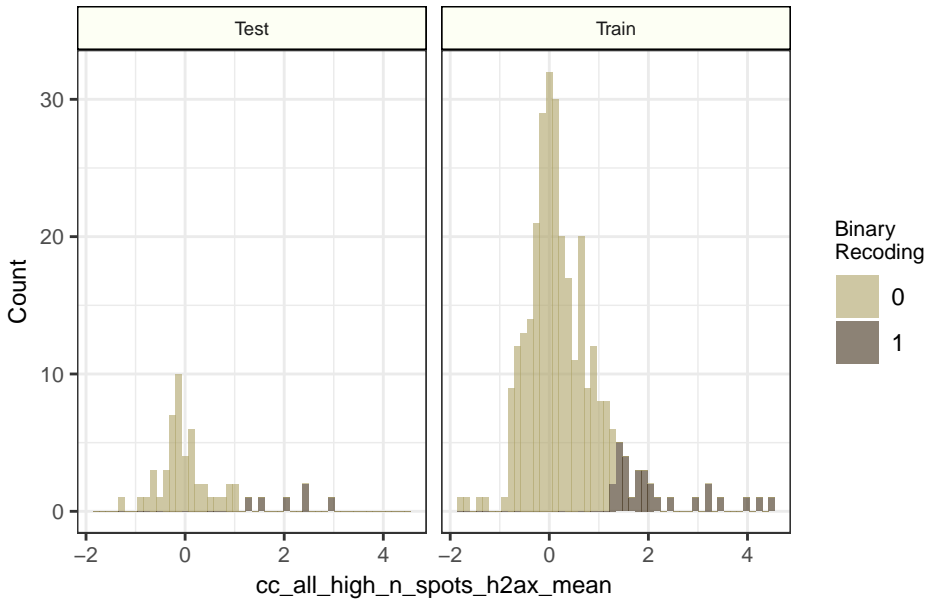
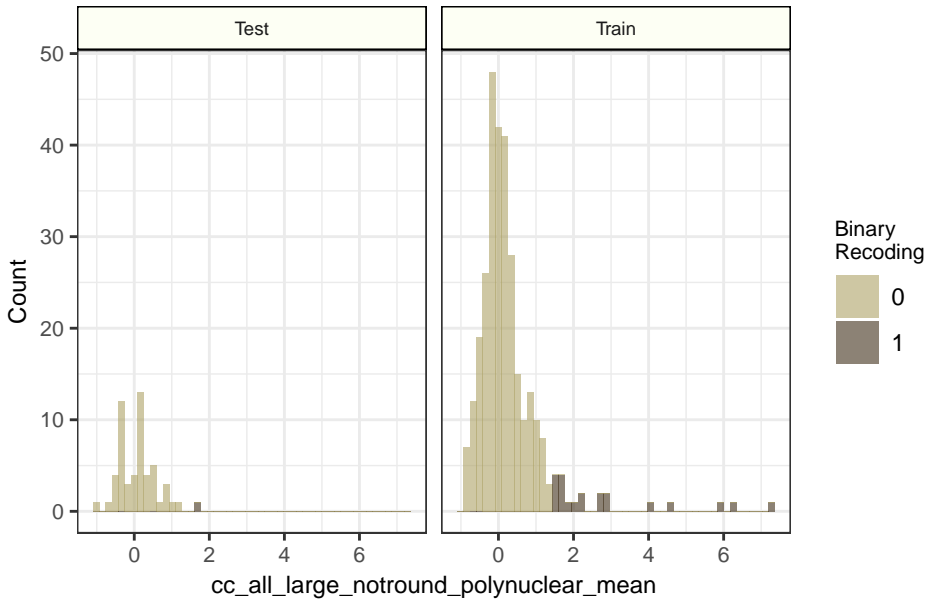


