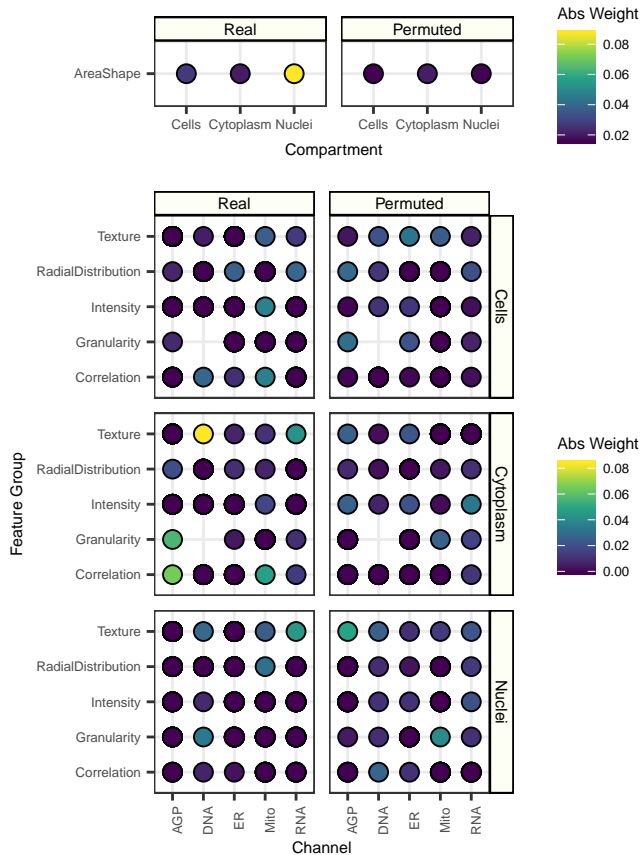
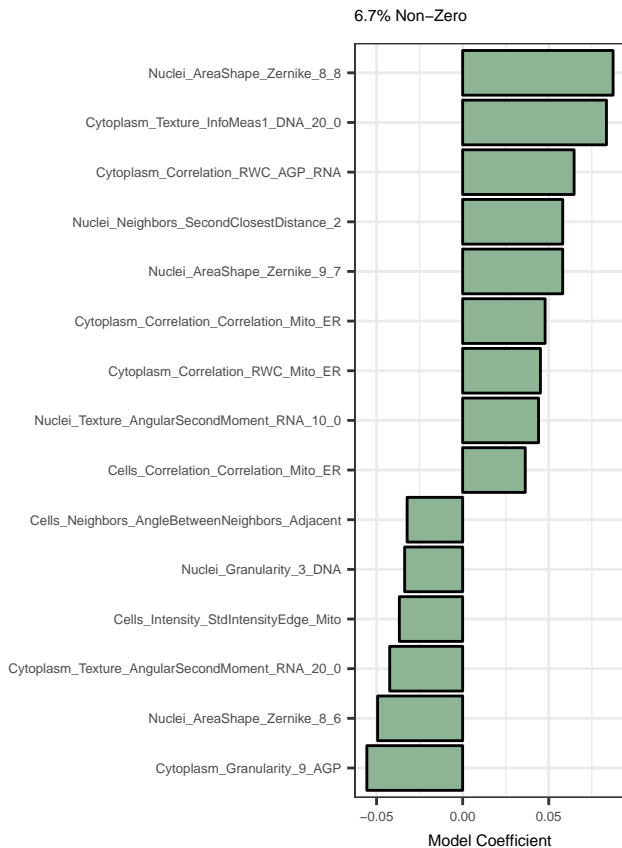
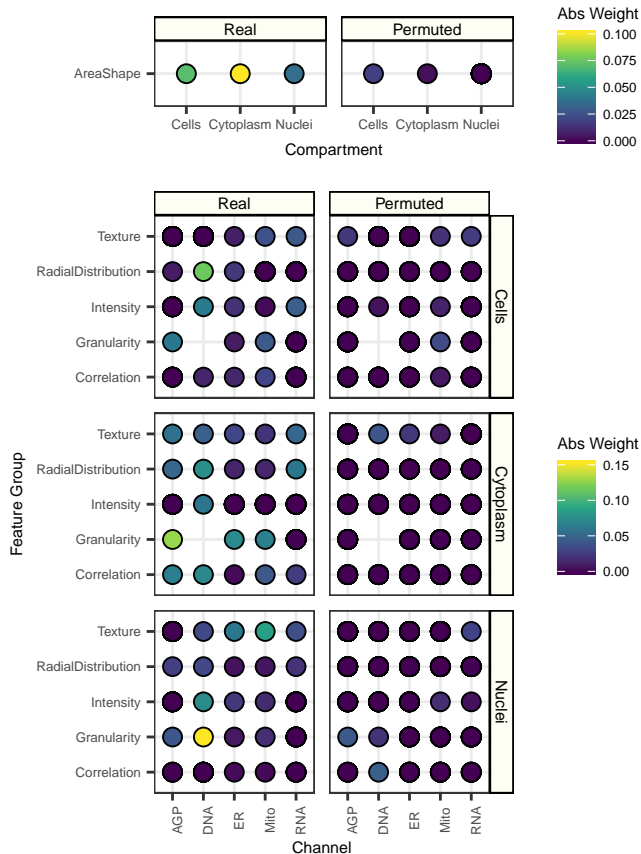
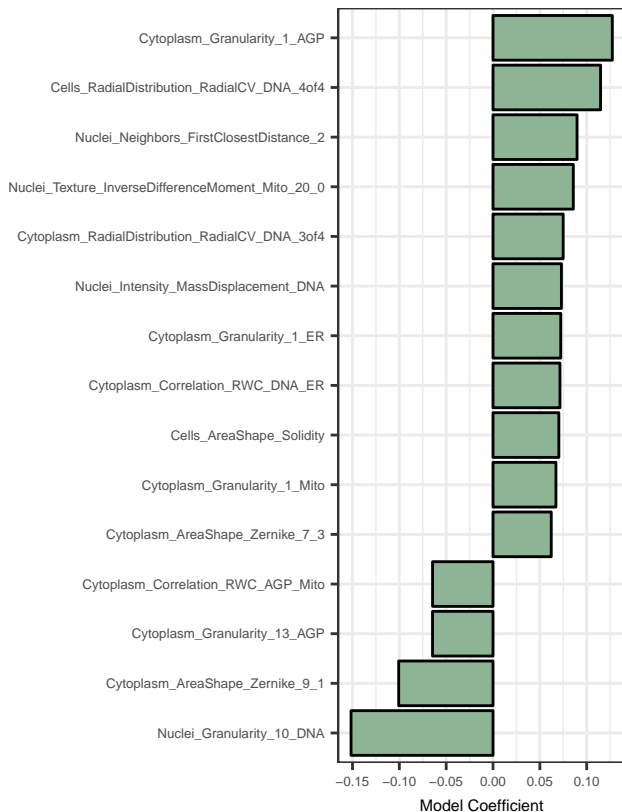


cc_all_high_n_spots_h2ax_mean



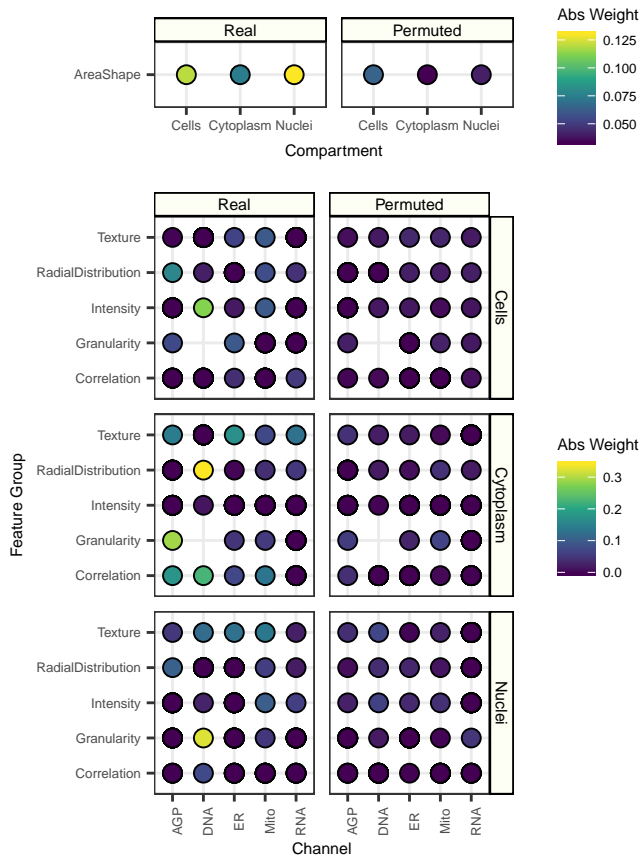
cc_all_large_notround_polynuclear_mean

14.33% Non-Zero



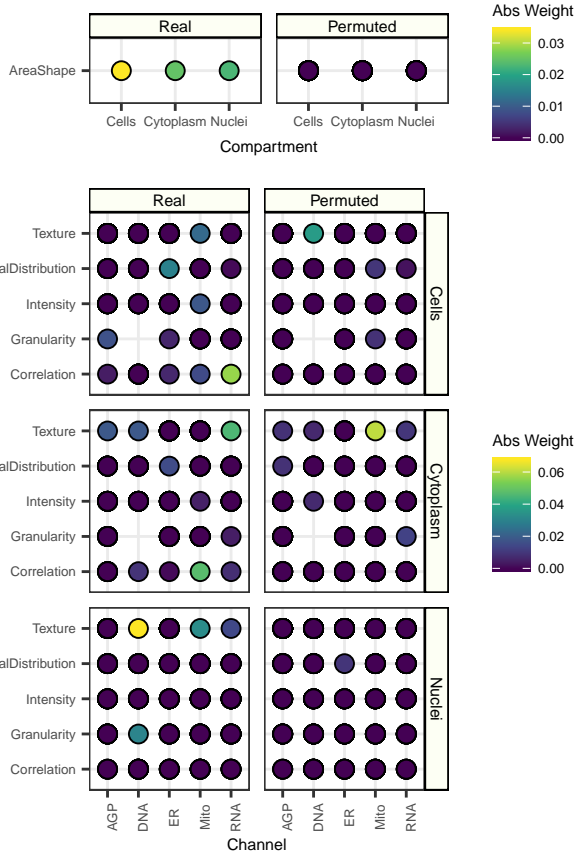
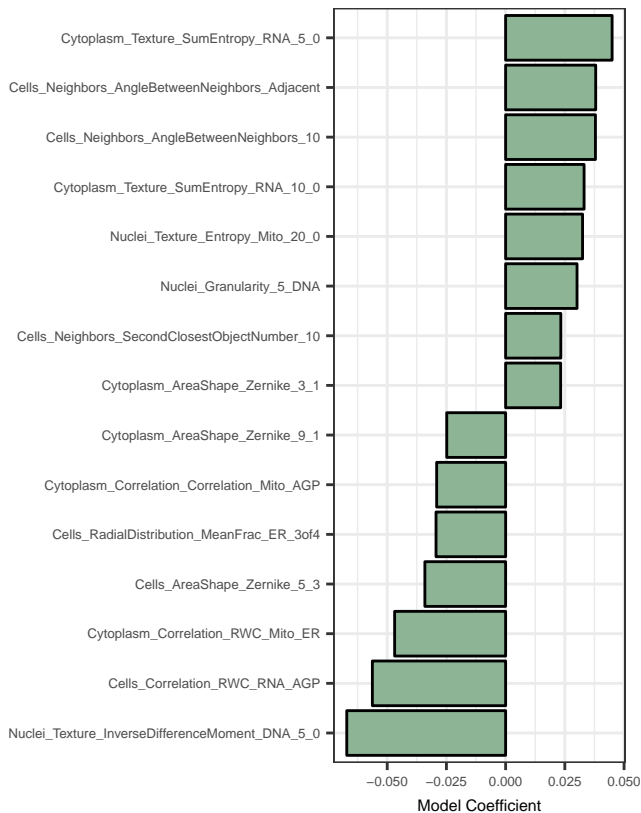
cc_all_large_round_polyploid_mean

7.07% Non-Zero

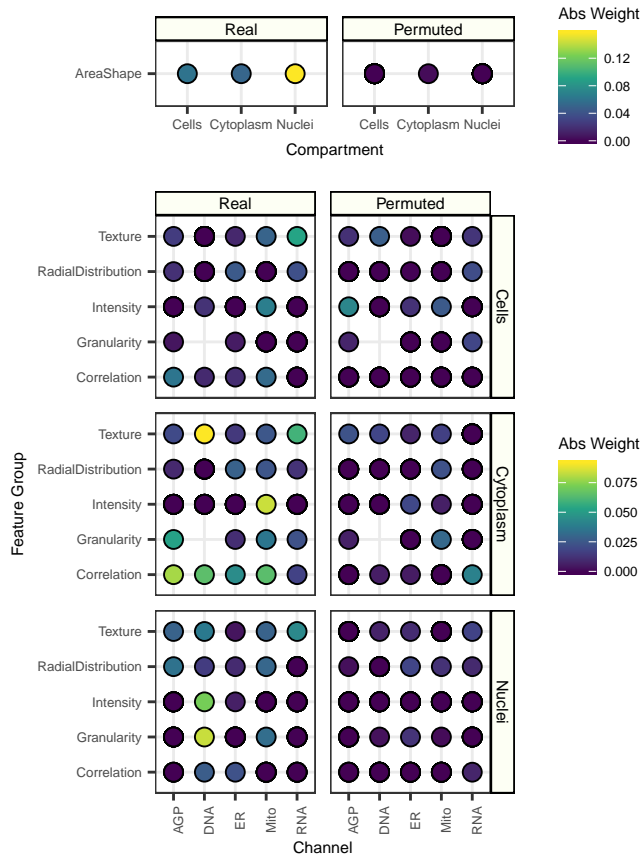
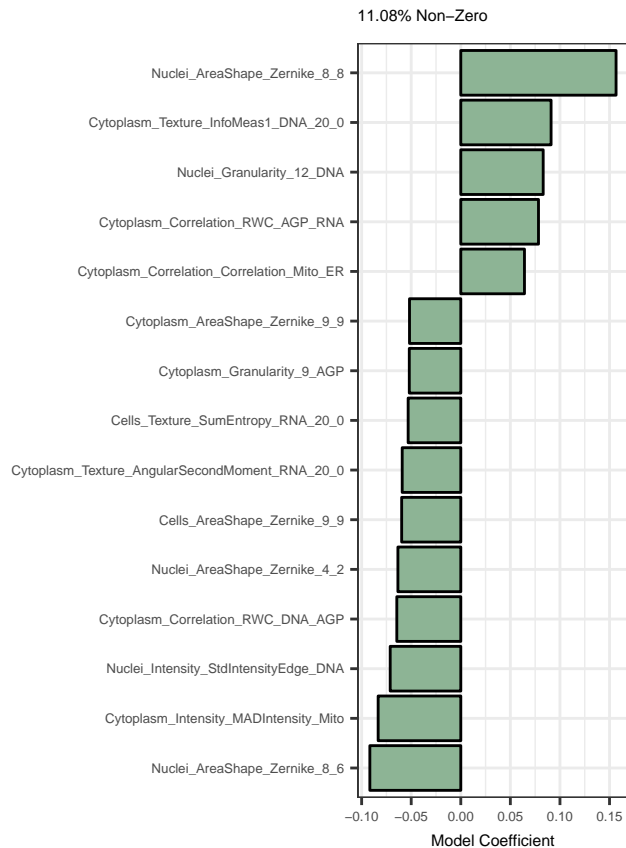


cc_all_n_objects

4.69% Non-Zero

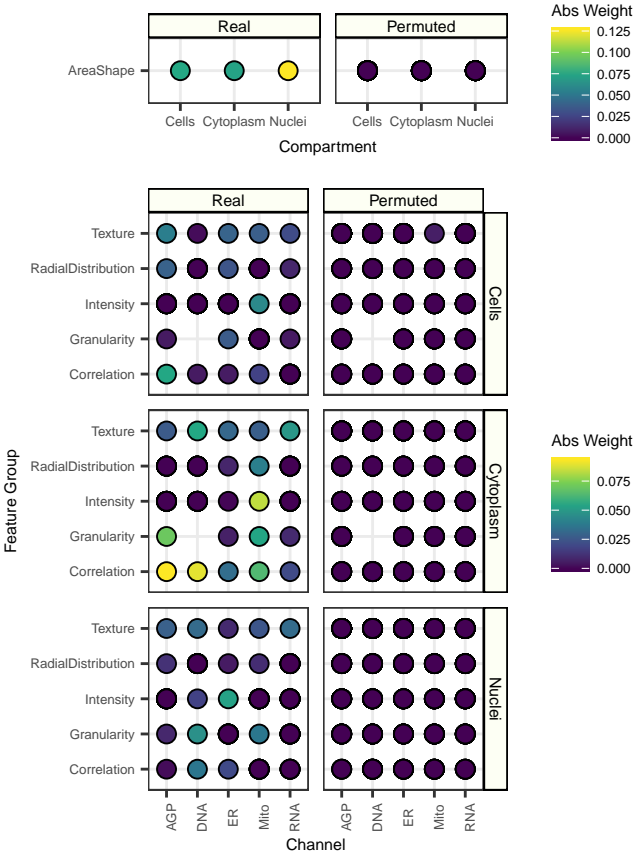


cc_all_n_spots_mean



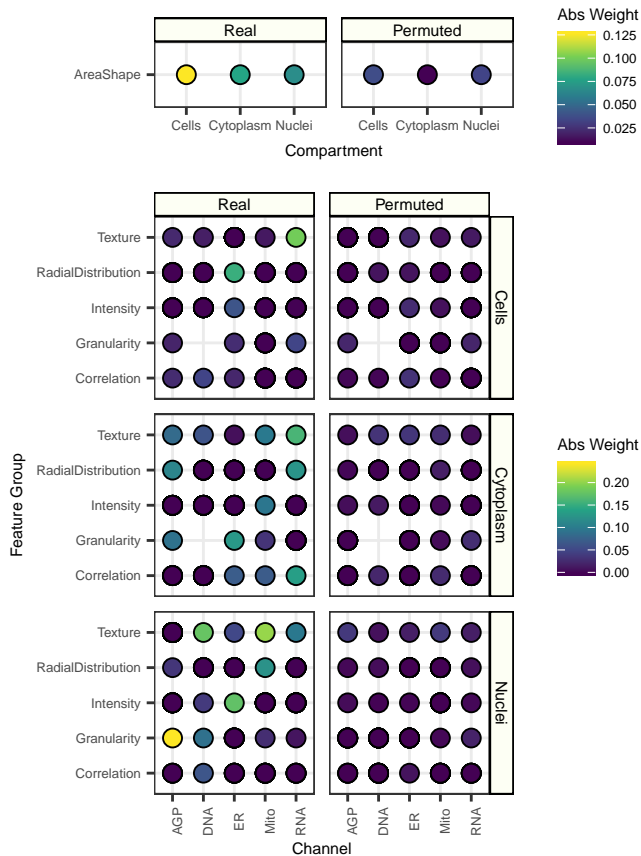
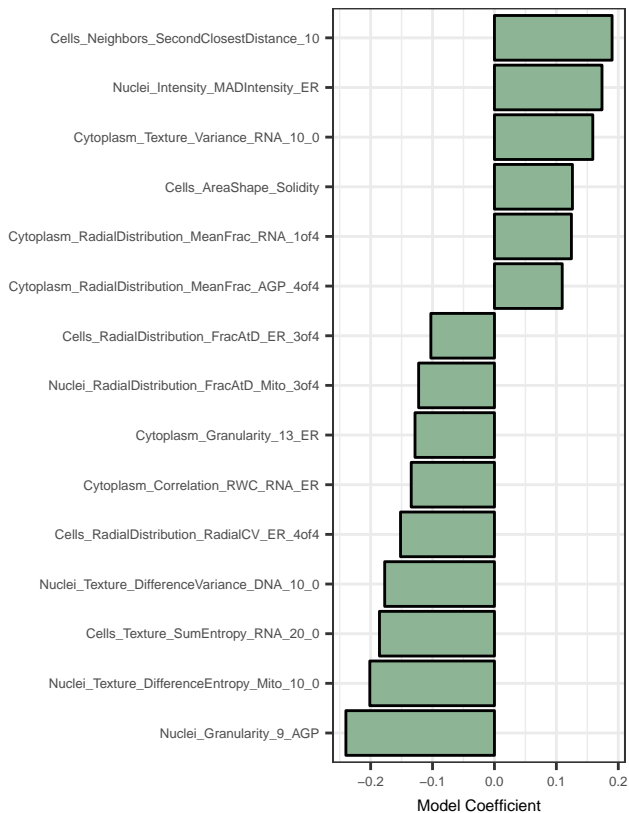
cc_all_n_spots_per_nucleus_area_mean

10.83% Non-Zero

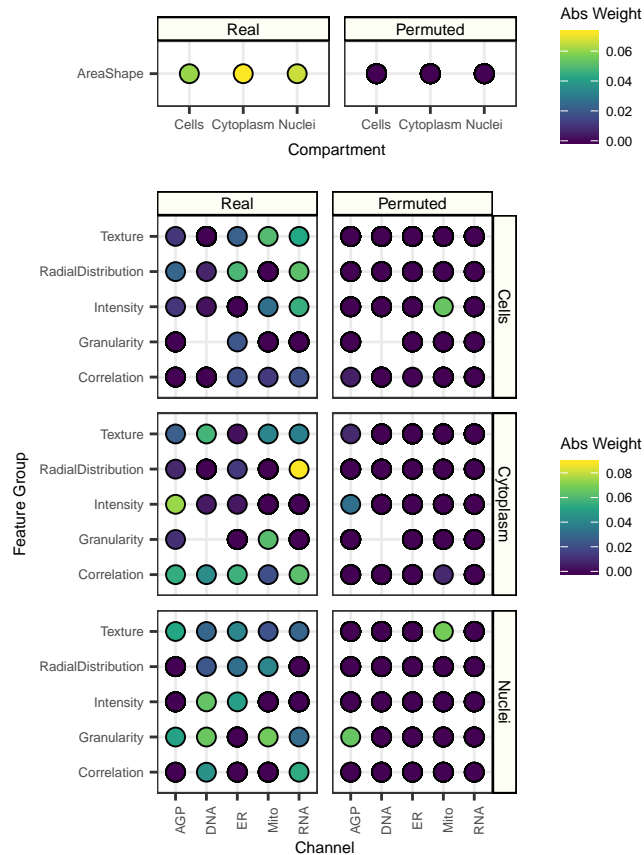
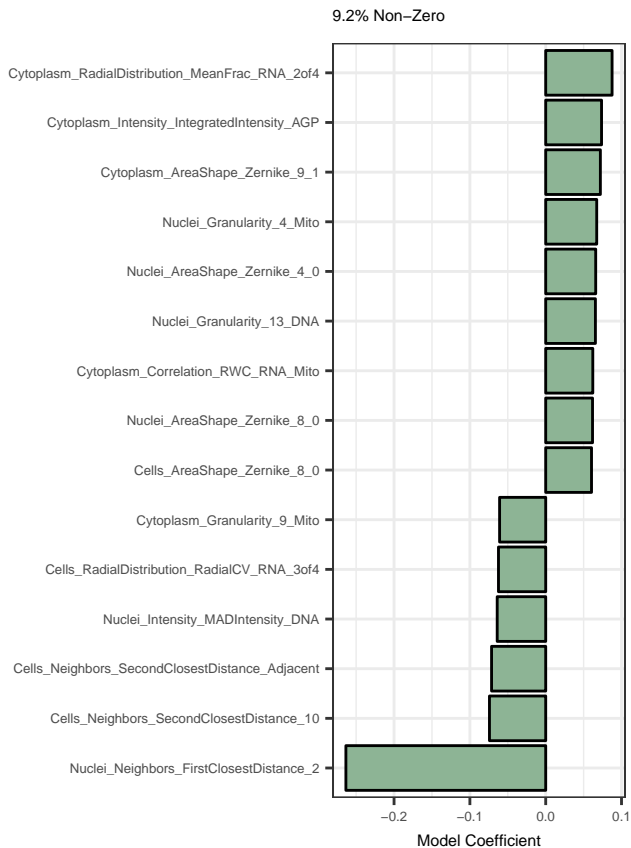


cc_all_nucleus_area_mean

4.51% Non-Zero

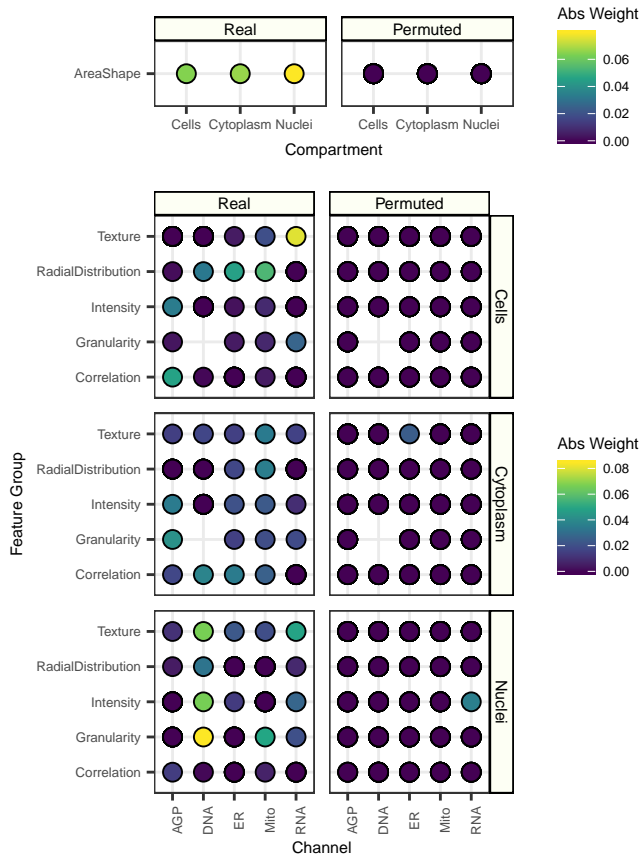


cc_all_nucleus_roundness_mean



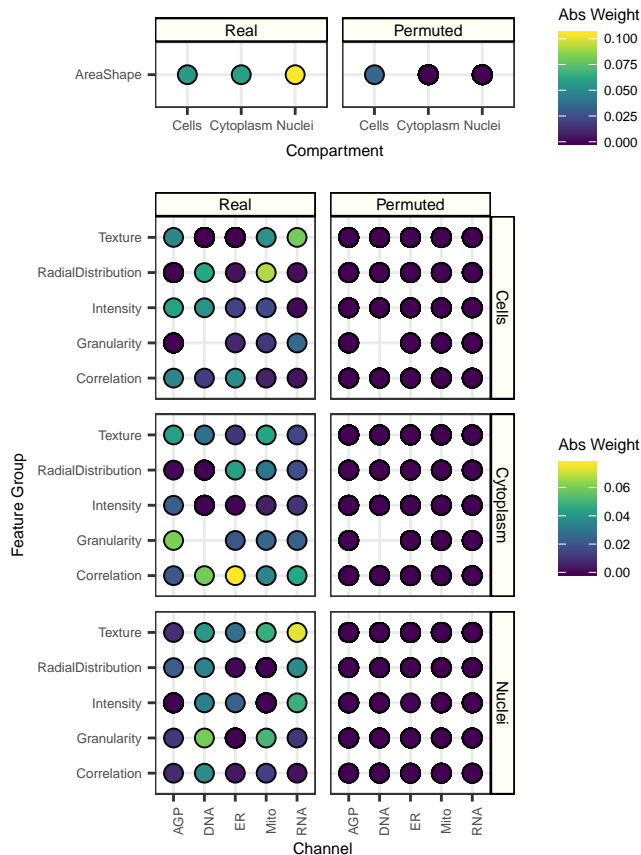
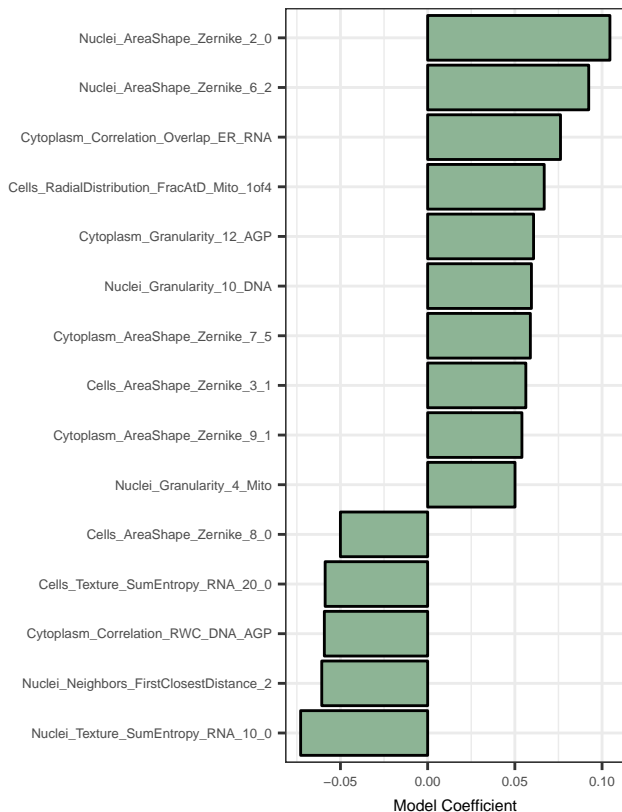
cc_cc_edu_pos_mean

