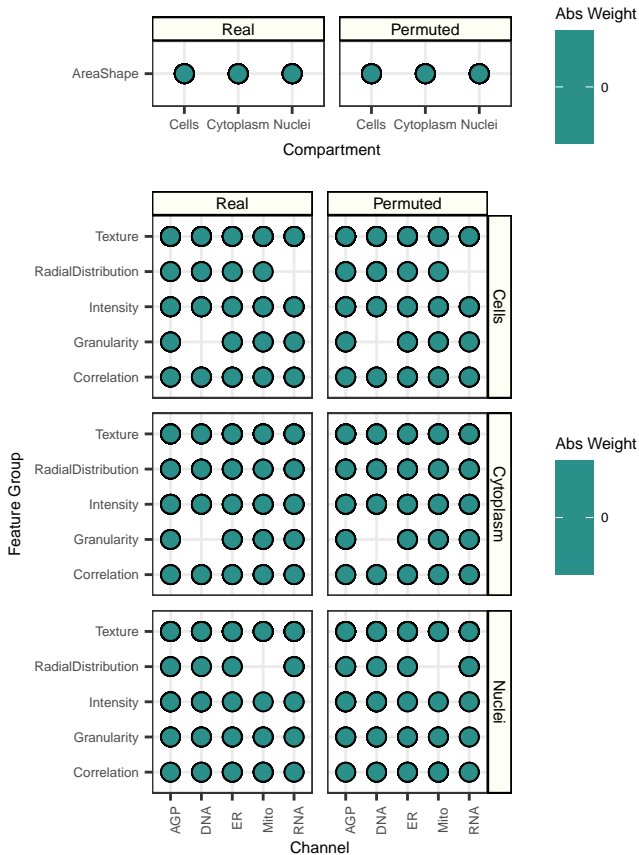
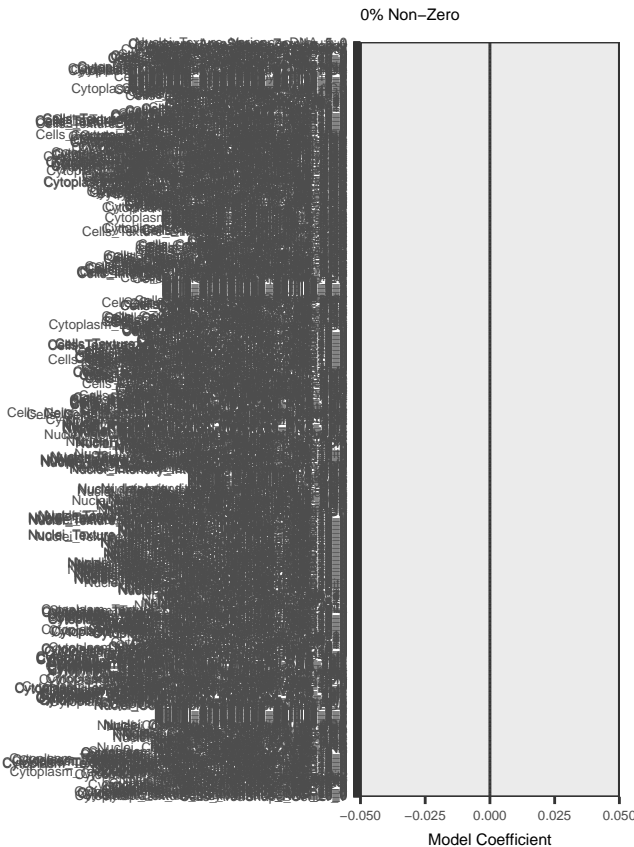
0.84% Non-Zero



cc_late_mitosis_n_spots_h2ax_per_nucleus_area_mean

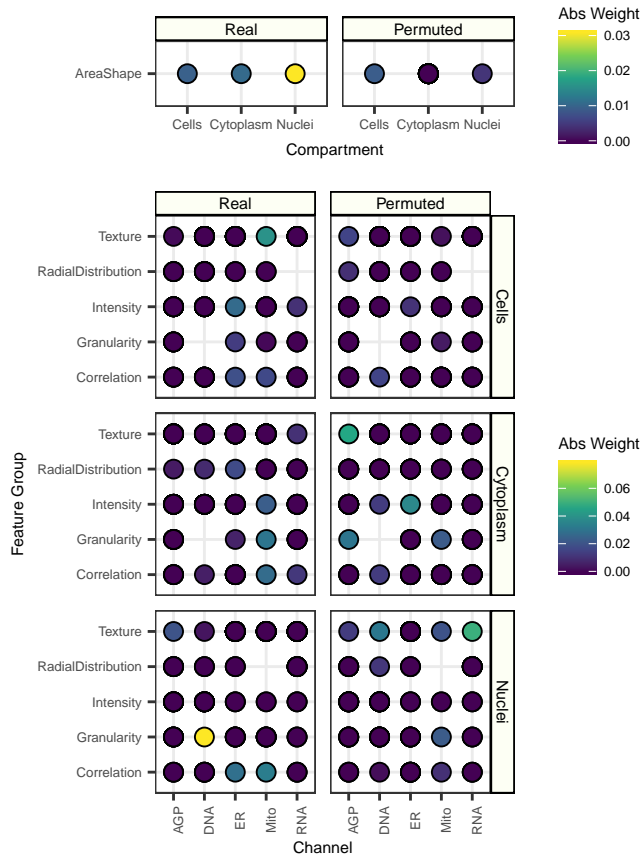
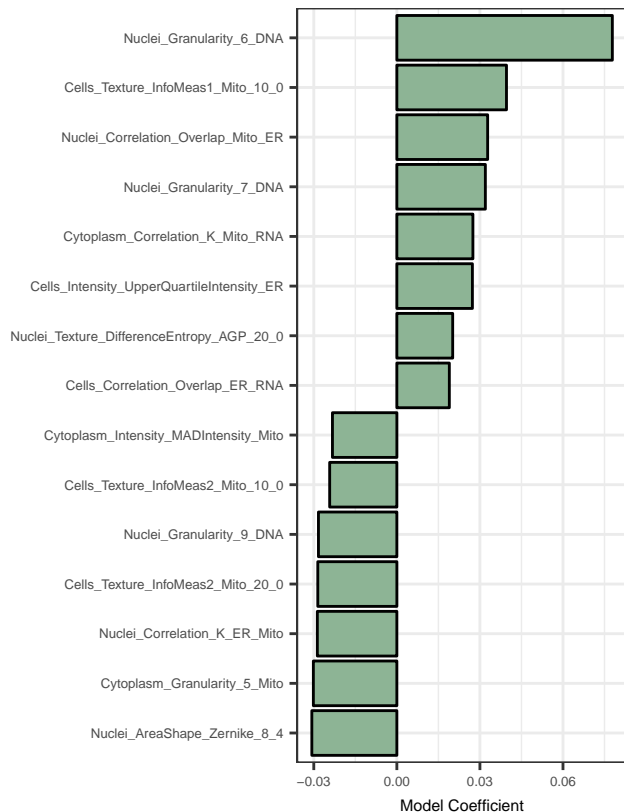


cc_mitosis_high_h2ax



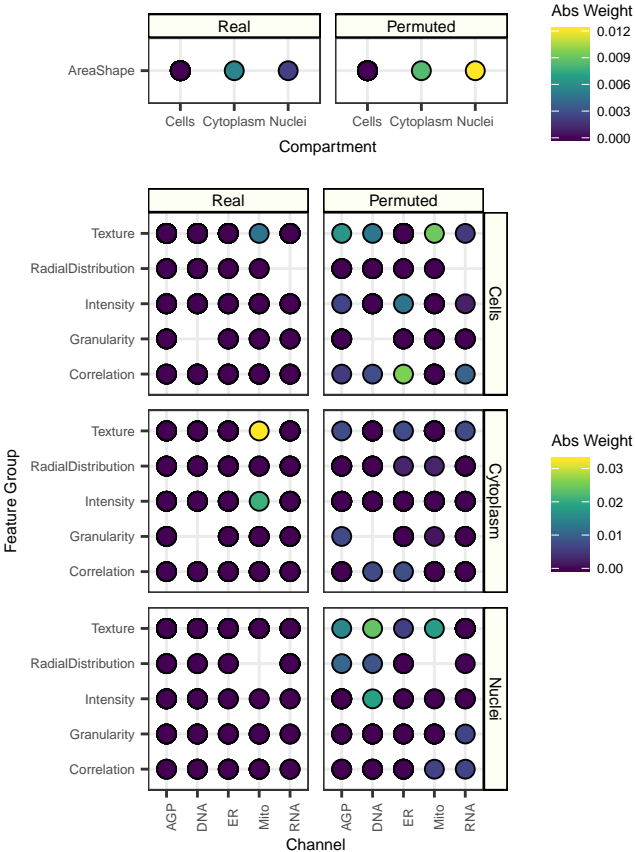
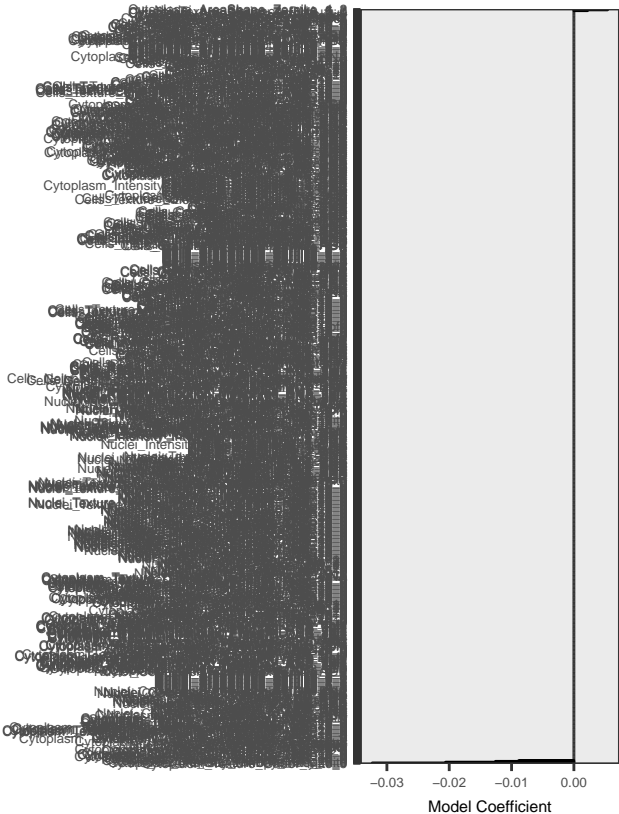
cc_mitosis_n_objects

6.11% Non-Zero



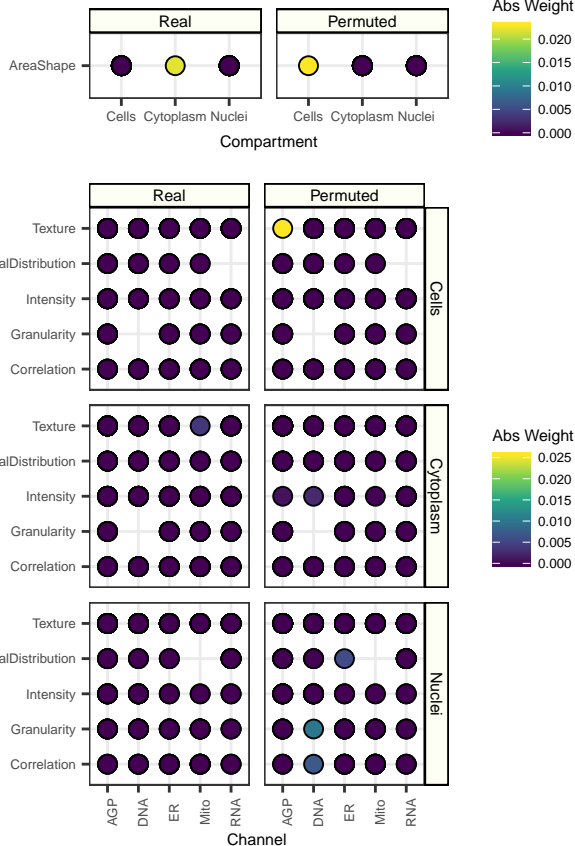
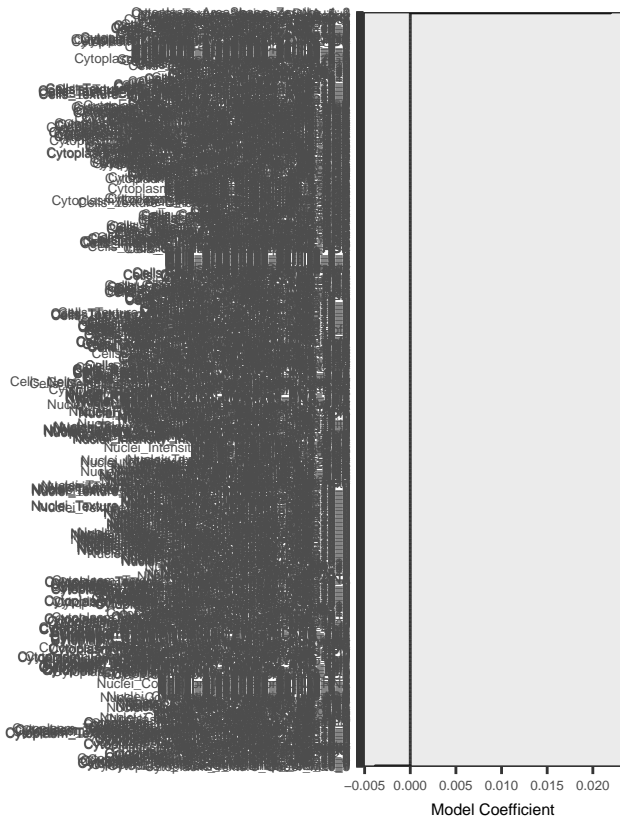
cc_mitosis_n_spots_h2ax_mean