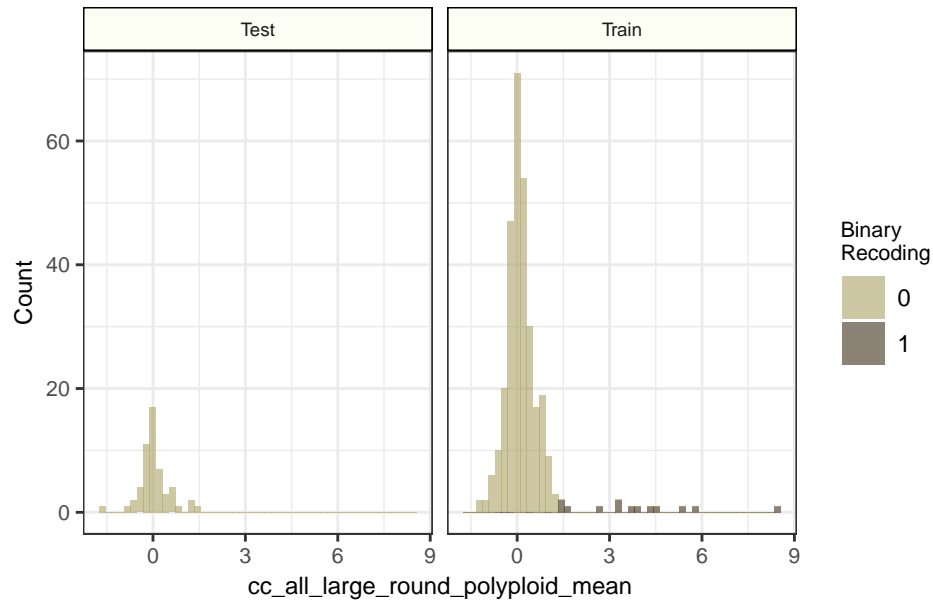
cc_all_high_n_spots_h2ax_mean



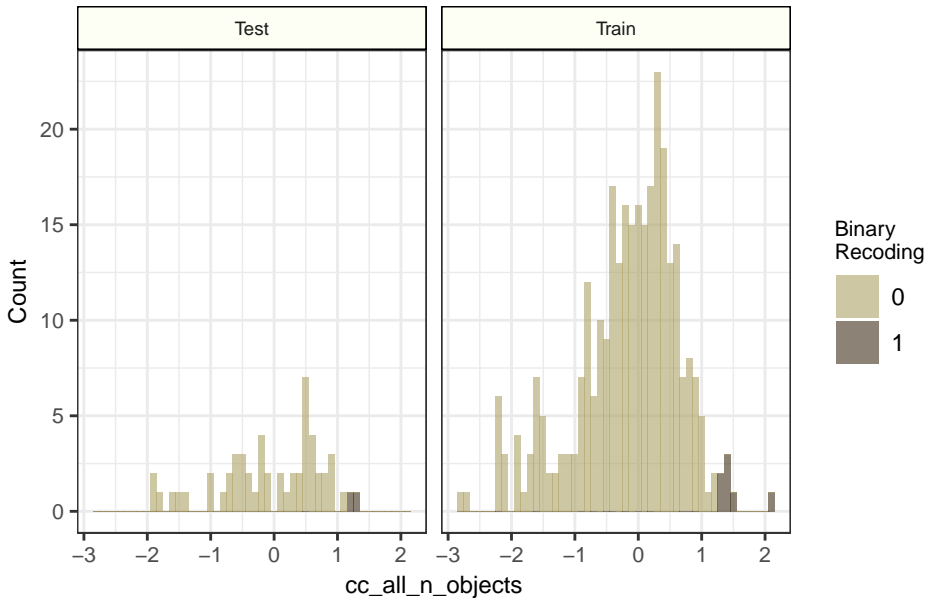
cc_all_large_notround_polynuclear_mean



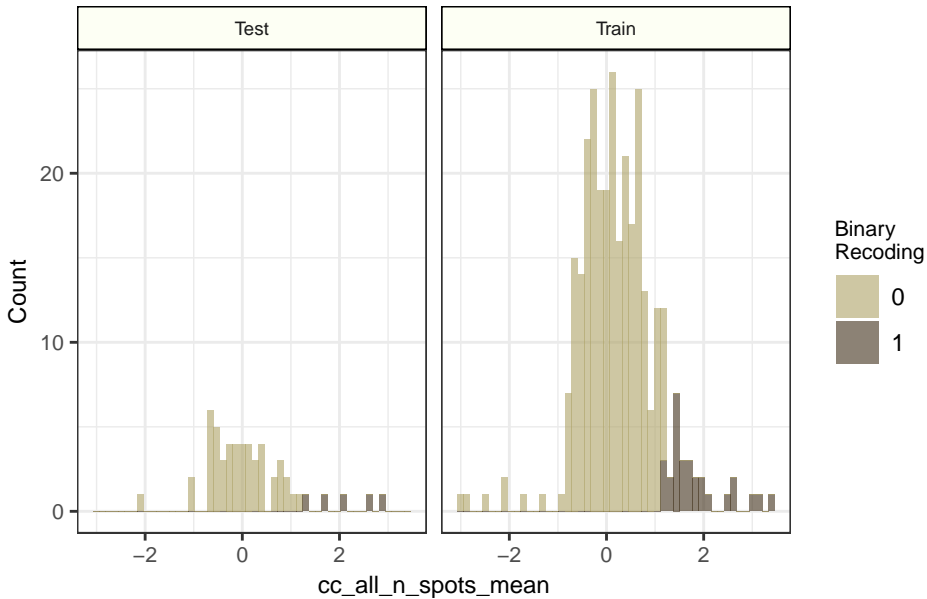
cc_all_large_round_polyploid_mean



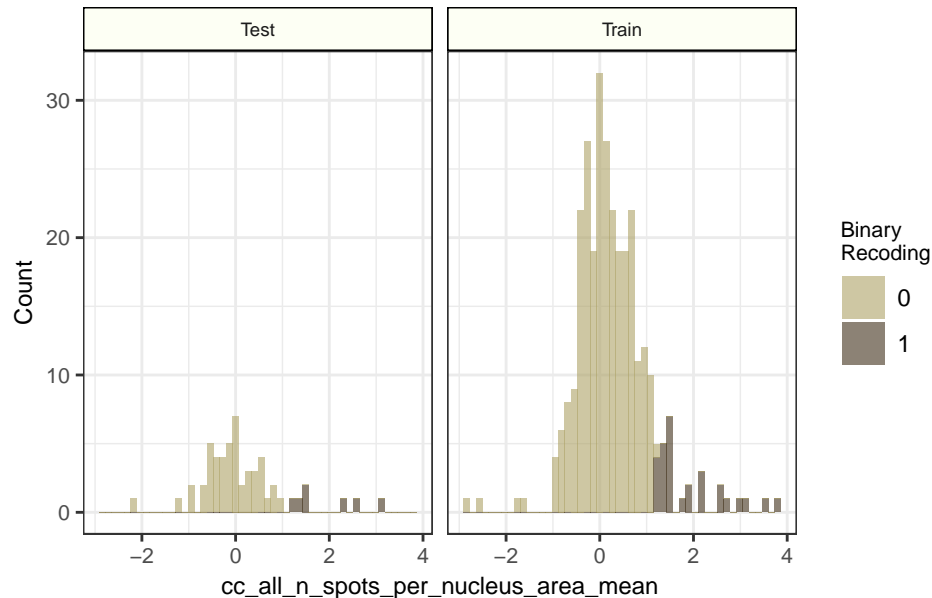
cc_all_n_objects



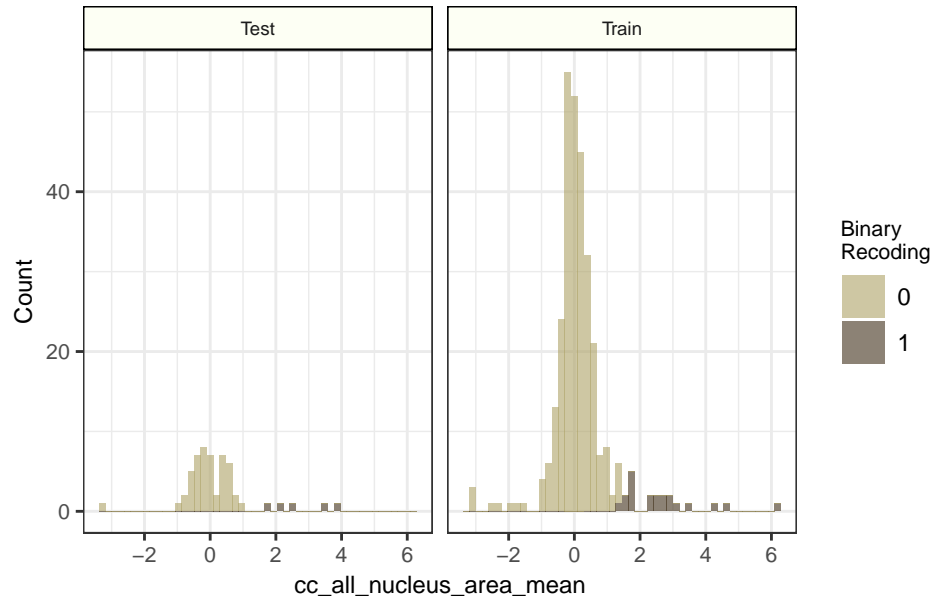
cc_all_n_spots_mean



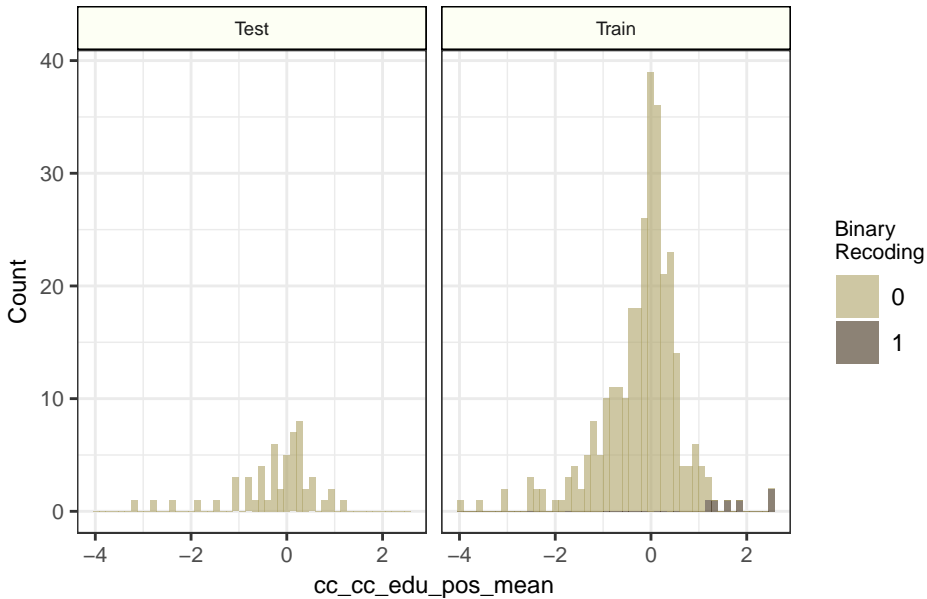
cc_all_n_spots_per_nucleus_area_mean



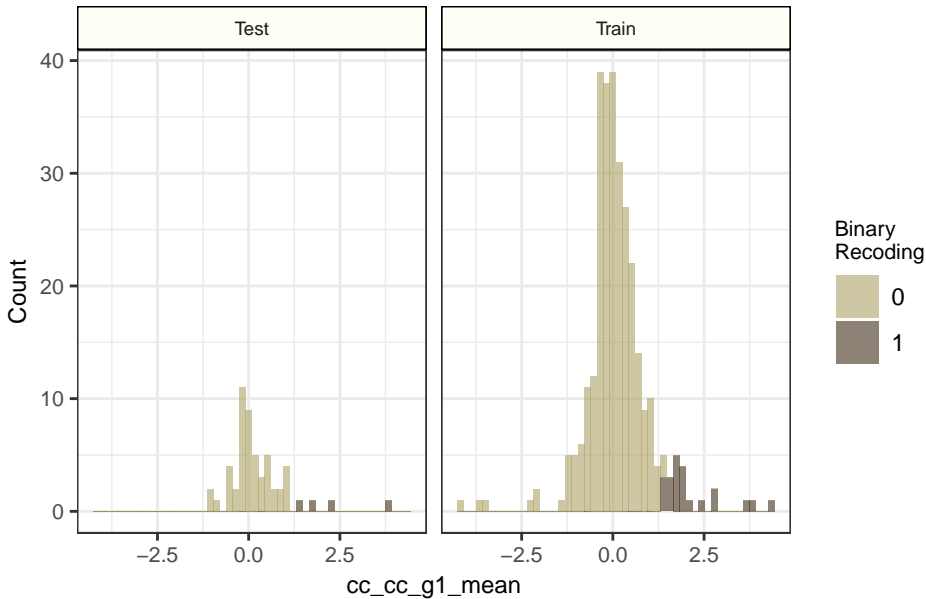
cc_all_nucleus_area_mean



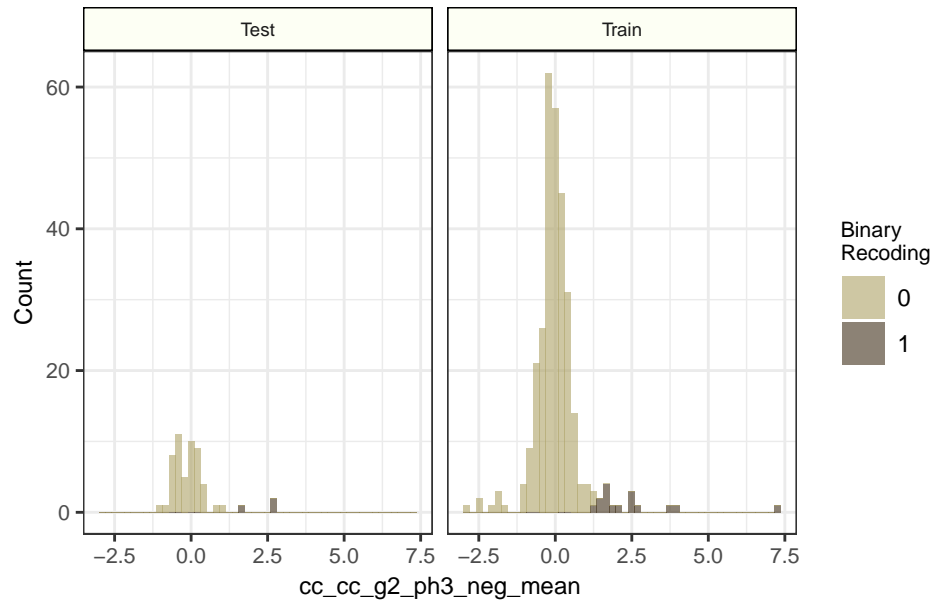