10.26% Non-Zero

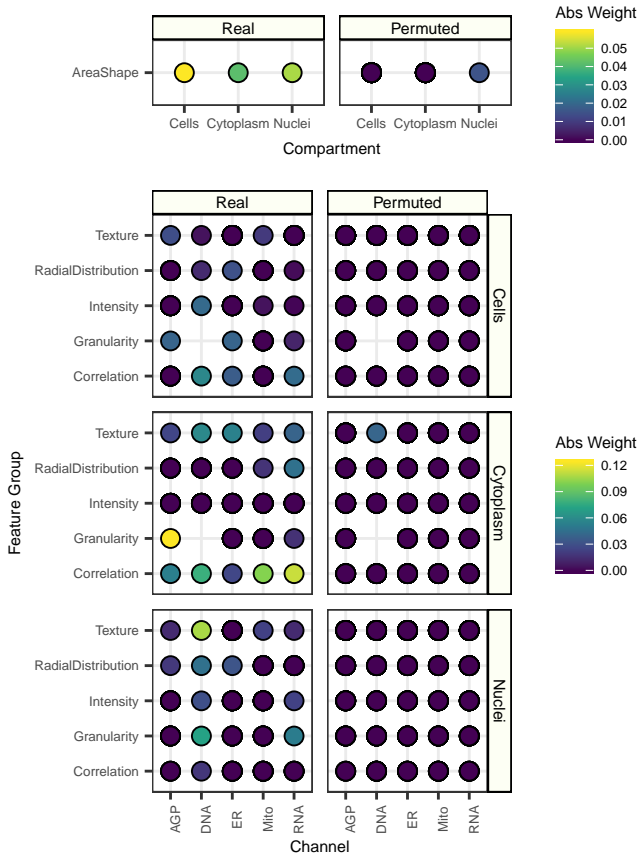
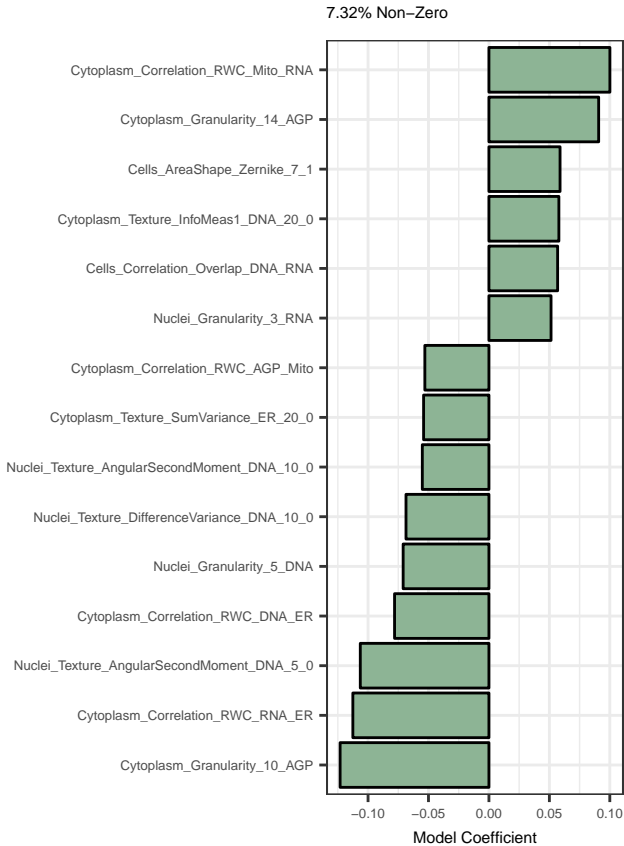


cc_cc_g1_mean

14.21% Non-Zero

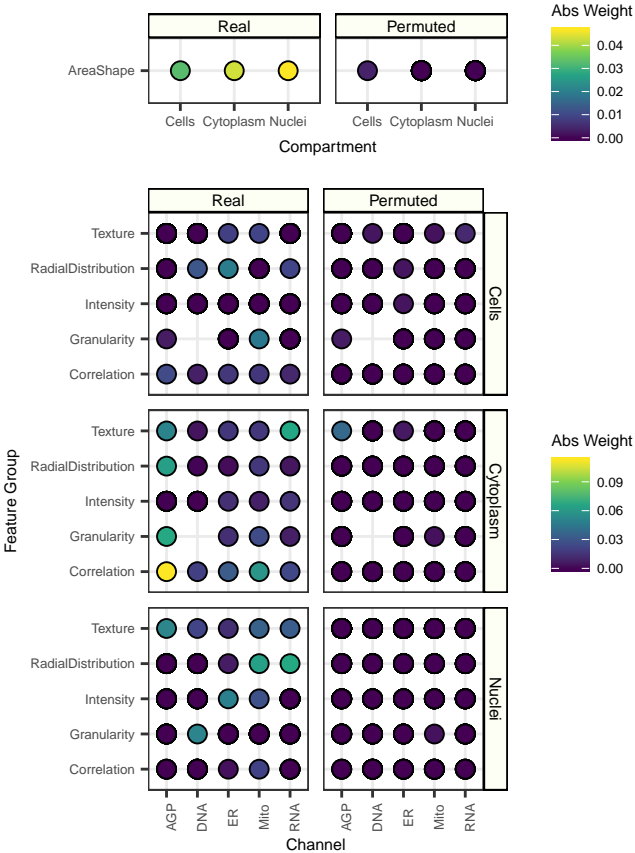
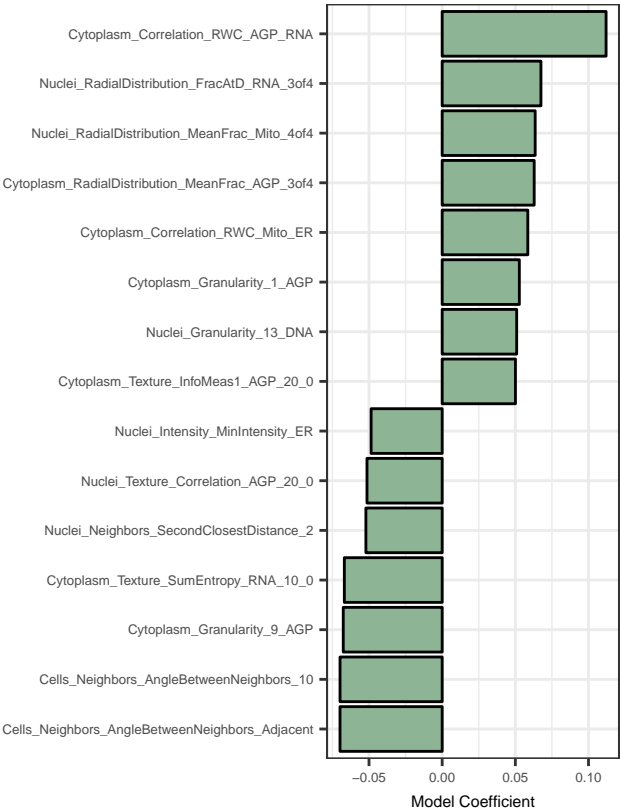


cc_cc_g2_ph3_neg_mean



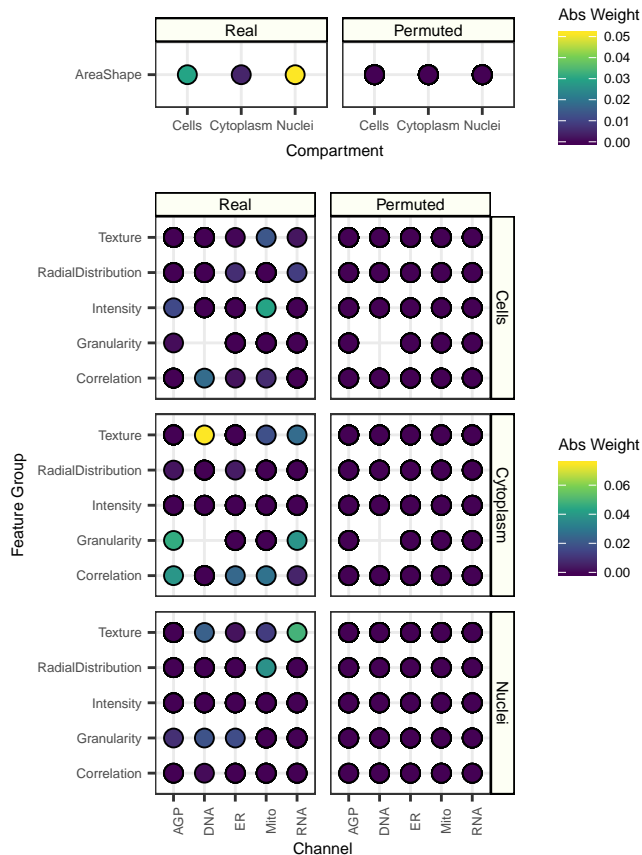
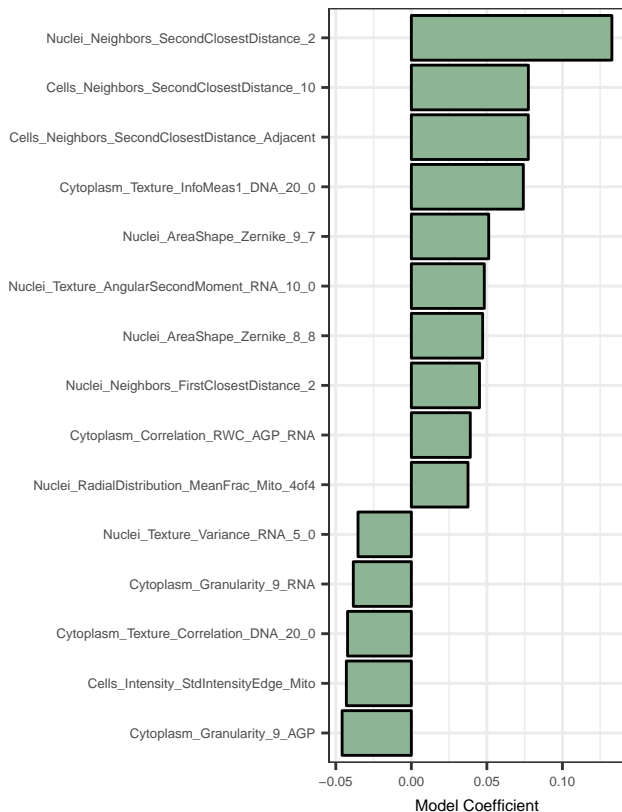
cc_cc_g2_ph3_pos_early_mitosis_mean

9.76% Non-Zero

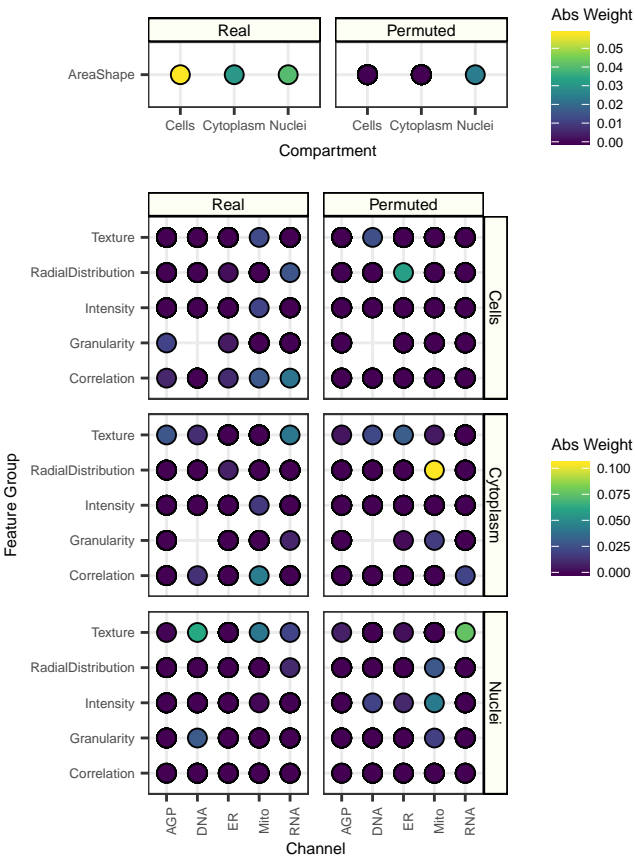
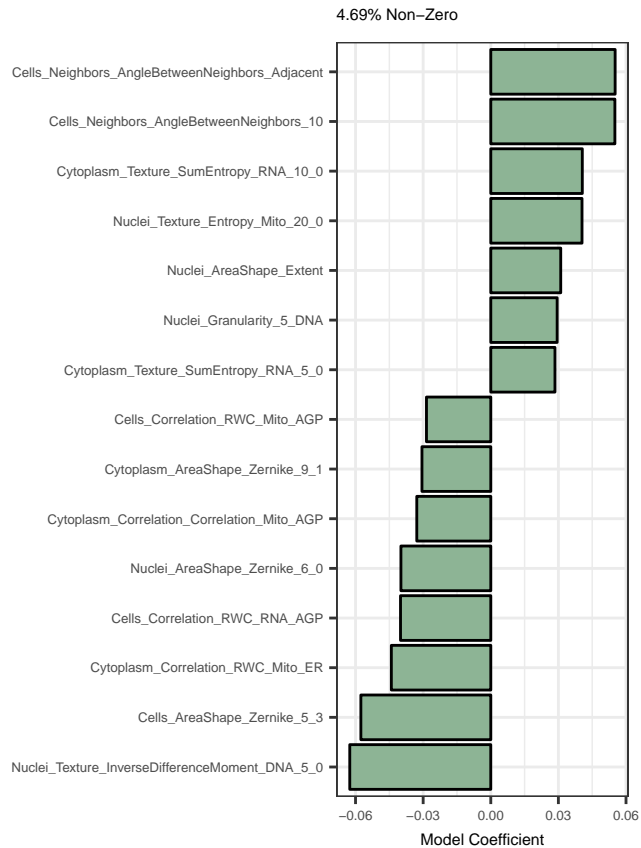


cc_cc_high_n_spots_h2ax_mean

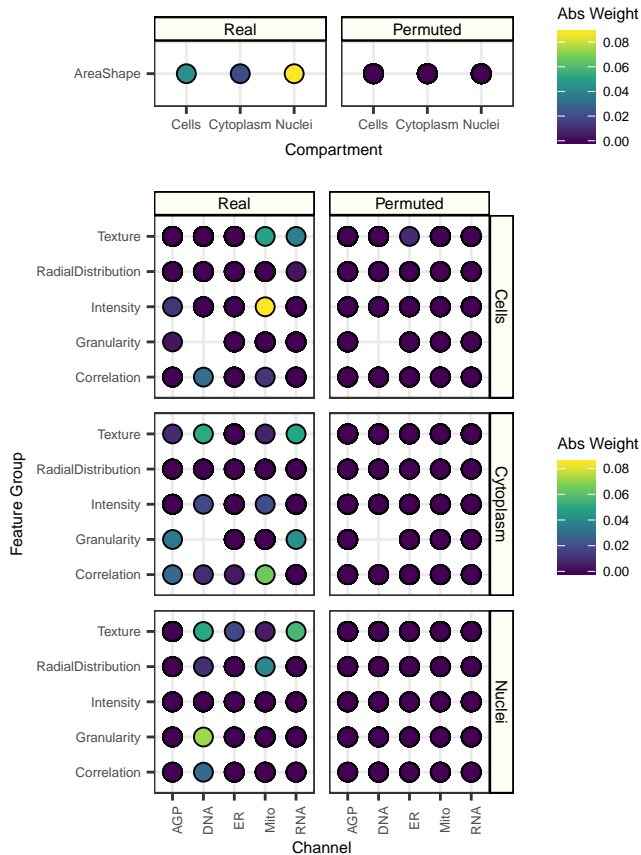
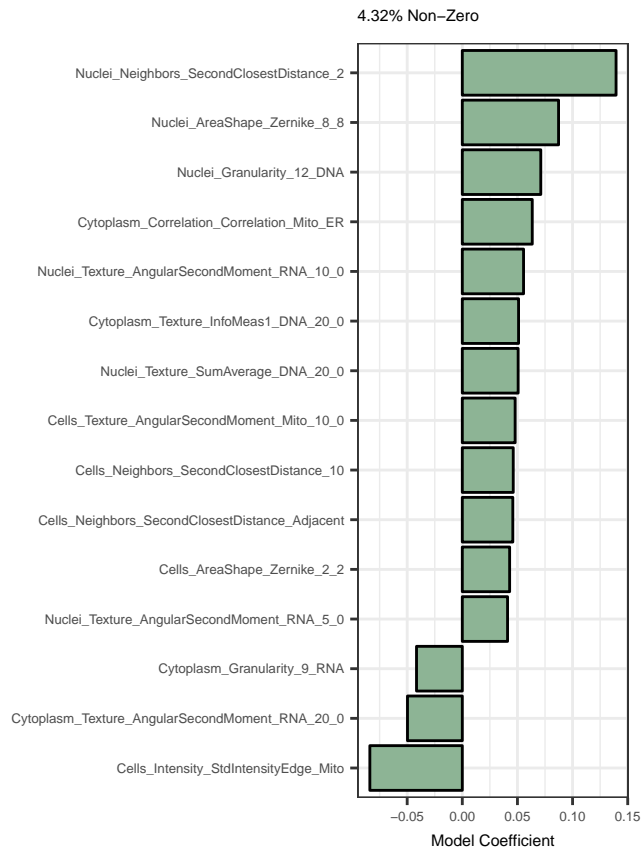
4.82% Non-Zero



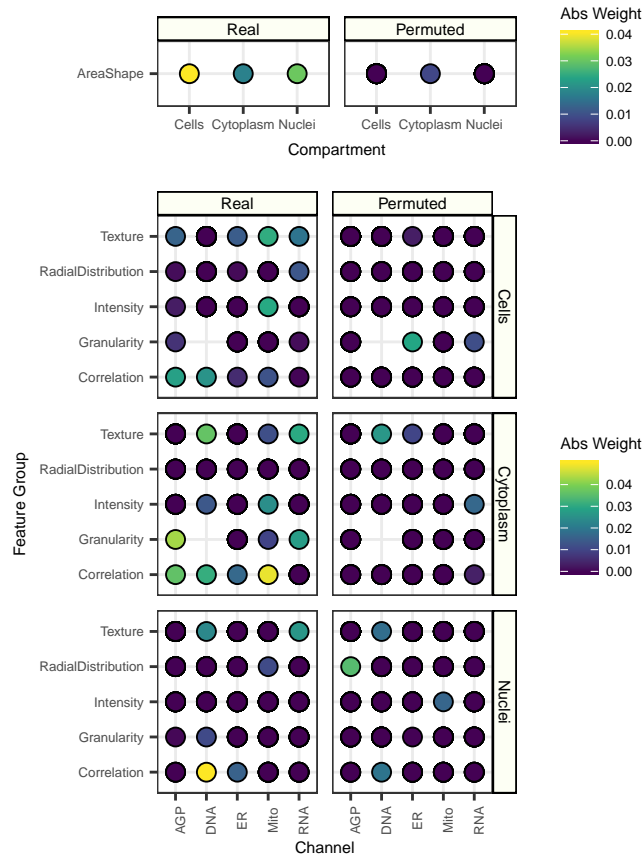
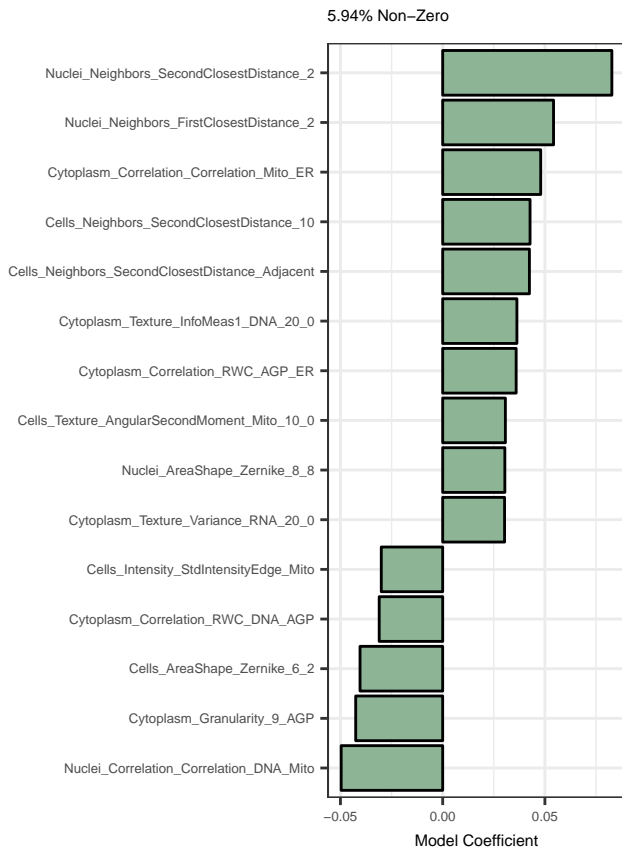
cc_cc_n_objects



cc_cc_n_spots_mean

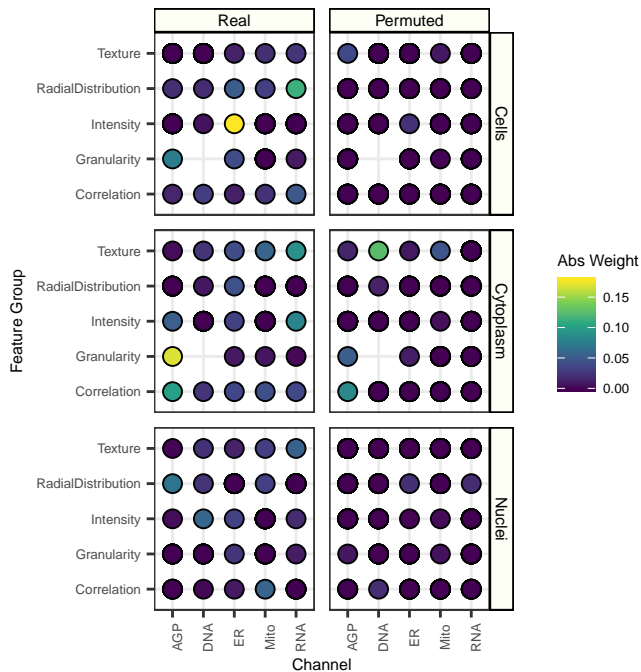
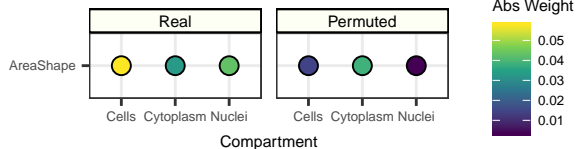