0.63% Non-Zero

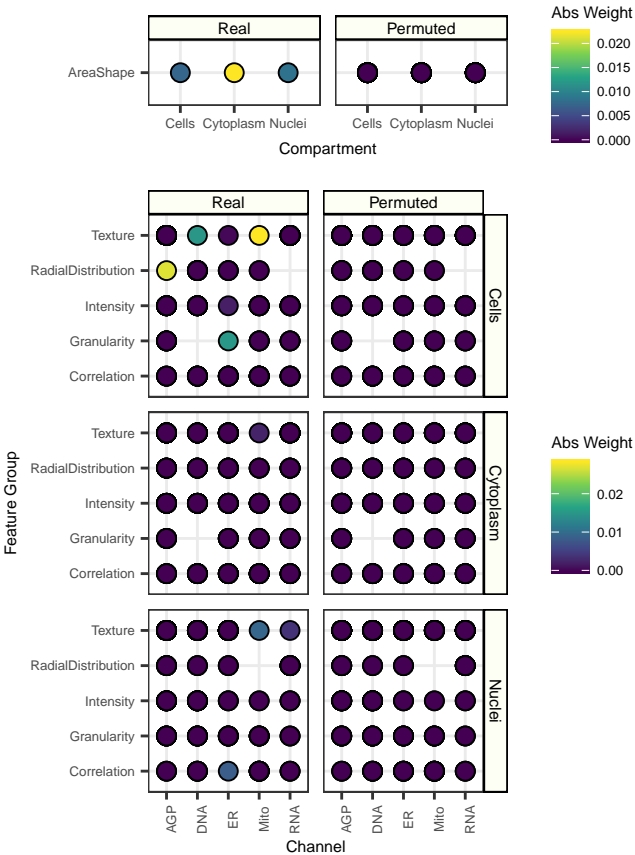
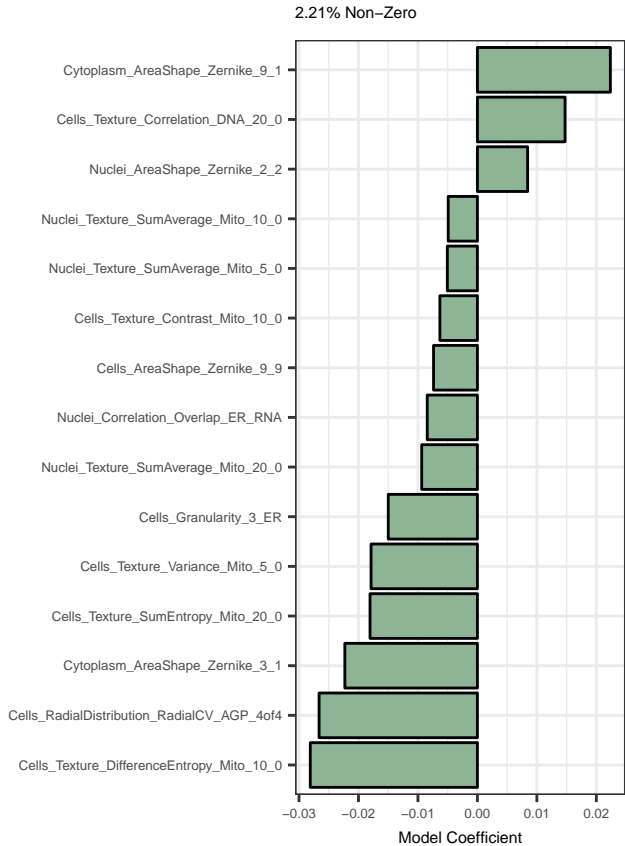


cc_mitosis_n_spots_h2ax_per_nucleus_area_mean

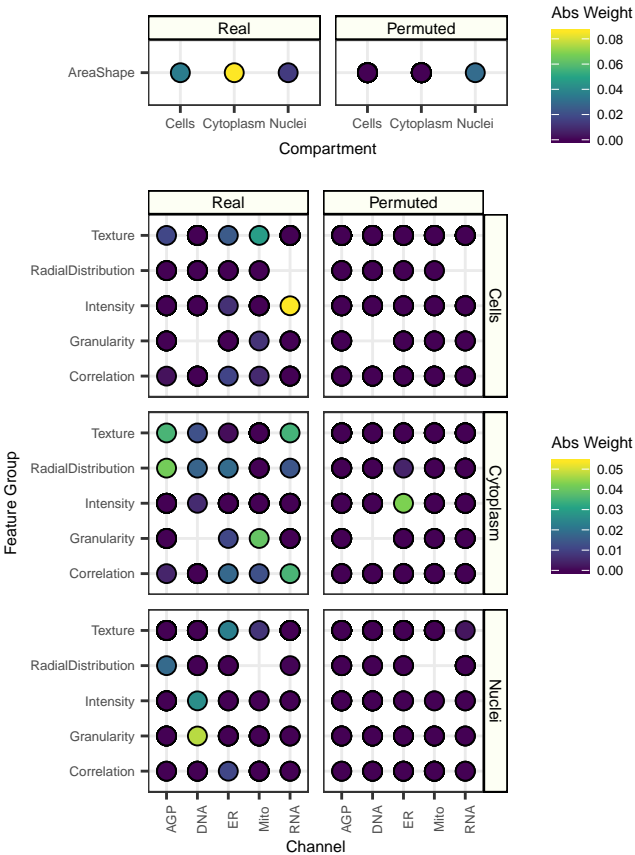
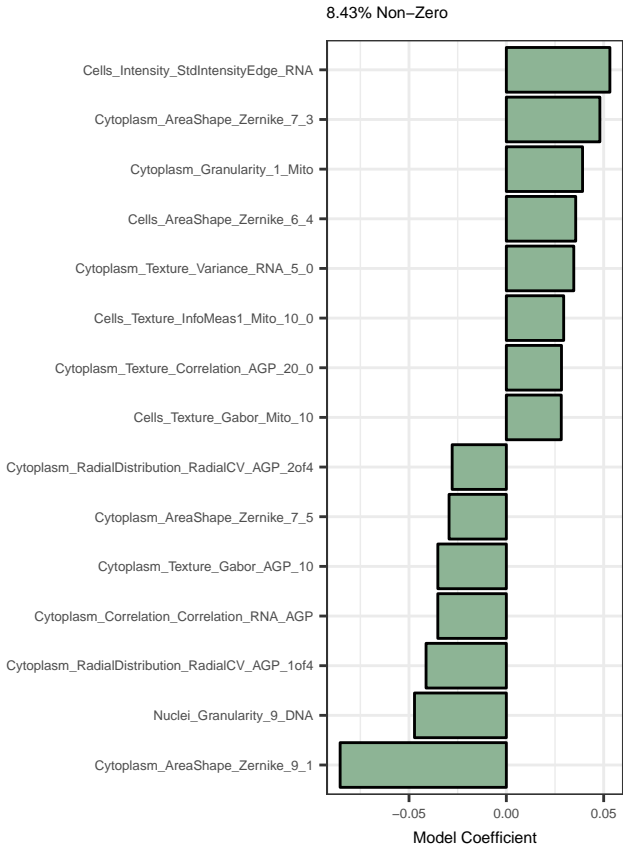
0.21% Non-Zero



cc_polynuclear_high_h2ax

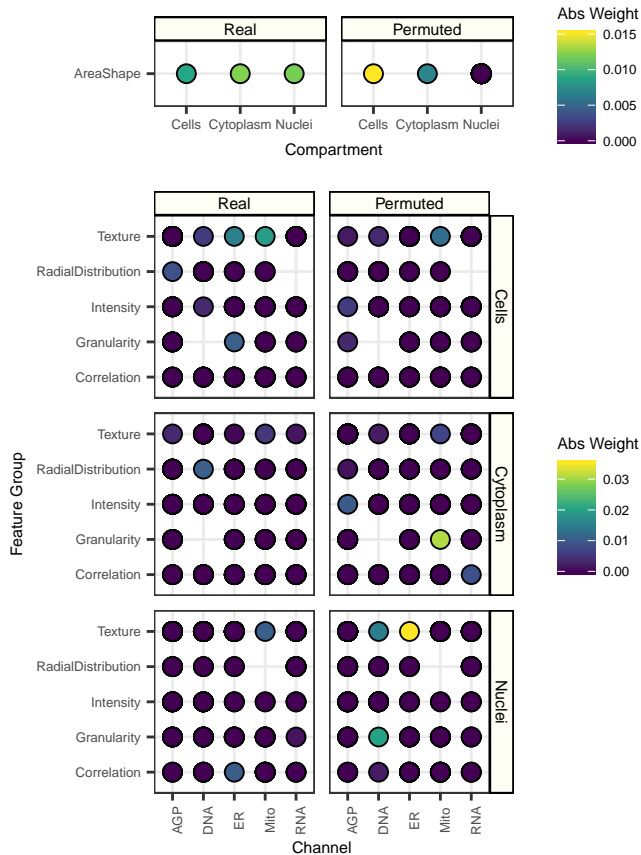
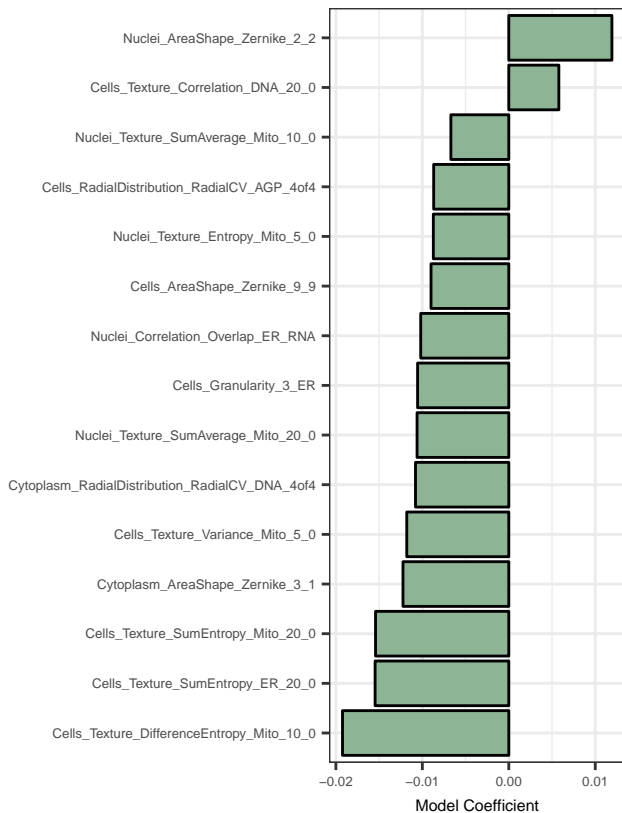