cc_cc_edu_pos_mean



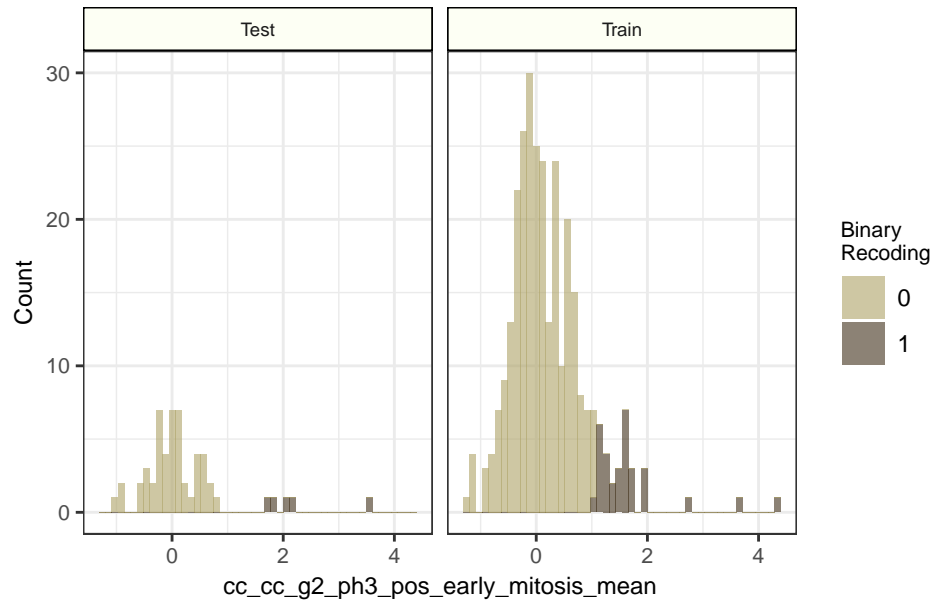
cc_cc_g1_mean



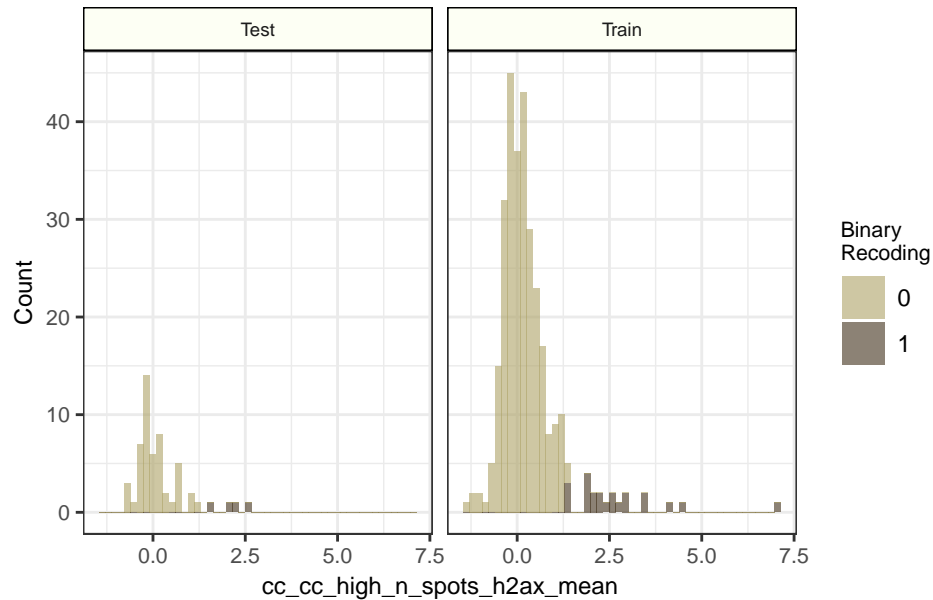
cc_cc_g2_ph3_neg_mean



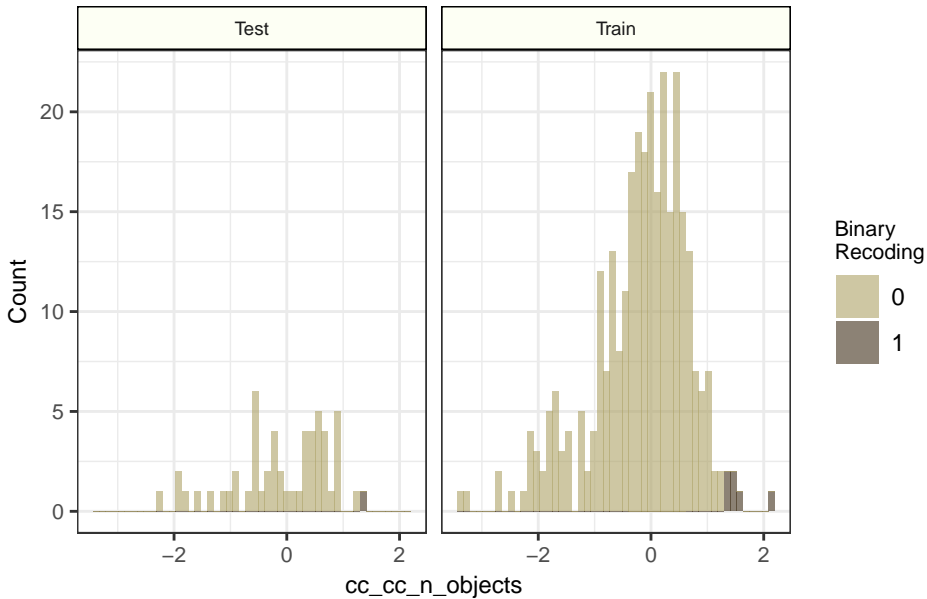
cc_cc_g2_ph3_pos_early_mitosis_mean



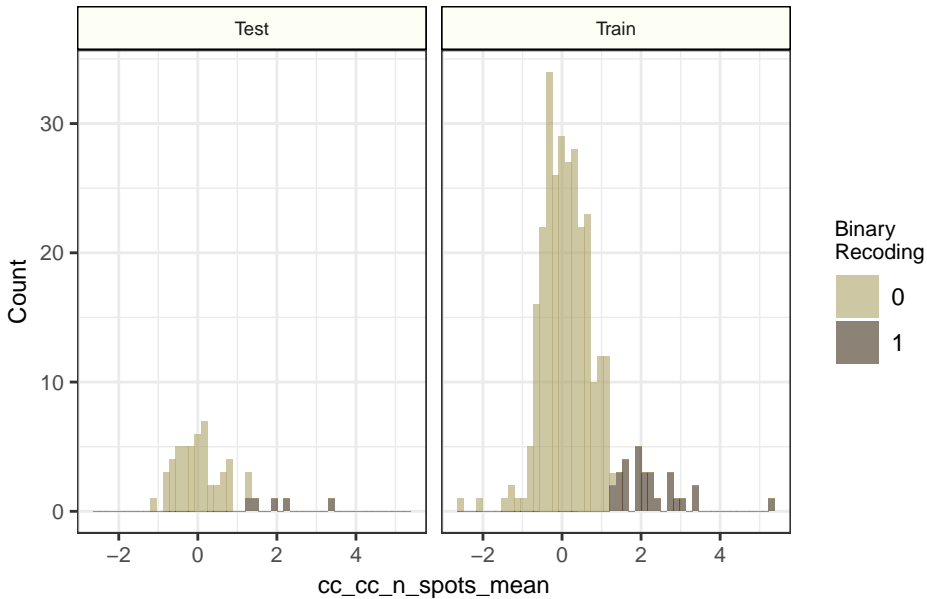
cc_cc_high_n_spots_h2ax_mean



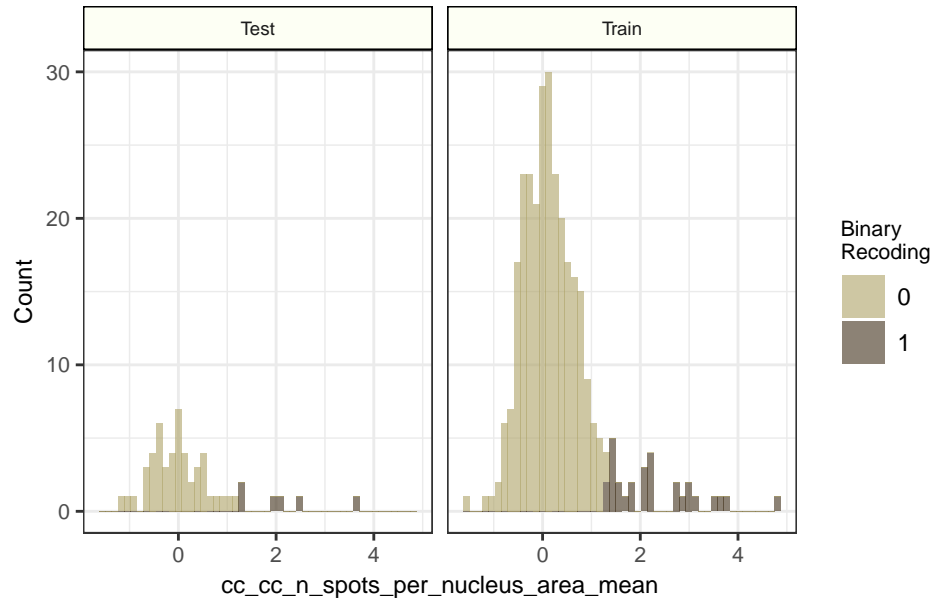
cc_cc_n_objects



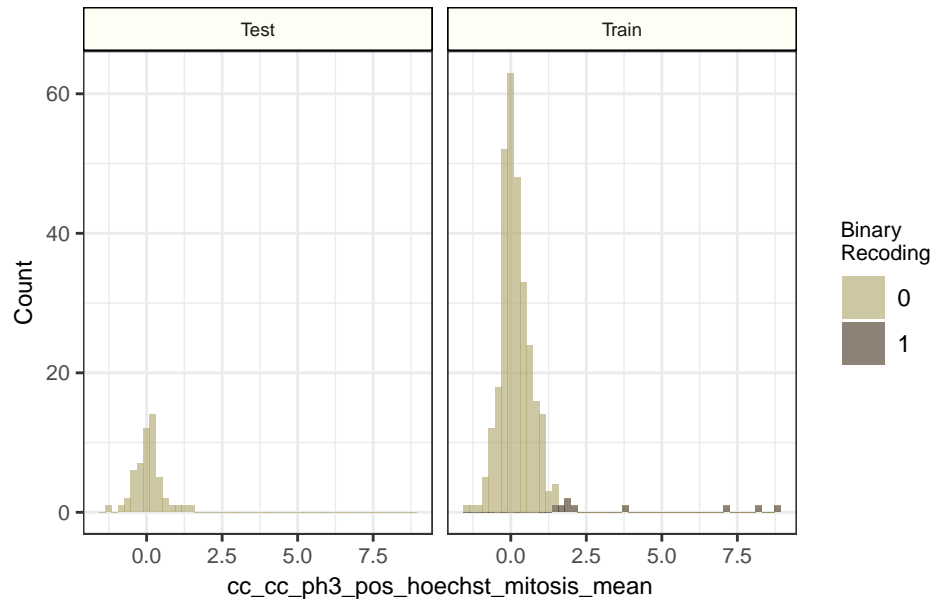
cc_cc_n_spots_mean



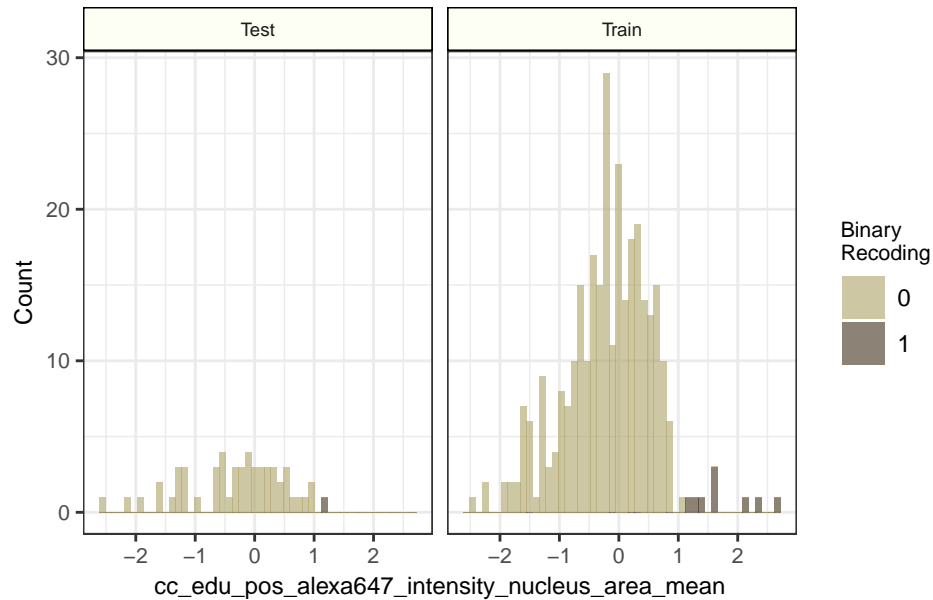
cc_cc_n_spots_per_nucleus_area_mean