cc_cc_n_spots_per_nucleus_area_mean

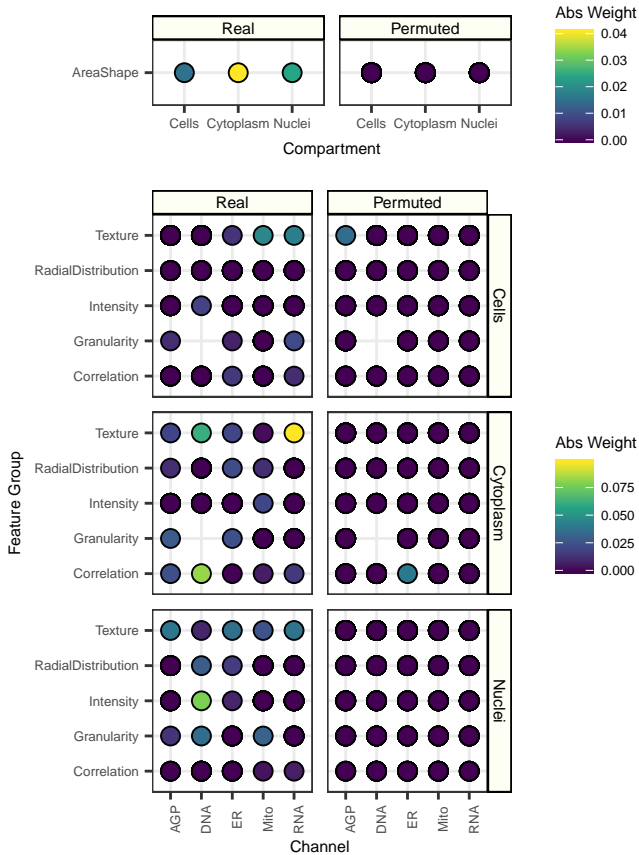


cc_cc_ph3_neg_hoechst_late_mitosis_mean

10.89% Non-Zero

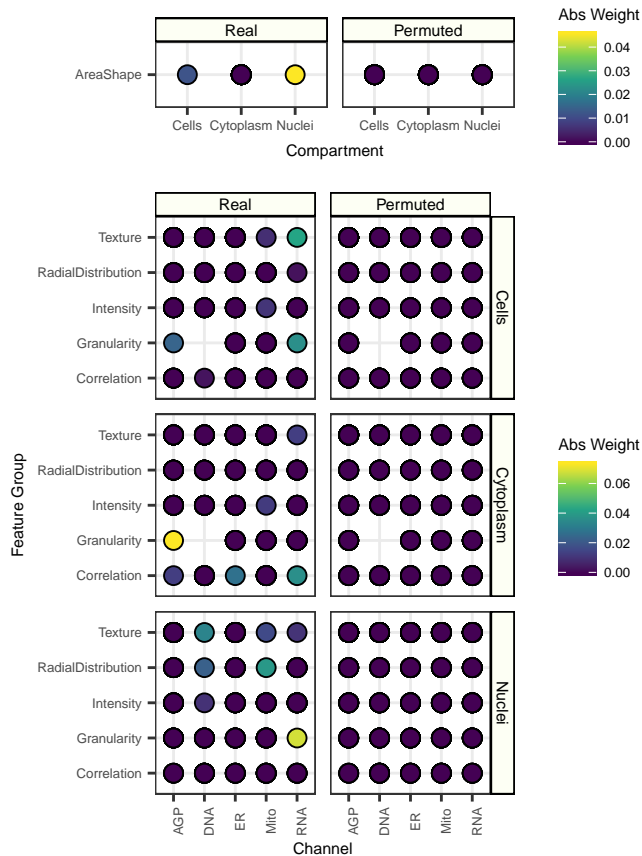
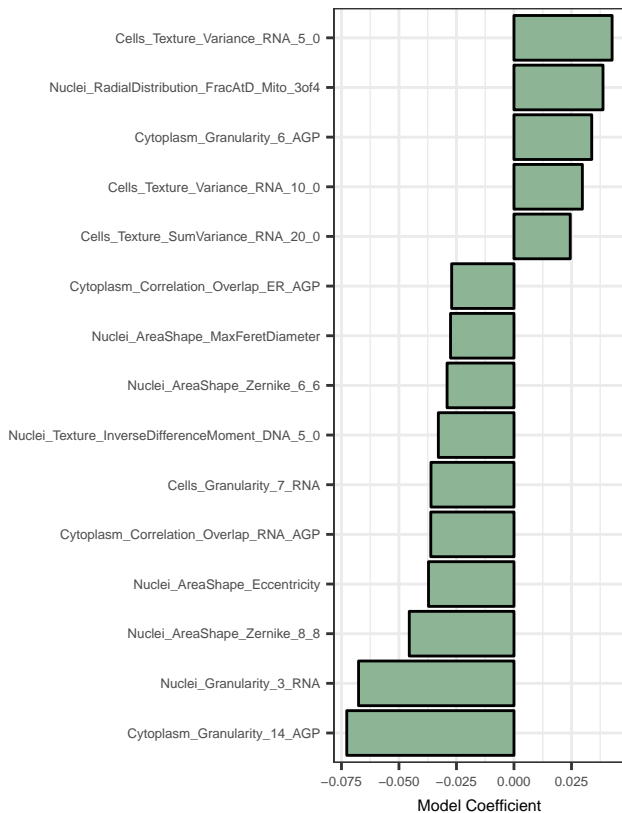


cc_cc_ph3_pos_hoechst_mitosis_mean



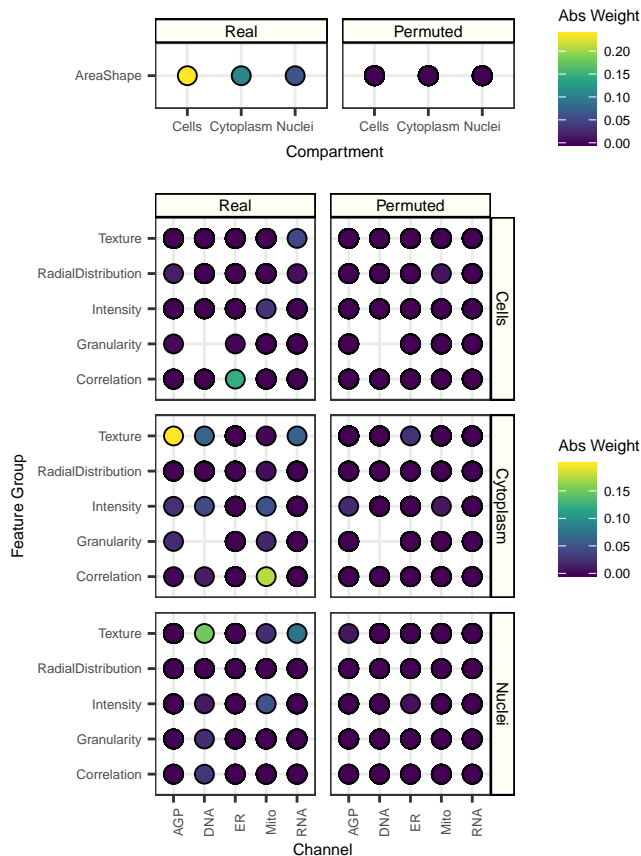
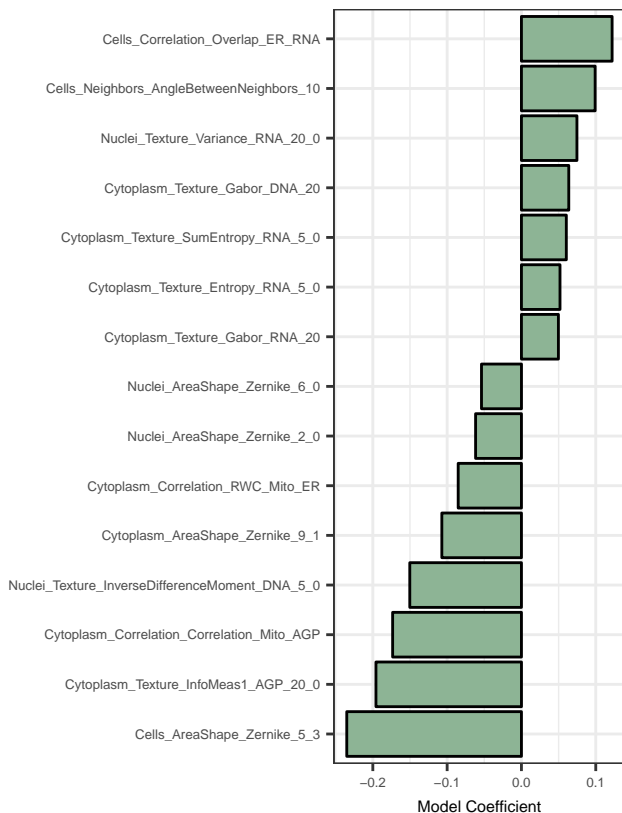
cc_edu_pos_alexa647_intensity_nucleus_area_mean

3.44% Non-Zero

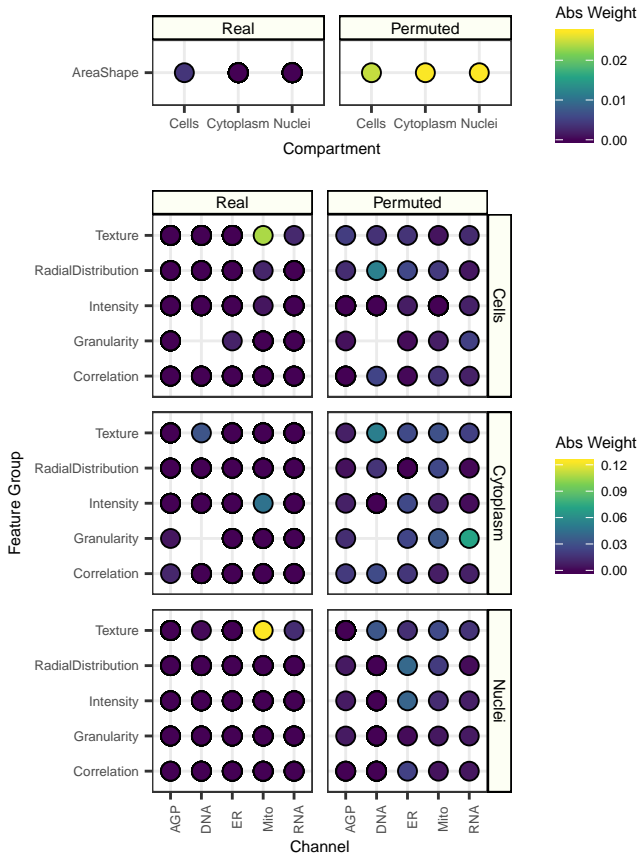


cc_edu_pos_alexa647_intensity_nucleus_area_sum

3.88% Non-Zero

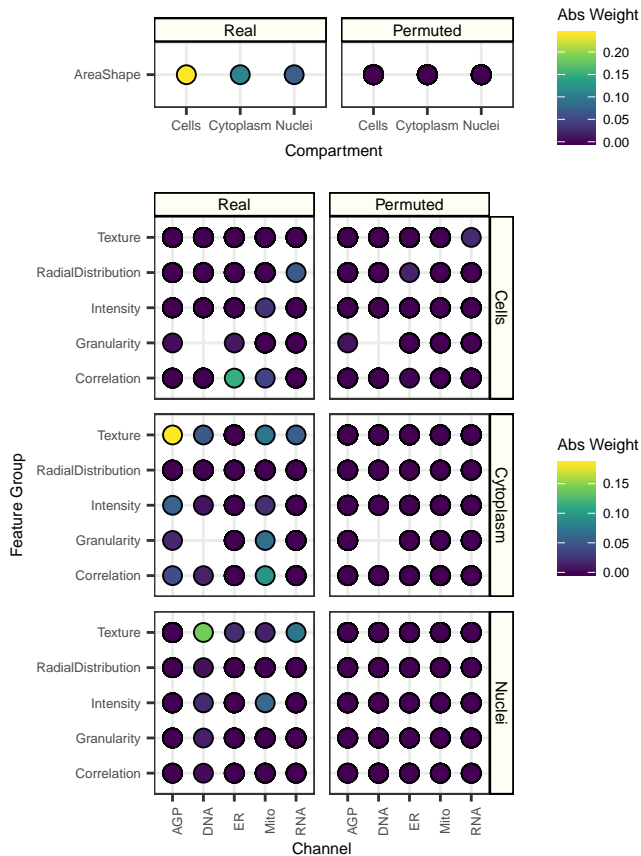
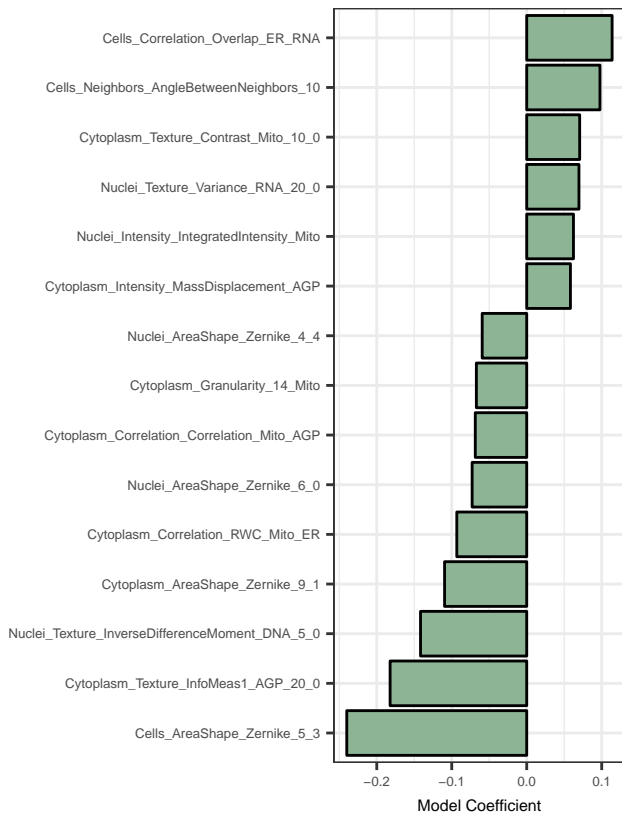


cc_edu_pos_high_n_spots_h2ax_mean



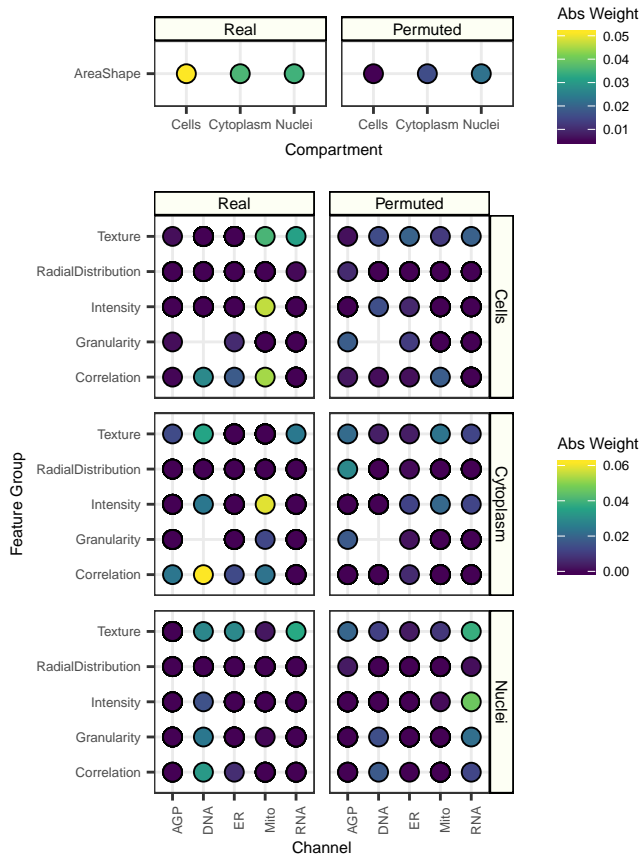
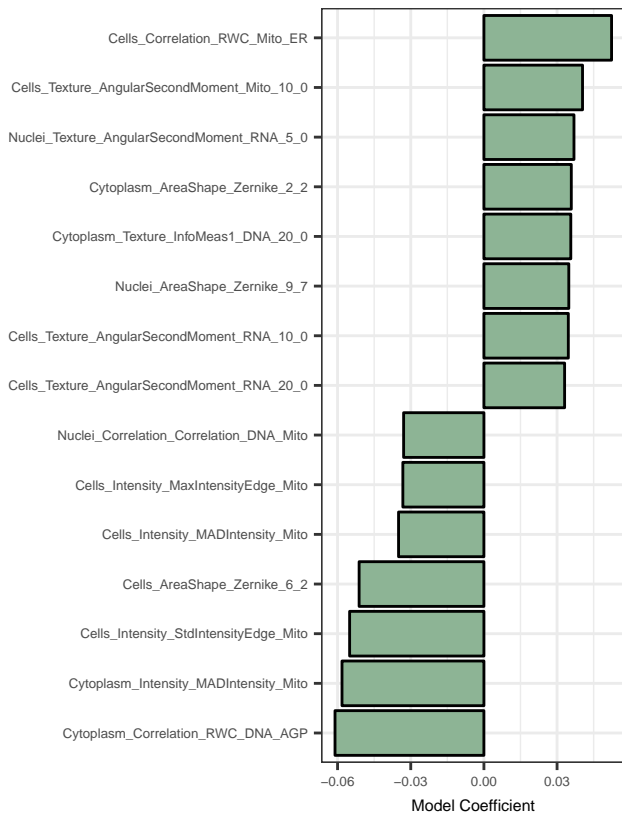
cc_edu_pos_n_objects

3.88% Non-Zero



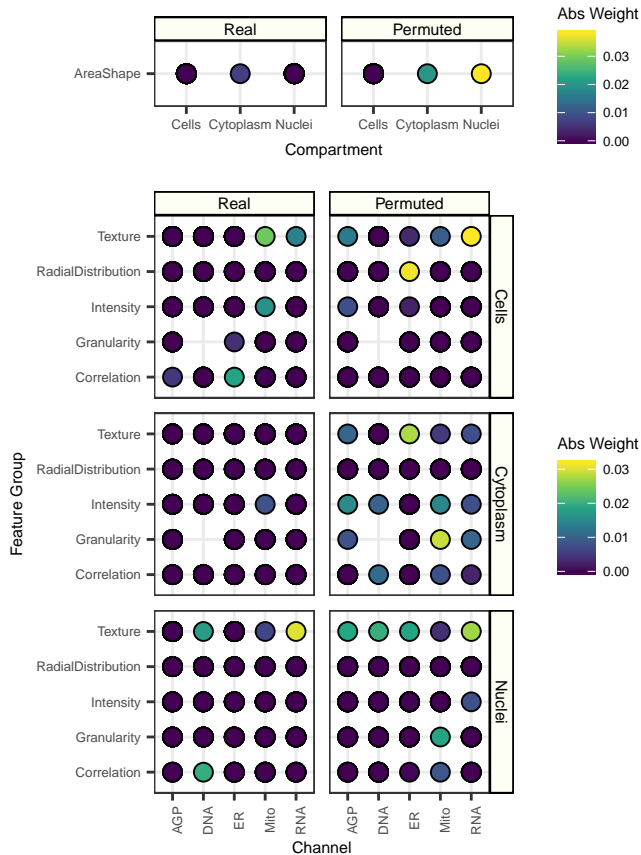
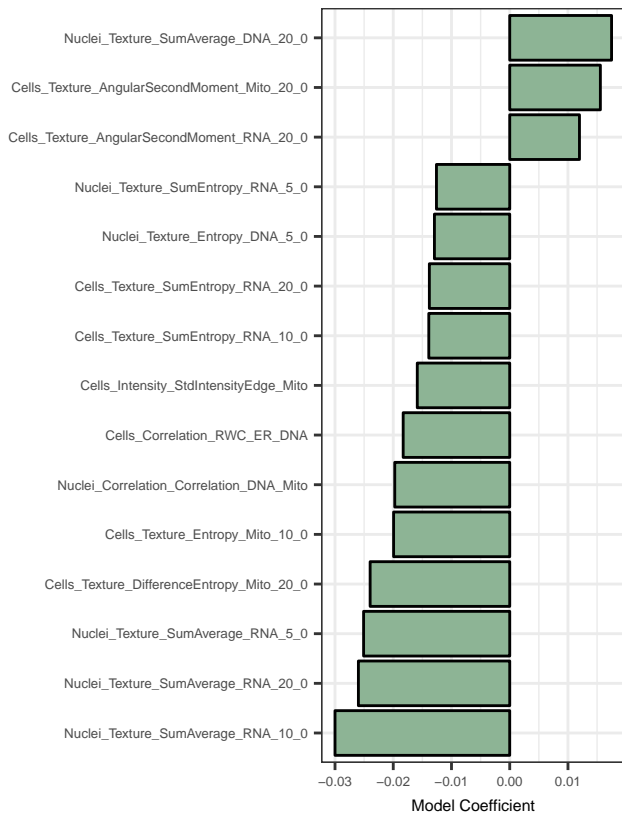
cc_edu_pos_n_spots_mean

6.45% Non-Zero



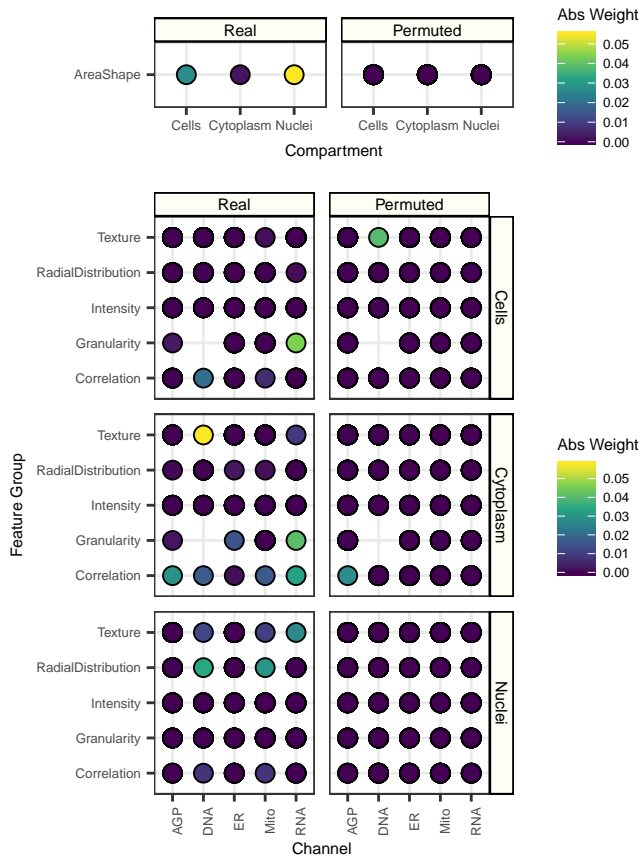
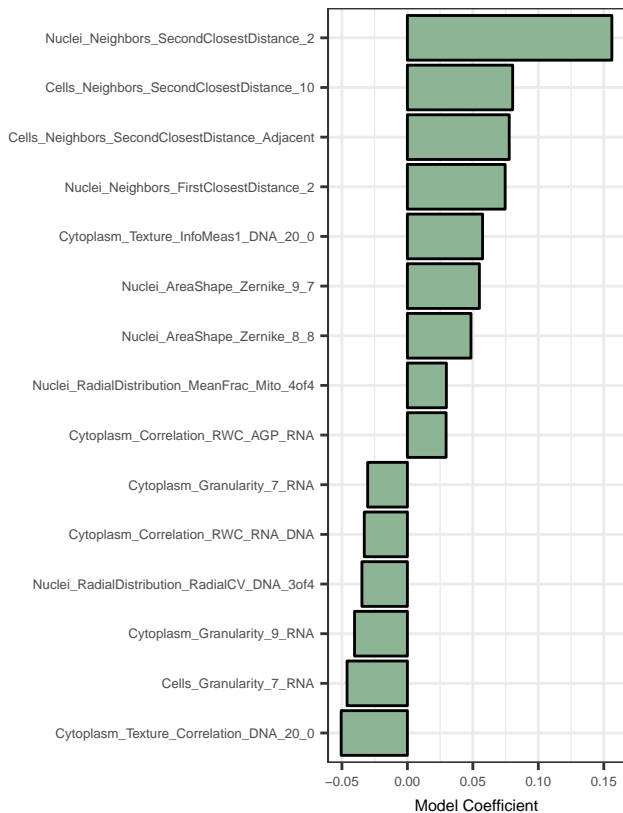
cc_edu_pos_n_spots_per_nucleus_area_mean