cc_polynuclear_n_objects



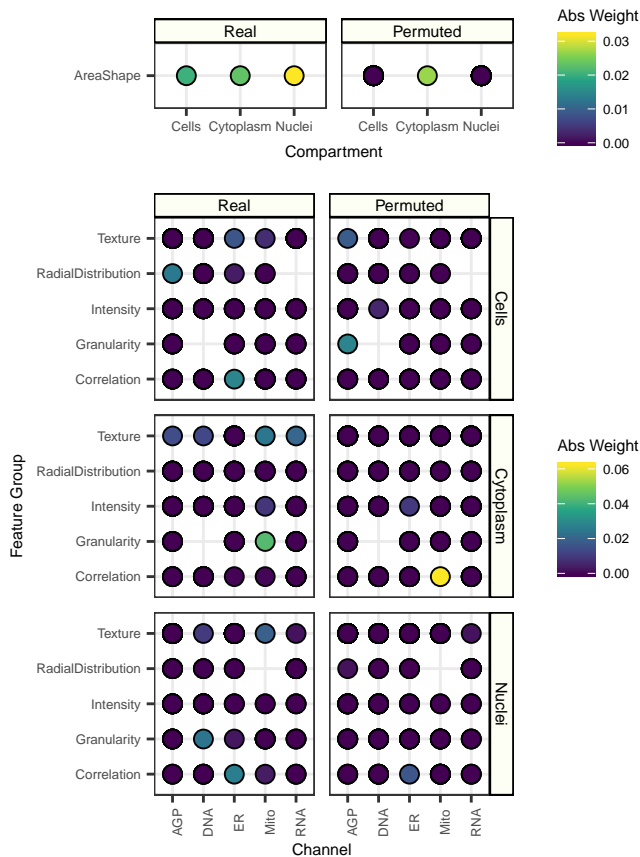
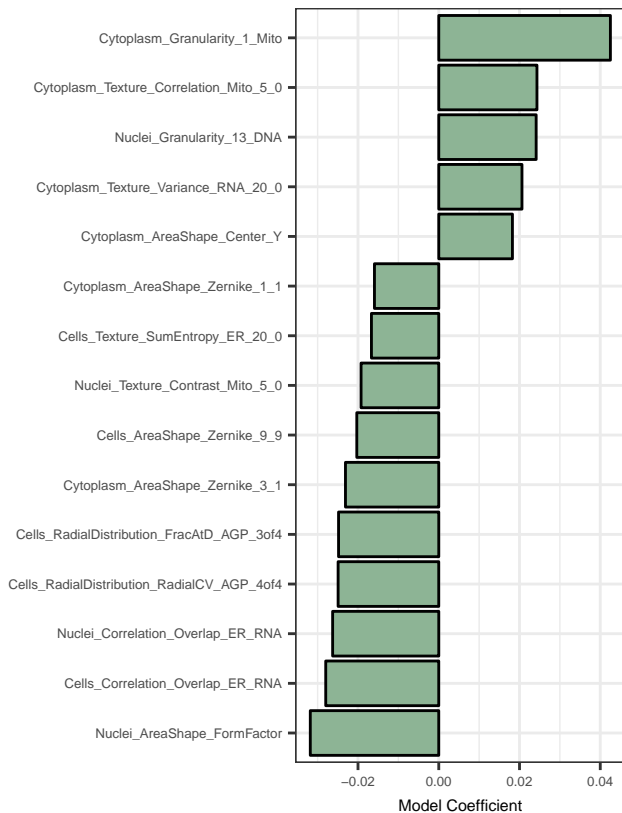
cc_polynuclear_n_spots_h2ax_mean

2.85% Non-Zero

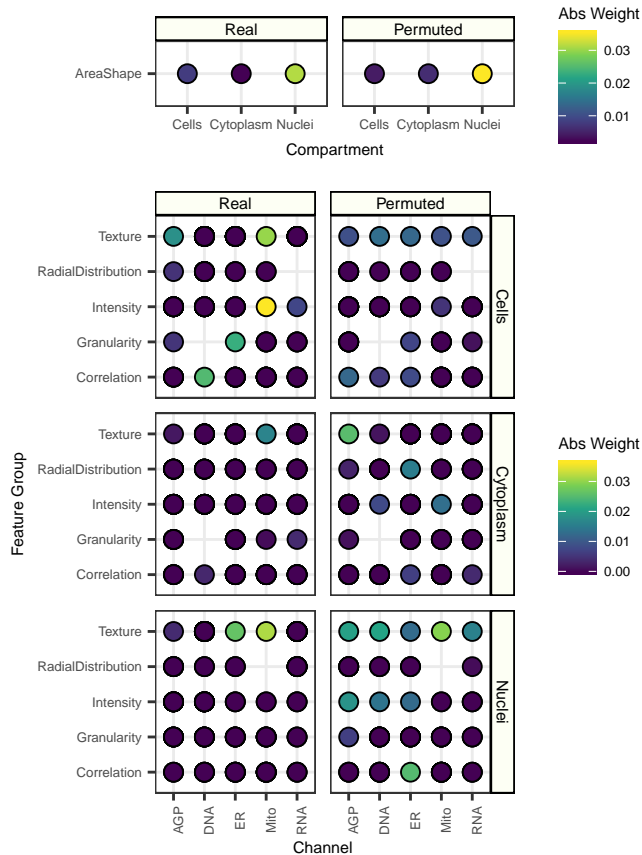
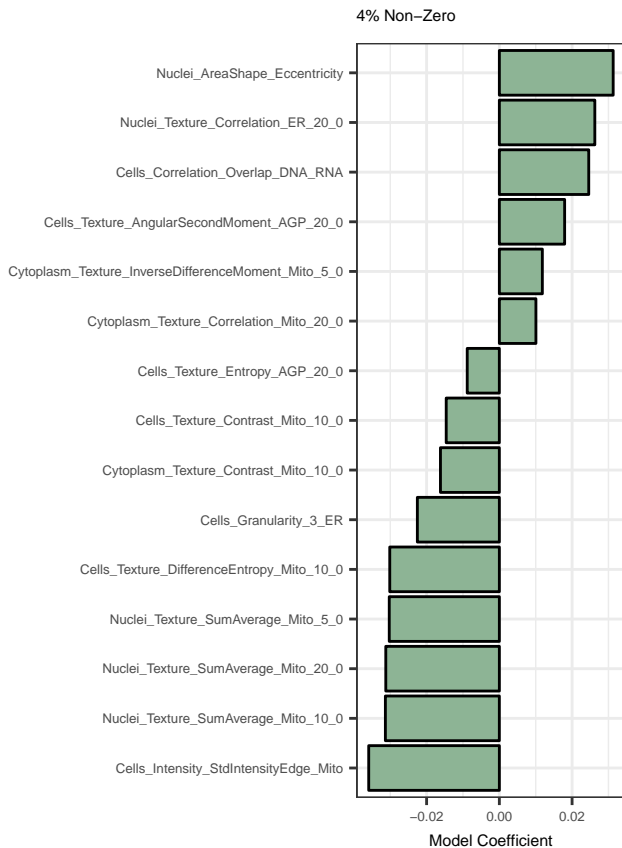


cc_polynuclear_n_spots_h2ax_per_nucleus_area_mean

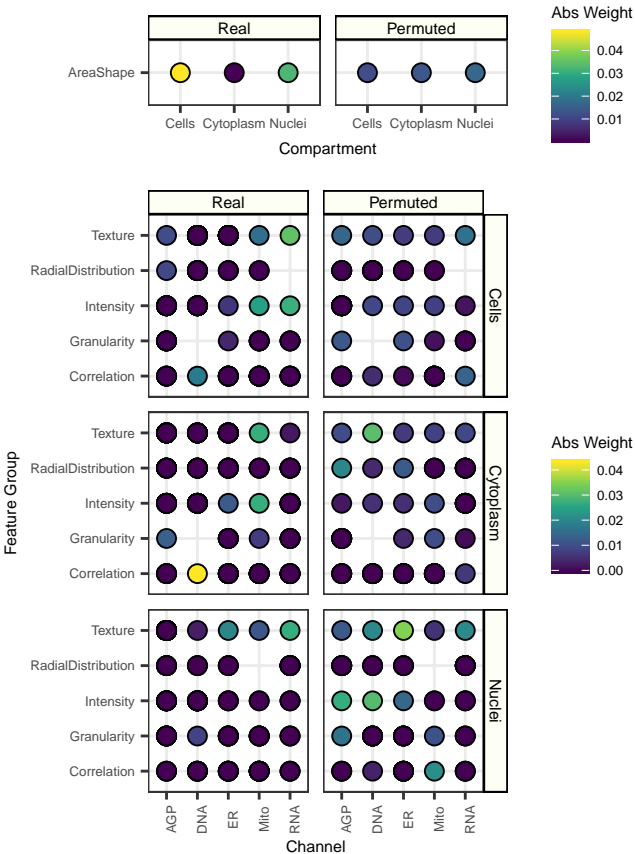
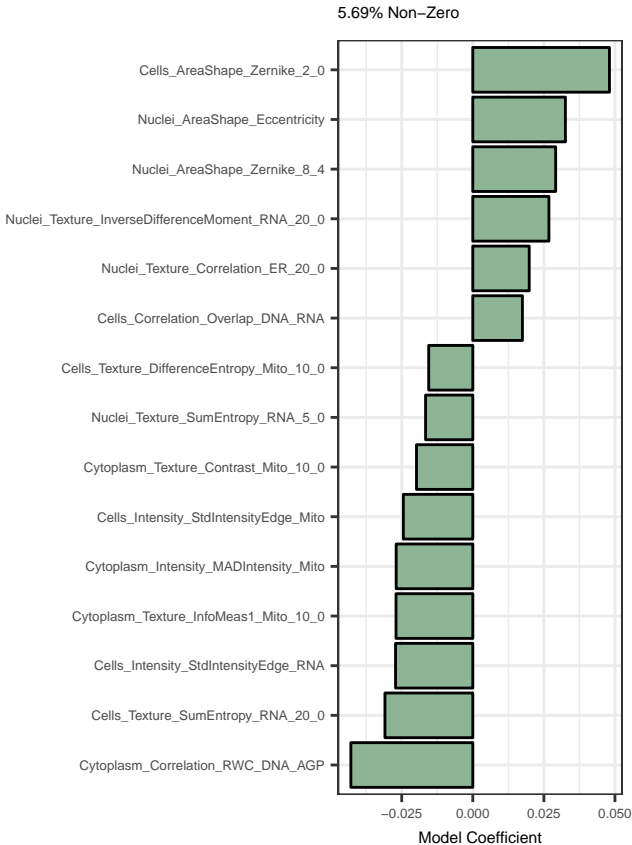
5.69% Non-Zero



cc_polyploid_high_h2ax

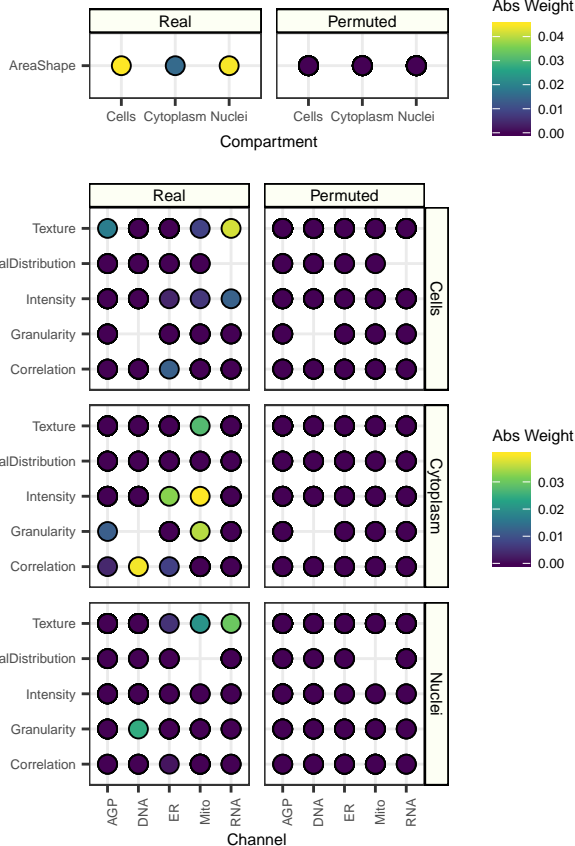
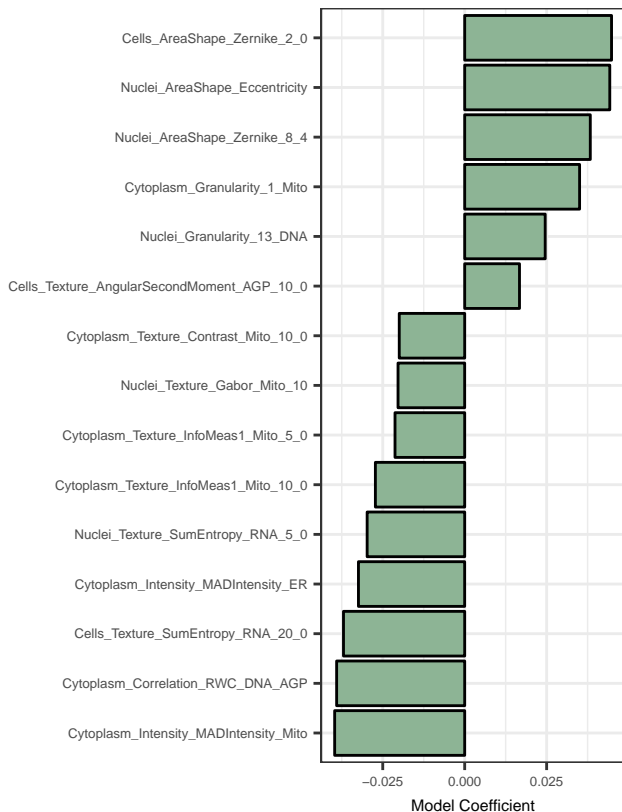