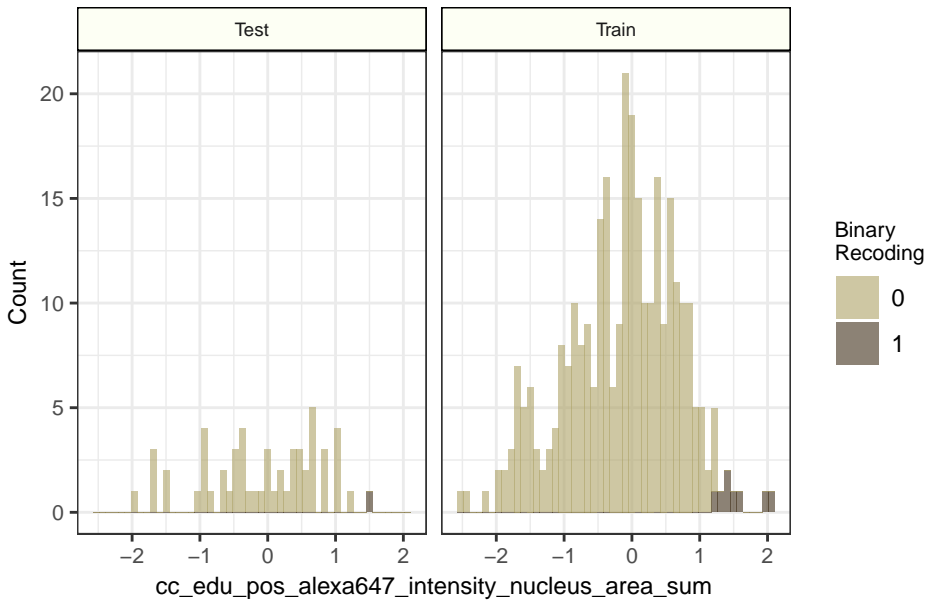
cc_cc_ph3_pos_hoechst_mitosis_mean



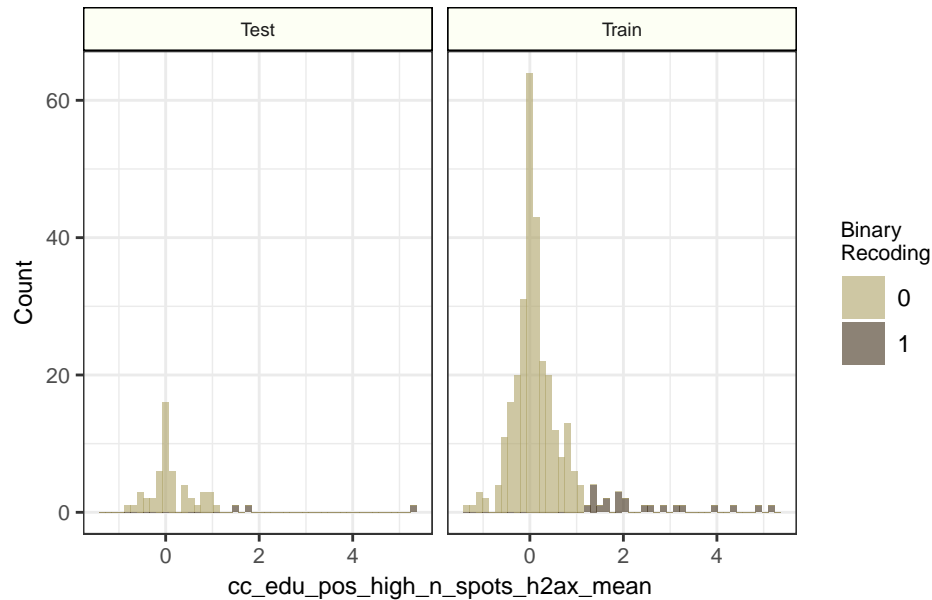
cc_edu_pos_alex647_intensity_nucleus_area_mean



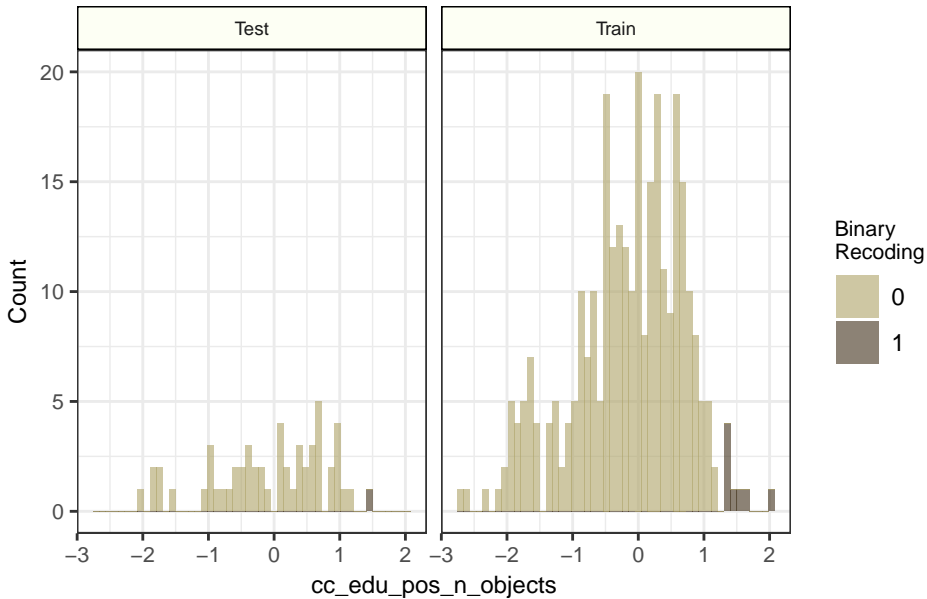
cc_edu_pos_alex647_intensity_nucleus_area_sum



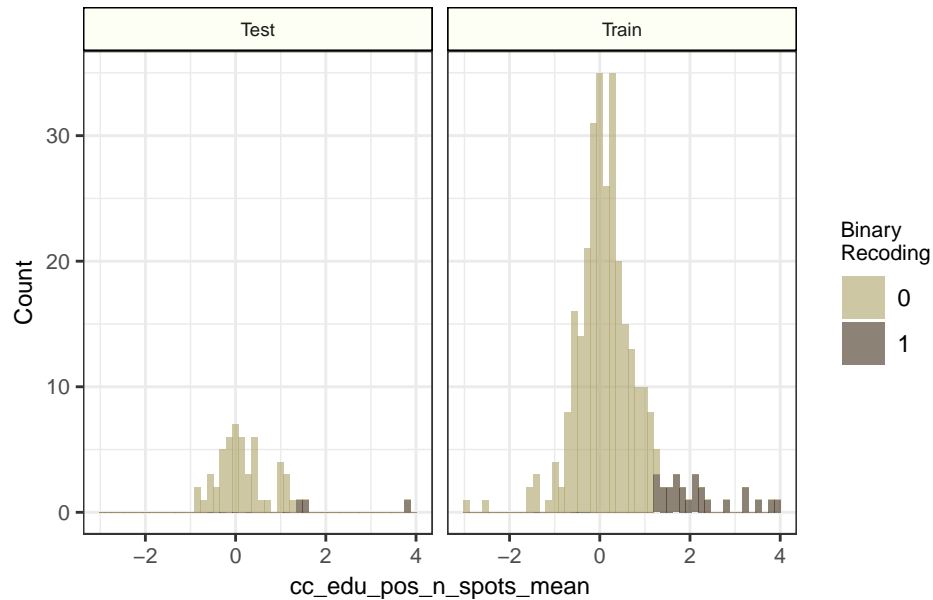
cc_edu_pos_high_n_spots_h2ax_mean



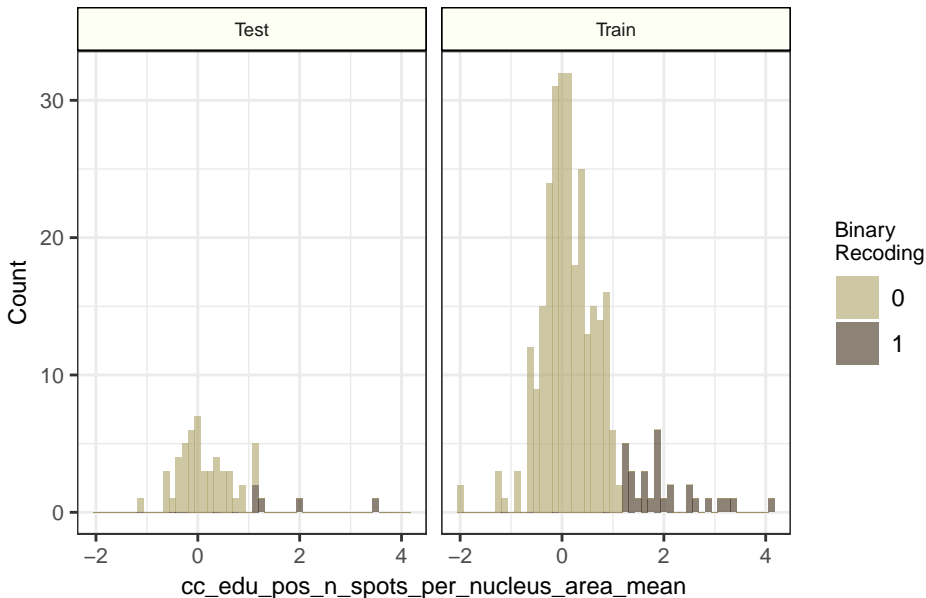
cc_edu_pos_n_objects



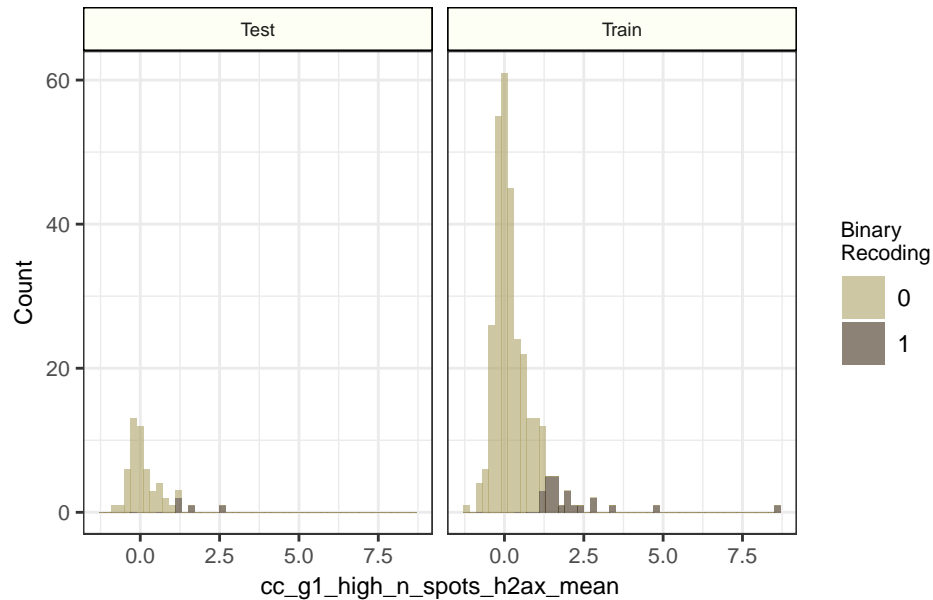
cc_edu_pos_n_spots_mean



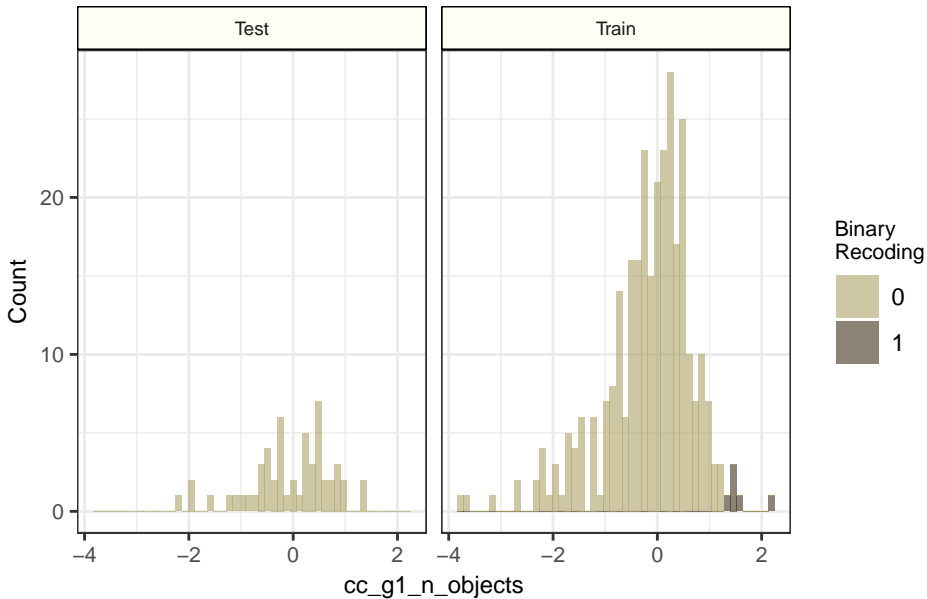
cc_edu_pos_n_spots_per_nucleus_area_mean