2.07% Non-Zero

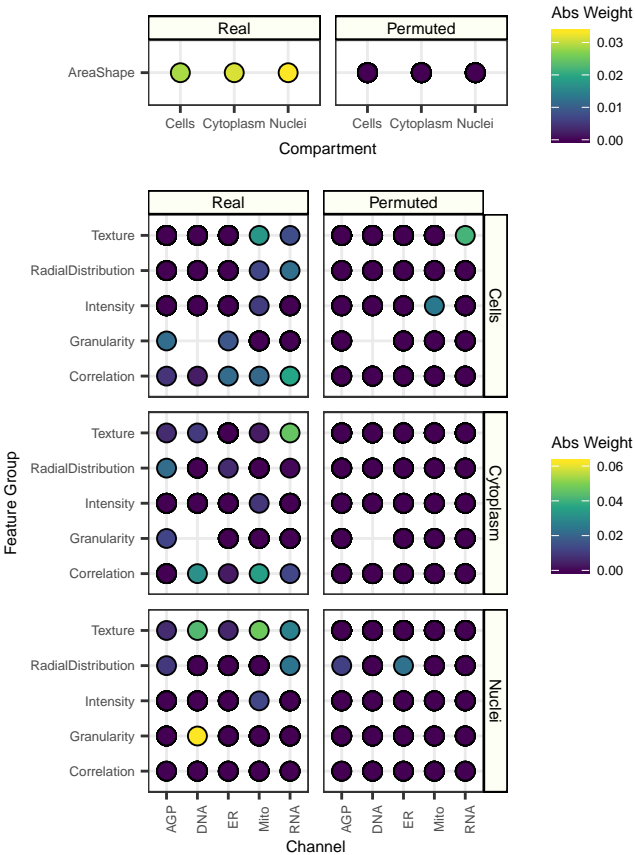
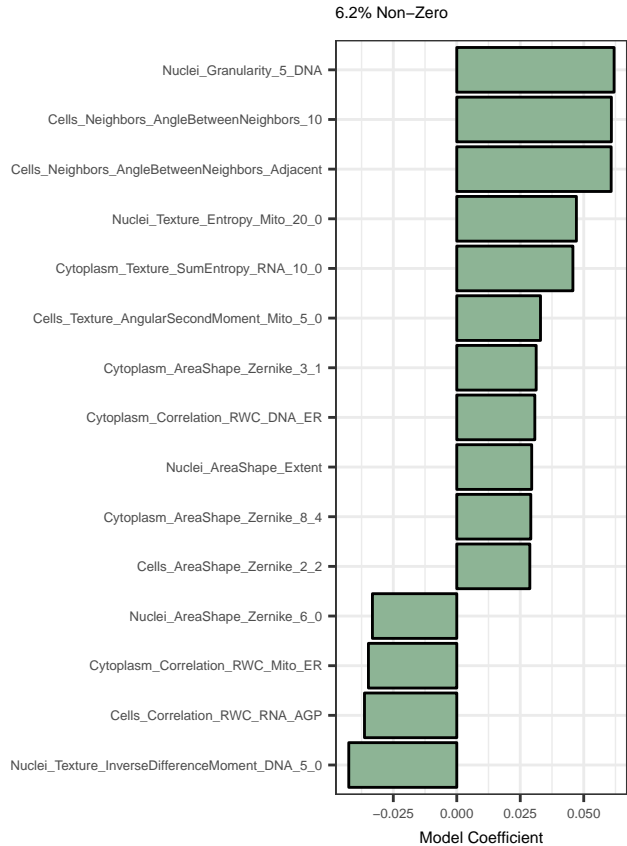


cc_g1_high_n_spots_h2ax_mean

3.25% Non-Zero

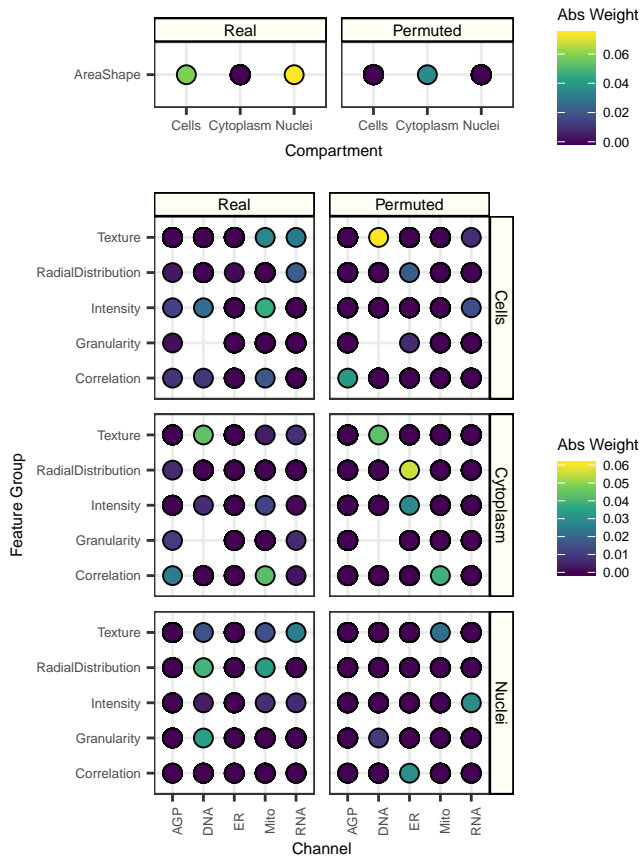
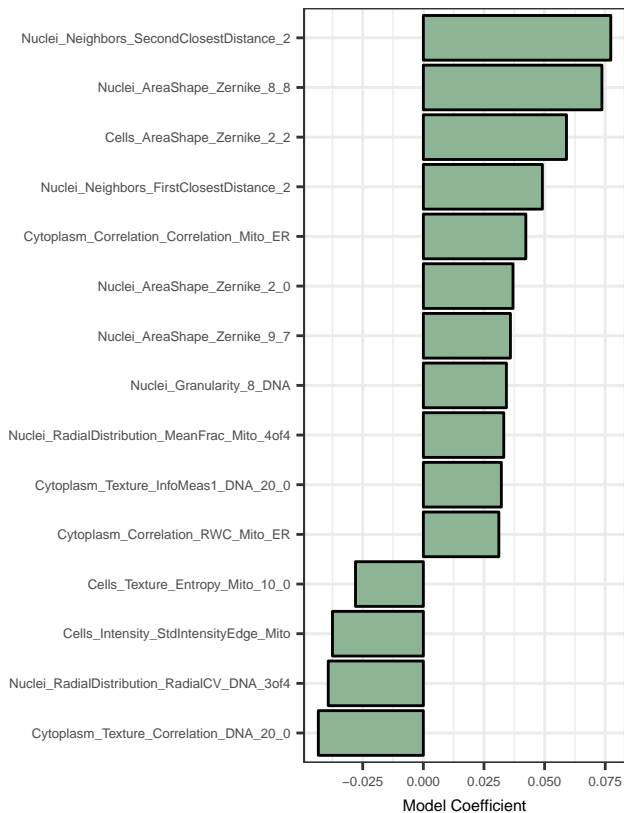


cc_g1_n_objects



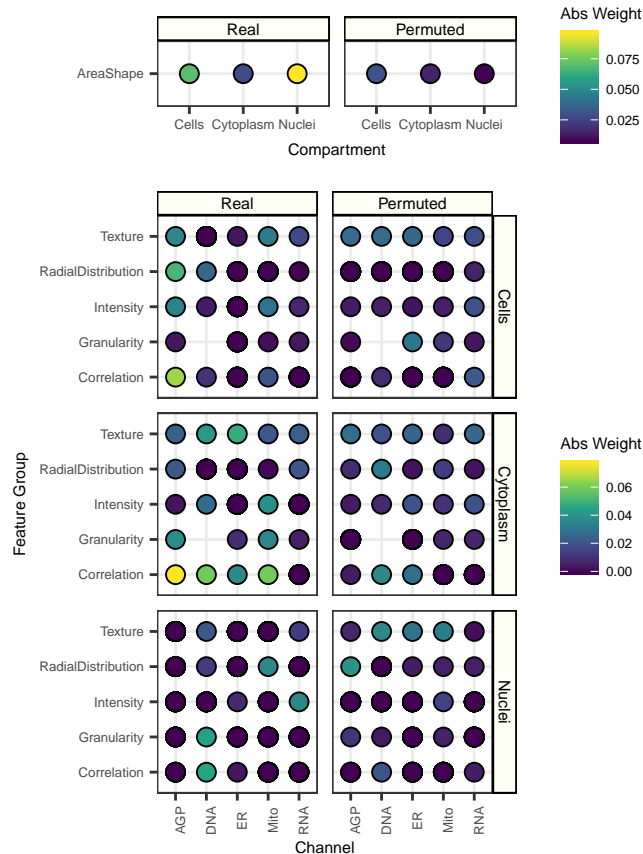
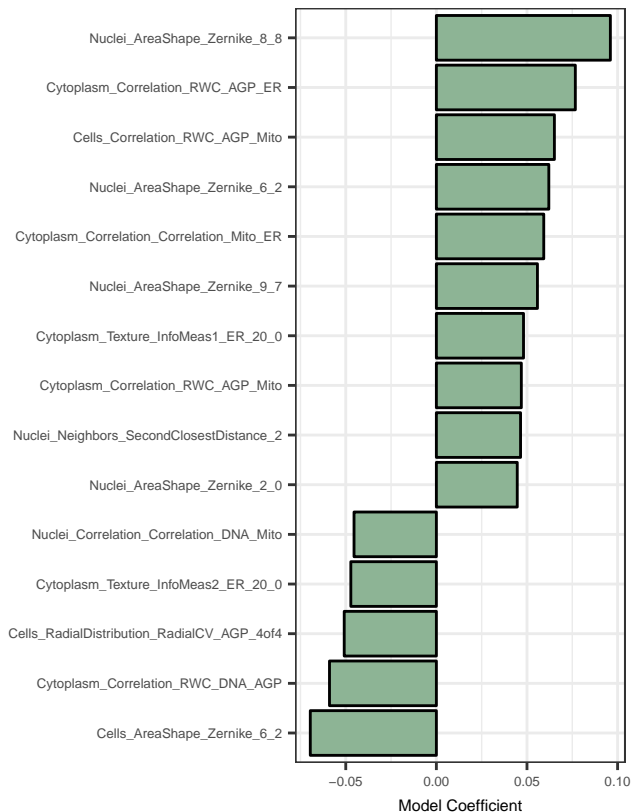
cc_g1_n_spots_mean

5.44% Non-Zero

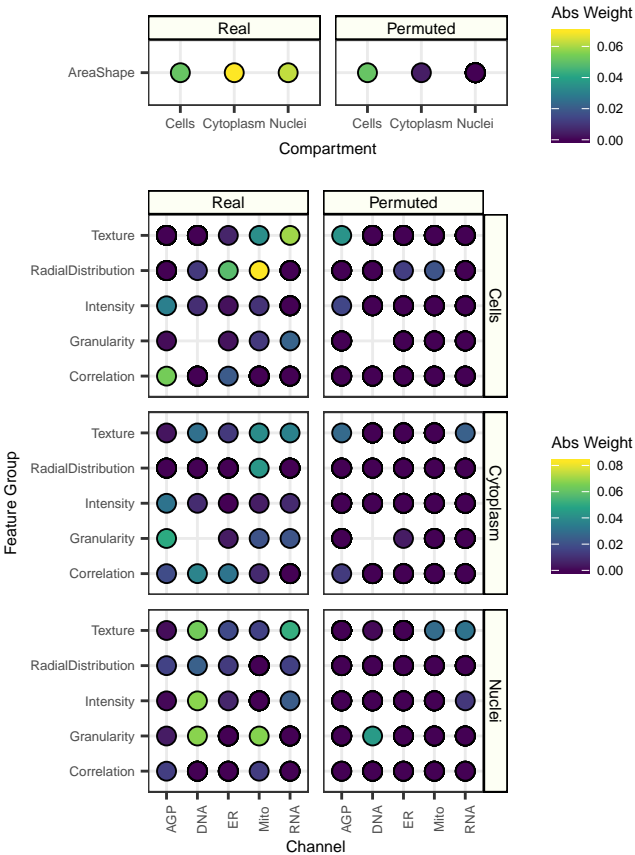
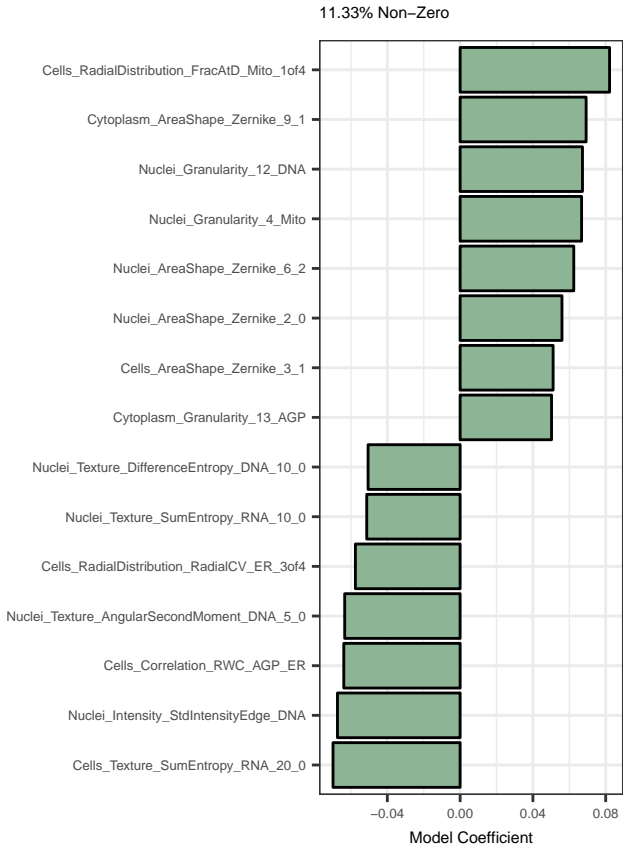


cc_g1_n_spots_per_nucleus_area_mean

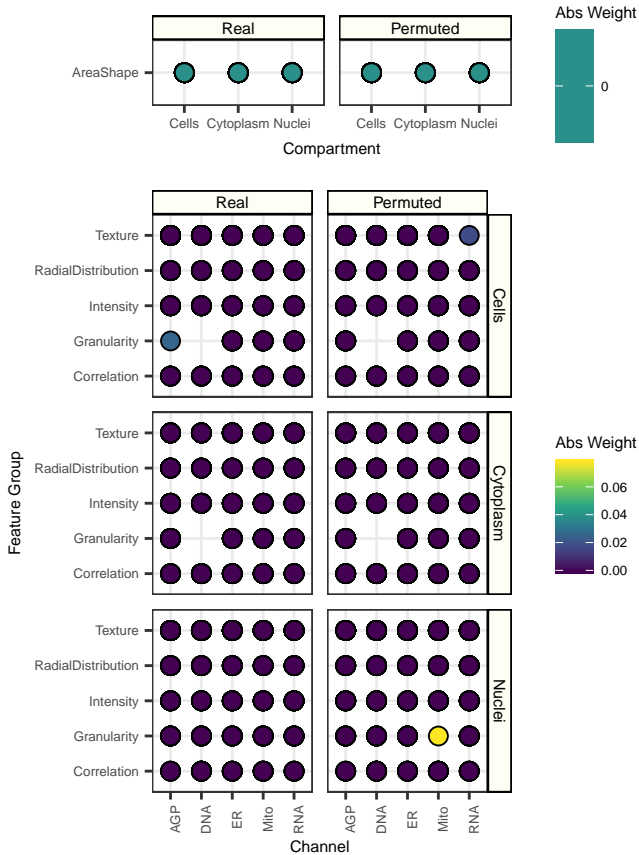
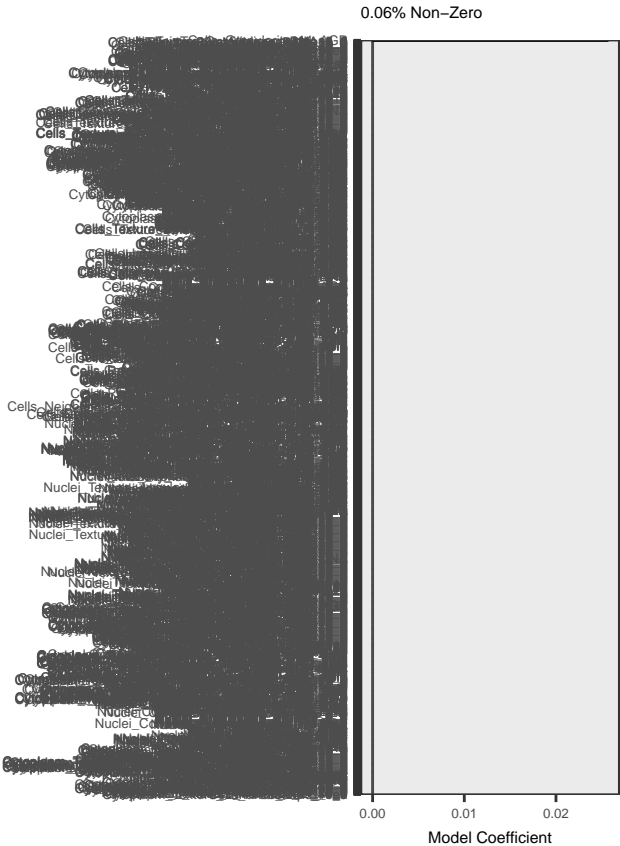
10.76% Non-Zero



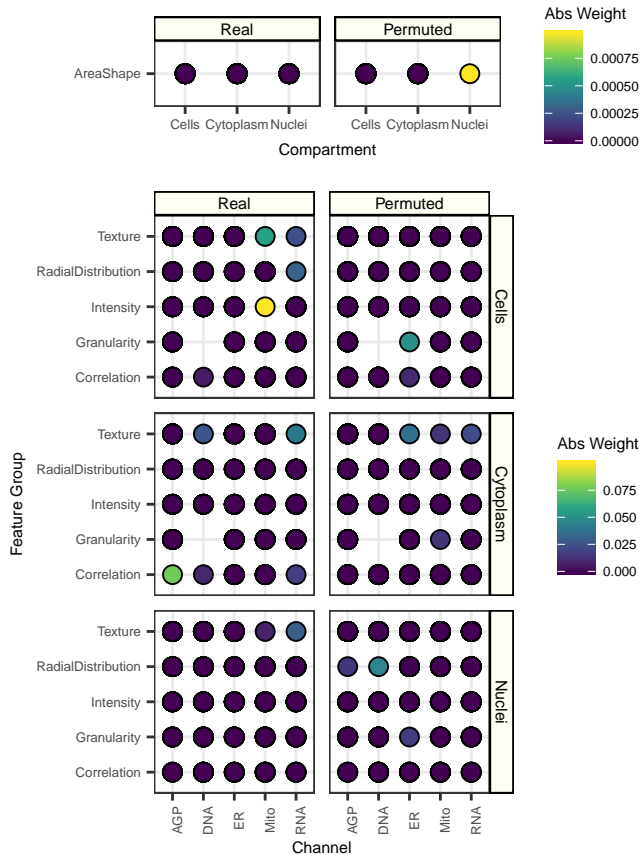
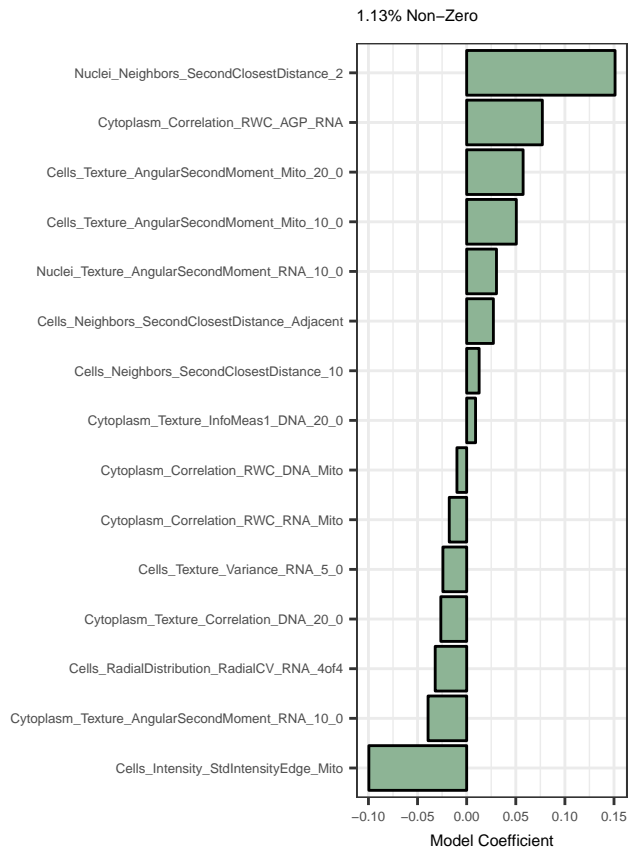
cc_g1_plus_g2



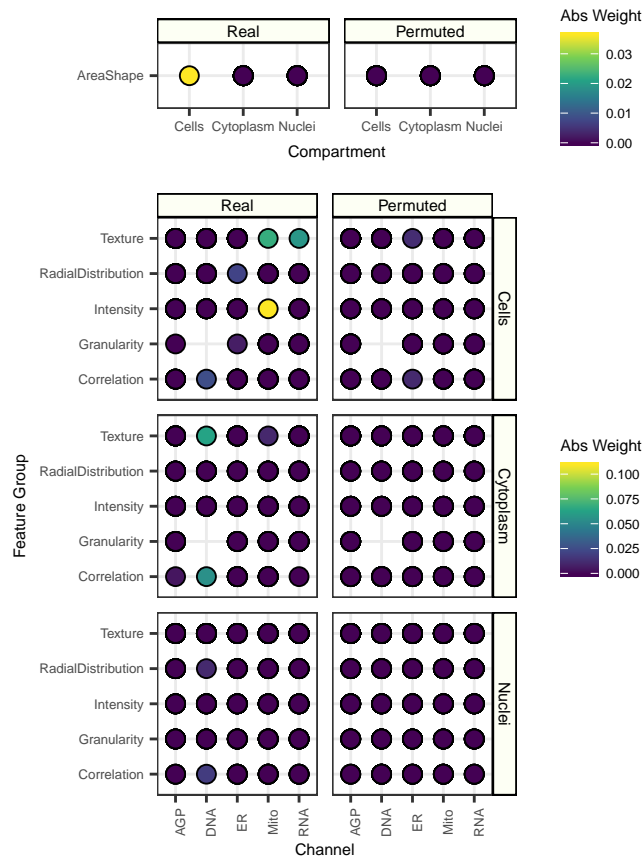
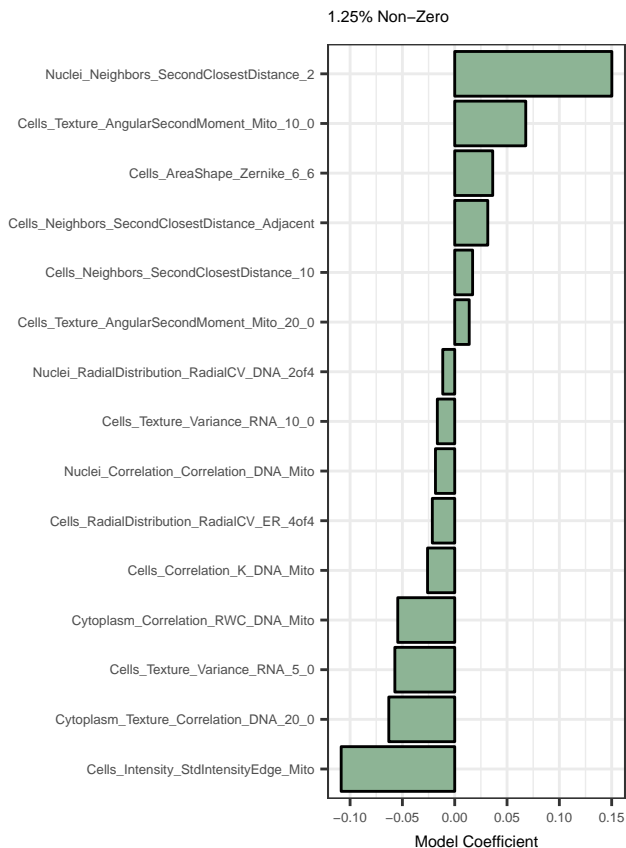
cc_g2_g1



cc_g2_ph3_neg_high_n_spots_h2ax_mean

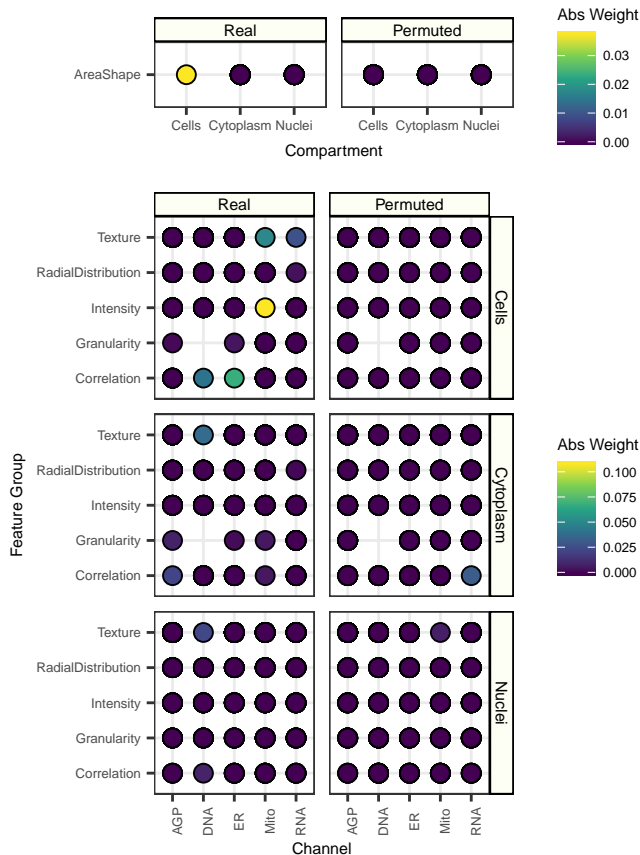
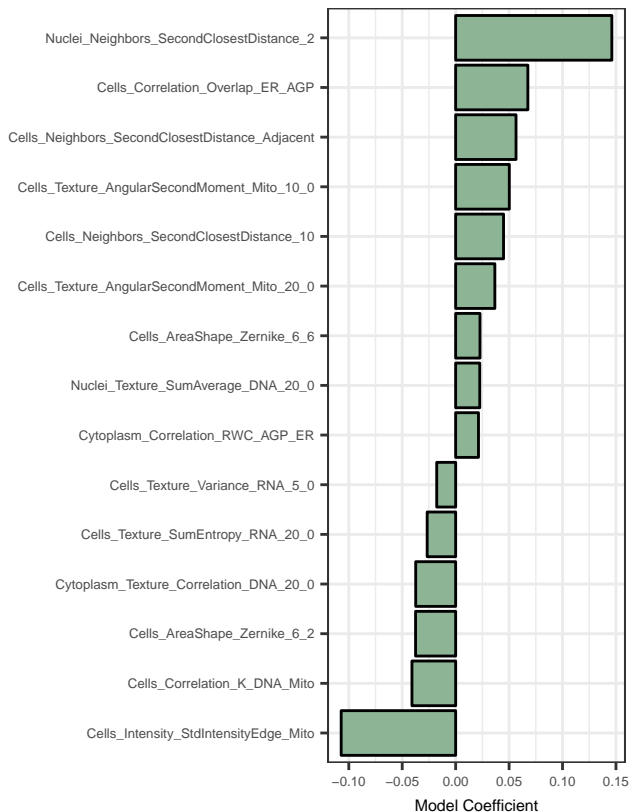


cc_g2_ph3_neg_n_spots_mean



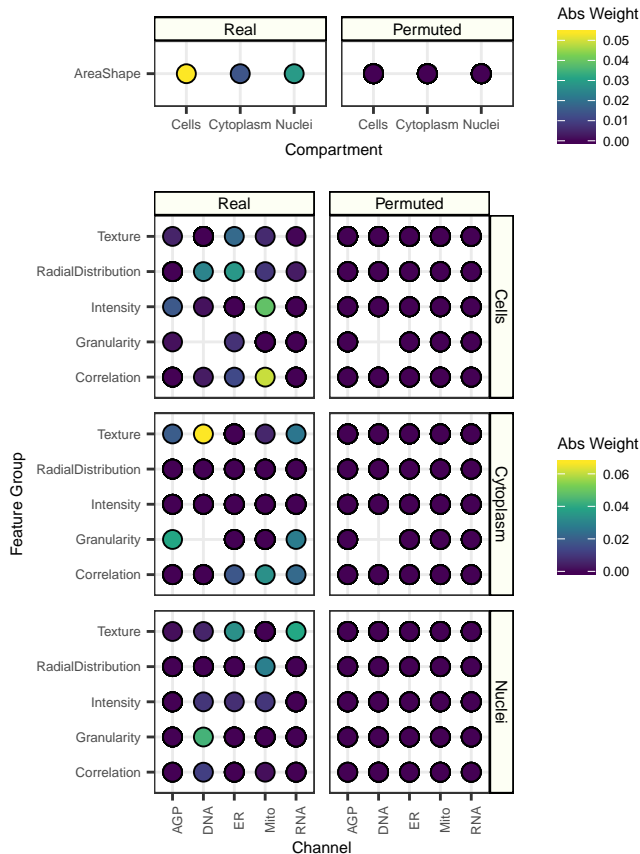
cc_g2_ph3_neg_n_spots_per_nucleus_area_mean