cc_polyloid_n_spots_h2ax_mean

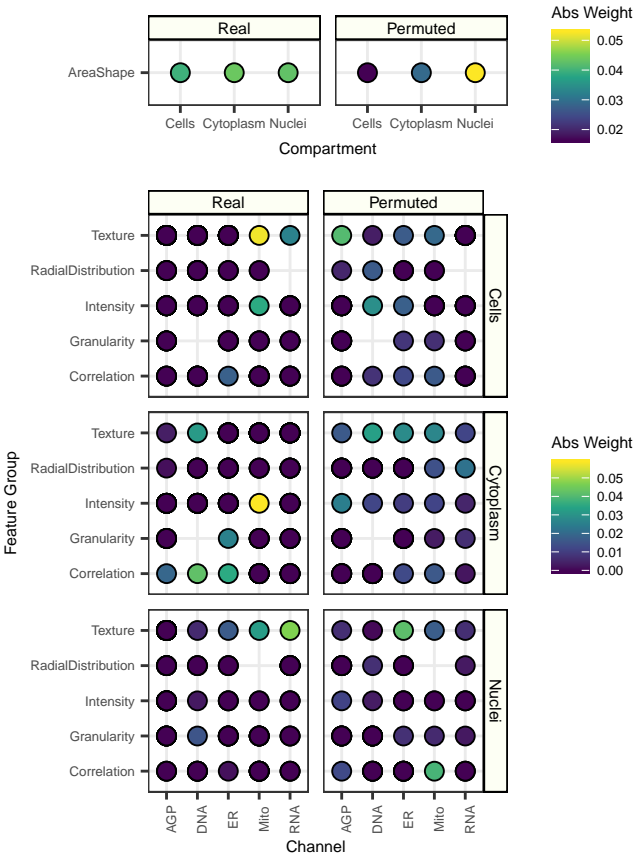


cc_polyploid_n_spots_h2ax_per_nucleus_area_mean

5.27% Non-Zero

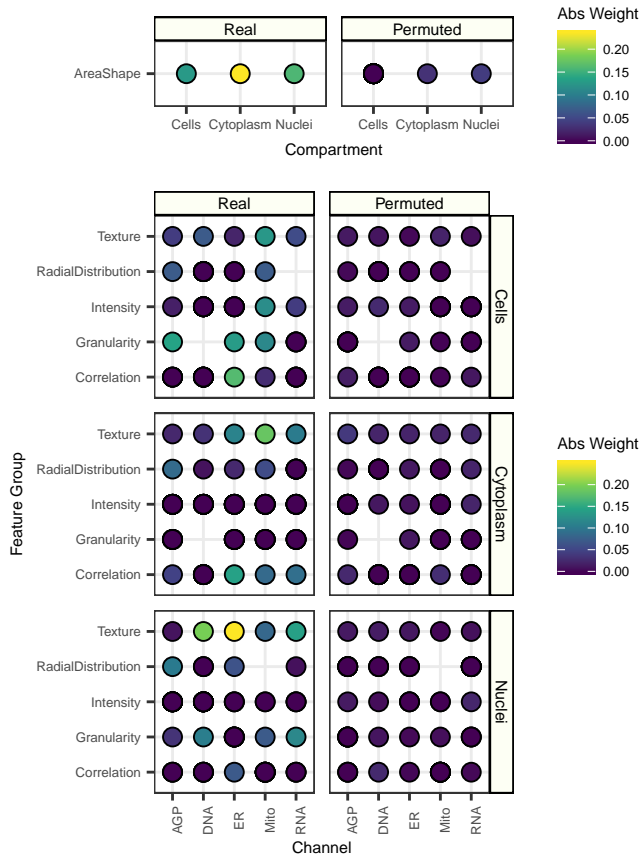
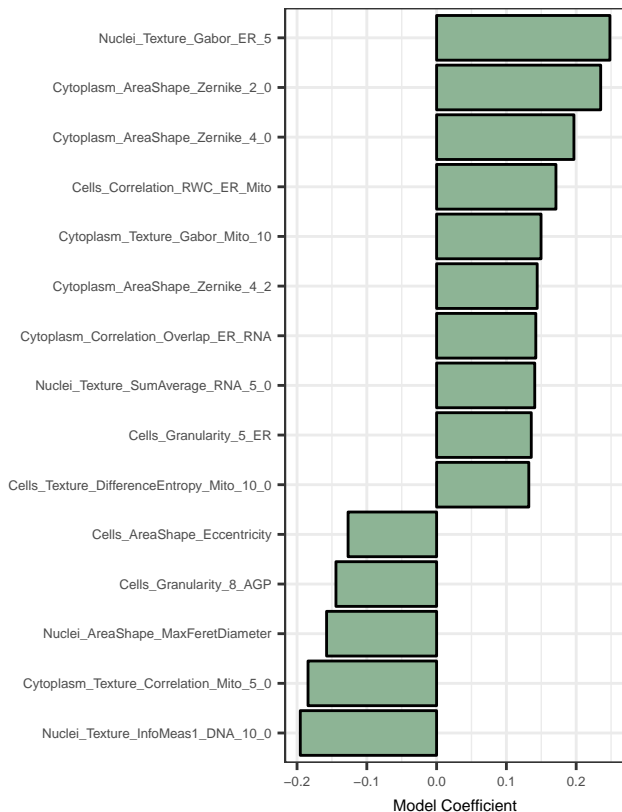


cc_s_high_h2ax



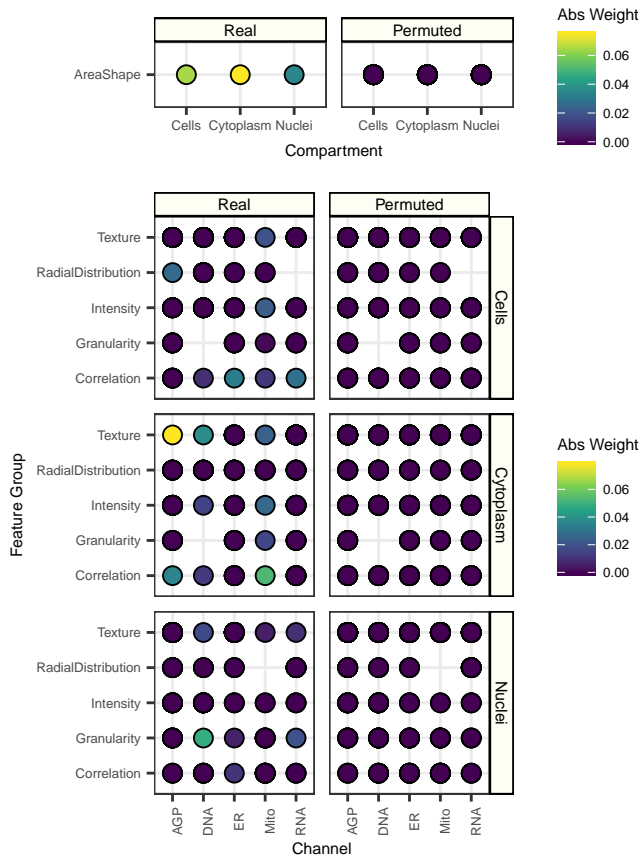
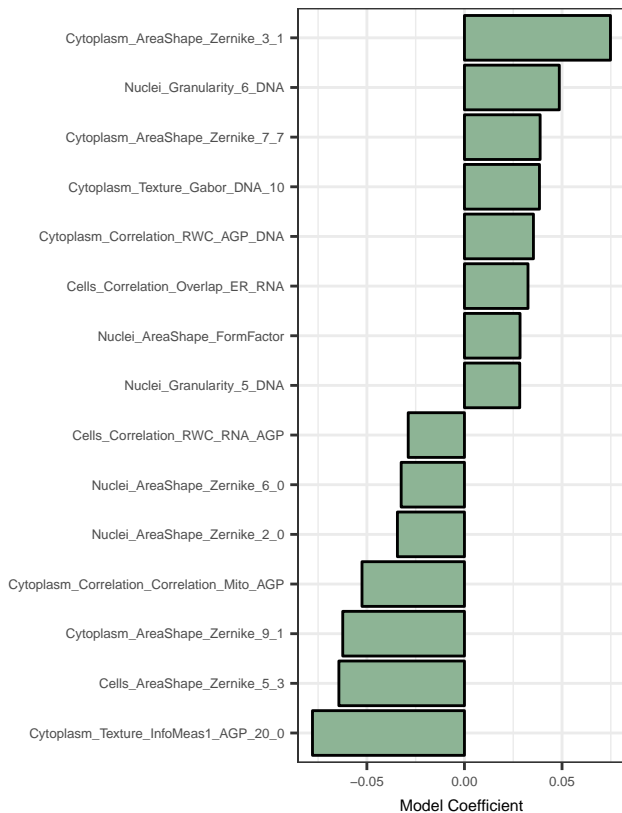
cc_s_intensity_nucleus_area_mean

11.28% Non-Zero



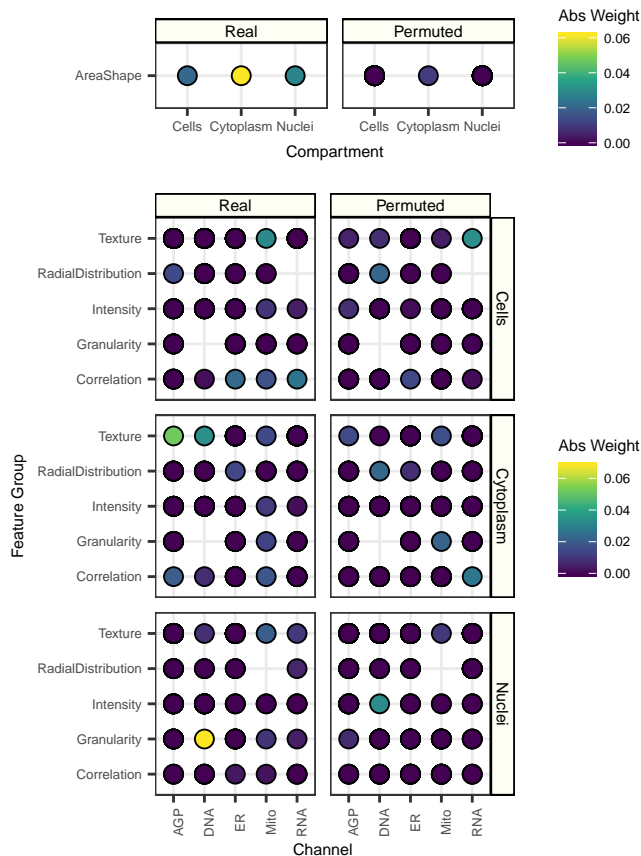
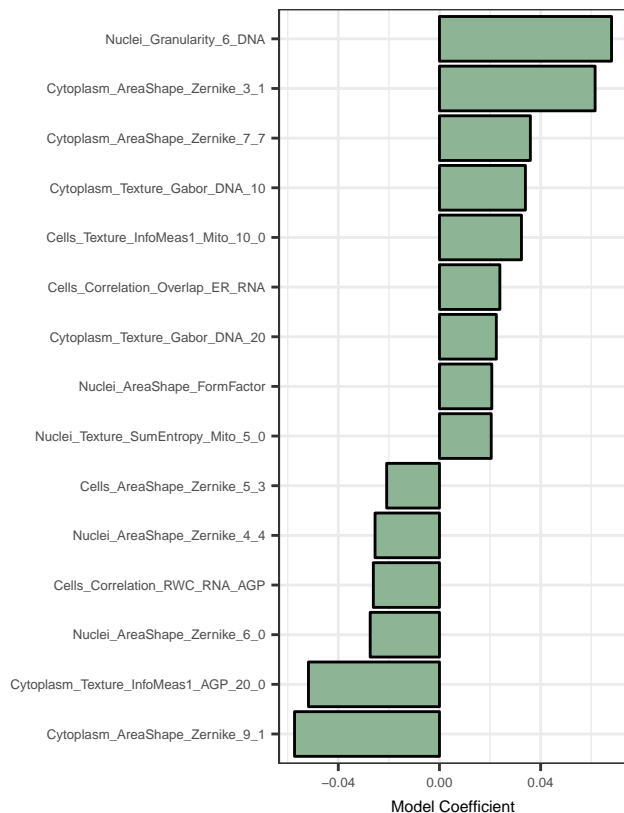
cc_s_intensity_nucleus_area_sum