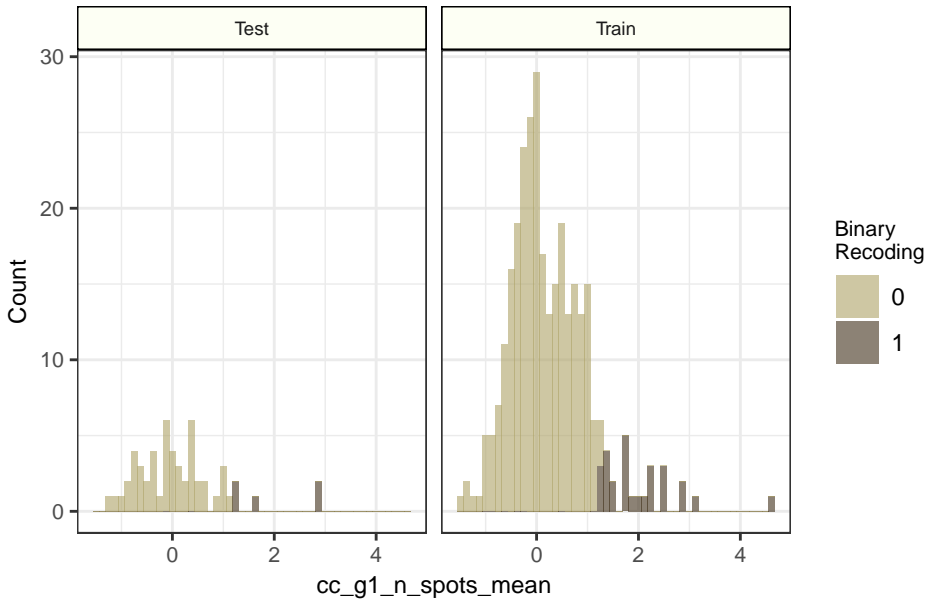
cc_g1_high_n_spots_h2ax_mean



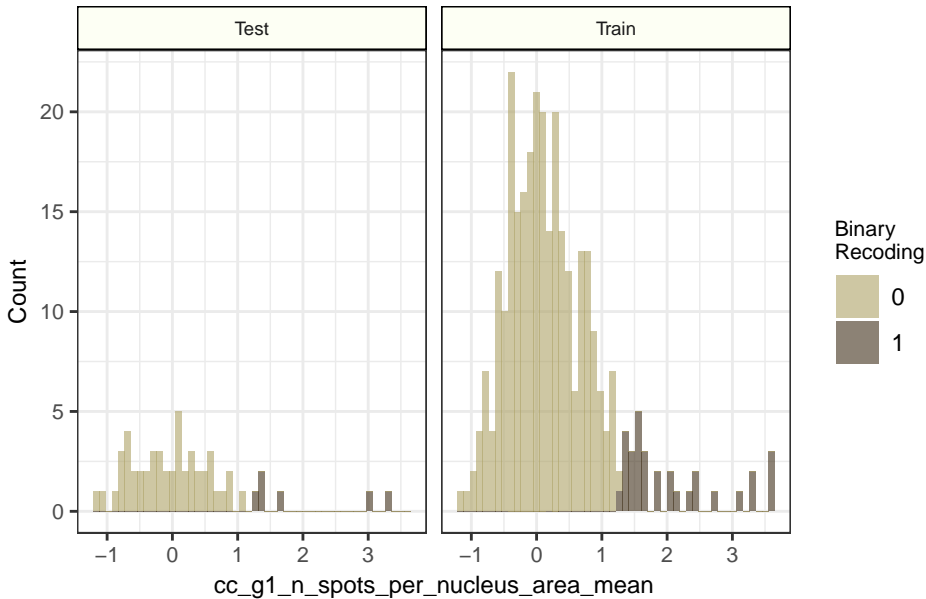
cc_g1_n_objects



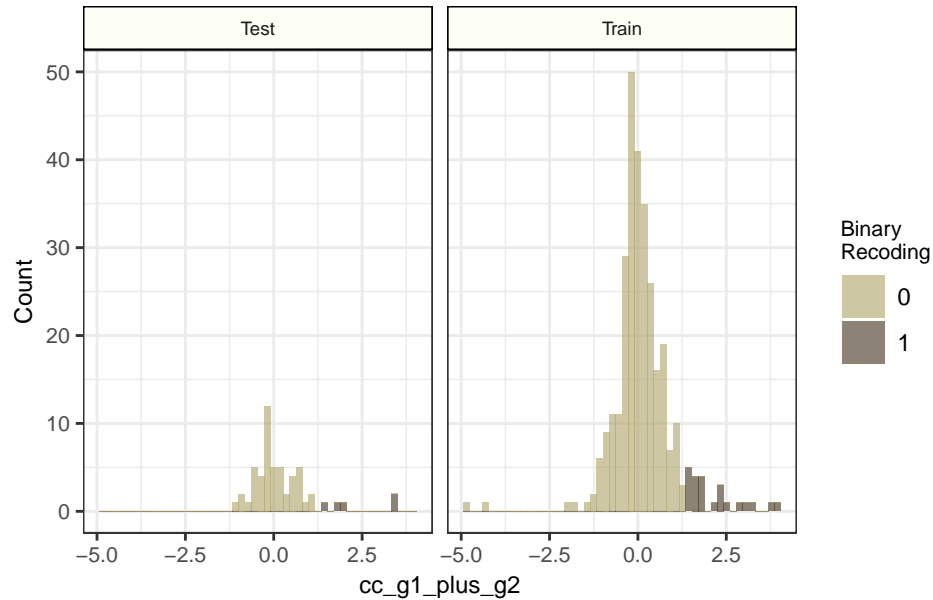
cc_g1_n_spots_mean



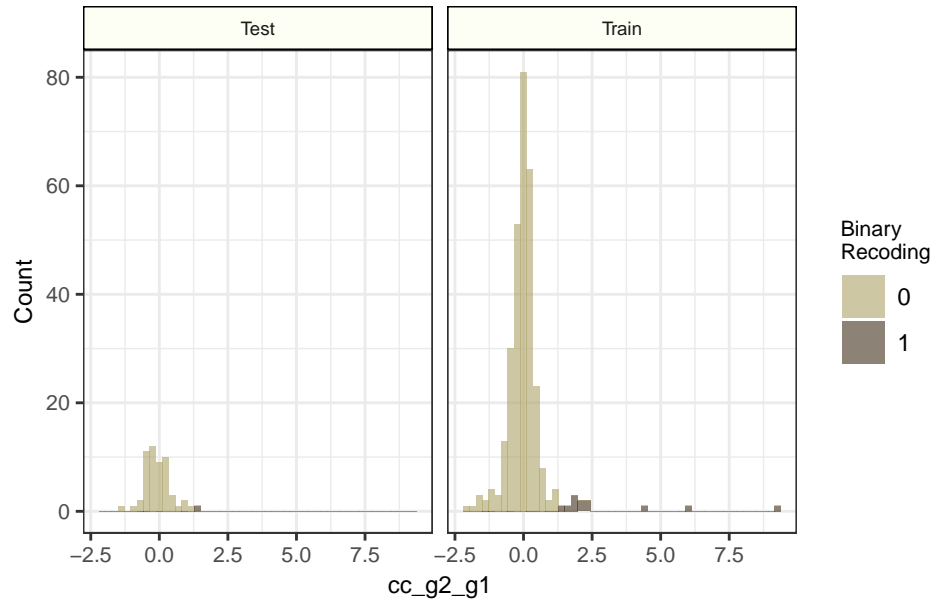
cc_g1_n_spots_per_nucleus_area_mean



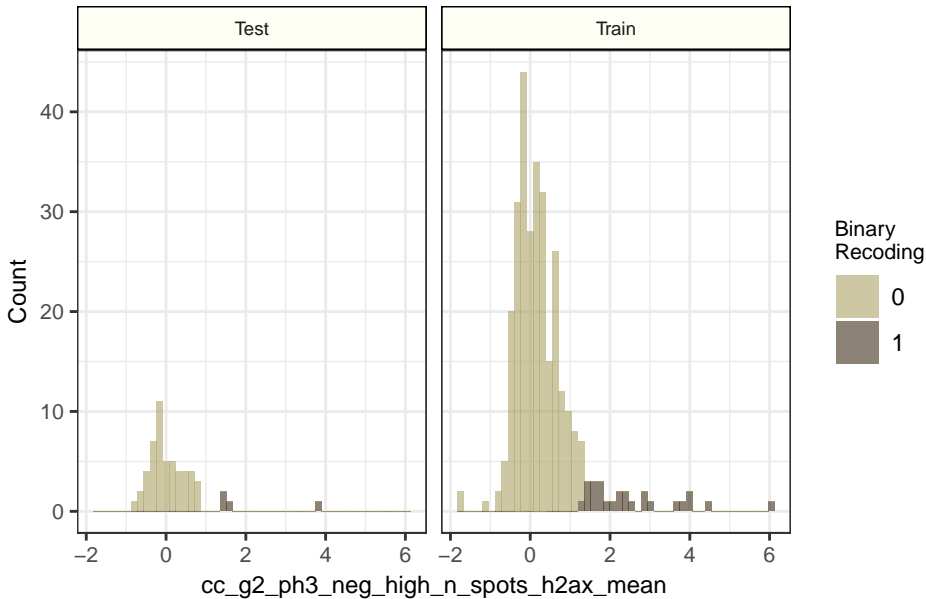
cc_g1_plus_g2



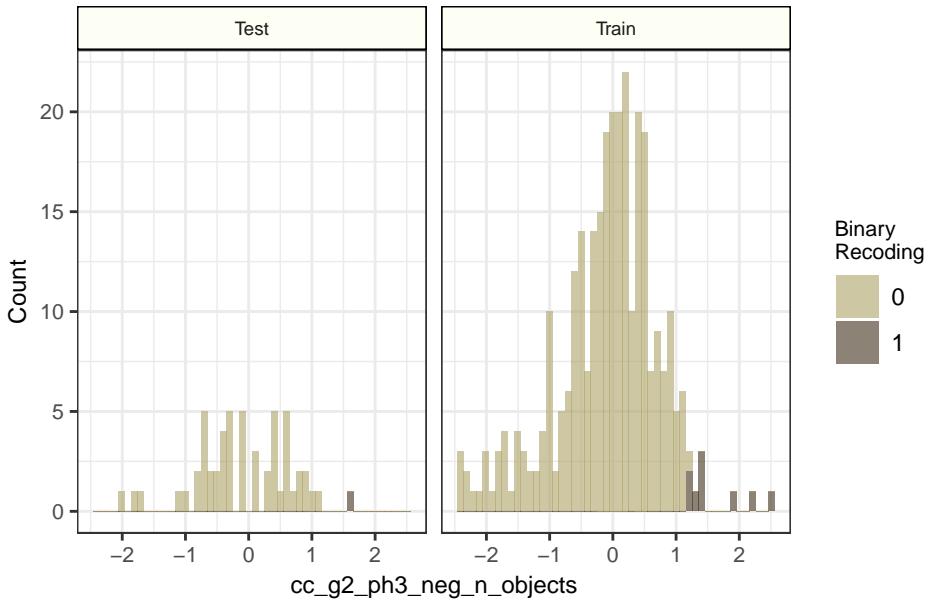
cc_g2_g1



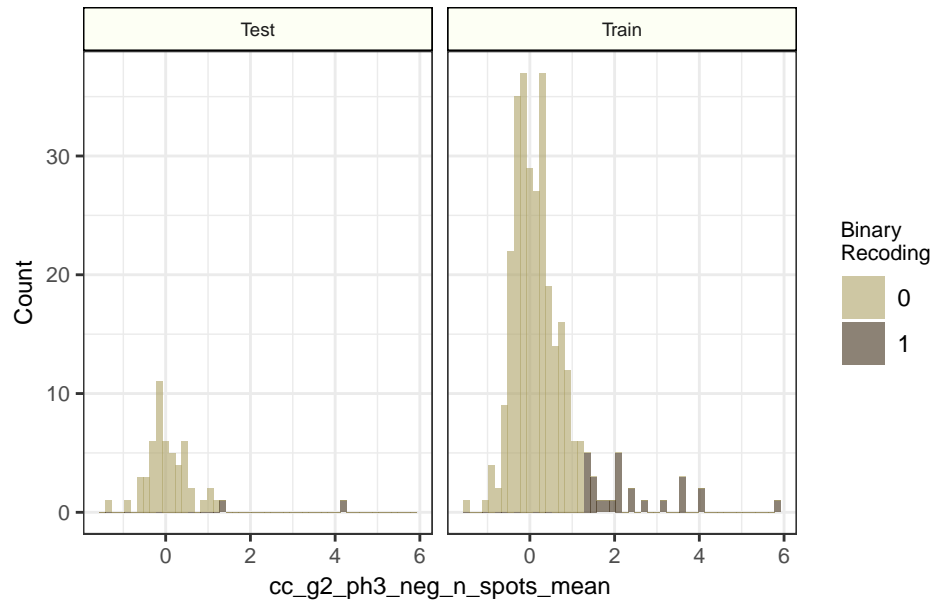
cc_g2_ph3_neg_high_n_spots_h2ax_mean



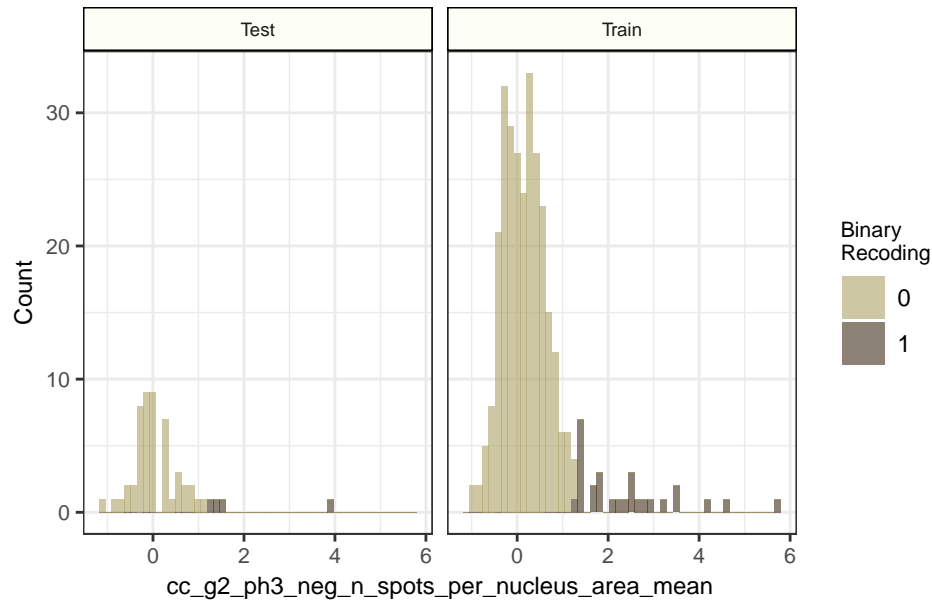
cc_g2_ph3_neg_n_objects



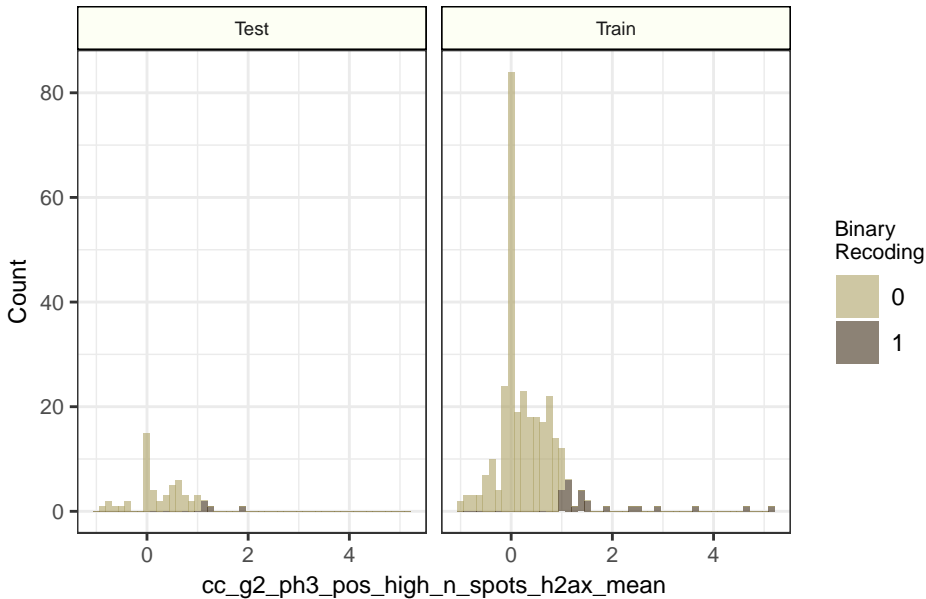
cc_g2_ph3_neg_n_spots_mean



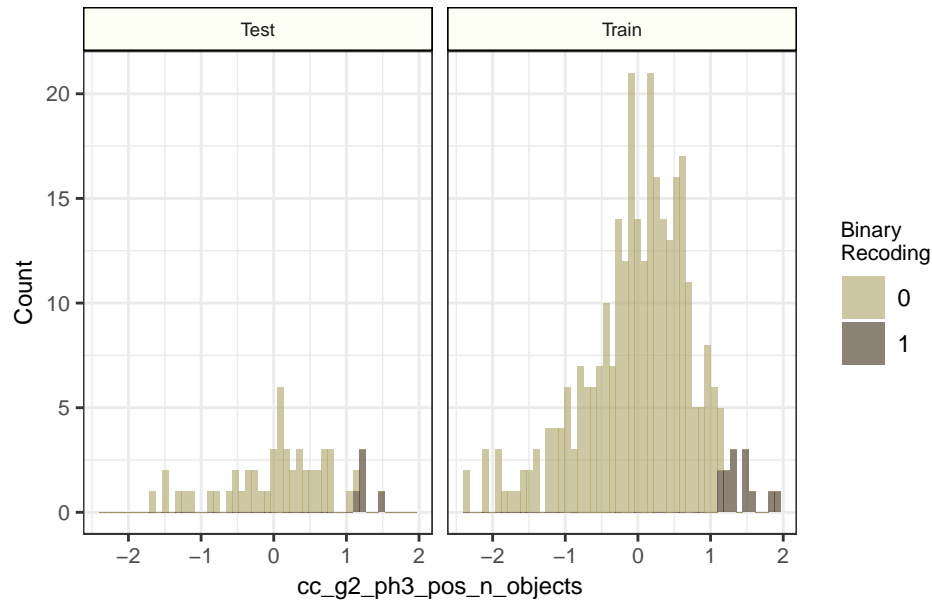
cc_g2_ph3_neg_n_spots_per_nucleus_area_mean



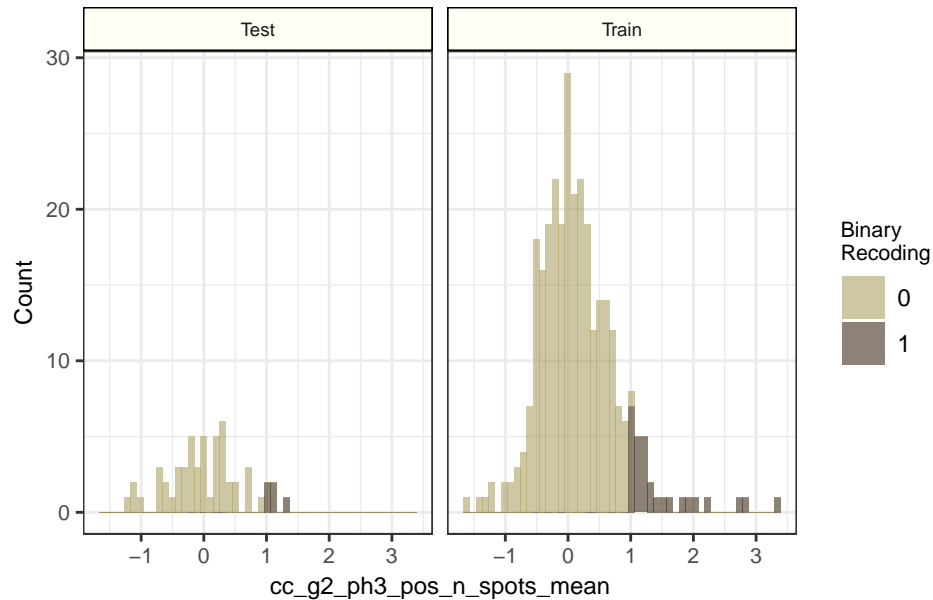
cc_g2_ph3_pos_high_n_spots_h2ax_mean



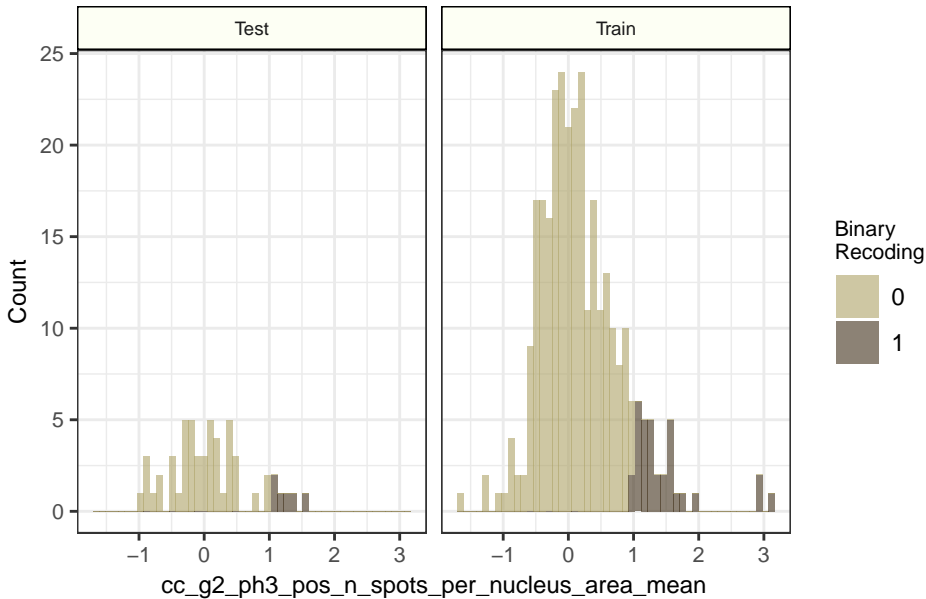
cc_g2_ph3_pos_n_objects



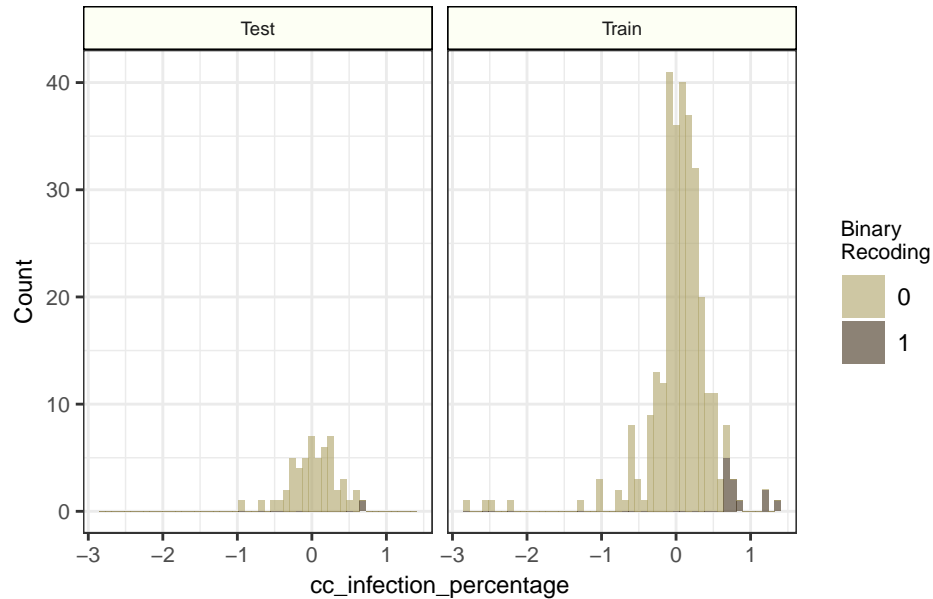
cc_g2_ph3_pos_n_spots_mean



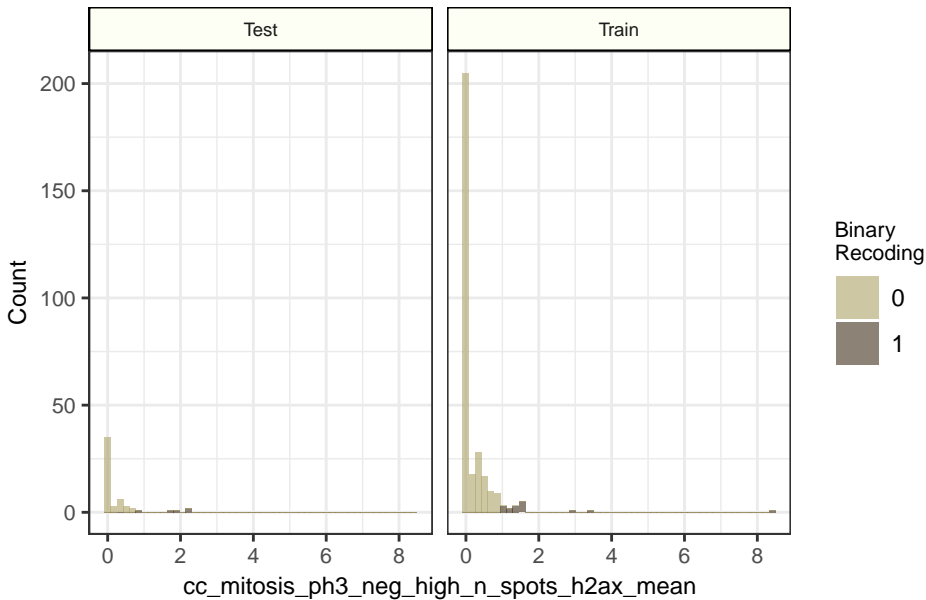
cc_g2_ph3_pos_n_spots_per_nucleus_area_mean