1.63% Non-Zero



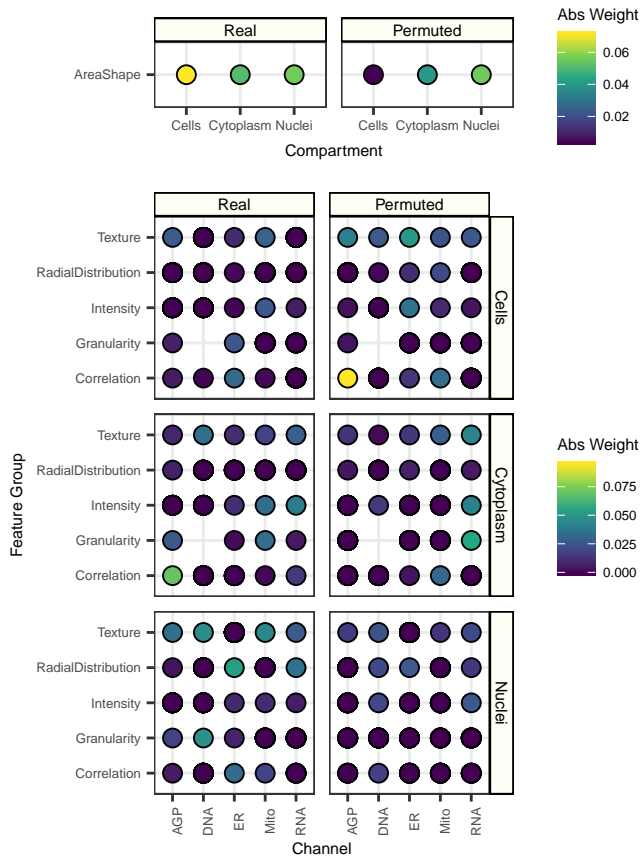
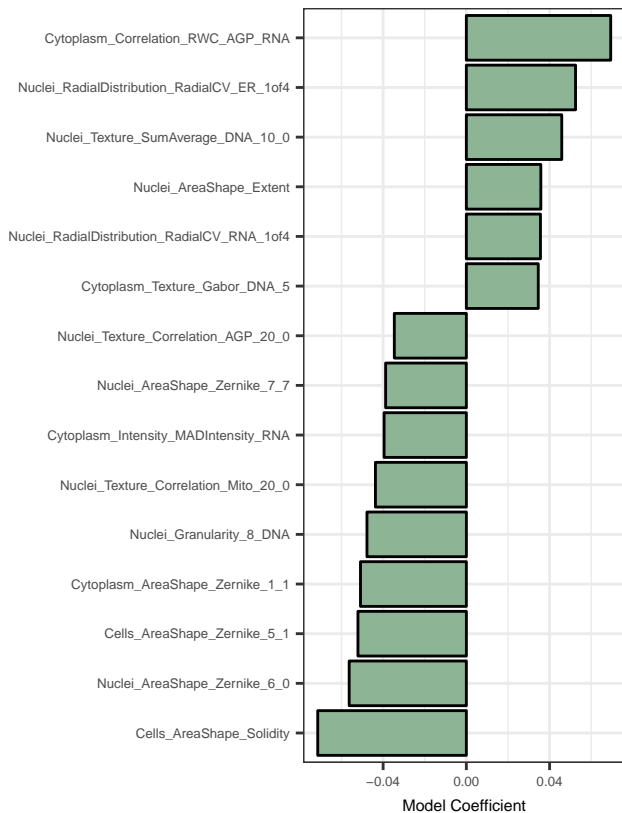
cc_g2_ph3_pos_high_n_spots_h2ax_mean

6.32% Non-Zero

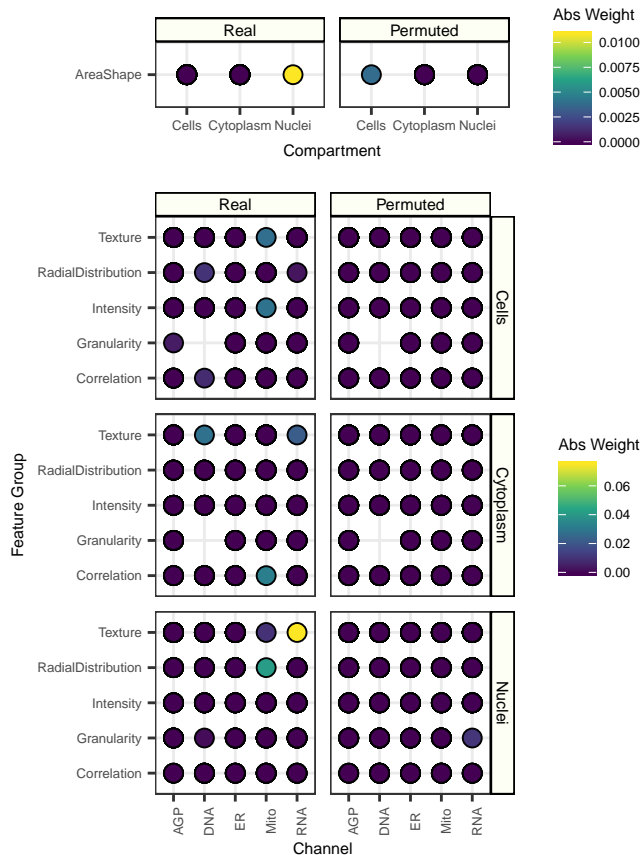
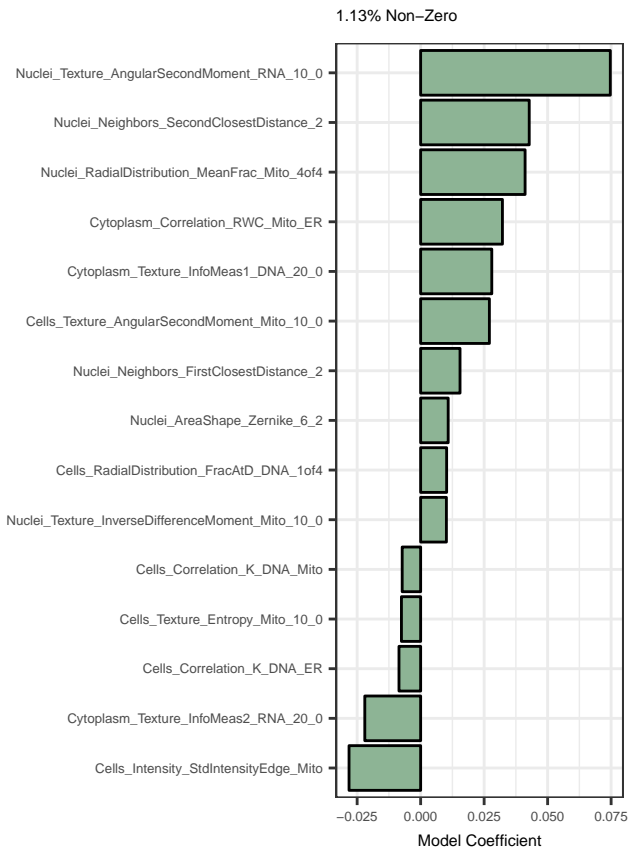


cc_g2_ph3_pos_n_objects

7.88% Non-Zero

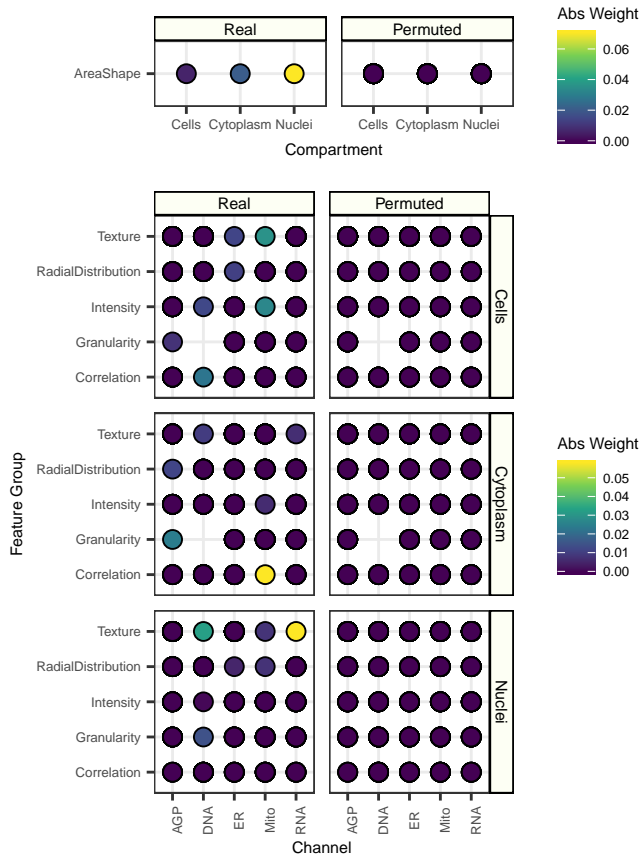
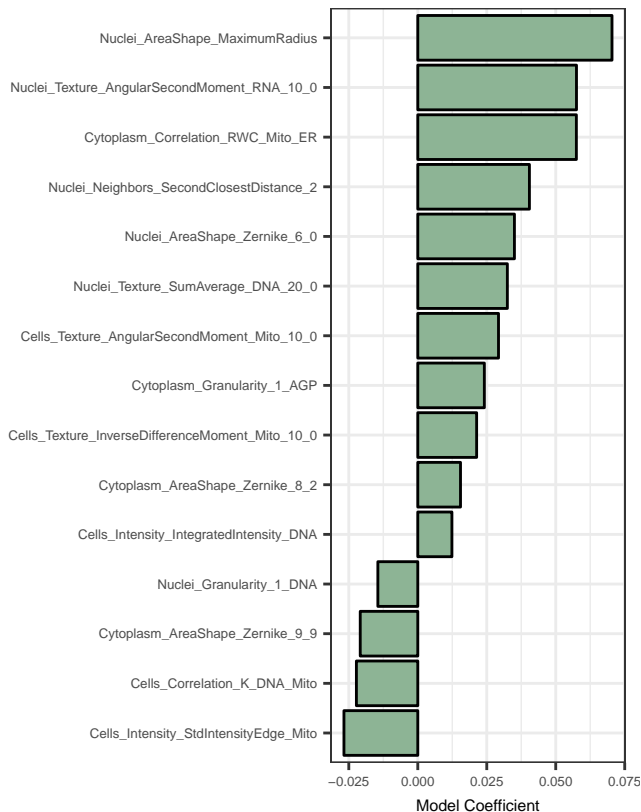


cc_g2_ph3_pos_n_spots_mean



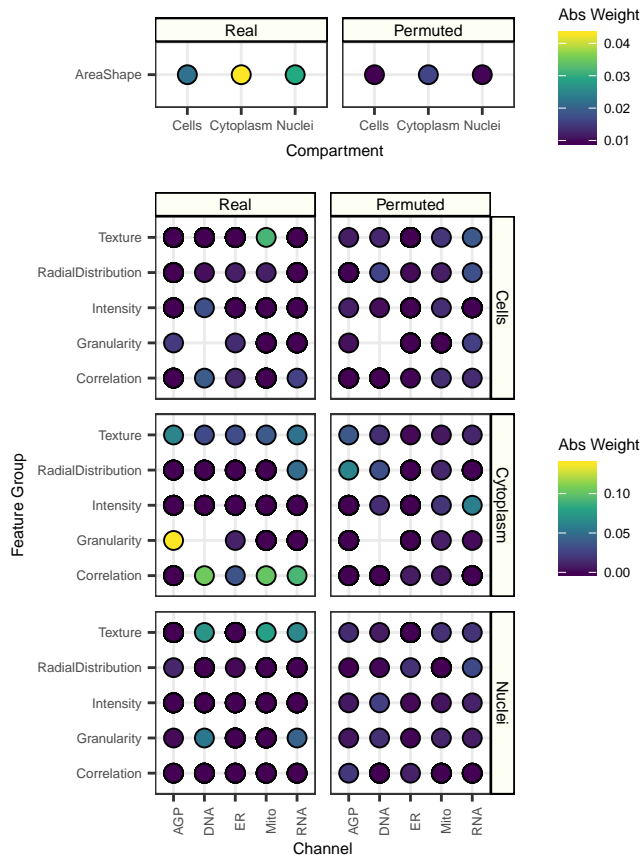
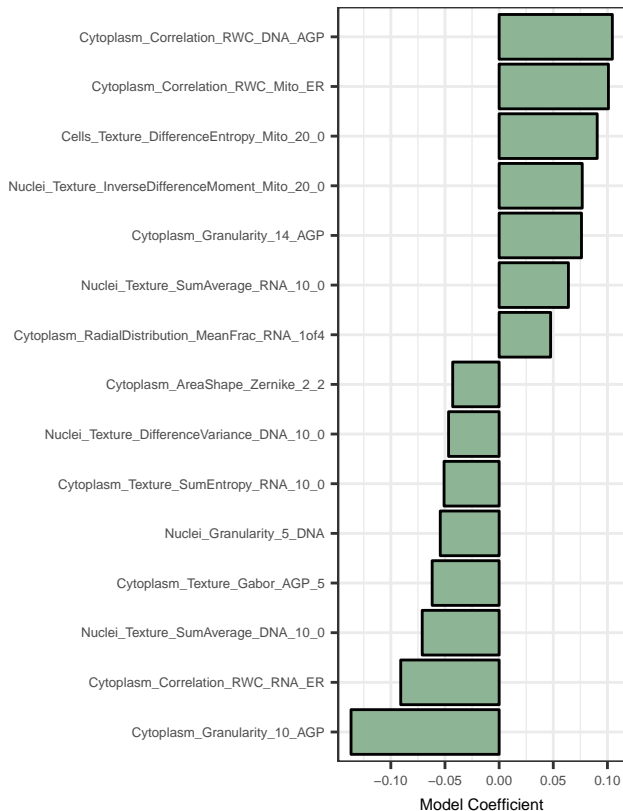
cc_g2_ph3_pos_n_spots_per_nucleus_area_mean

2.13% Non-Zero



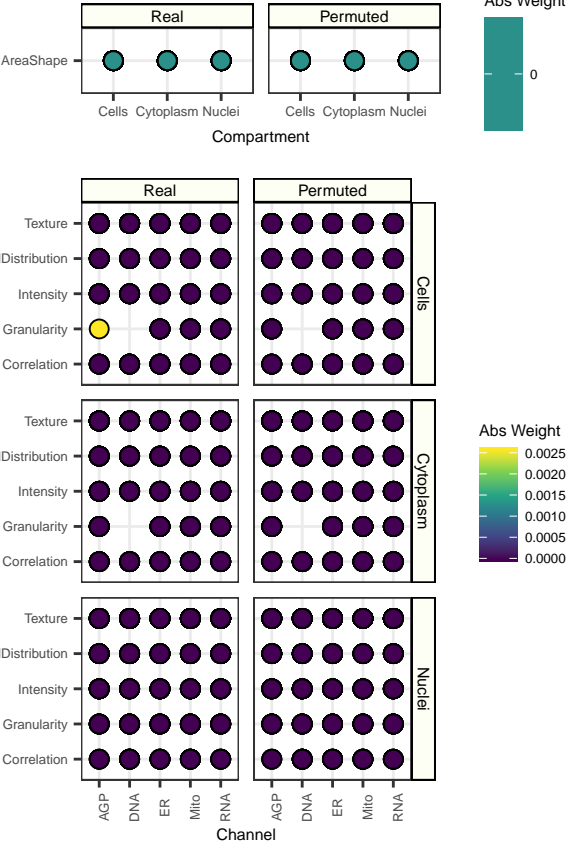
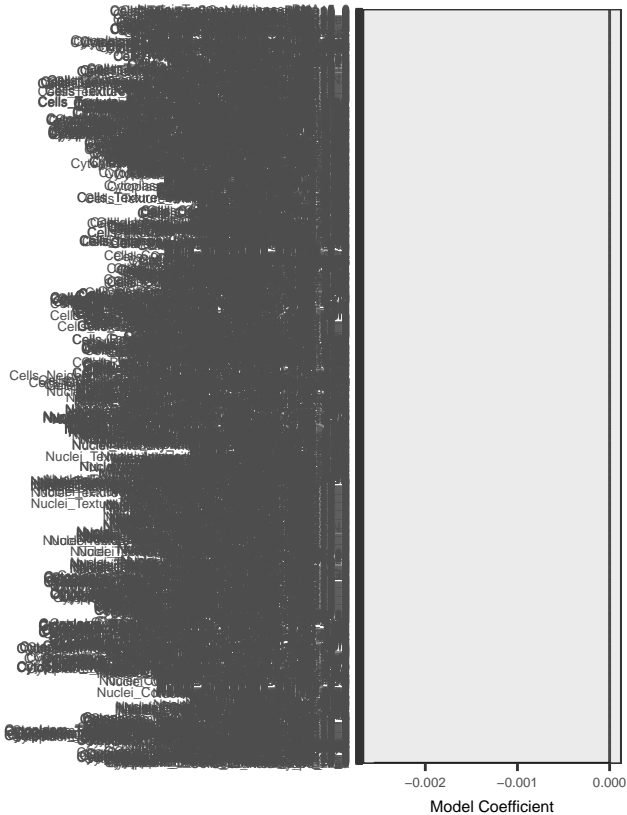
cc_g2_plus_all_m

5.69% Non-Zero



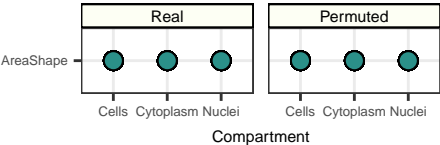
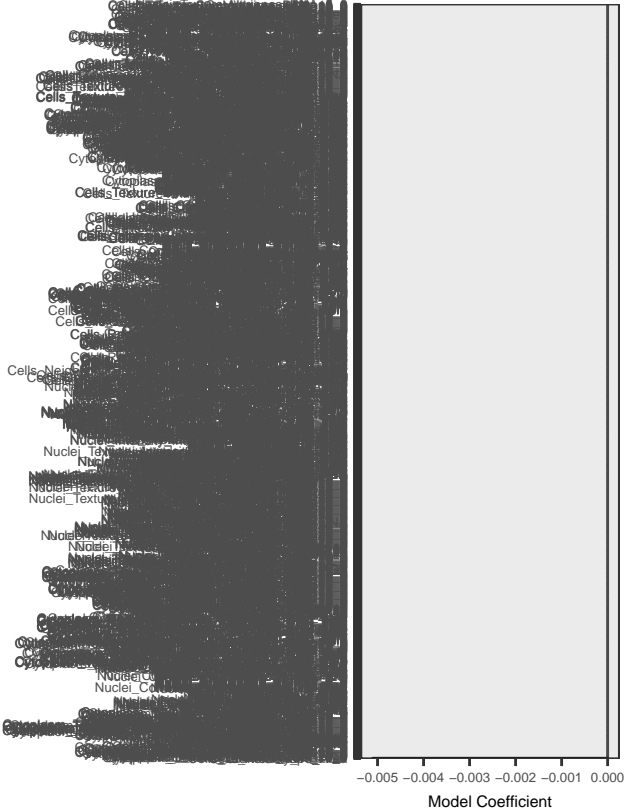
cc_infection_percentage

0.06% Non-Zero

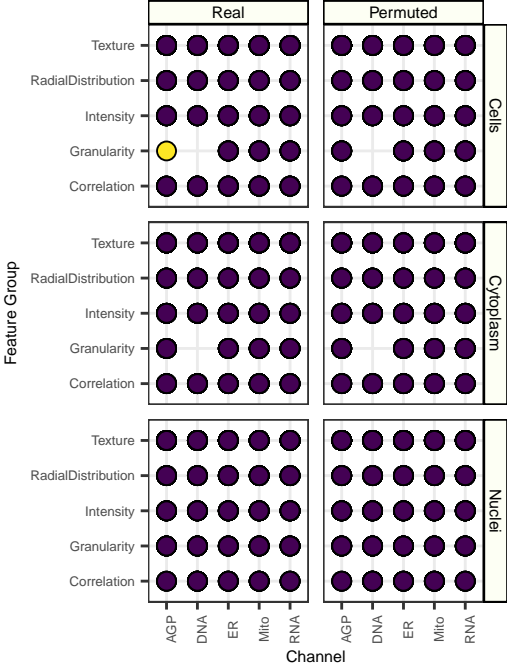


cc_mitosis_ph3_neg_high_n_spots_h2ax_mean

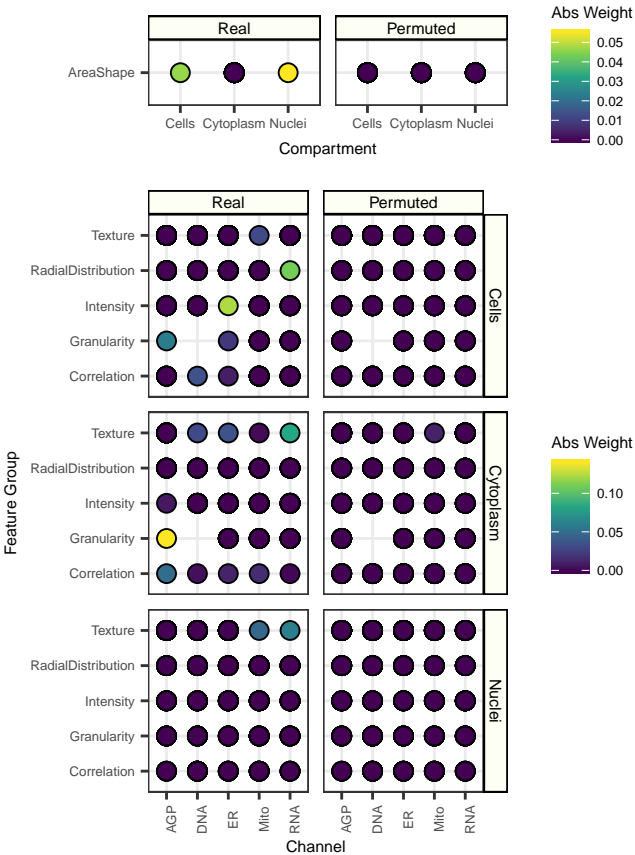
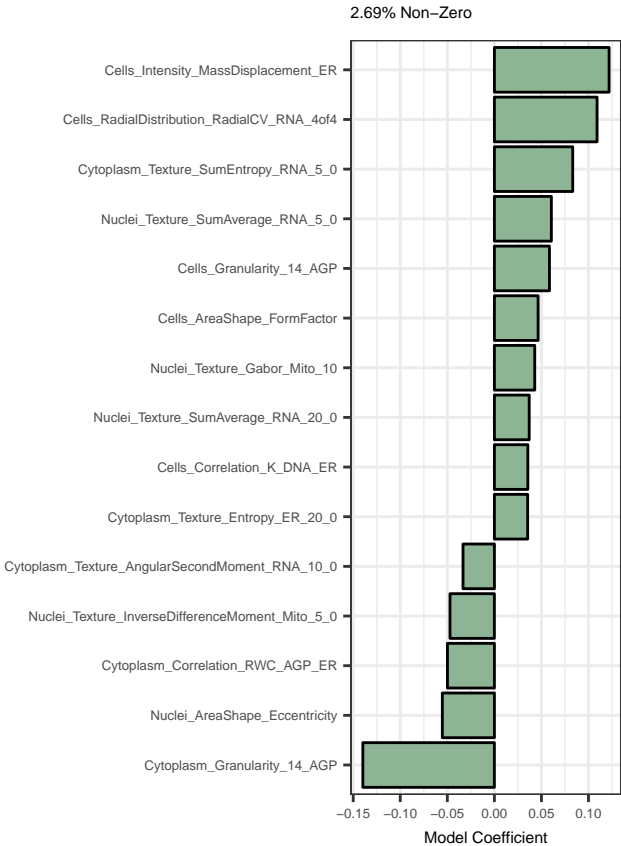
0.06% Non-Zero



Abs Weight

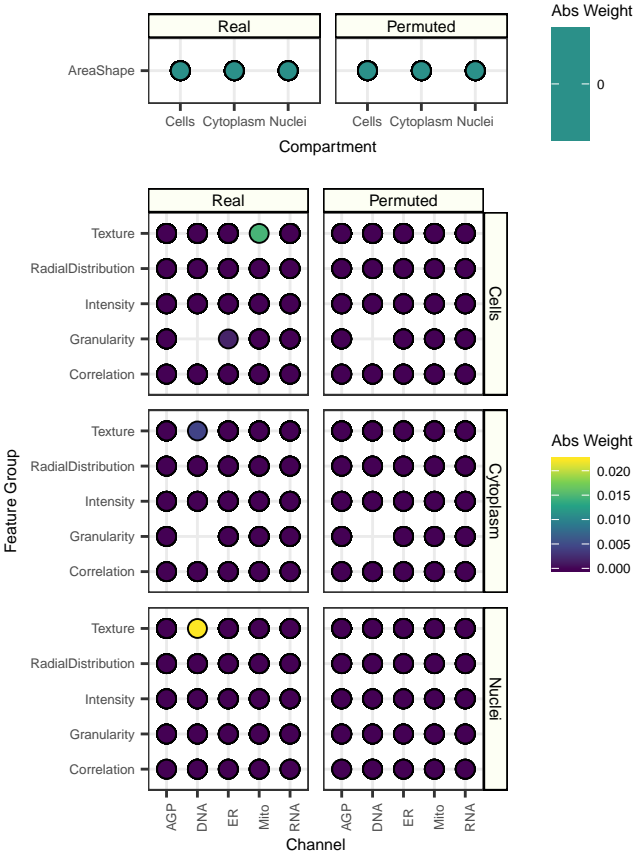


cc_mitosis_ph3_neg_n_objects



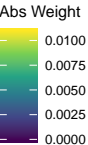
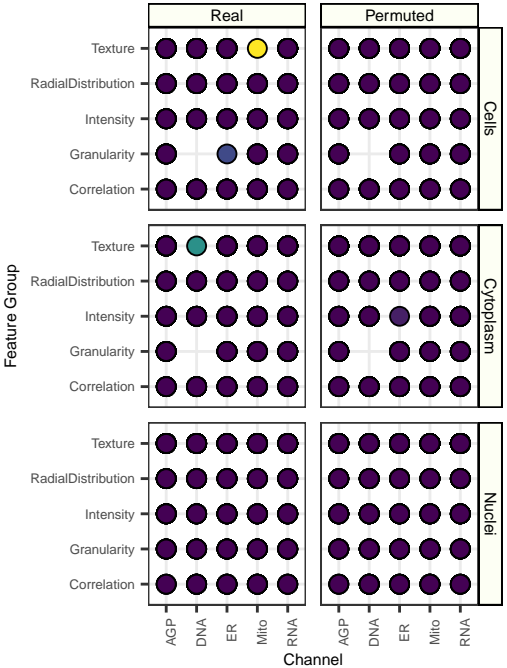
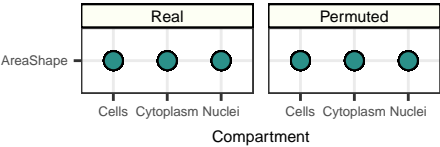
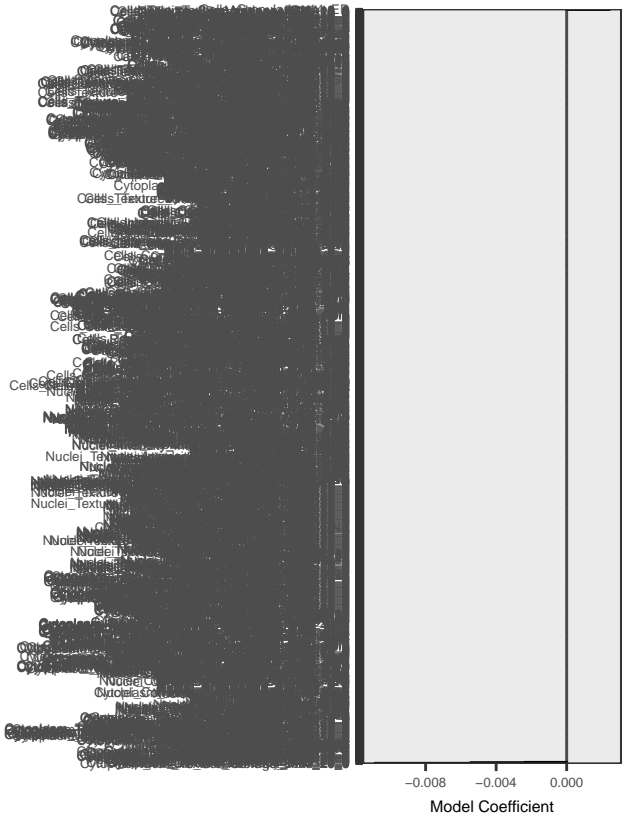
cc_mitosis_ph3_neg_n_spots_mean