7.9% Non-Zero



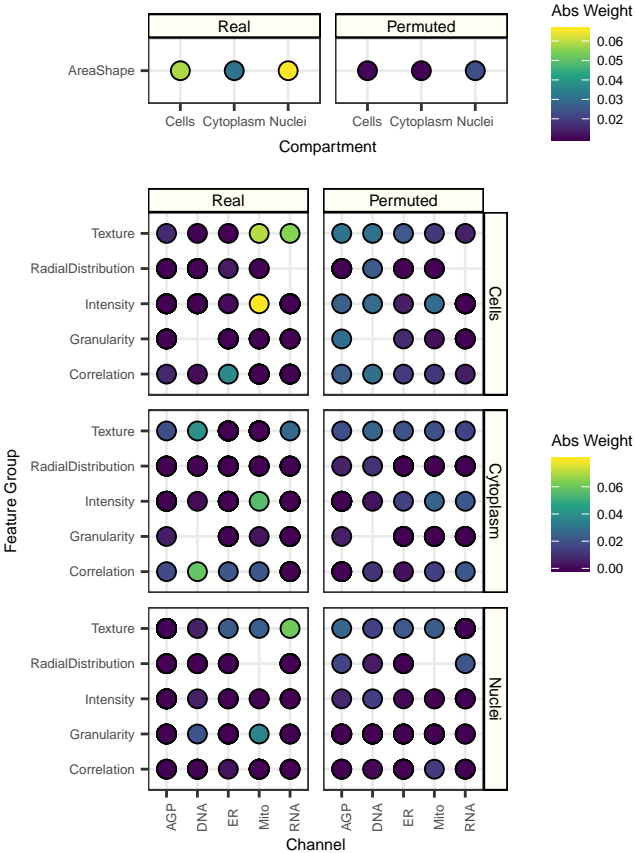
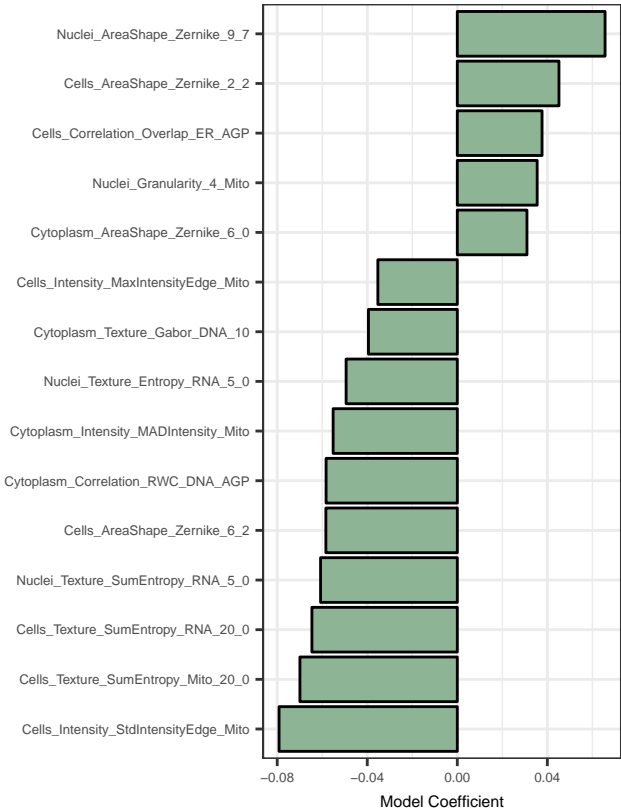
cc_s_n_objects

8.85% Non-Zero



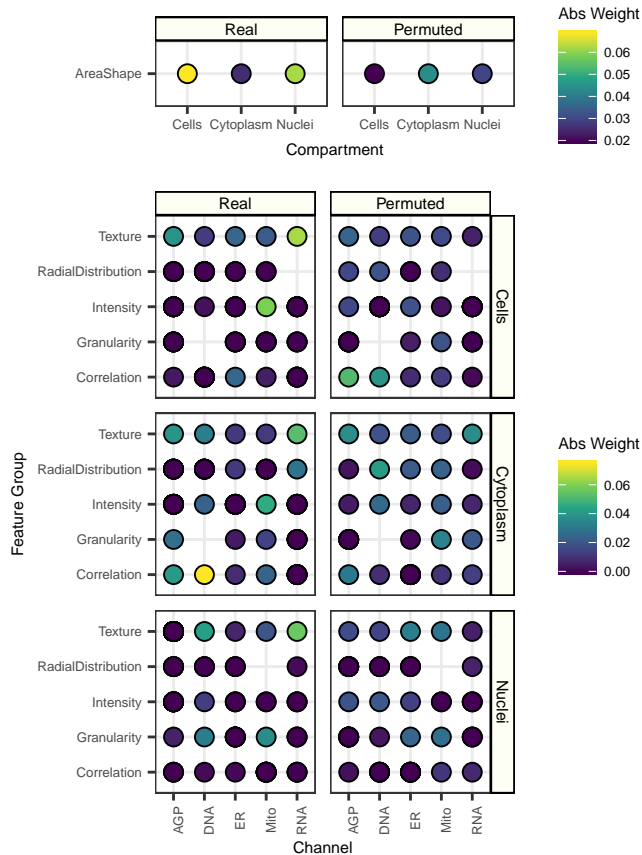
cc_s_n_spots_h2ax_mean

7.9% Non-Zero

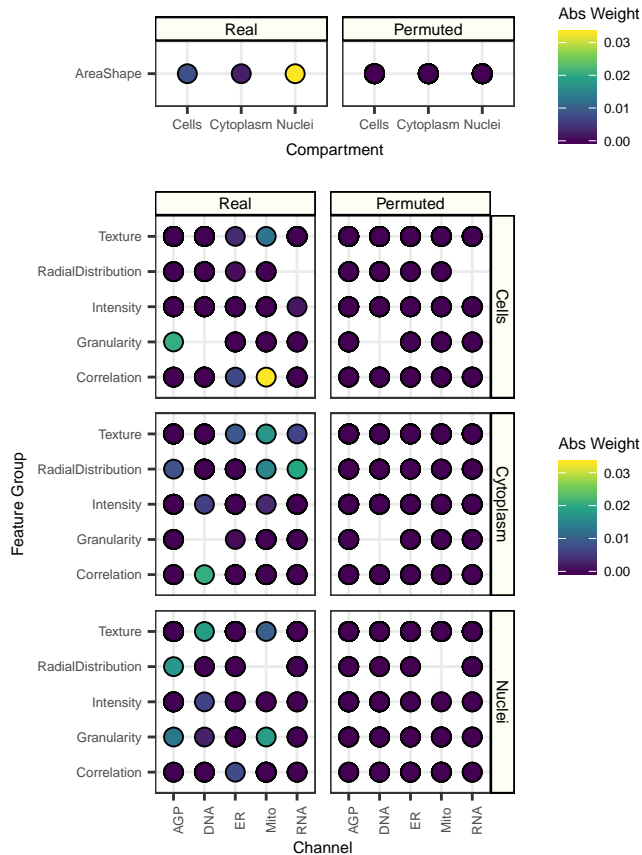
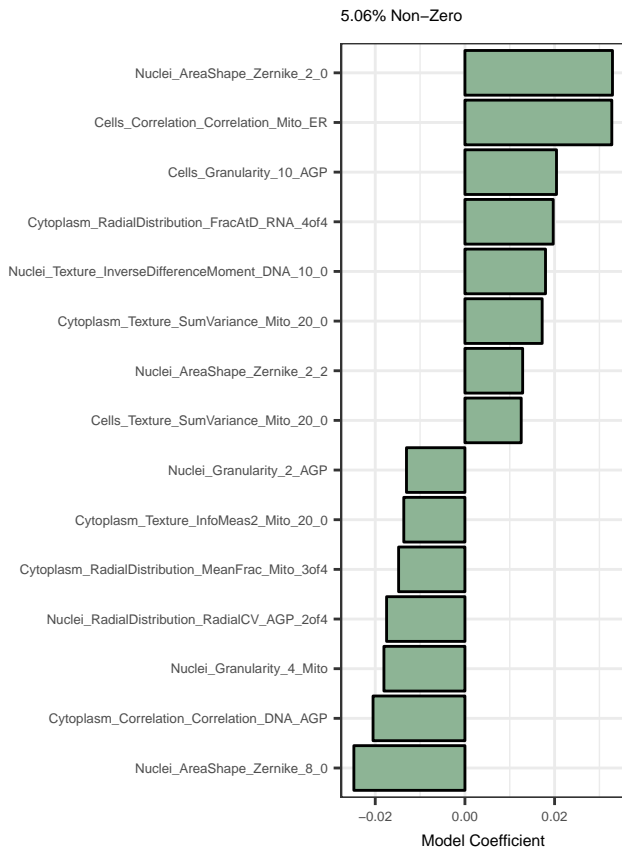


cc_s_n_spots_h2ax_per_nucleus_area_mean

10.12% Non-Zero

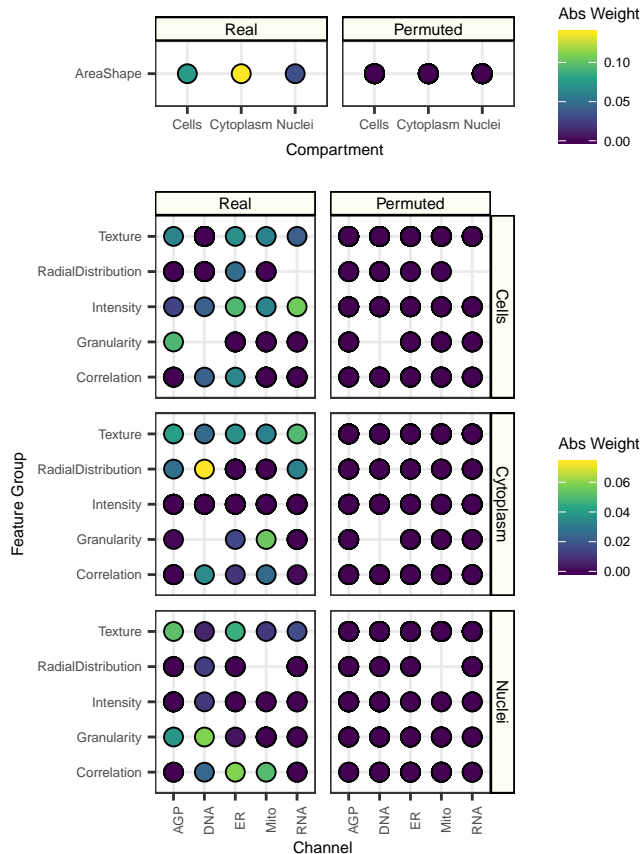
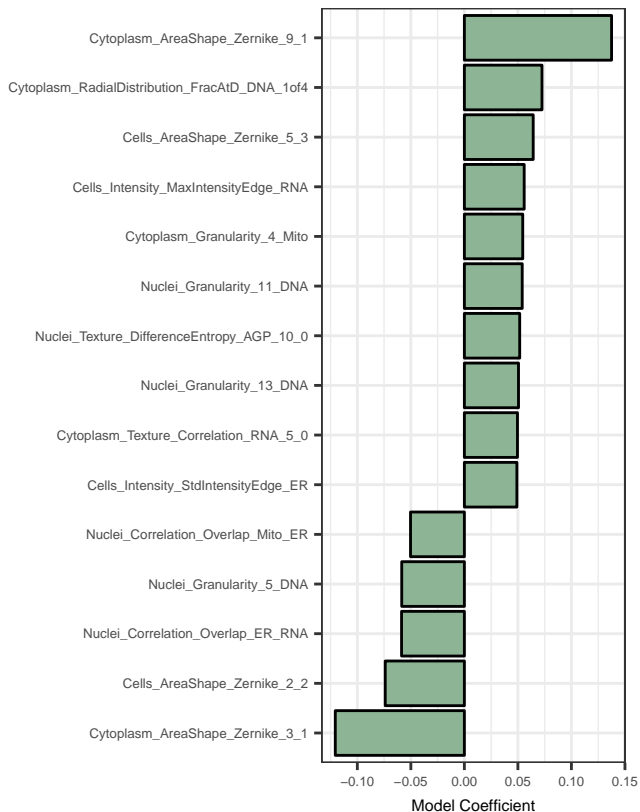