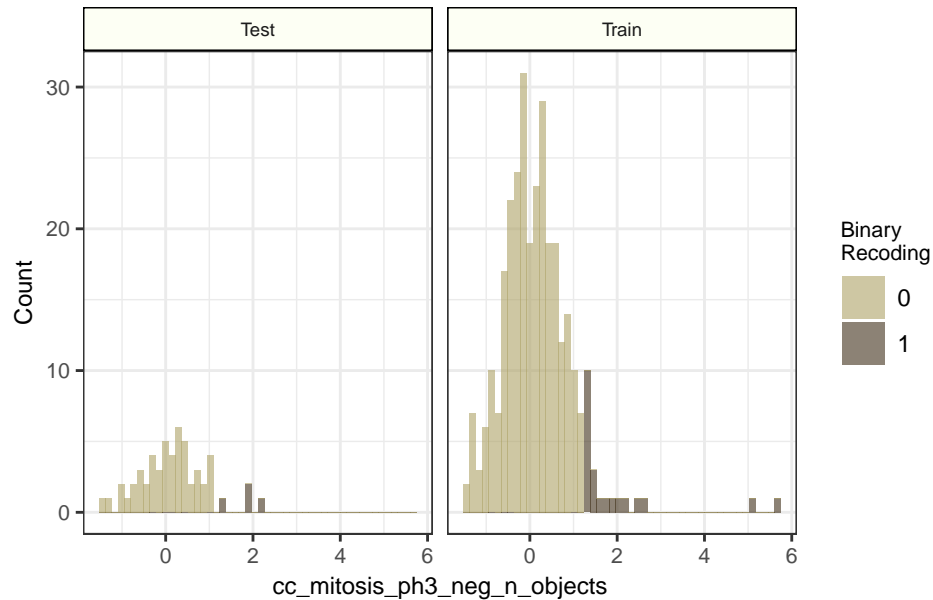
cc_infection_percentage



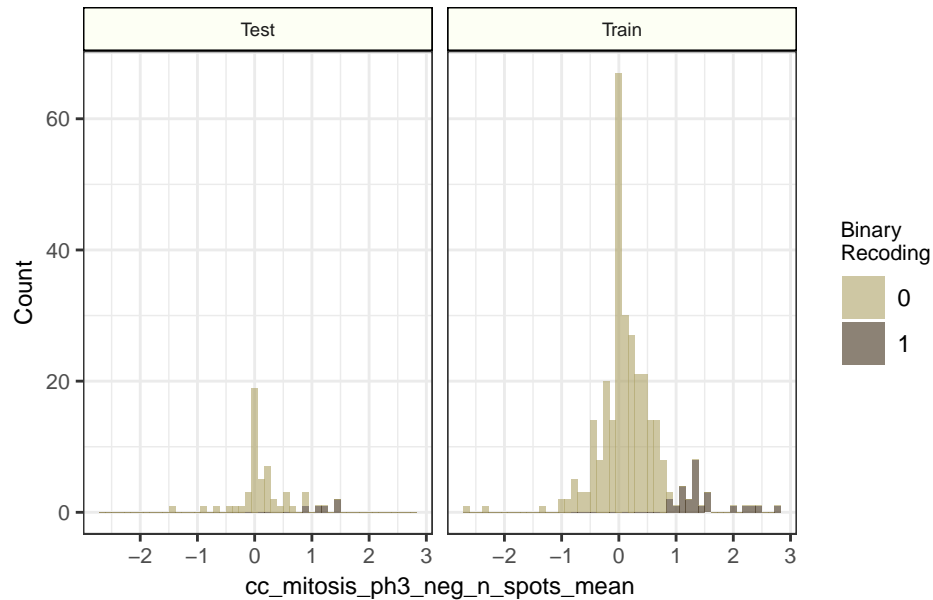
cc_mitosis_ph3_neg_high_n_spots_h2ax_mean



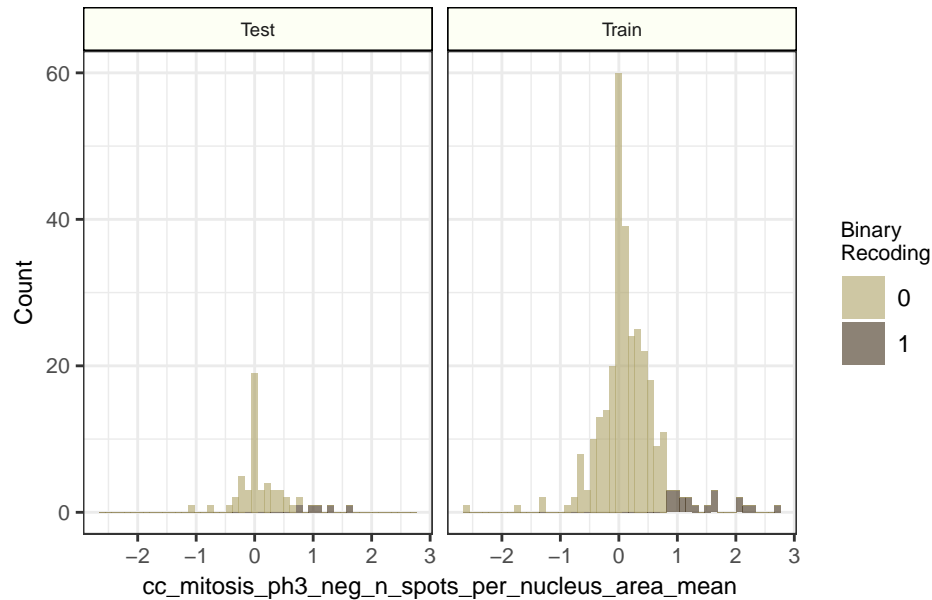
cc_mitosis_ph3_neg_n_objects



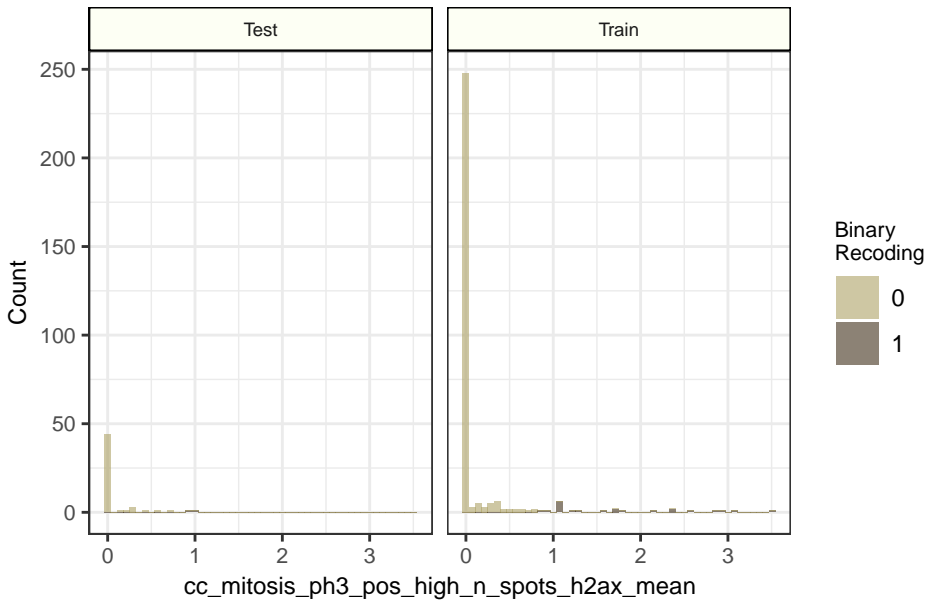
cc_mitosis_ph3_neg_n_spots_mean



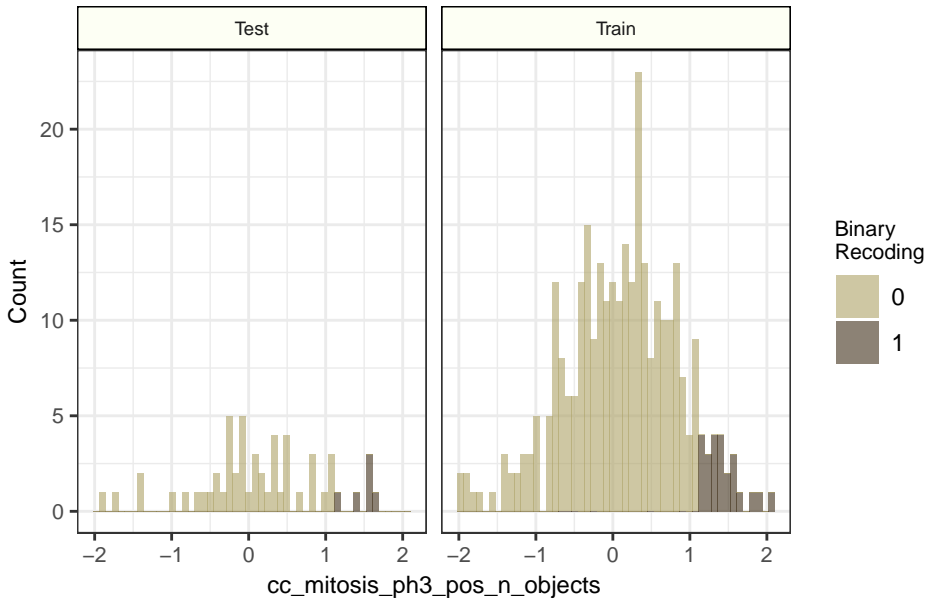
cc_mitosis_ph3_neg_n_spots_per_nucleus_area_me



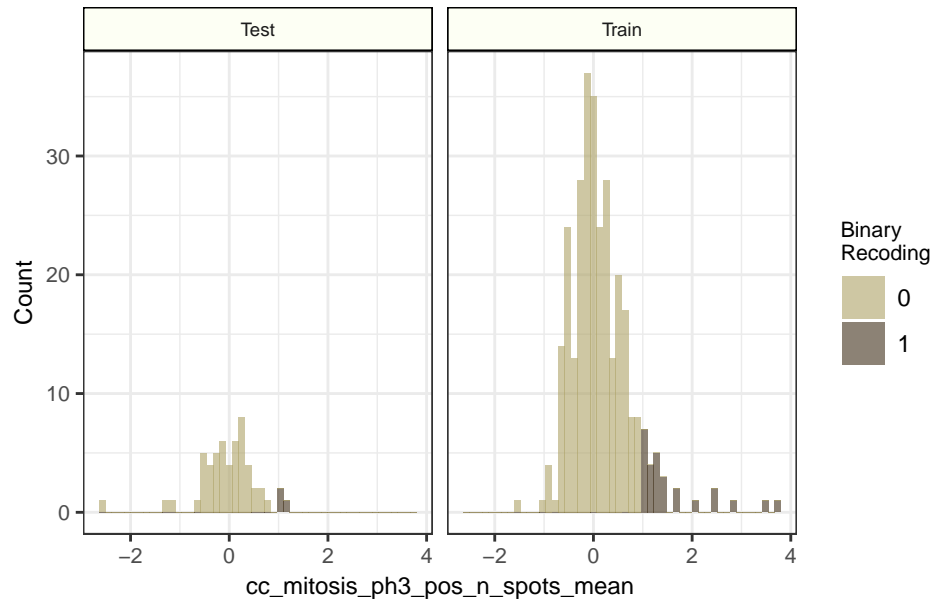
cc_mitosis_ph3_pos_high_n_spots_h2ax_mean