0.38% Non-Zero

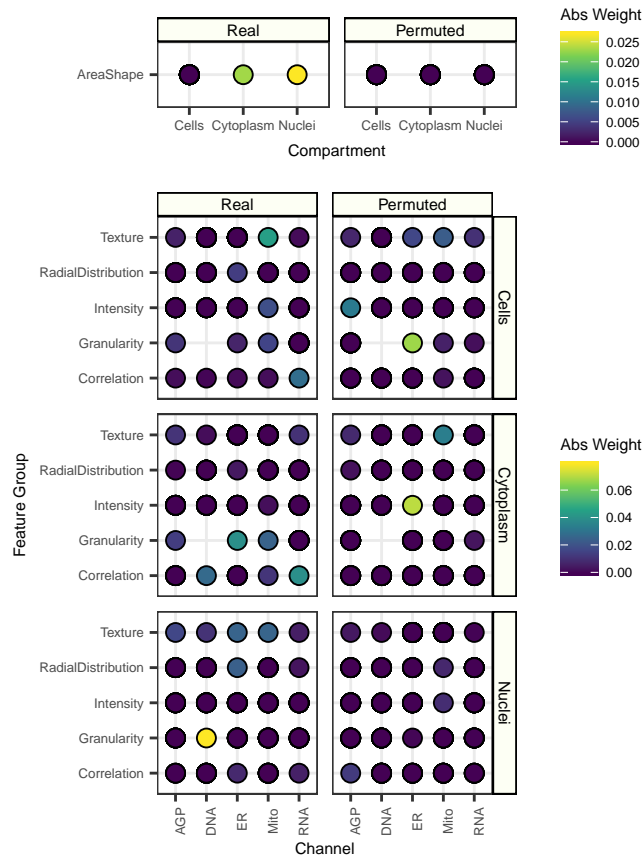
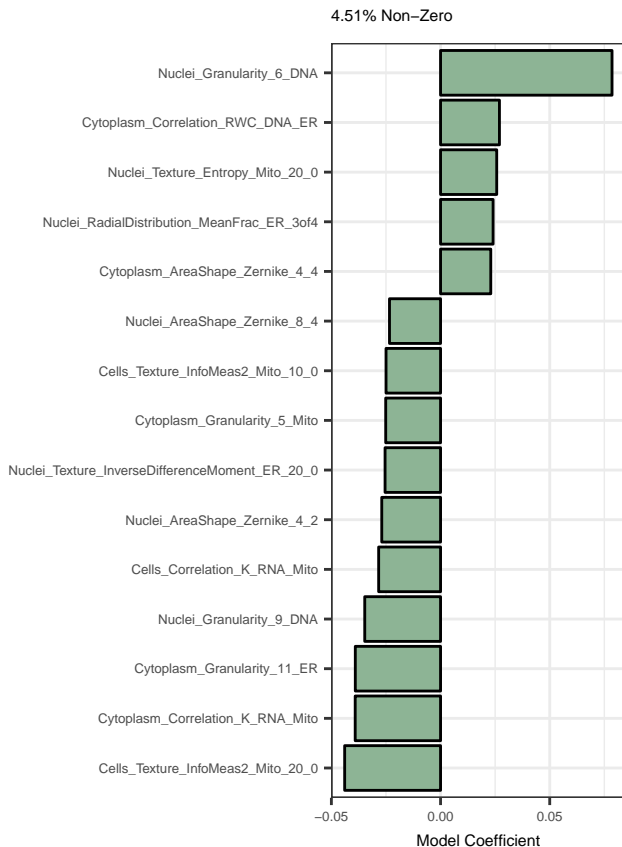


cc_mitosis_ph3_neg_n_spots_per_nucleus_area_mean

0.25% Non-Zero

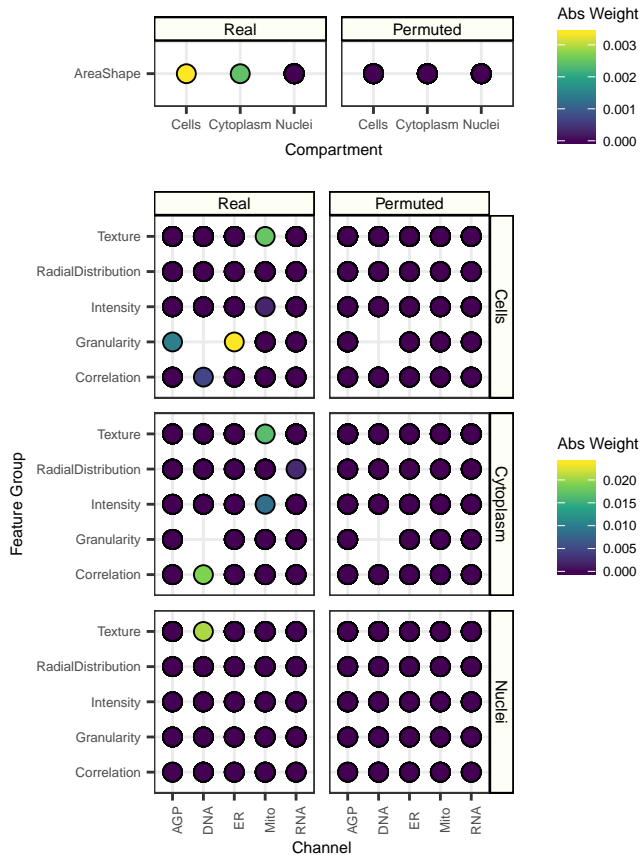
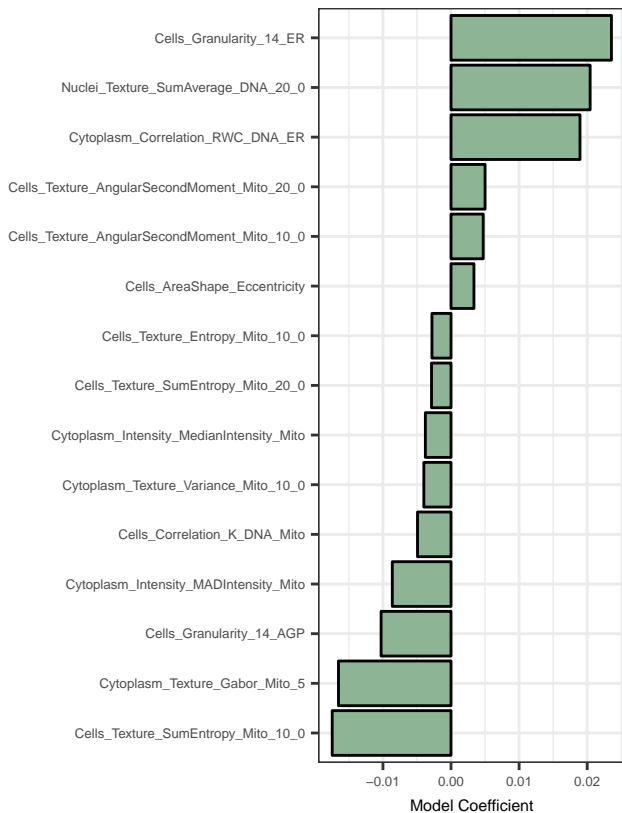


cc_mitosis_ph3_pos_n_objects

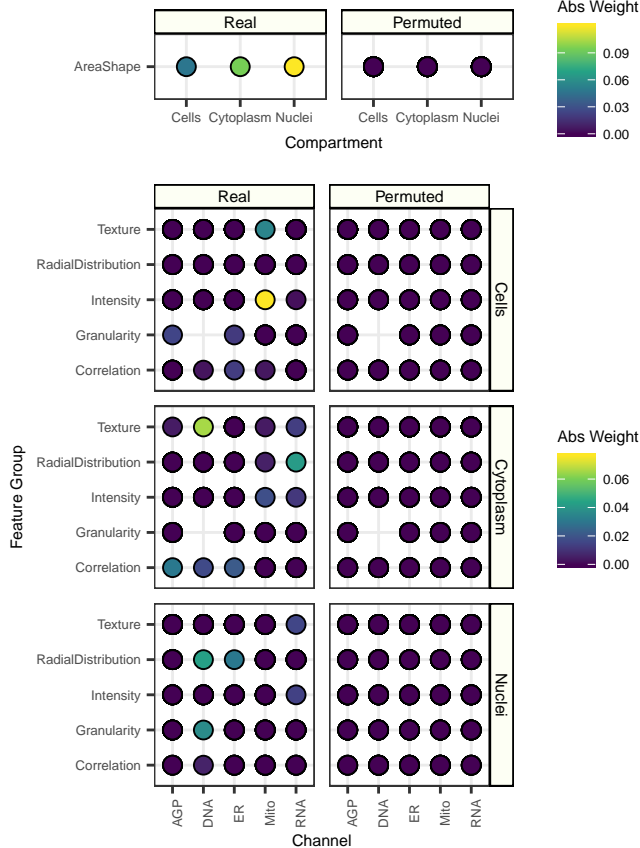
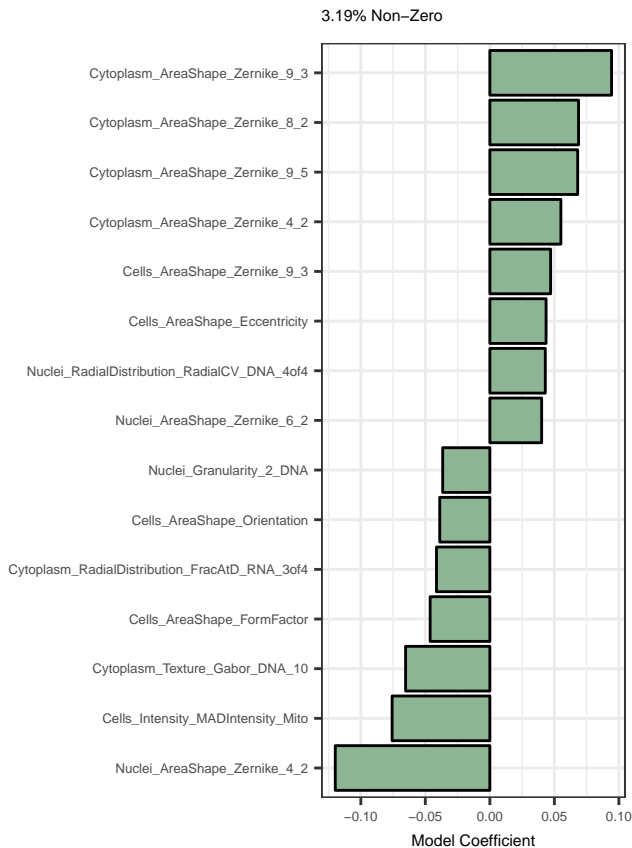


cc_mitosis_ph3_pos_n_spots_mean

1.44% Non-Zero

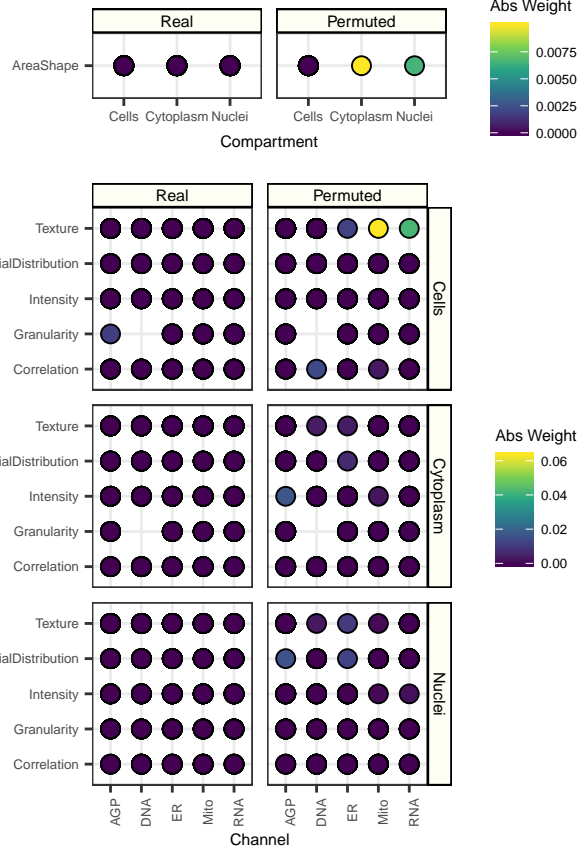


cc_mitosis_ph3_pos_n_spots_per_nucleus_area_mean

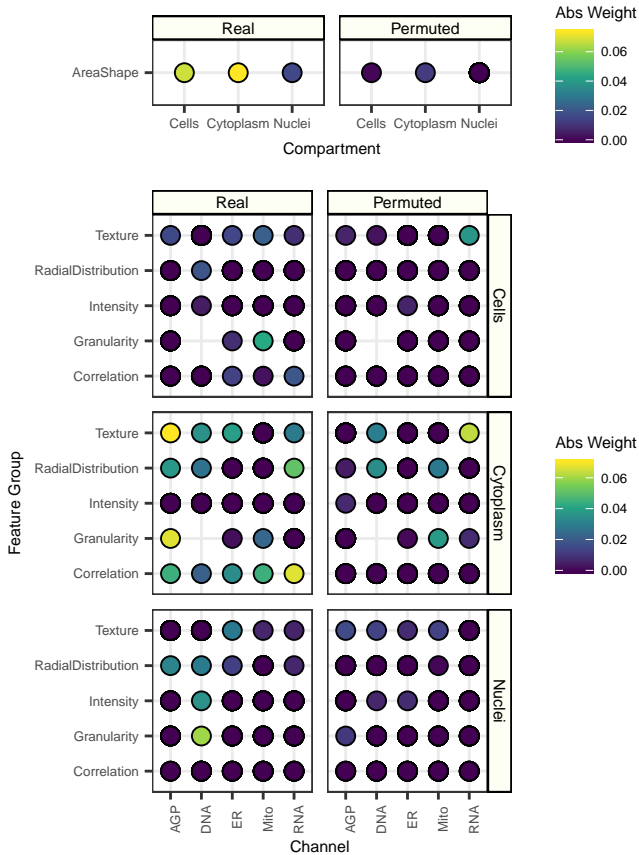
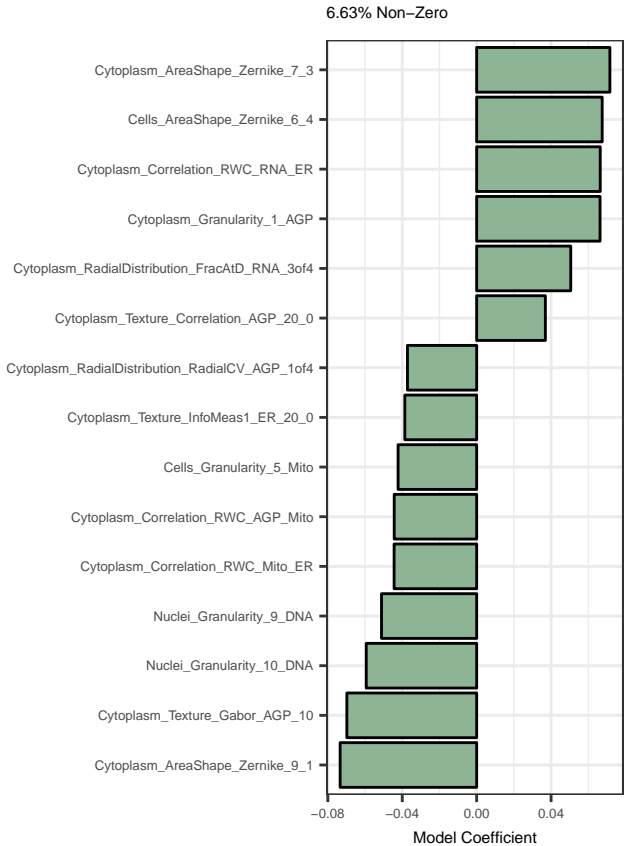


cc_polynuclear_high_n_spots_h2ax_mean

0.06% Non-Zero

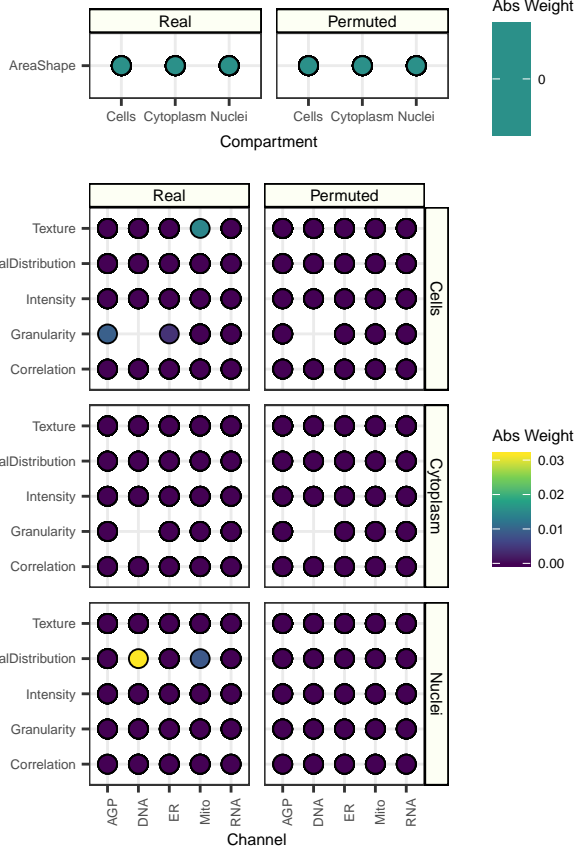


cc_polynuclear_n_objects



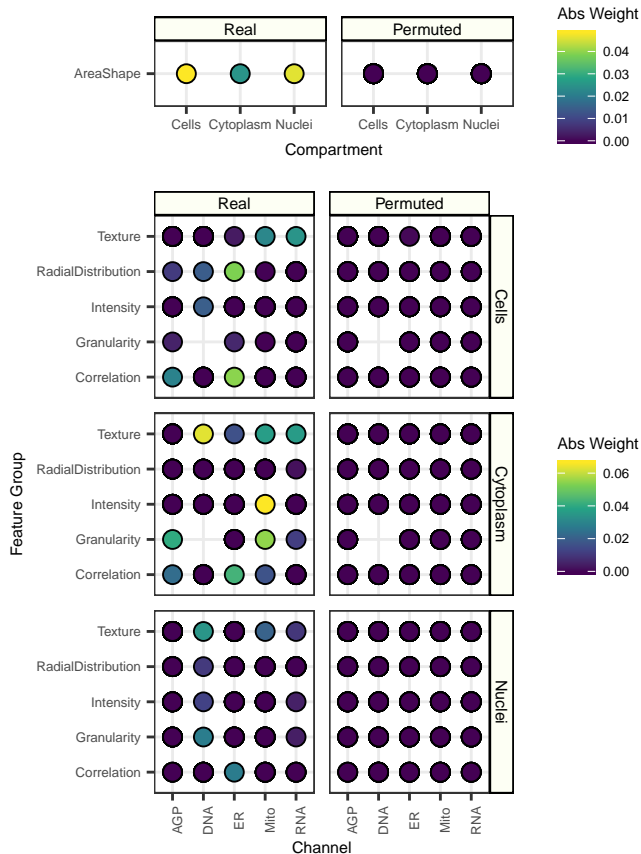
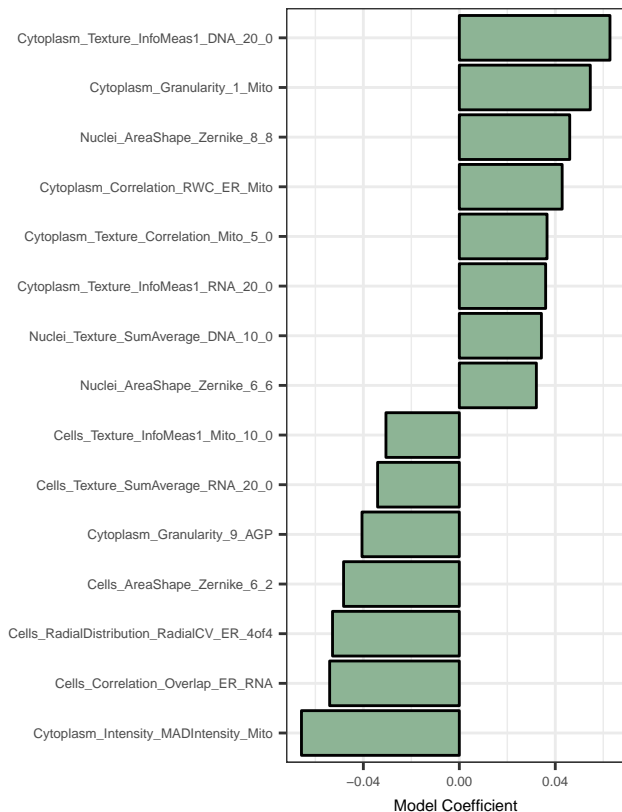
cc_polynuclear_n_spots_mean

0.38% Non-Zero



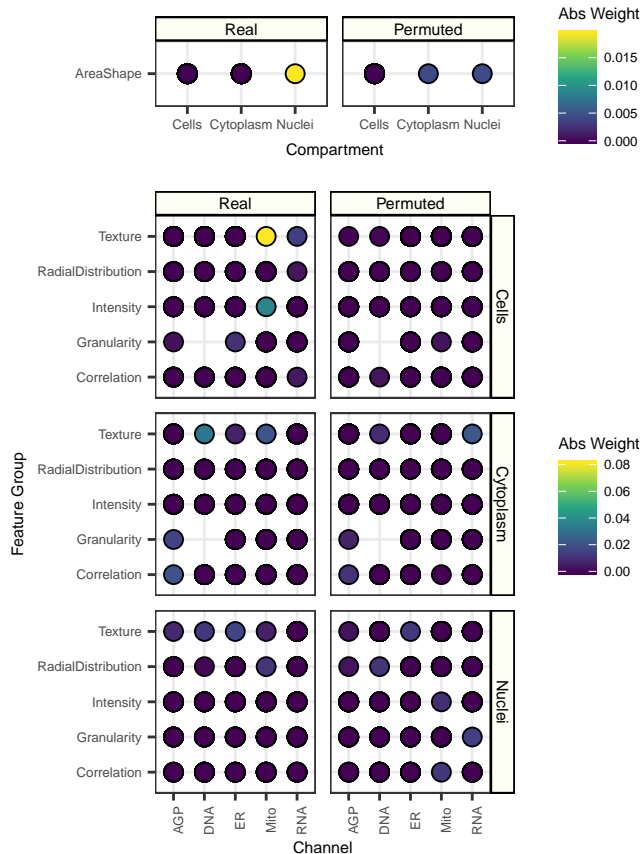
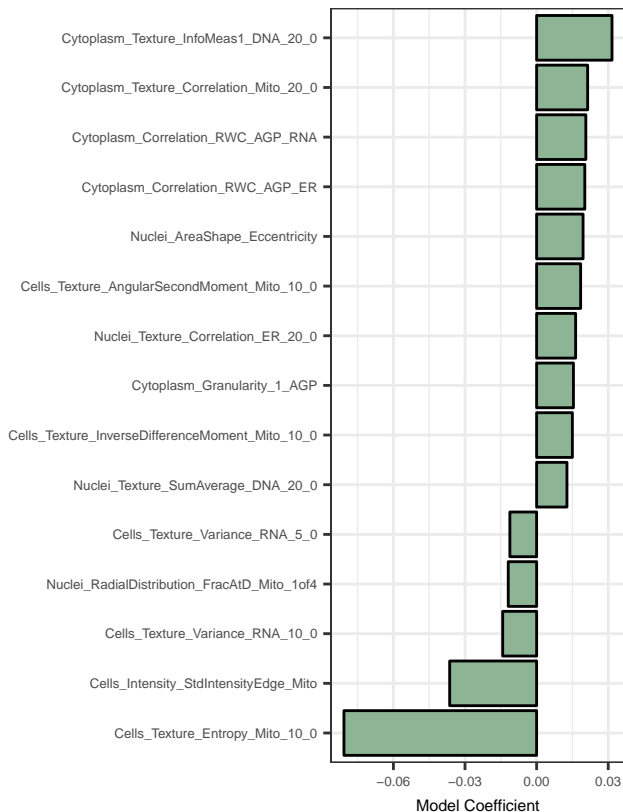
cc_polynuclear_n_spots_per_nucleus_area_mean