vb_infection_percentage



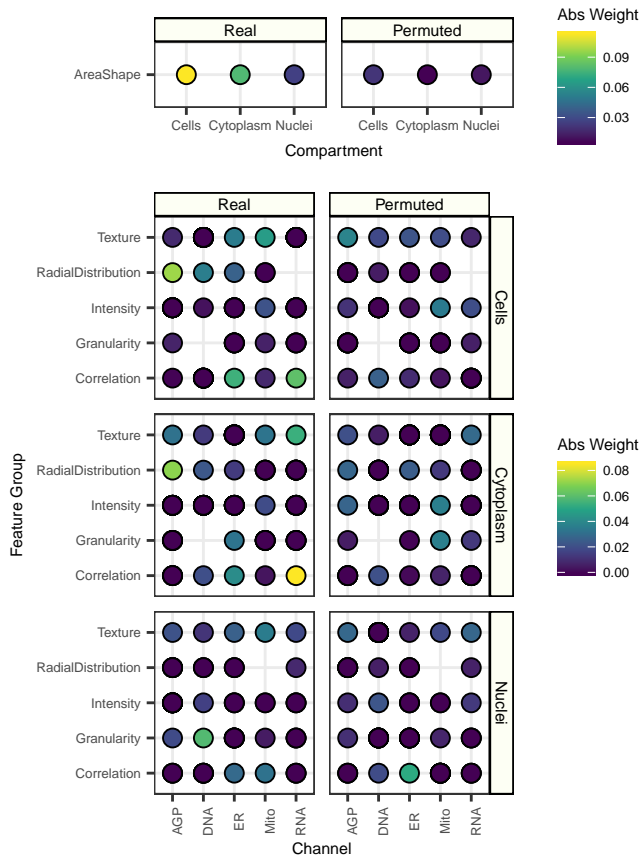
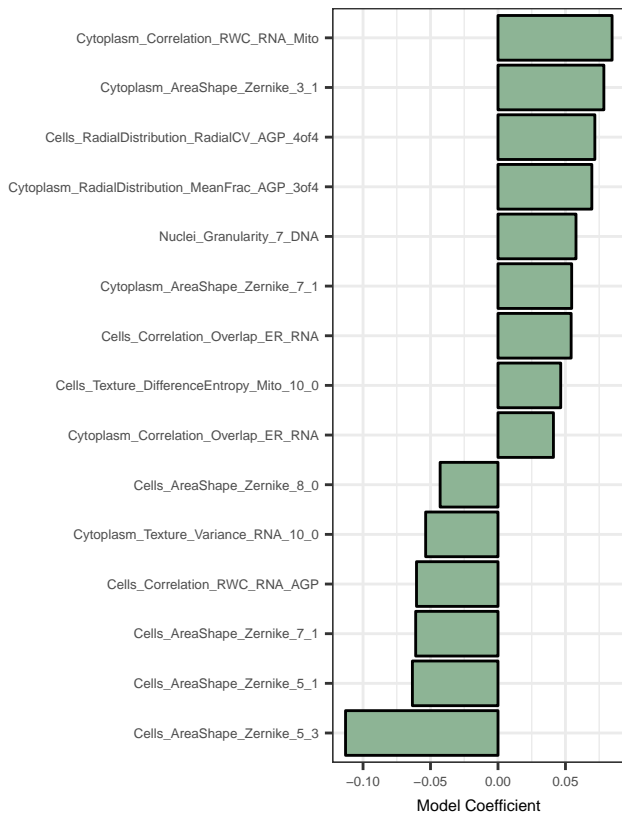
vb_live_cell_area

13.17% Non-Zero

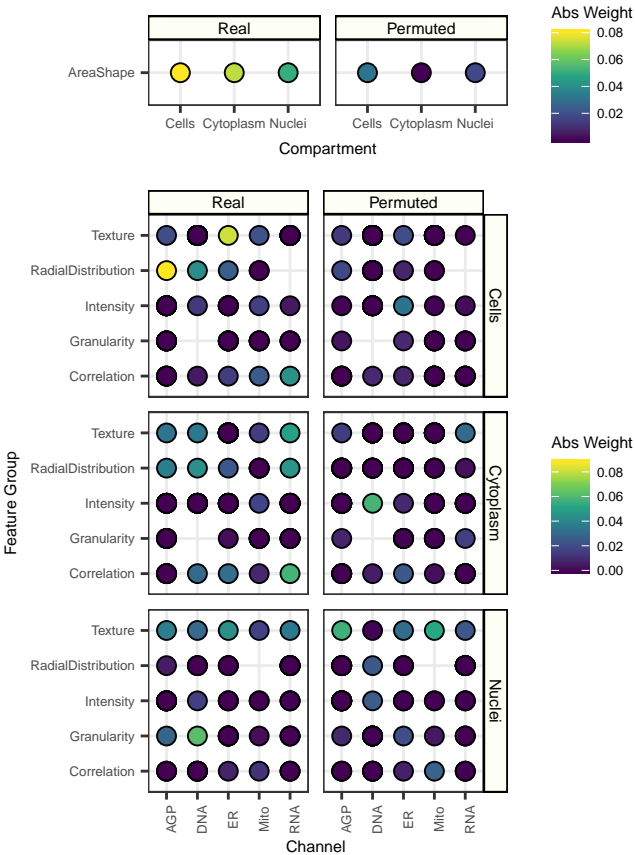
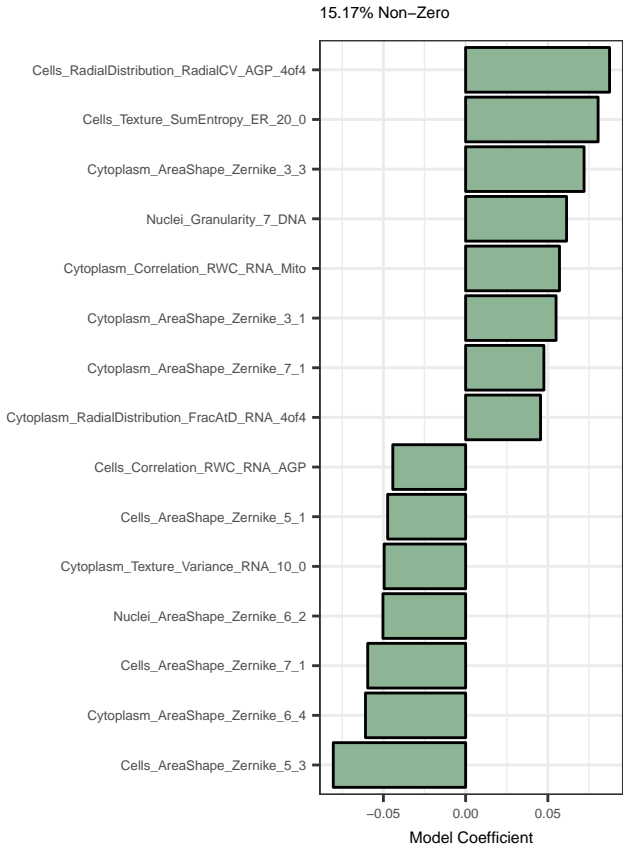


vb_live_cell_roundness

14.33% Non-Zero

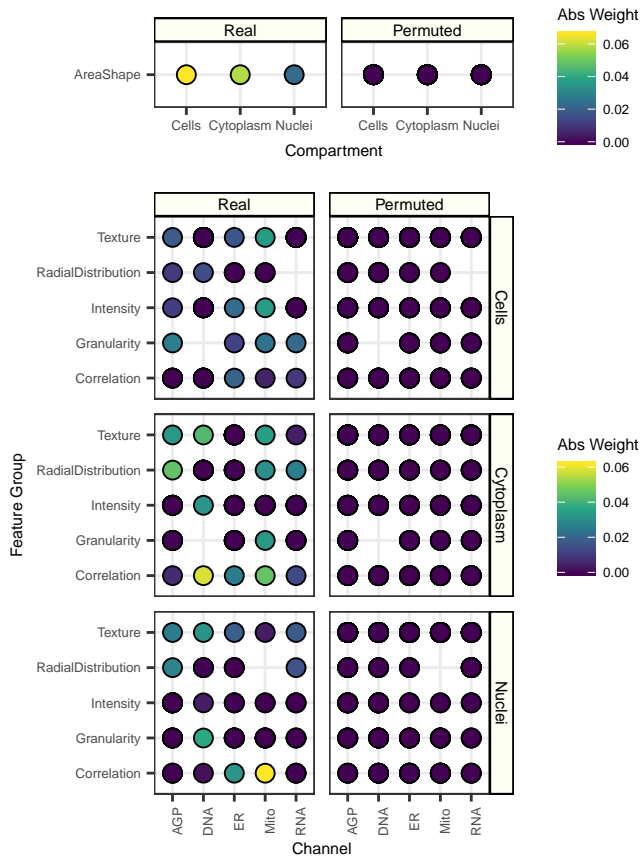
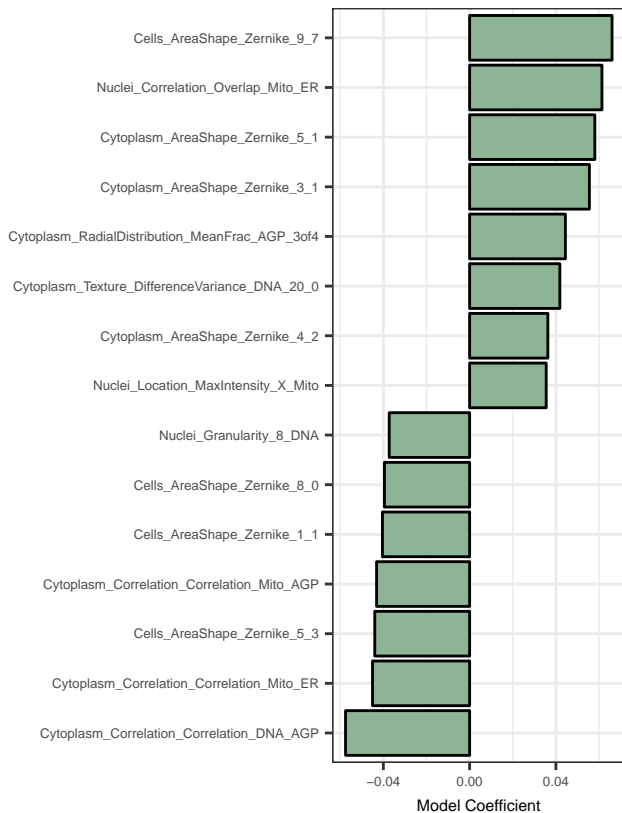