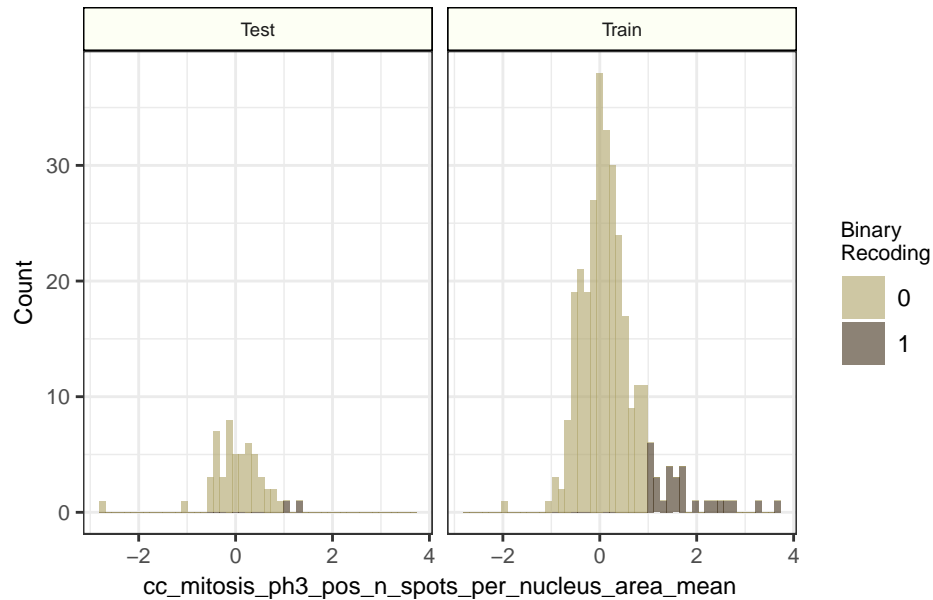
cc_mitosis_ph3_pos_n_objects



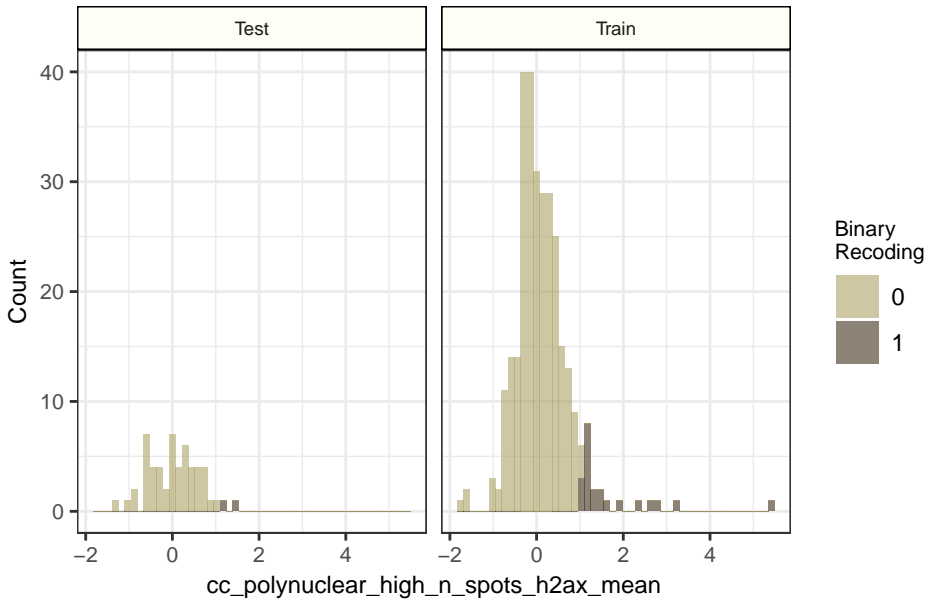
cc_mitosis_ph3_pos_n_spots_mean



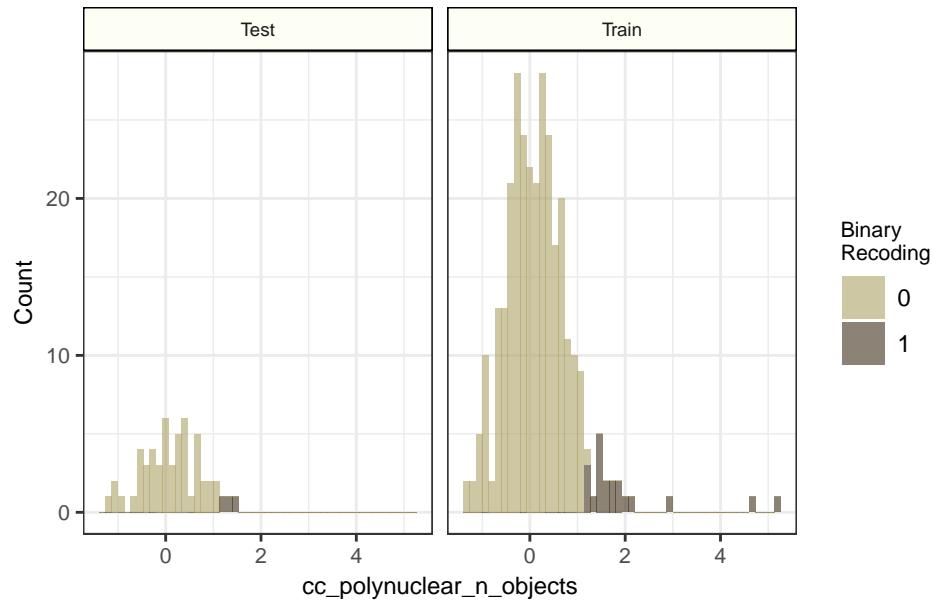
cc_mitosis_ph3_pos_n_spots_per_nucleus_area_me



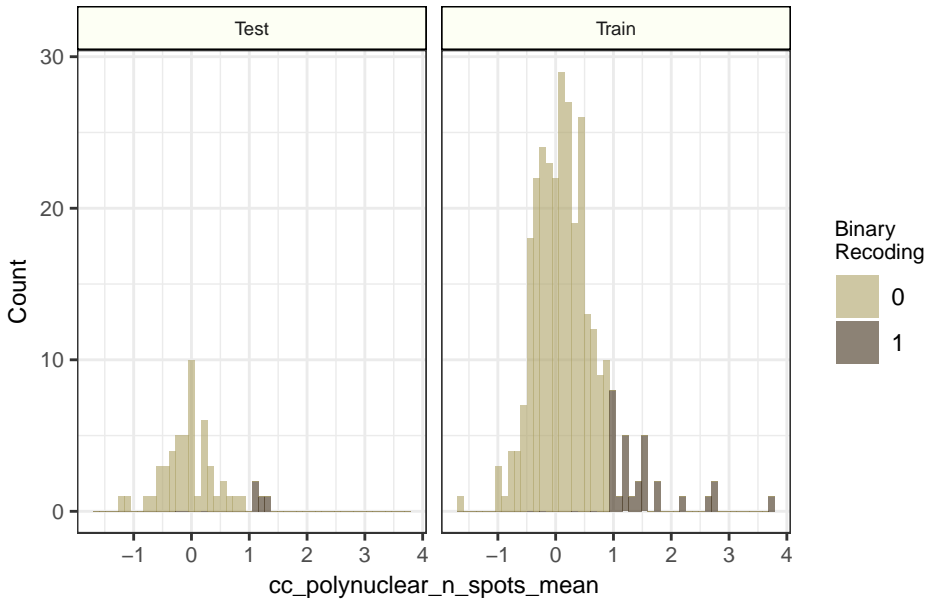
cc_polynuclear_high_n_spots_h2ax_mean



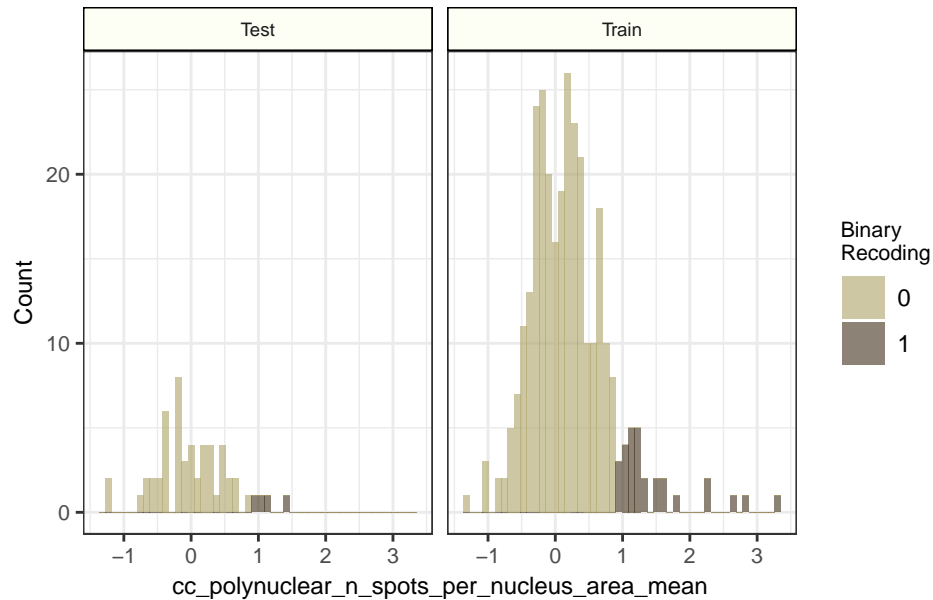
cc_polynuclear_n_objects



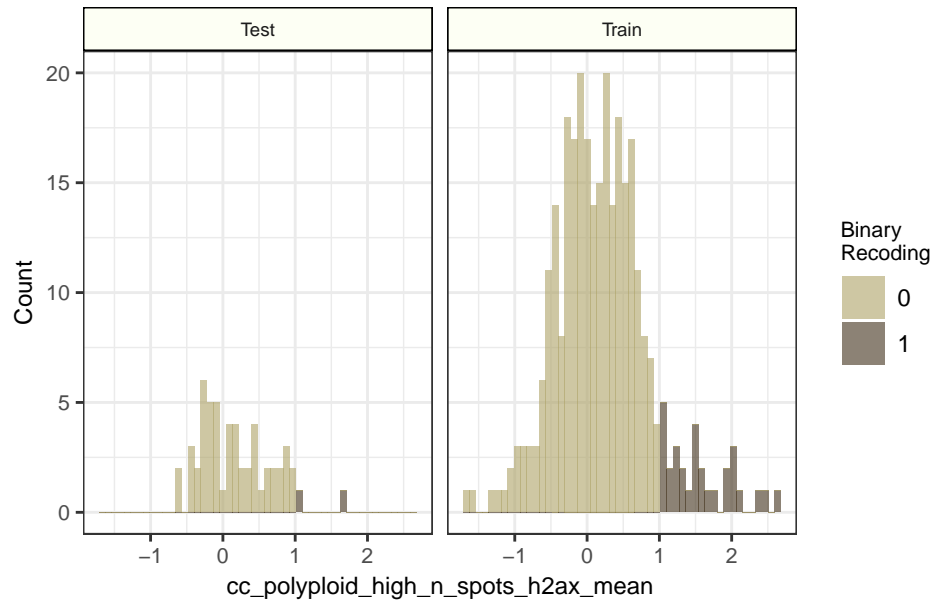
cc_polynuclear_n_spots_mean



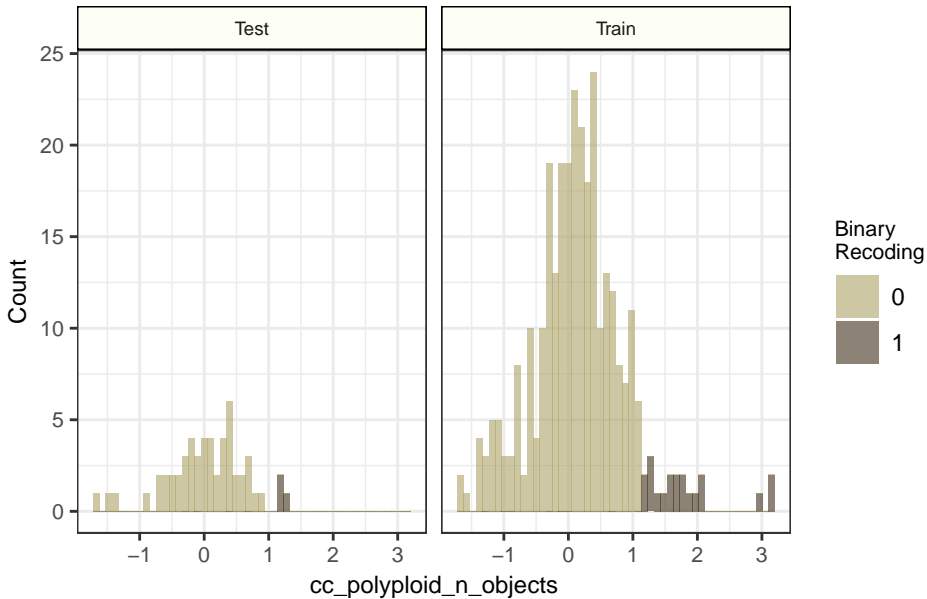
cc_polynuclear_n_spots_per_nucleus_area_mean