4.94% Non-Zero



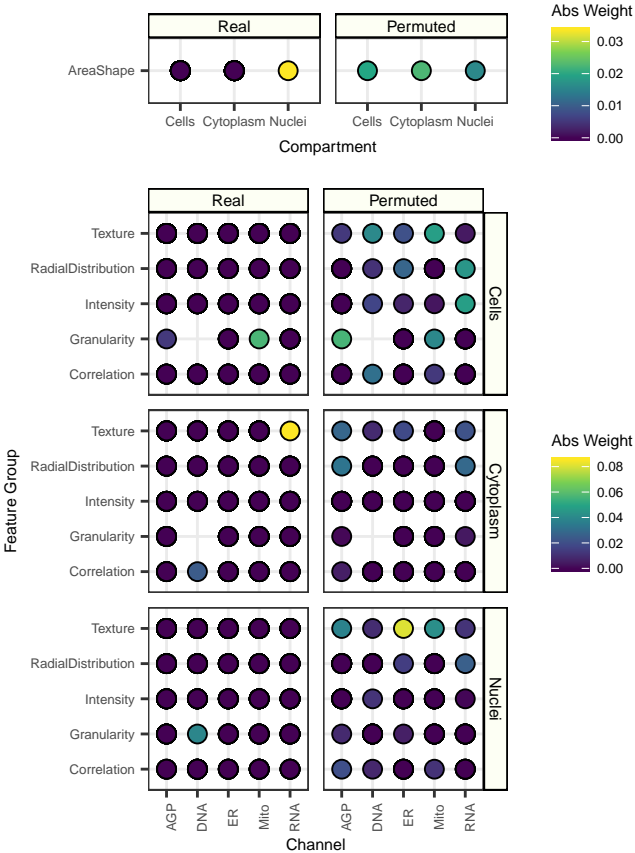
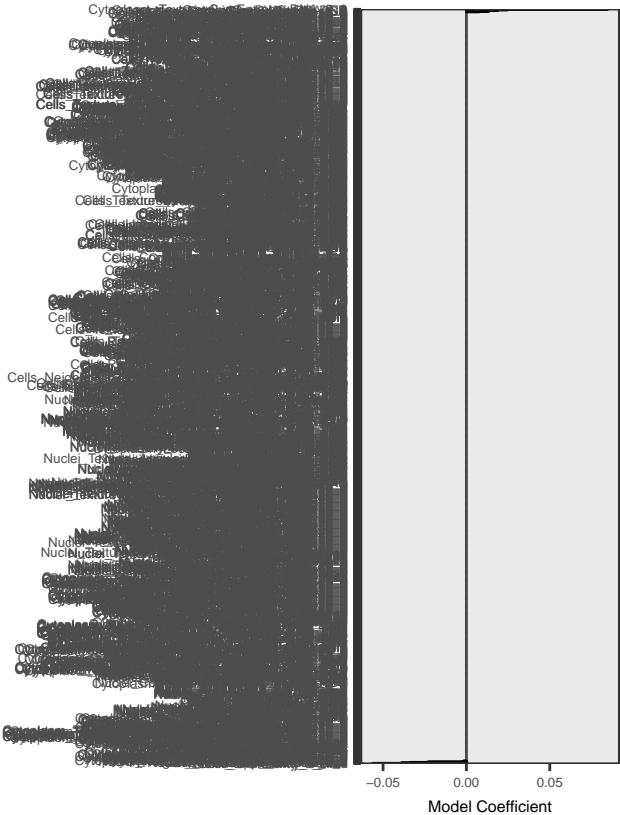
cc_polyploid_high_n_spots_h2ax_mean

2.13% Non-Zero

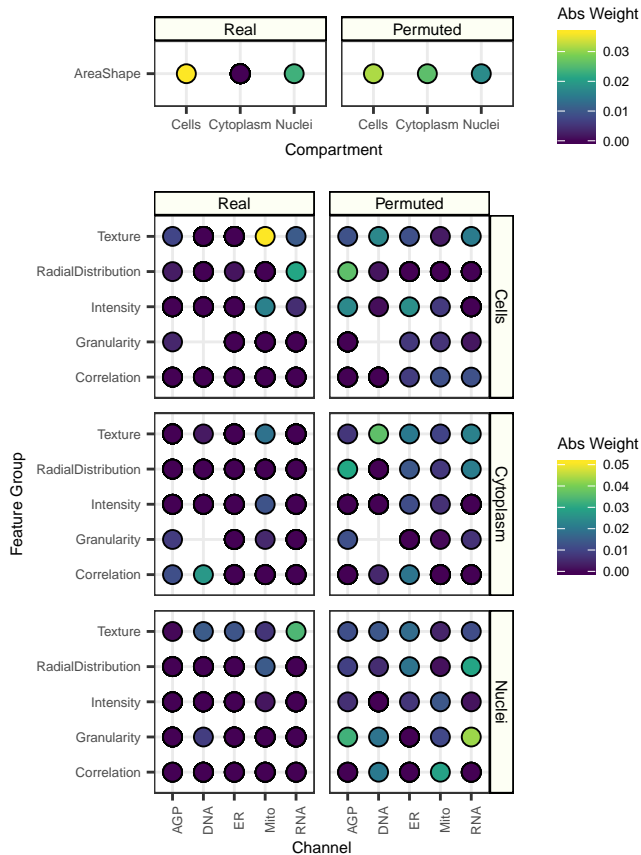
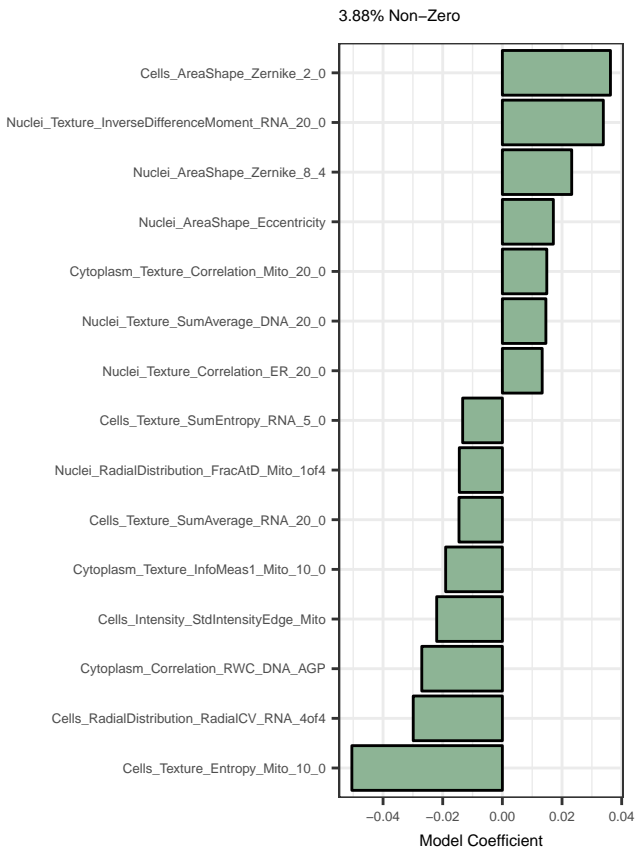


cc_polyploid_n_objects

0.75% Non-Zero

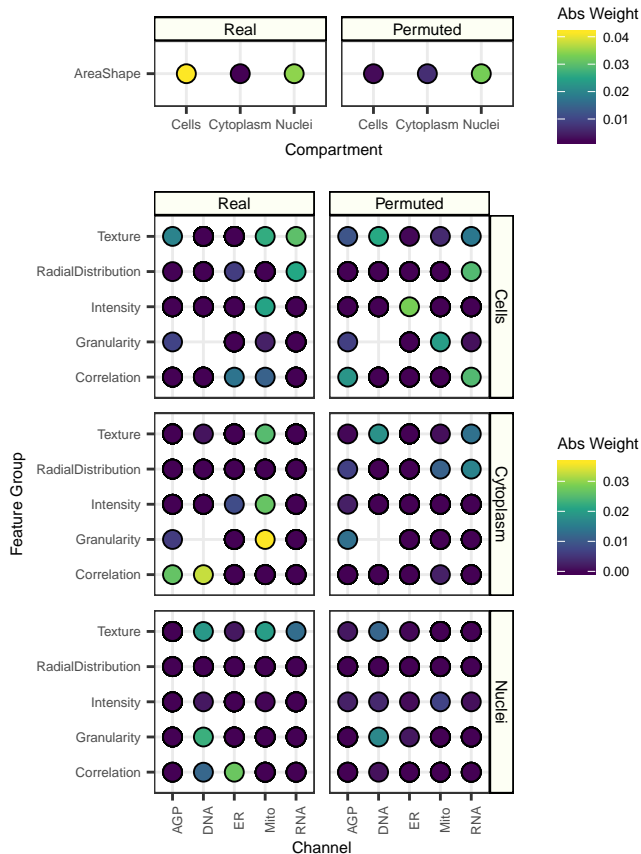
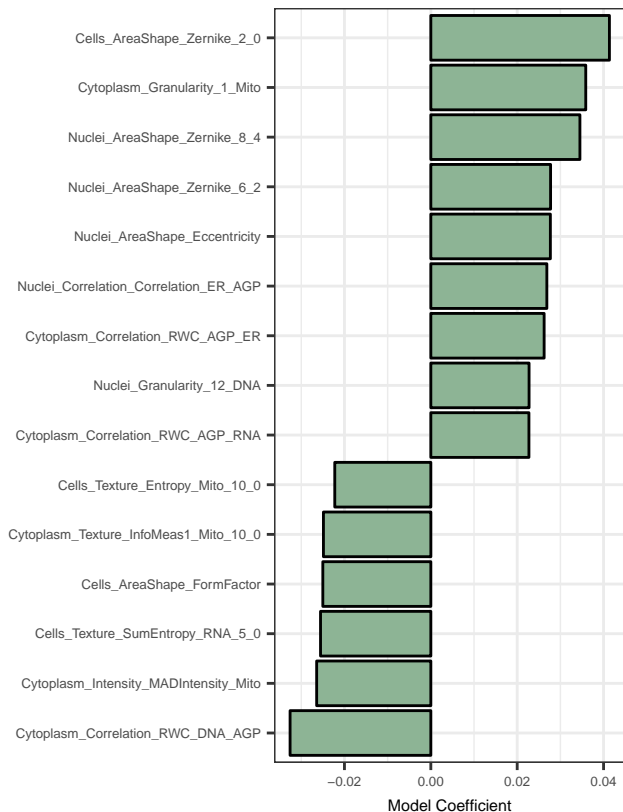


cc_polyloid_n_spots_mean



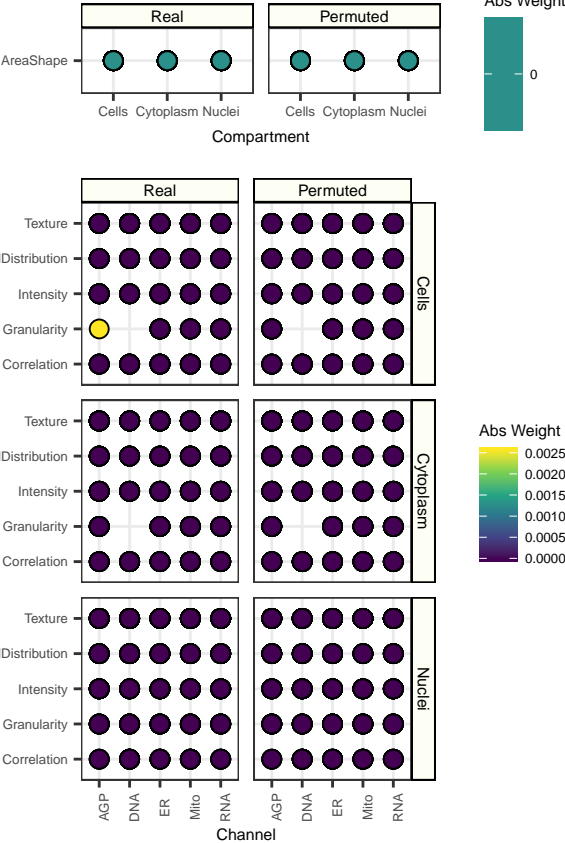
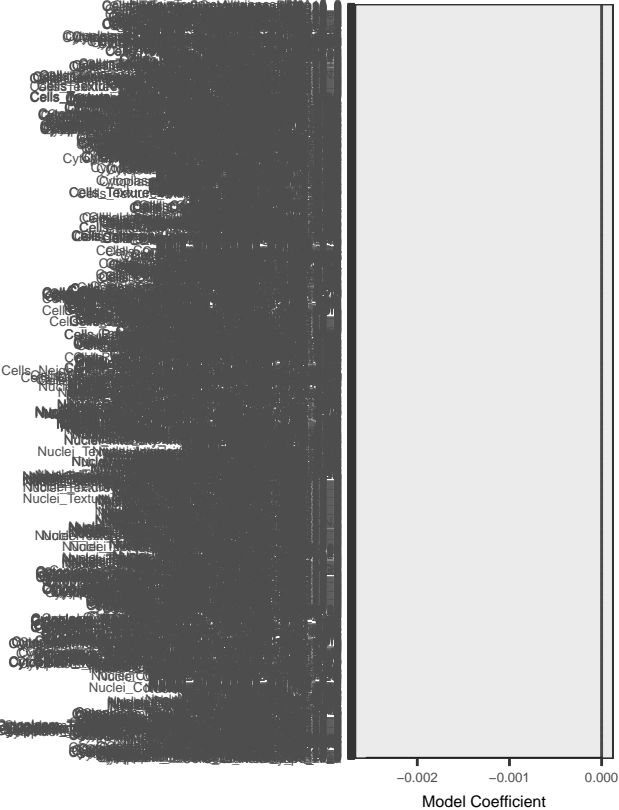
cc_polyloid_n_spots_per_nucleus_area_mean

5.01% Non-Zero

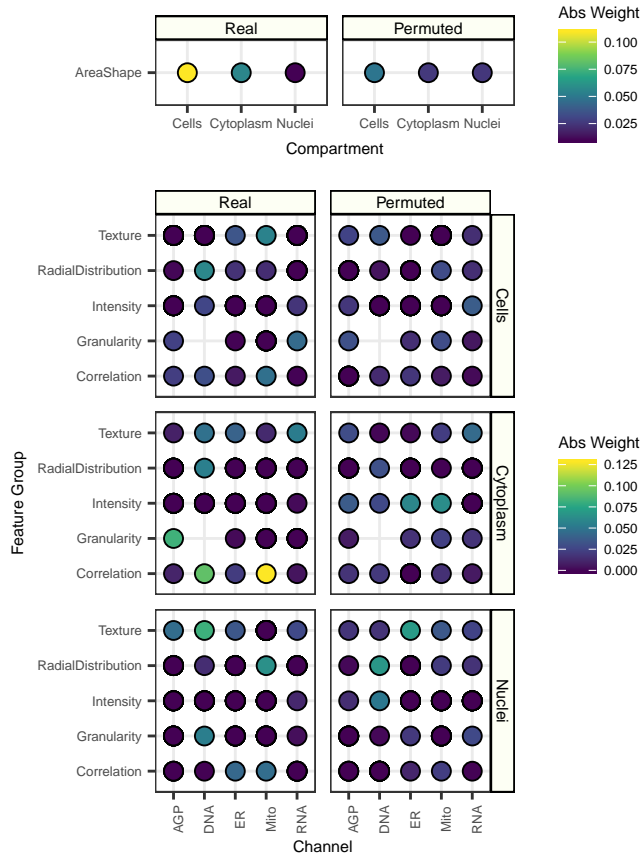


vb_infection_percentage

0.06% Non-Zero

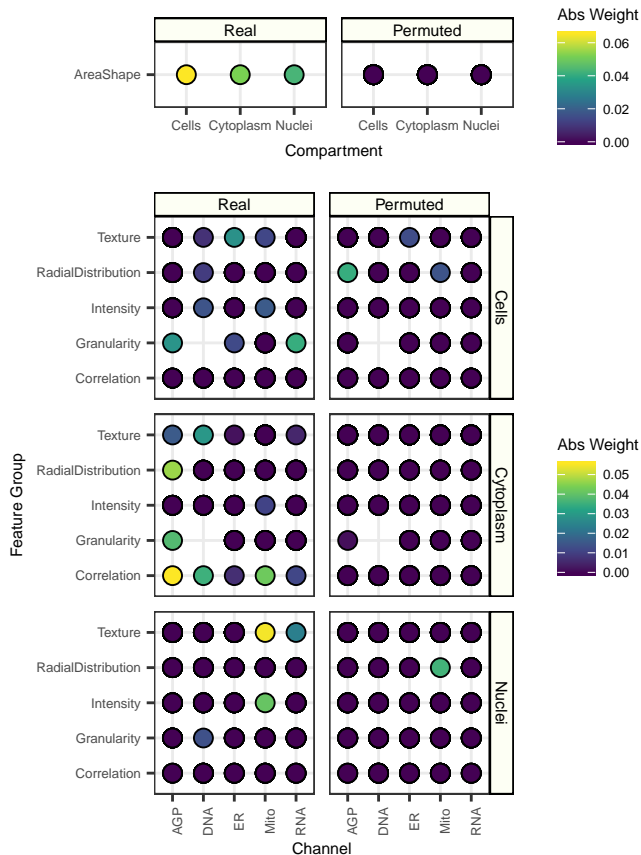
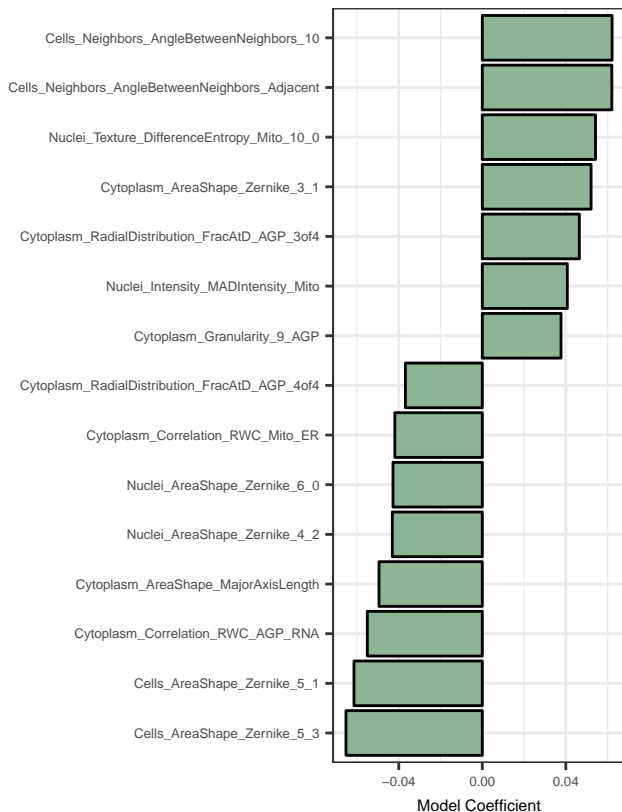


vb_live_cell_area



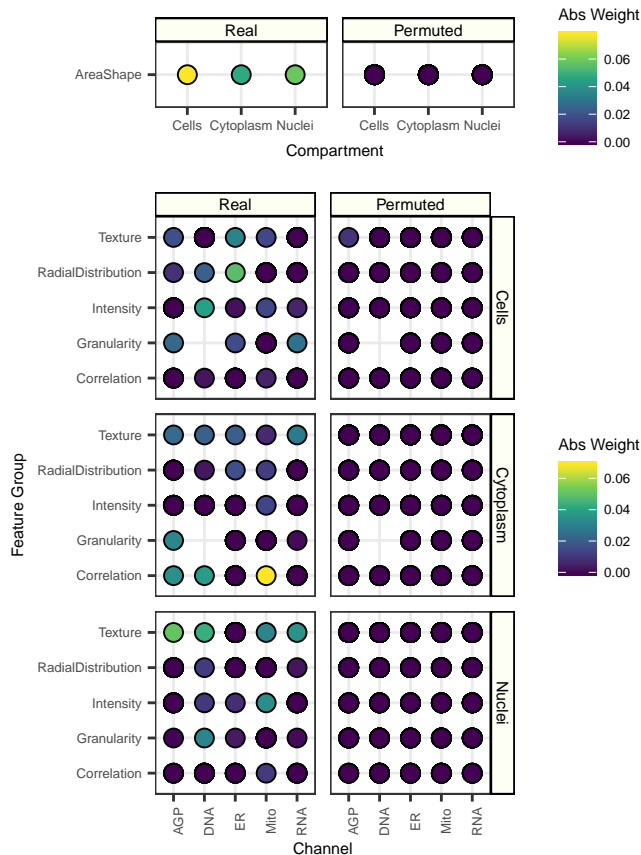
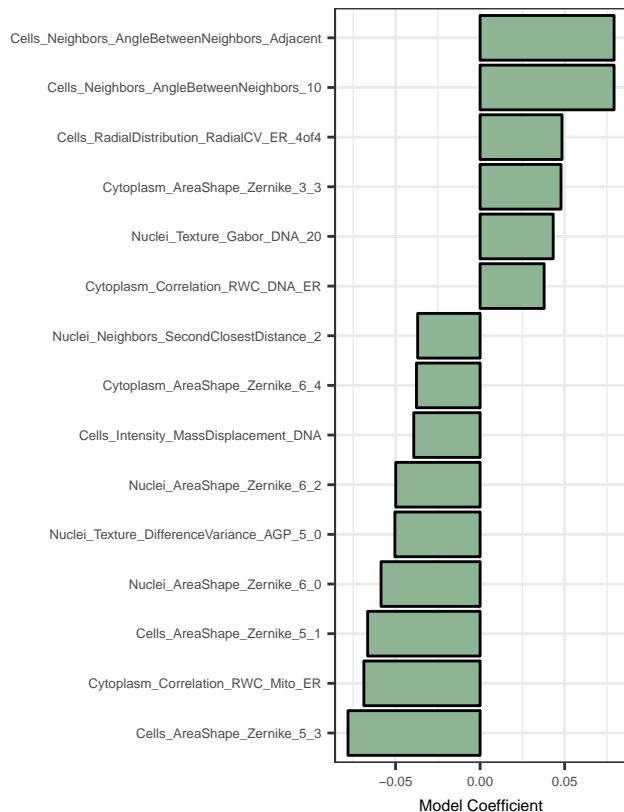
vb_live_cell_roundness

4.07% Non-Zero

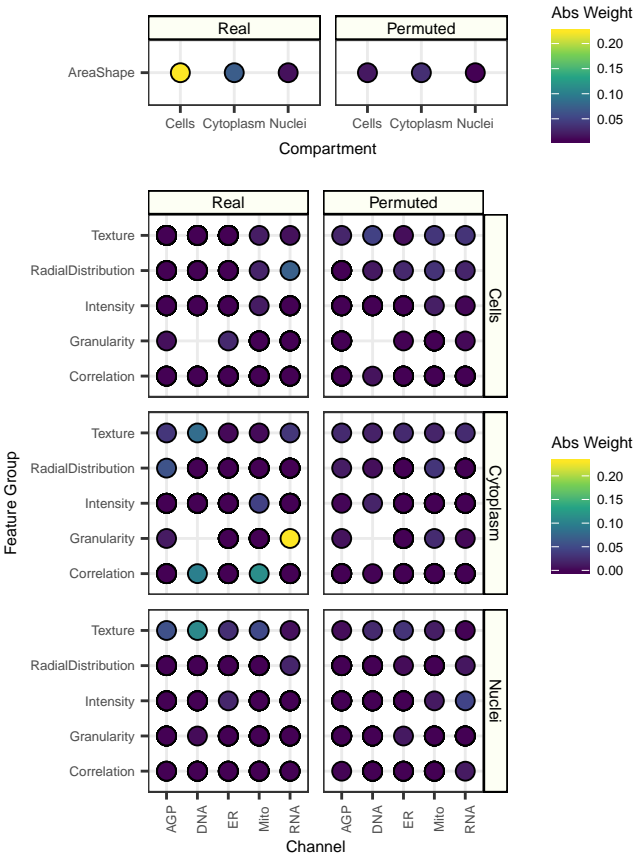
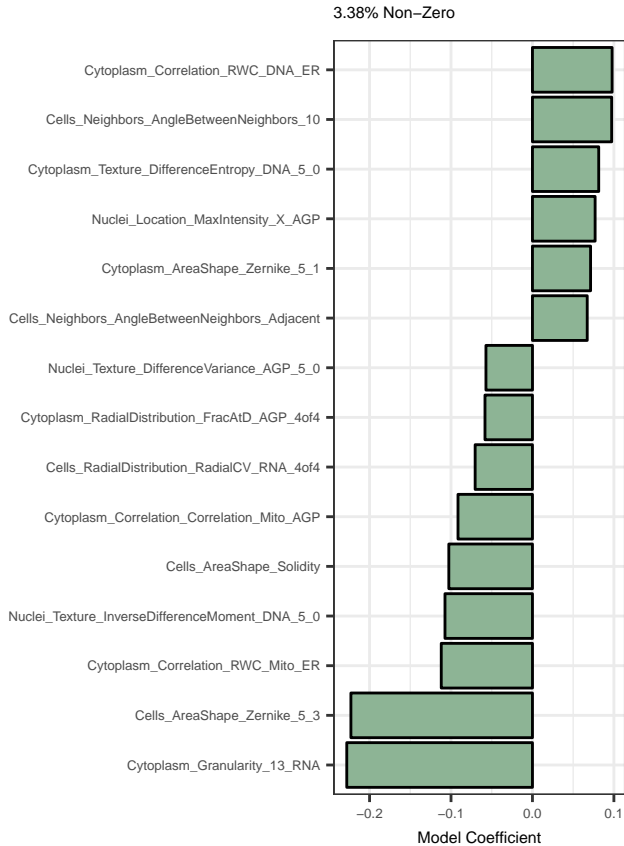