vb_live_cell_width_length

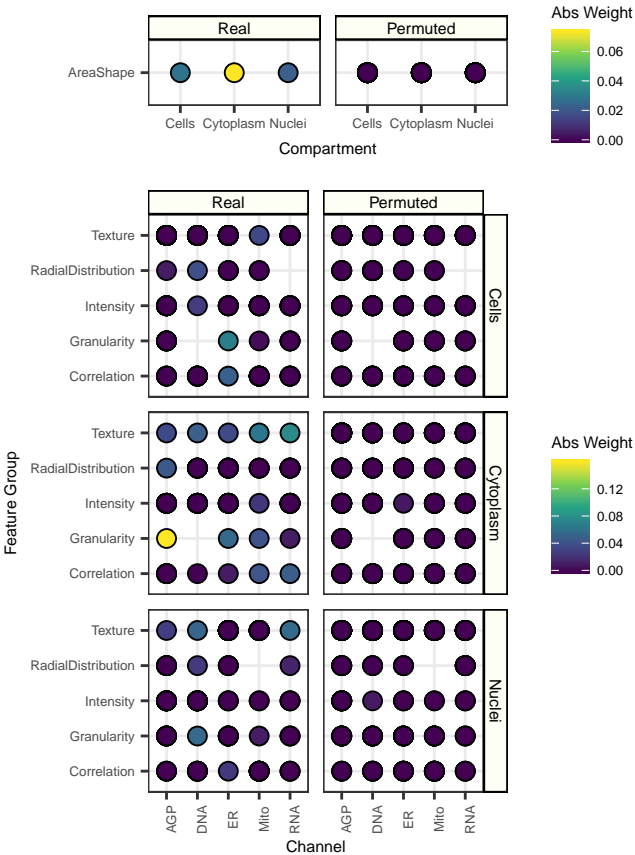
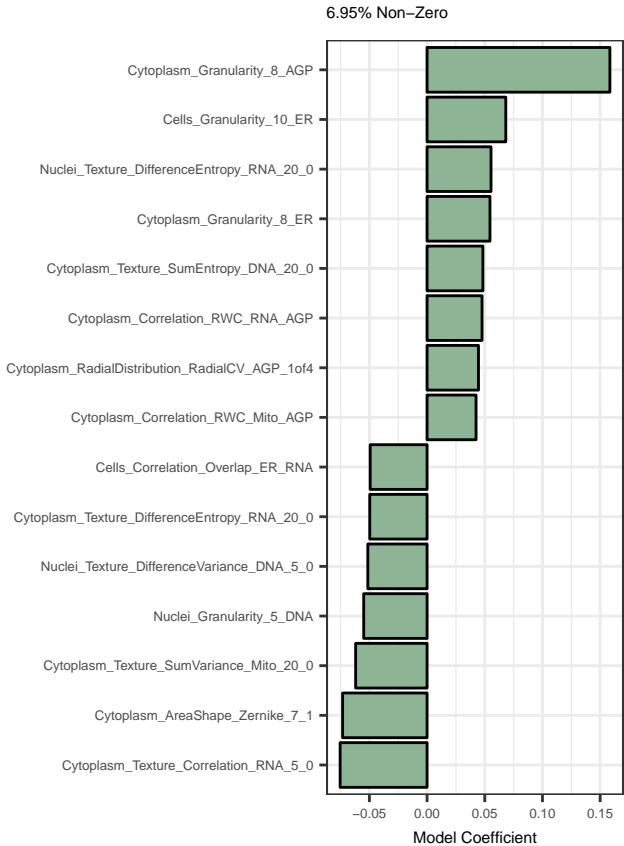


vb_num_live_cells

12.96% Non-Zero

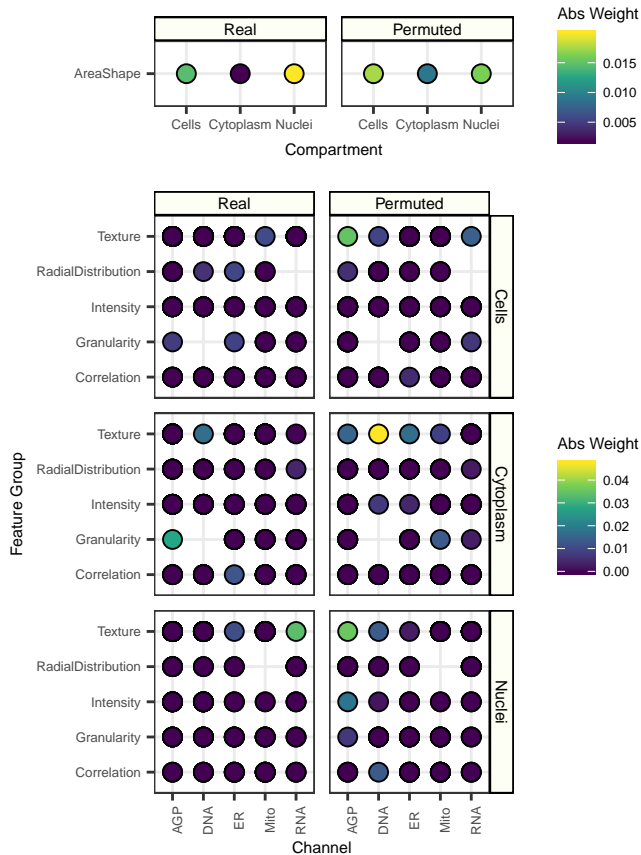
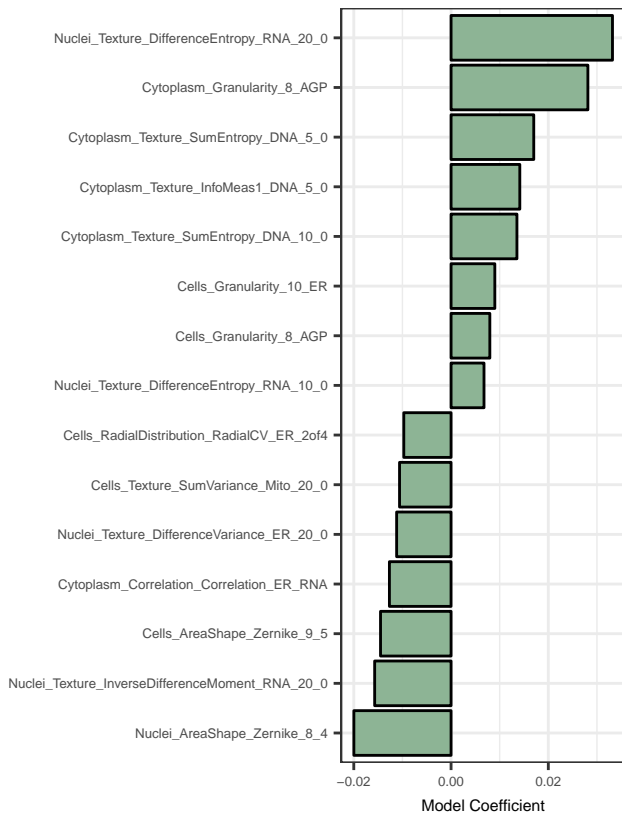


vb_percent_all_apoptosis



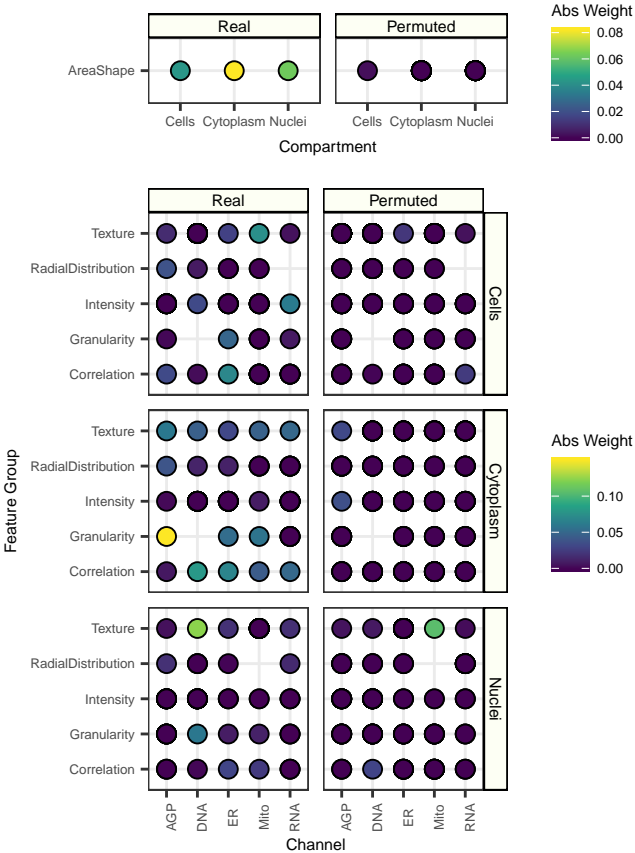
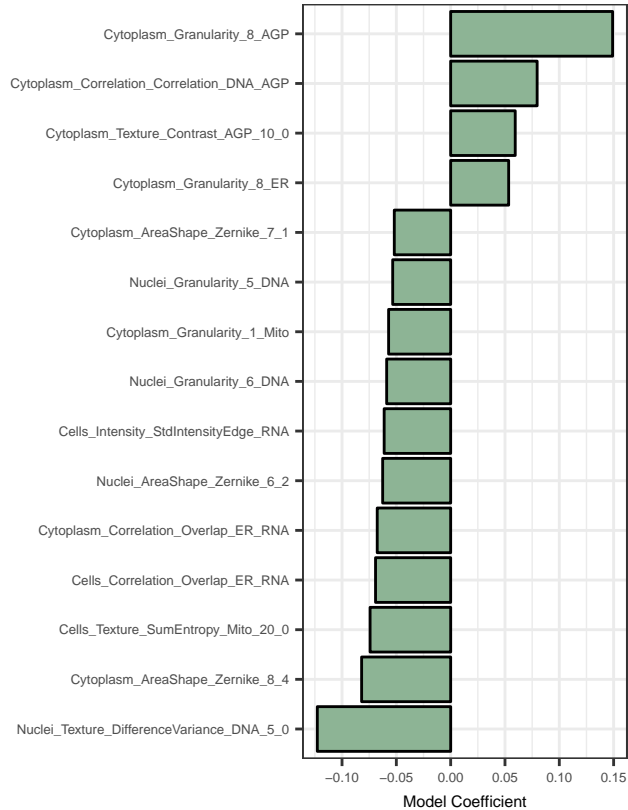
vb_percent_caspase_dead_only

2.63% Non-Zero



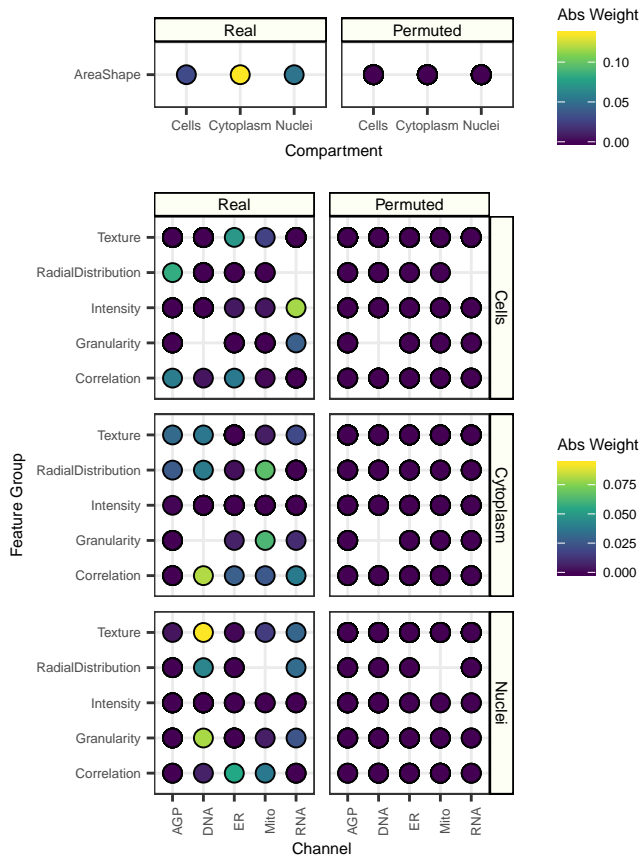
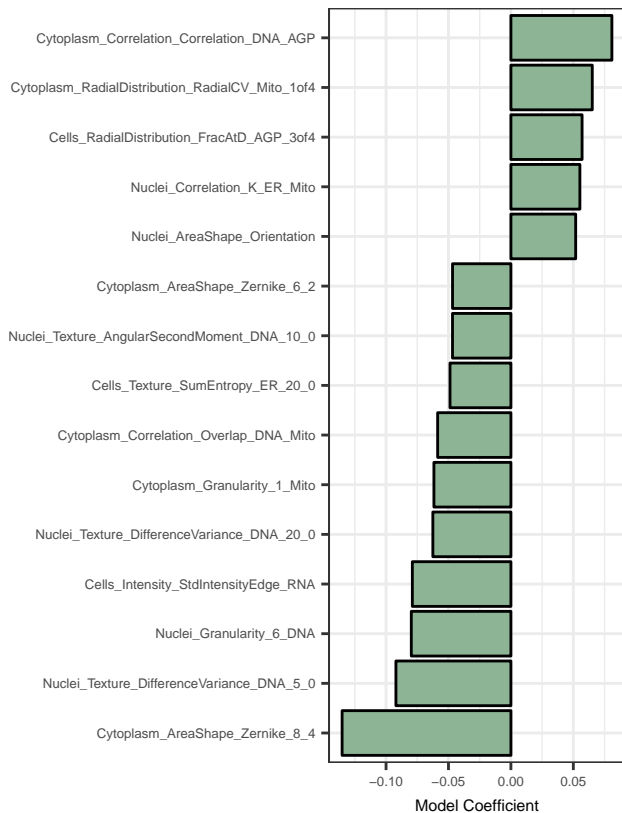
vb_percent_dead