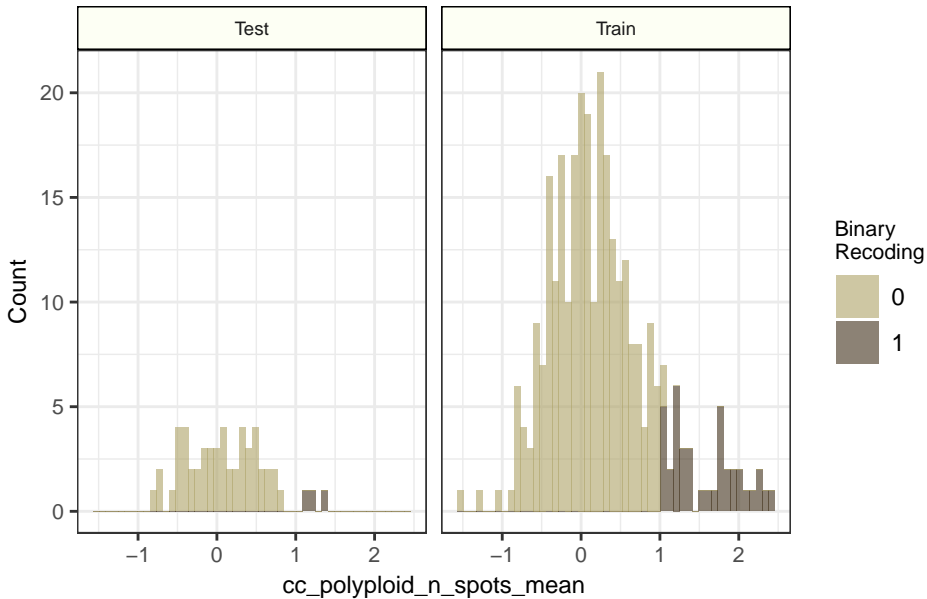
cc_polyploid_high_n_spots_h2ax_mean



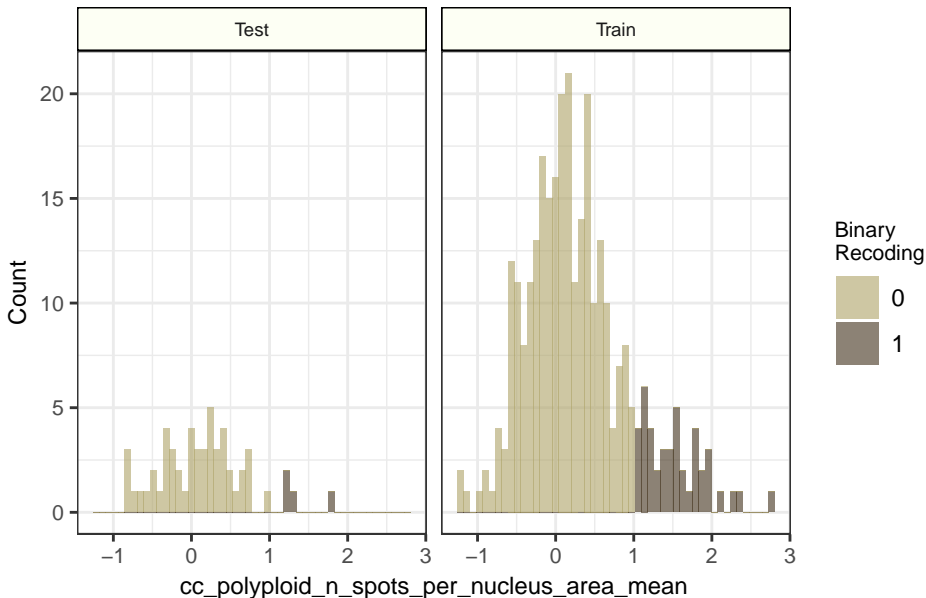
cc_polyploid_n_objects



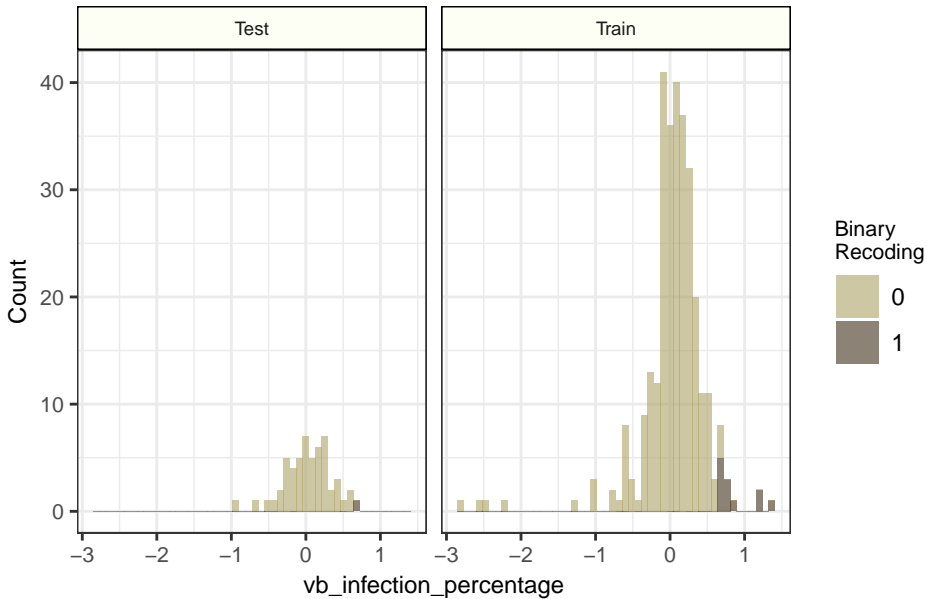
cc_polyploid_n_spots_mean



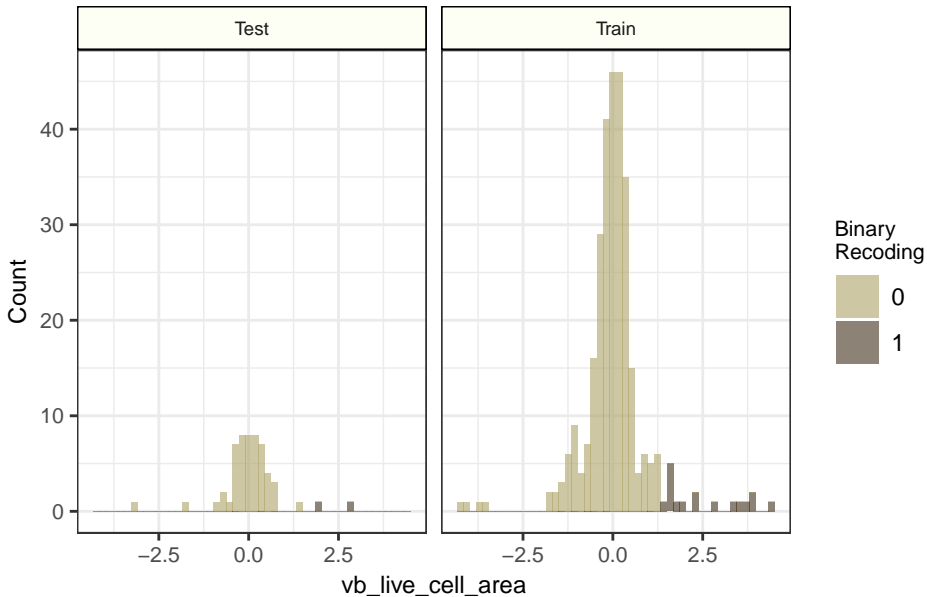
cc_polyploid_n_spots_per_nucleus_area_mean



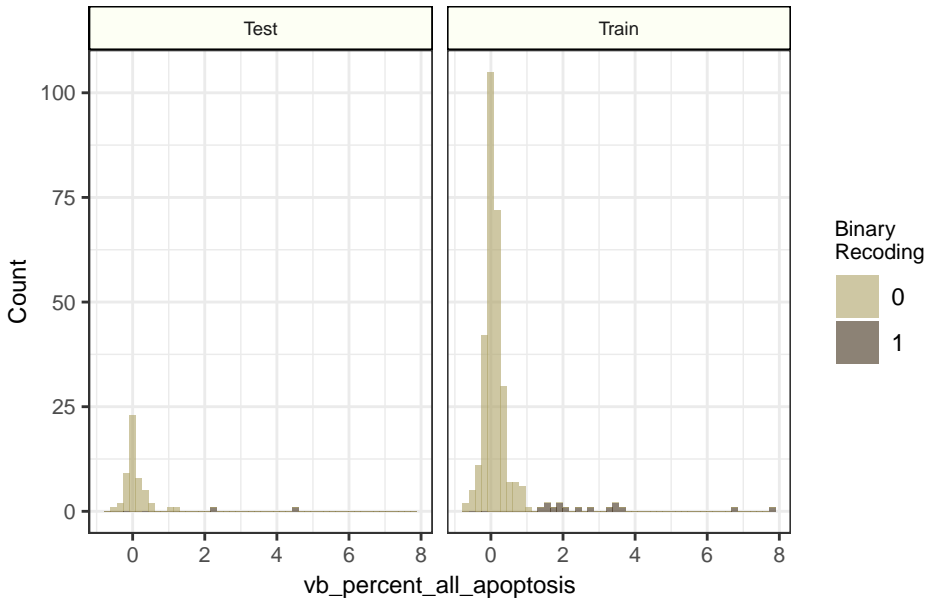
vb_infection_percentage



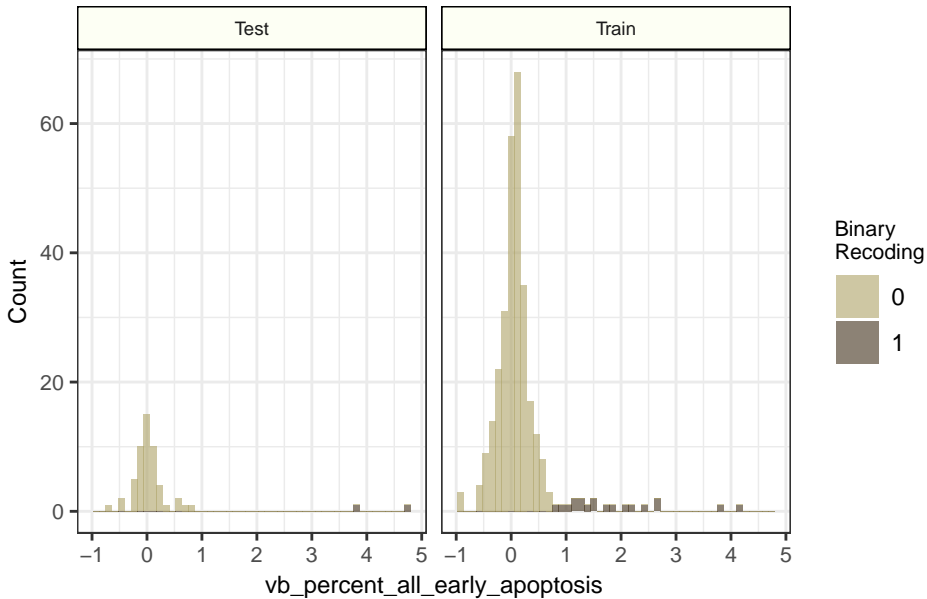
vb_live_cell_area



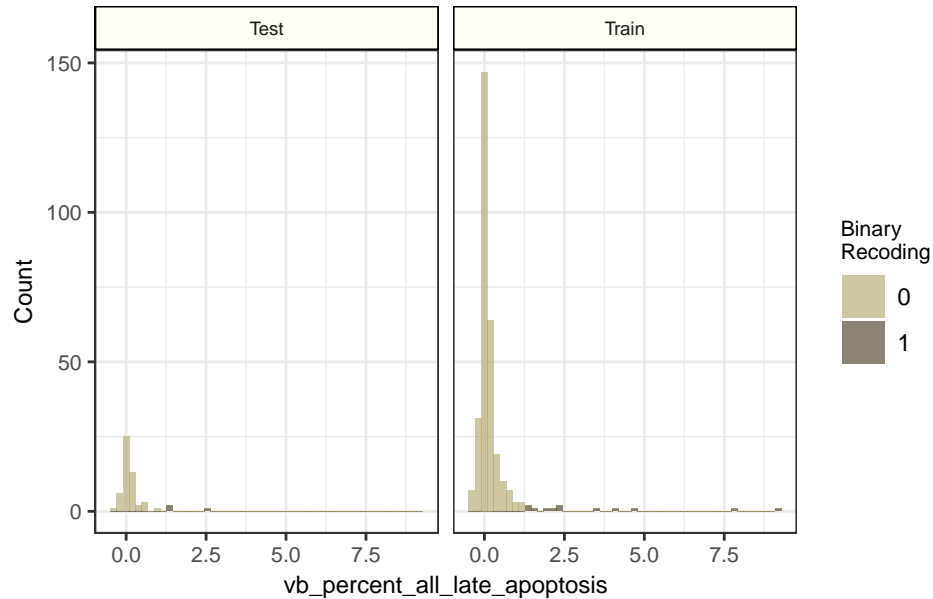
vb_percent_all_apoptosis



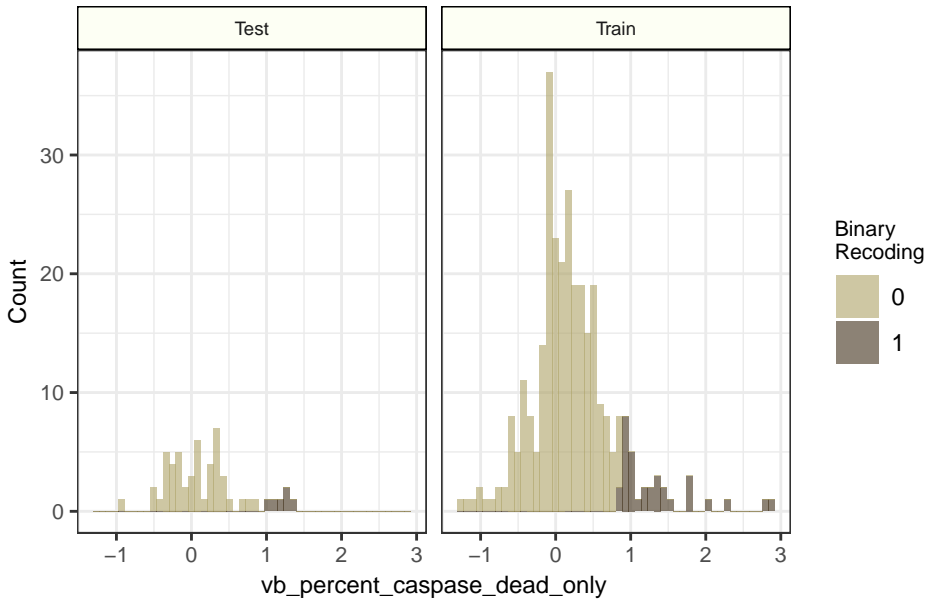
vb_percent_all_early_apoptosis



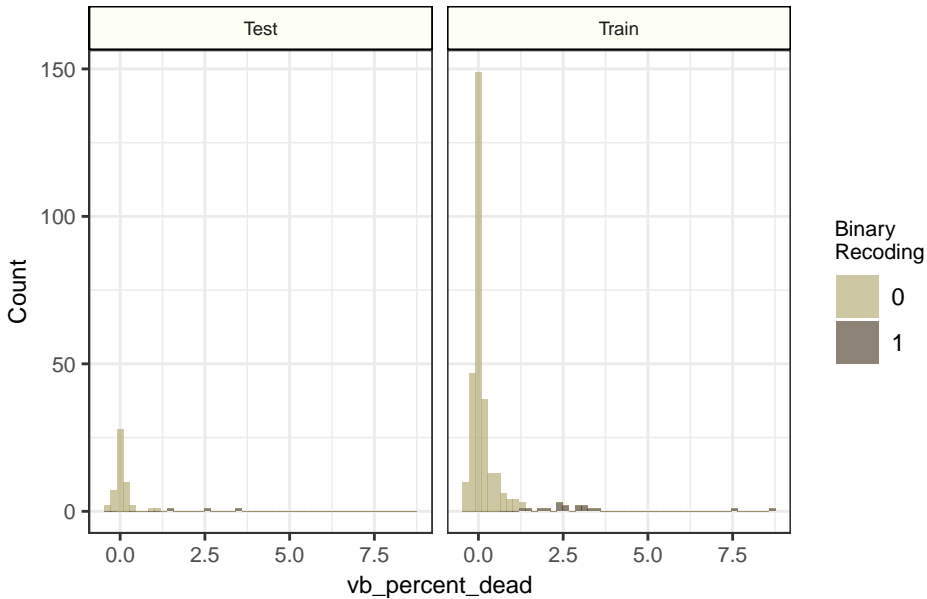
vb_percent_all_late_apoptosis



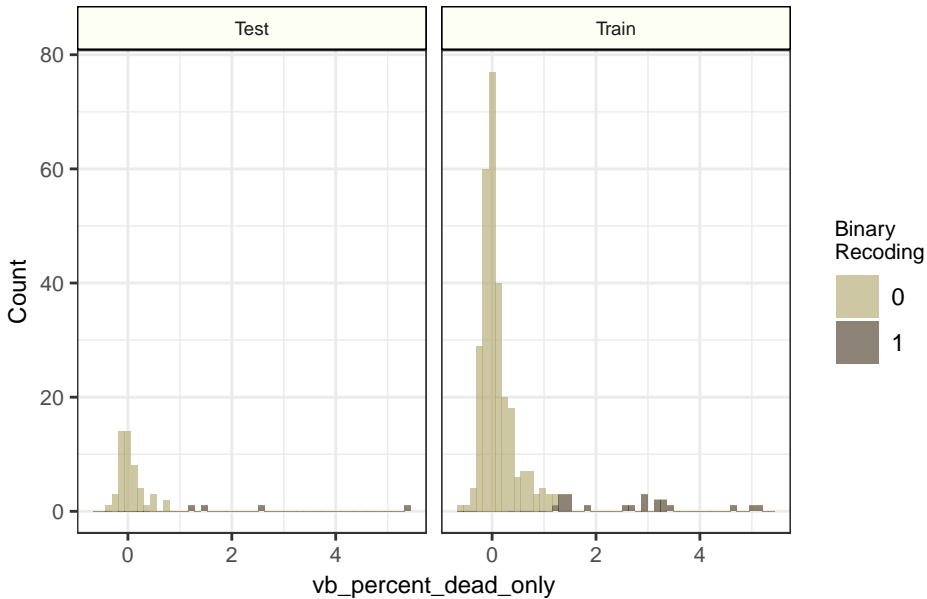
vb_percent_caspase_dead_only



vb_percent_dead



vb_percent_dead_only



vb_ros_mean