vb_live_cell_width_length

10.01% Non-Zero

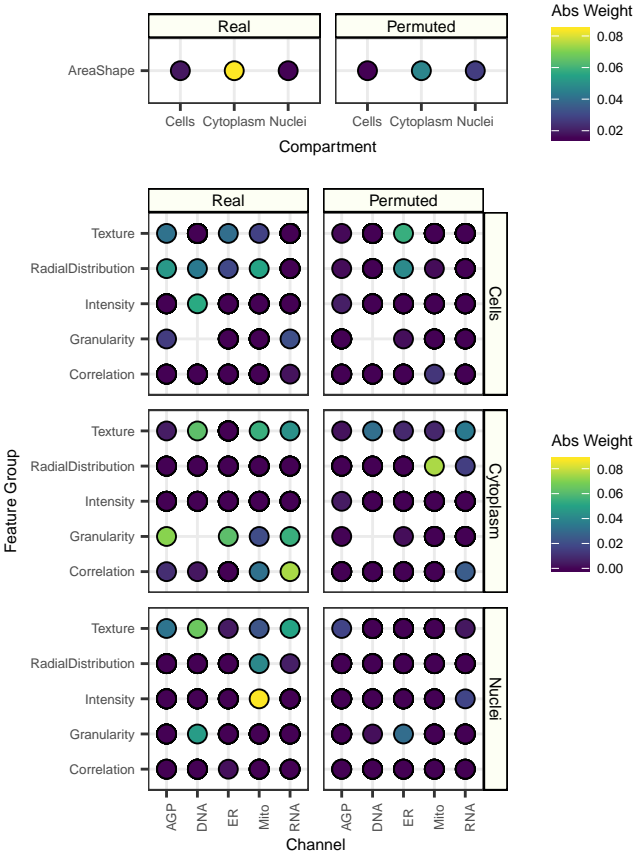
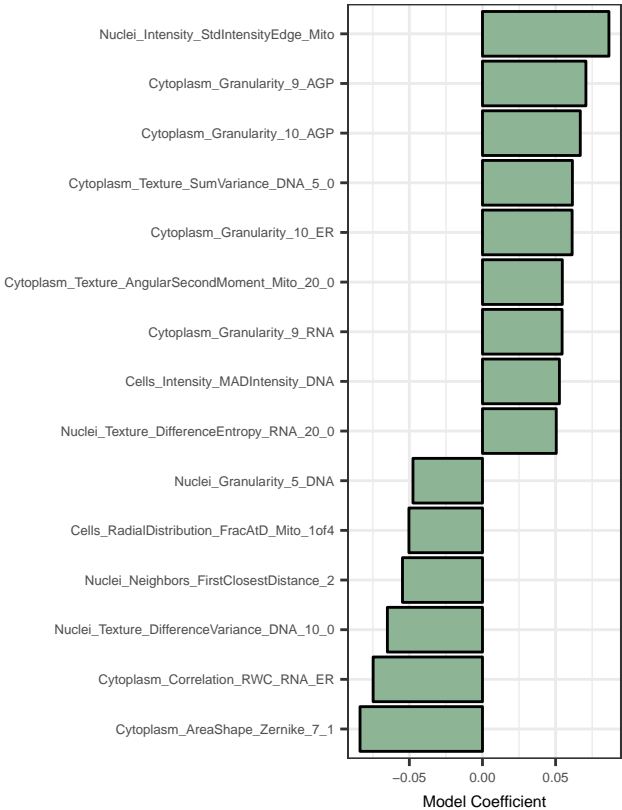


vb_num_live_cells



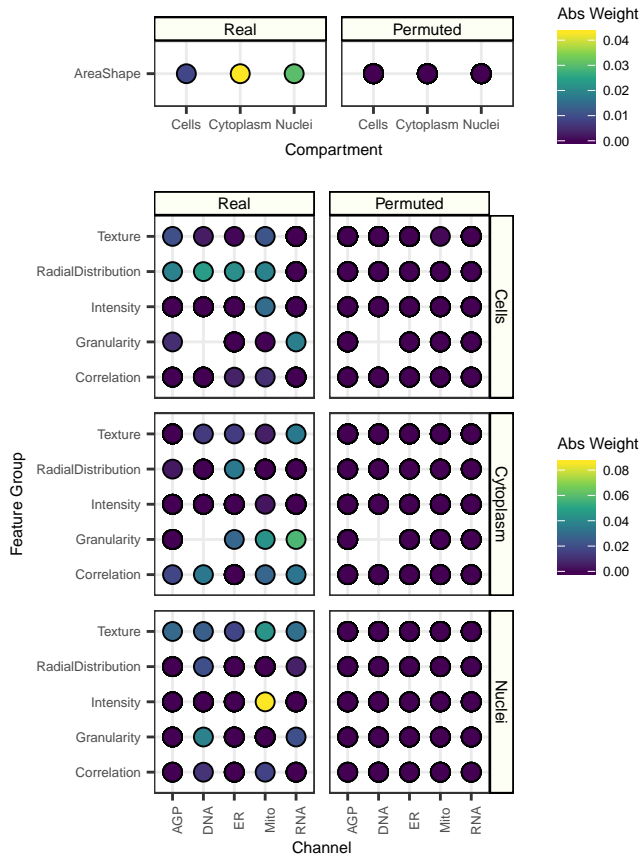
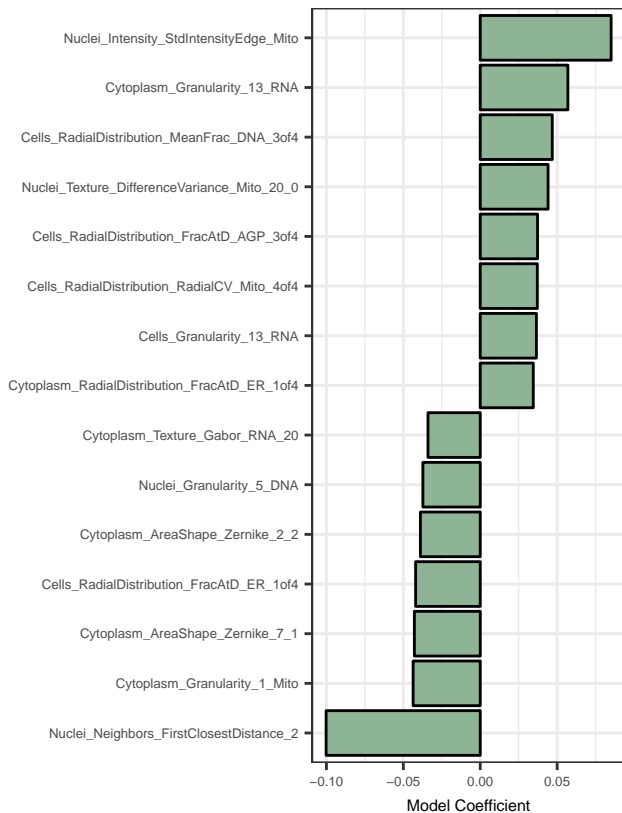
vb_percent_all_apoptosis

4.63% Non-Zero



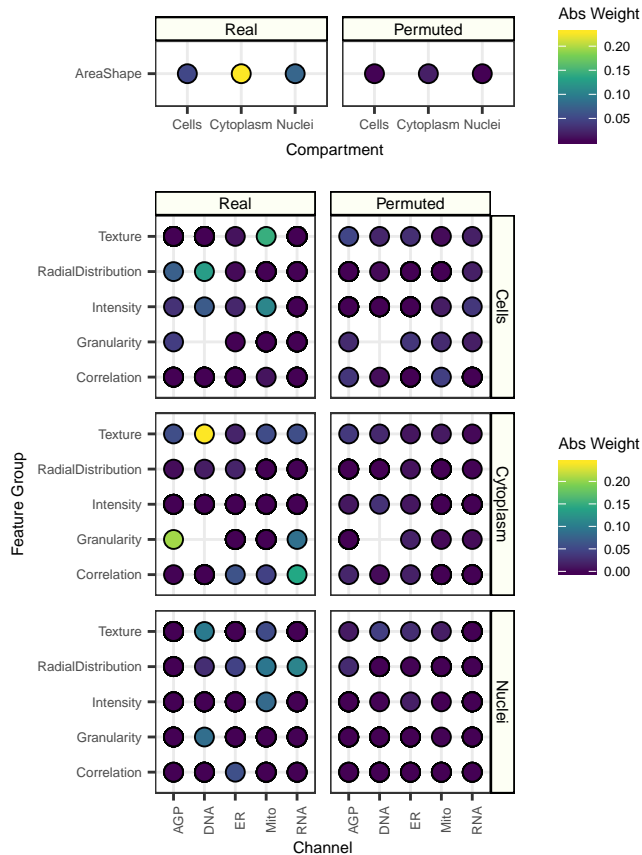
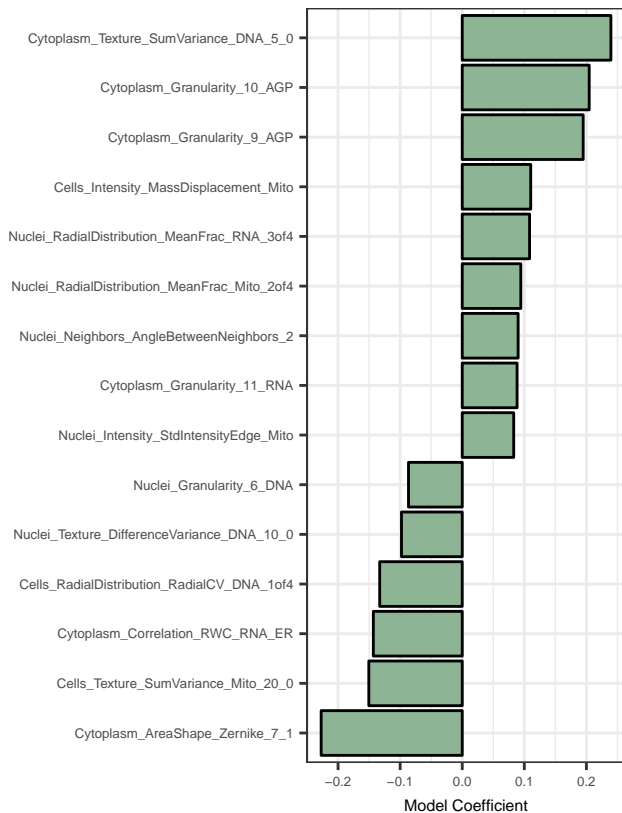
vb_percent_all_early_apoptosis

7.01% Non-Zero



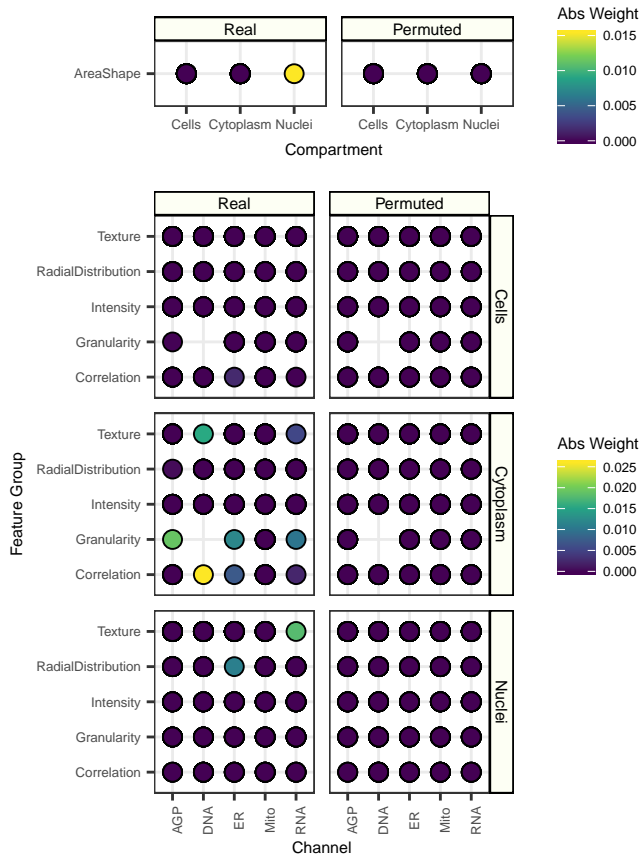
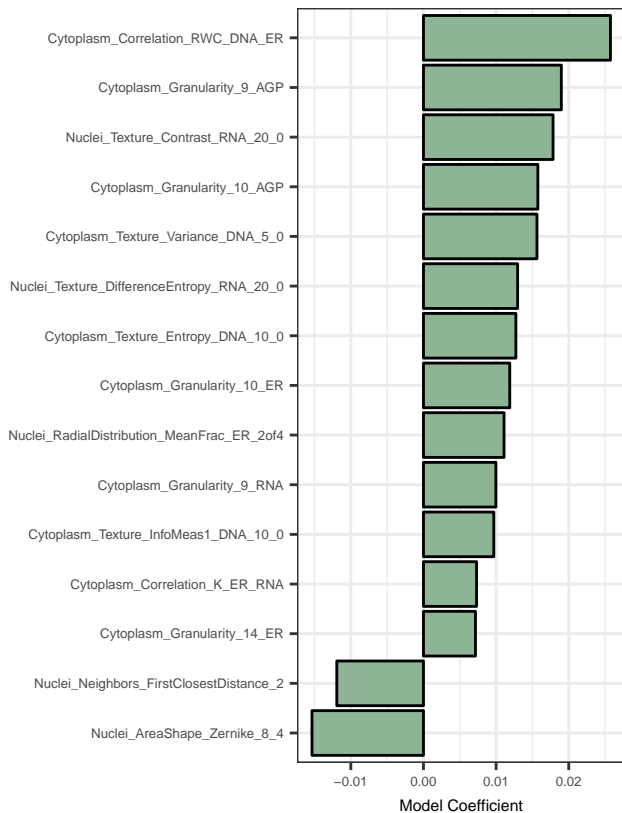
vb_percent_all_late_apoptosis

4.32% Non-Zero

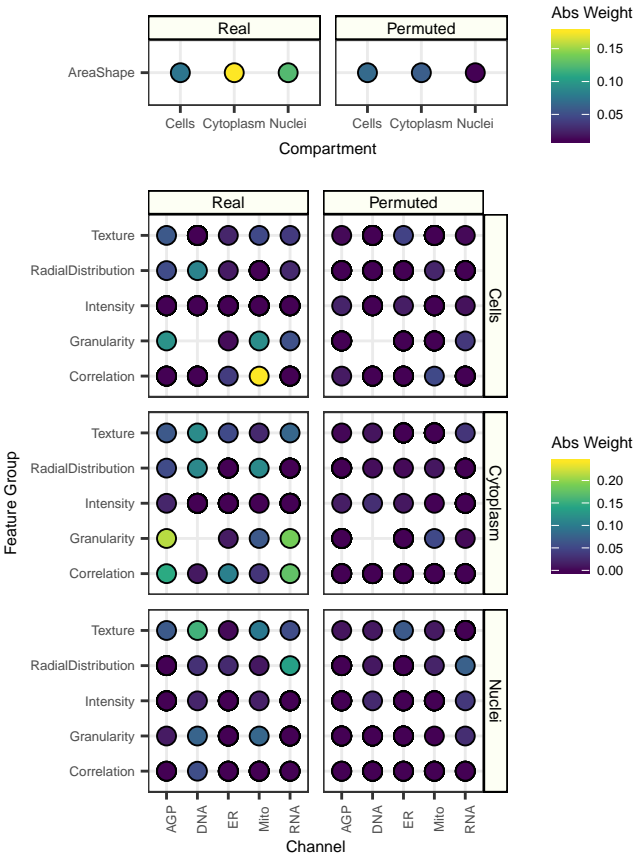
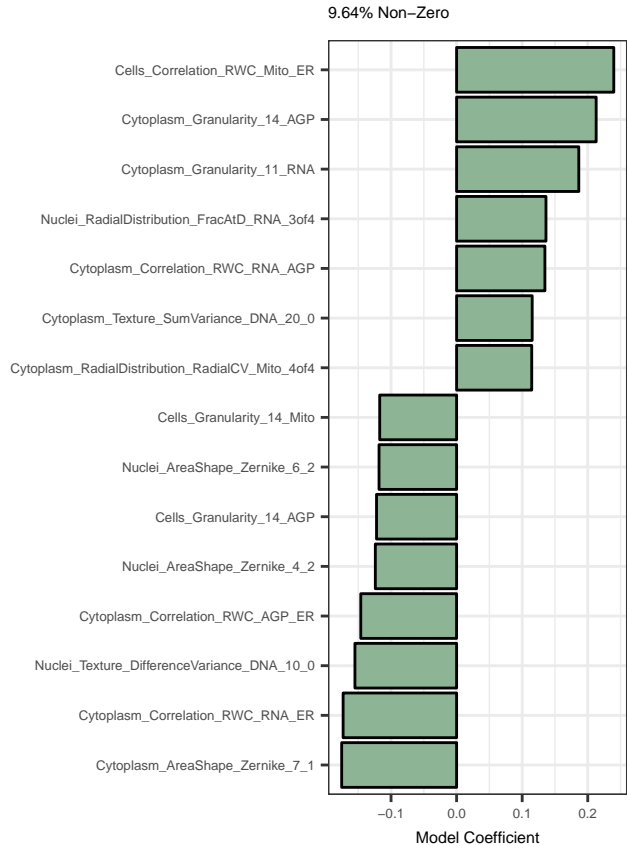


vb_percent_caspase_dead_only

1.94% Non-Zero

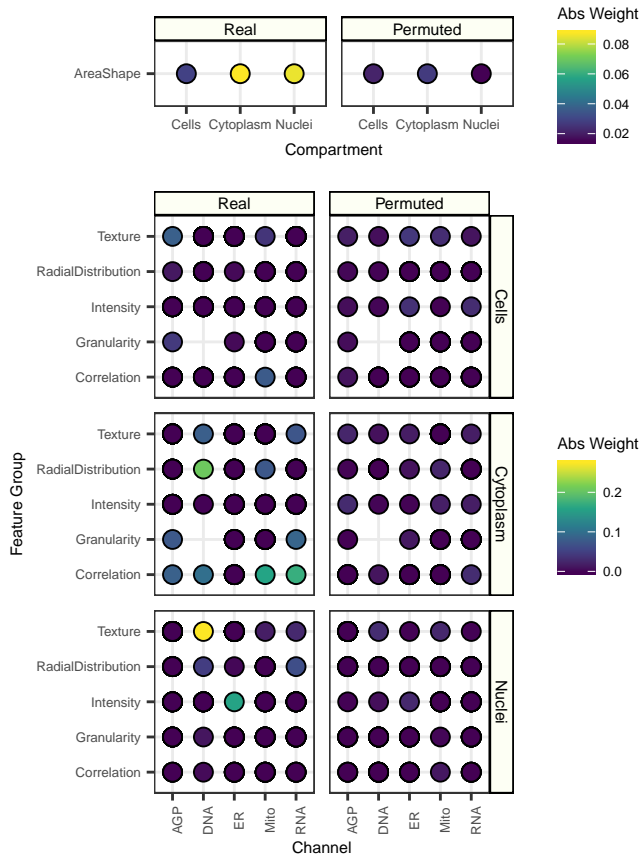
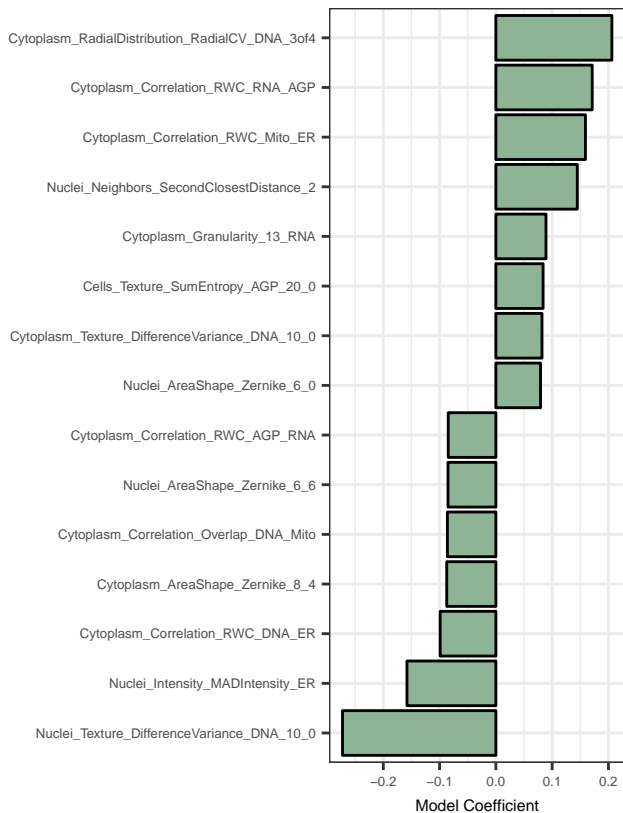


vb_percent_dead

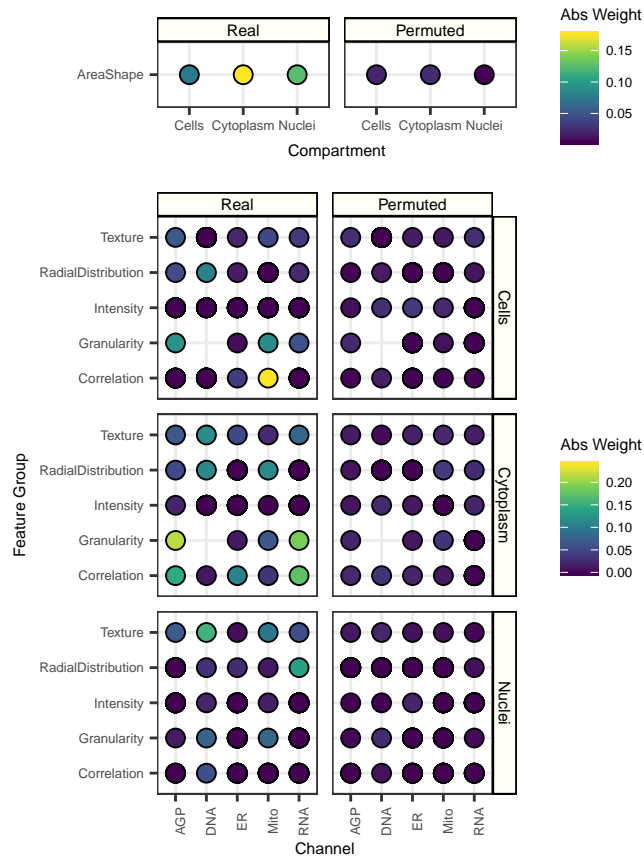
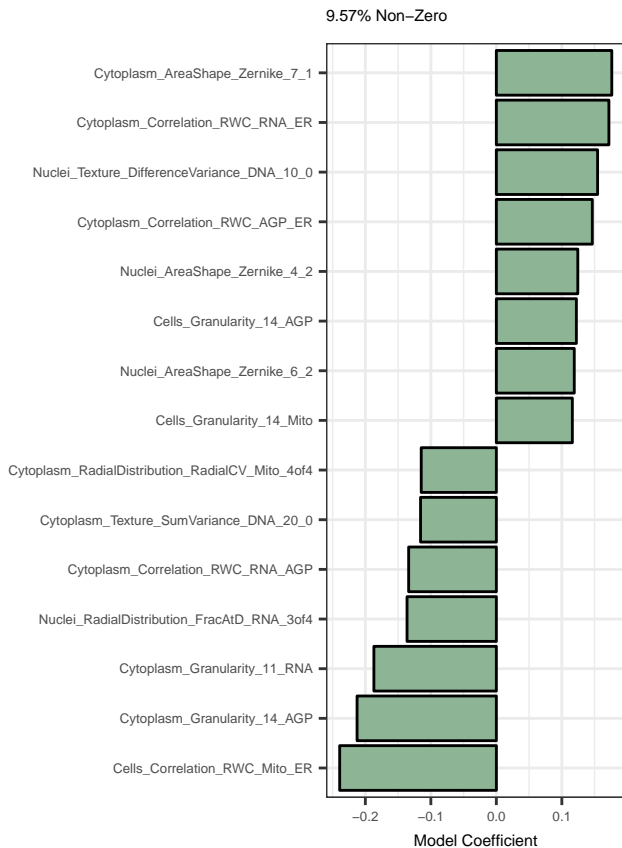


vb_percent_dead_only

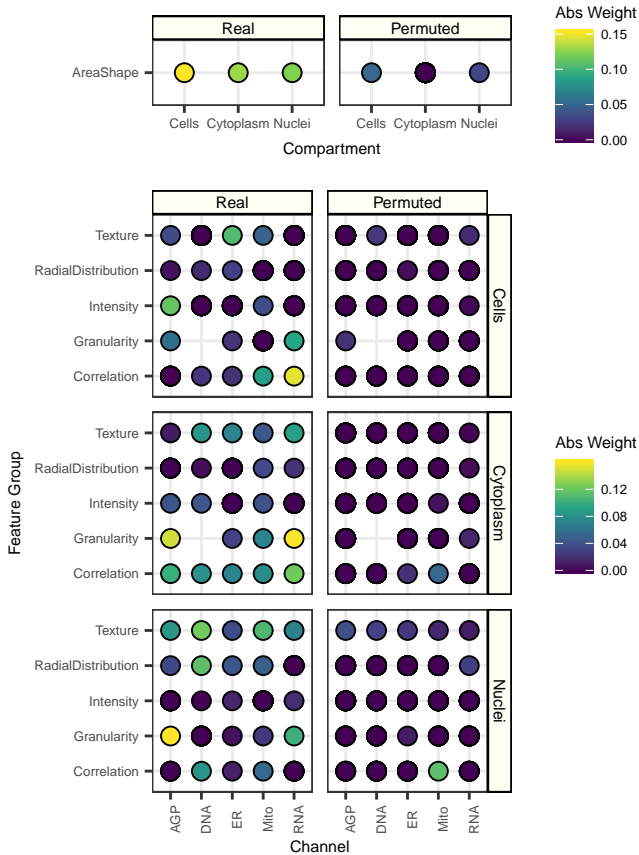
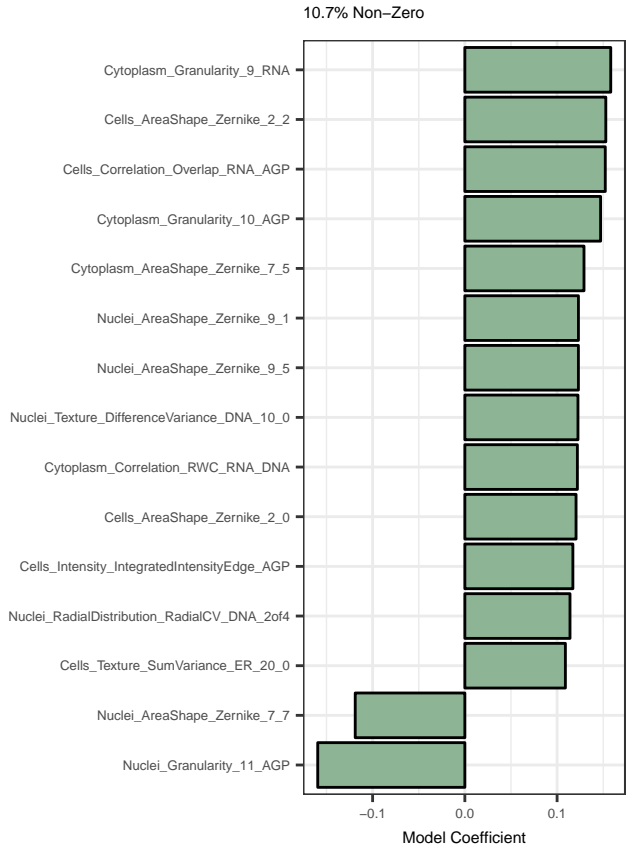
3.88% Non-Zero



vb_percent_live



vb_ros_back_mean



vb_ros_mean