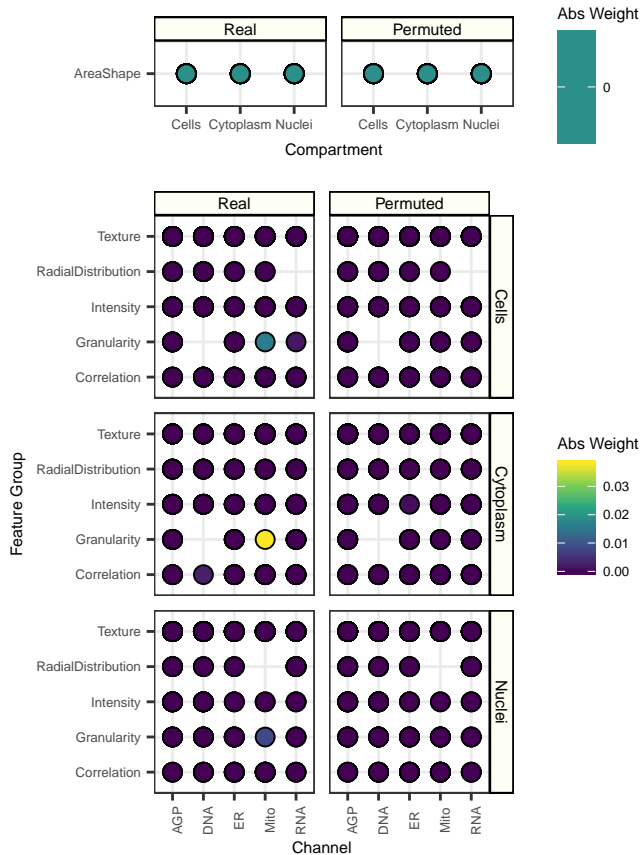
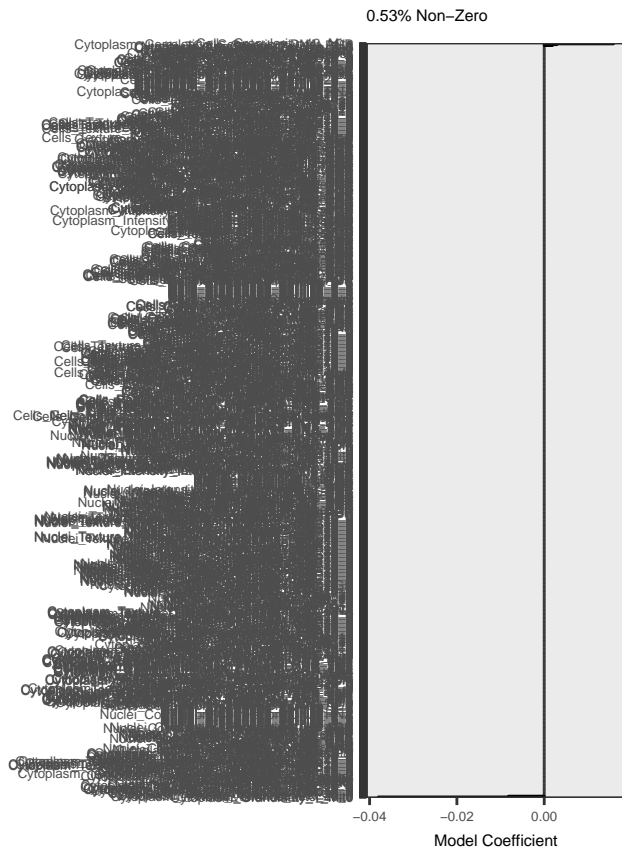
12.43% Non-Zero



vb_percent_dead_only

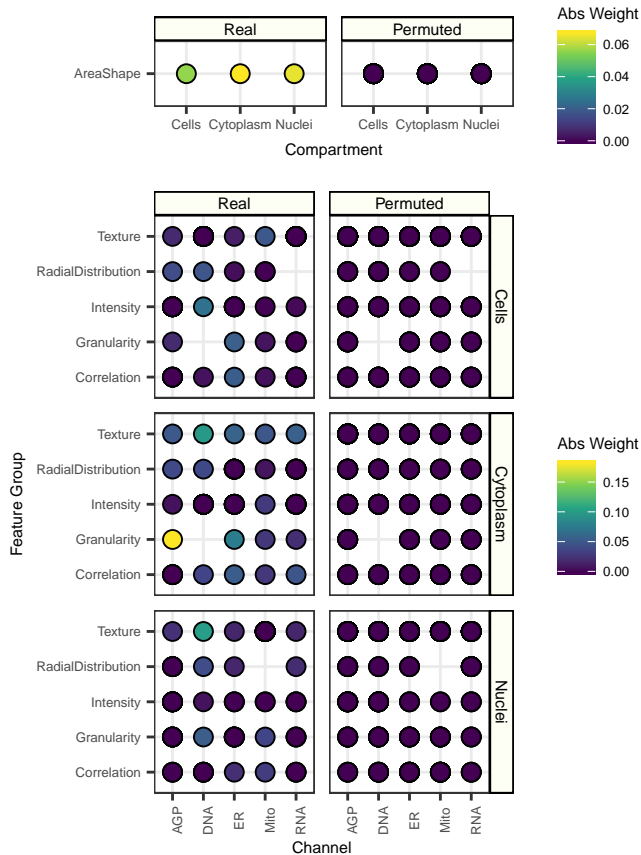
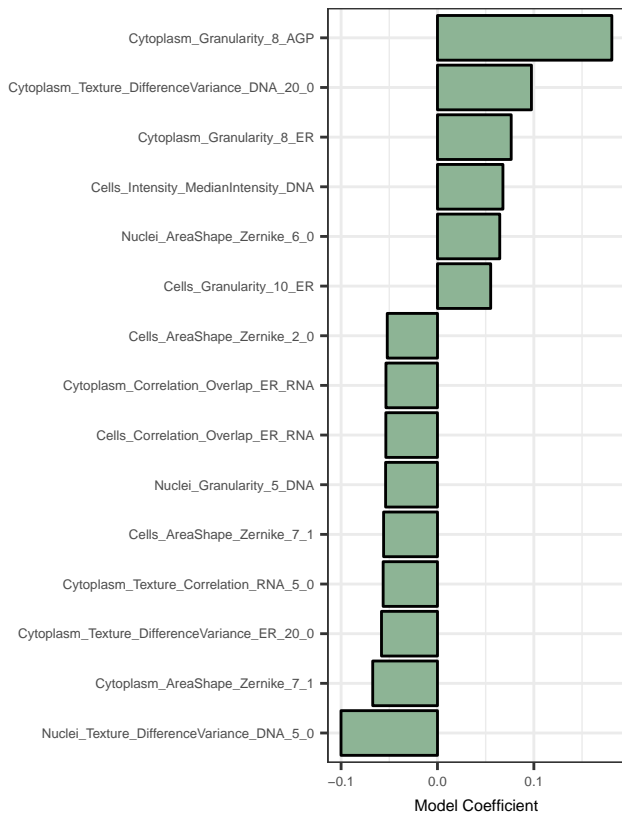
11.59% Non-Zero



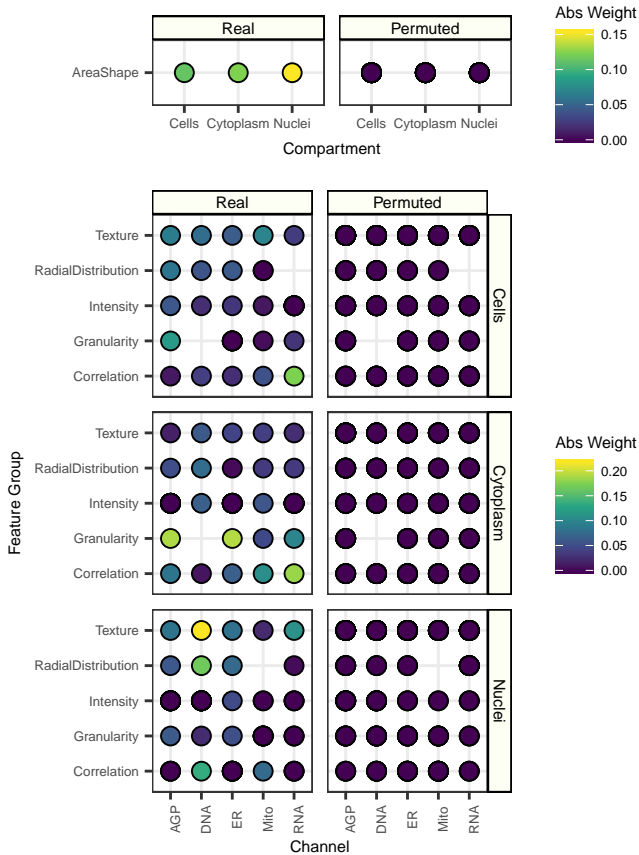
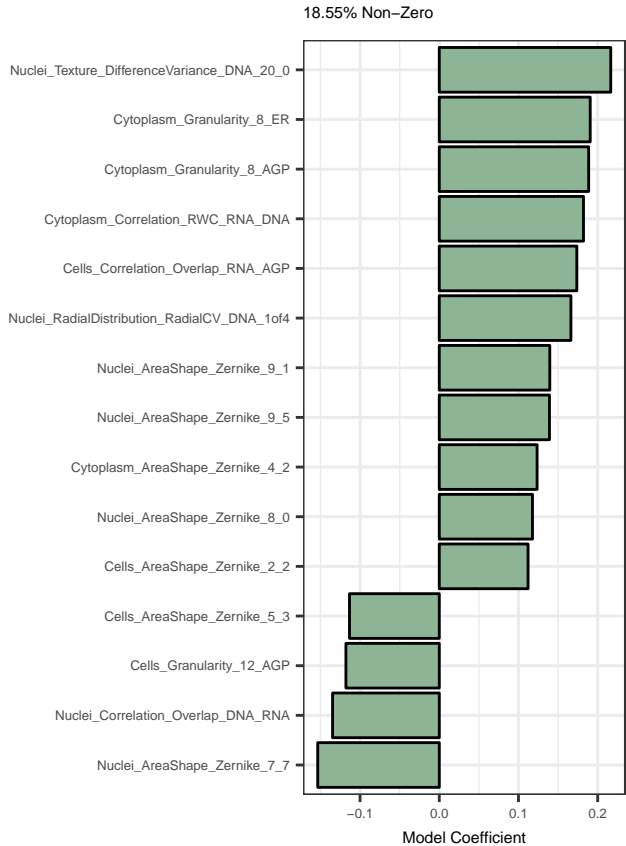
vb_percent_early_apoptosis

vb_percent_late_apoptosis

13.28% Non-Zero



vb_ros_back_mean



vb_ros_mean

