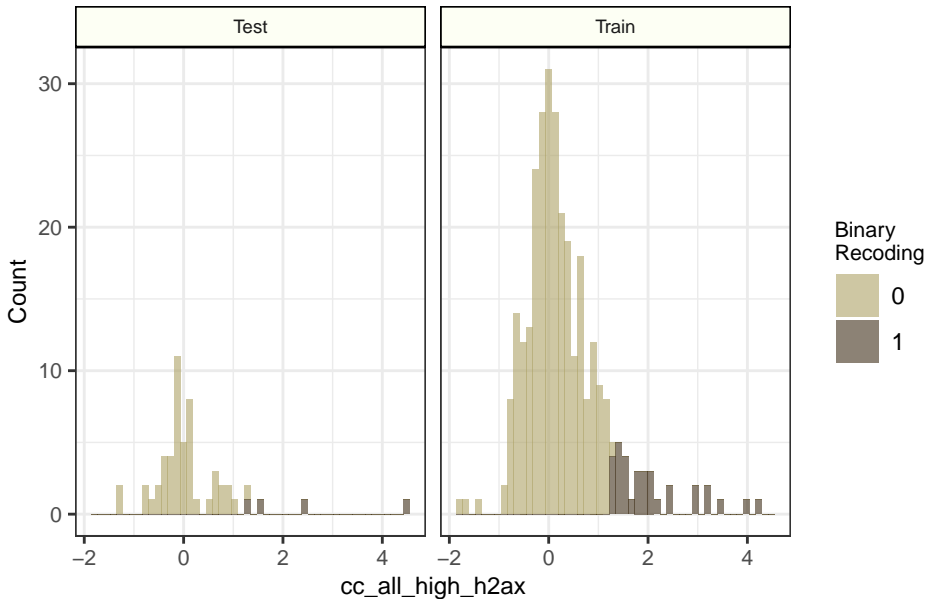
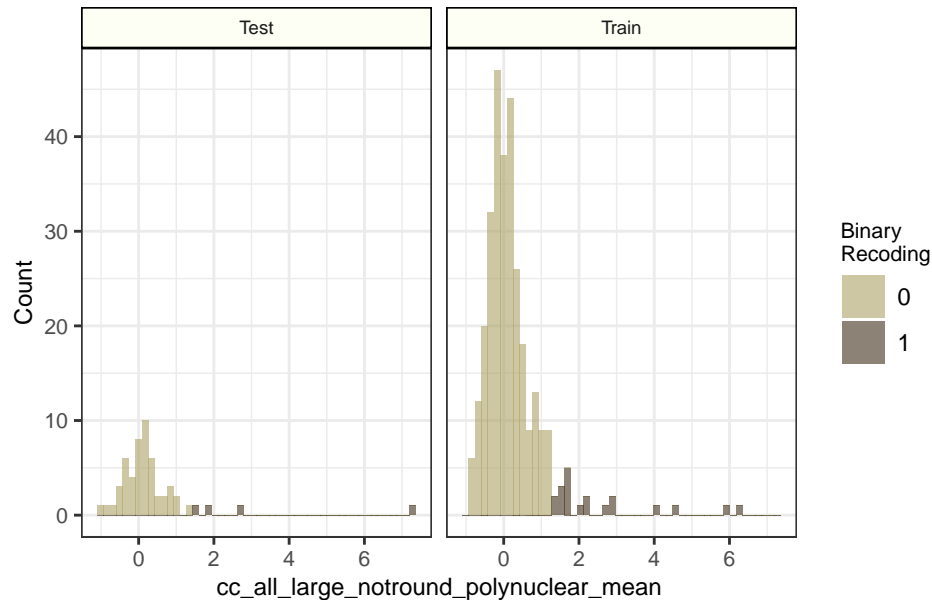


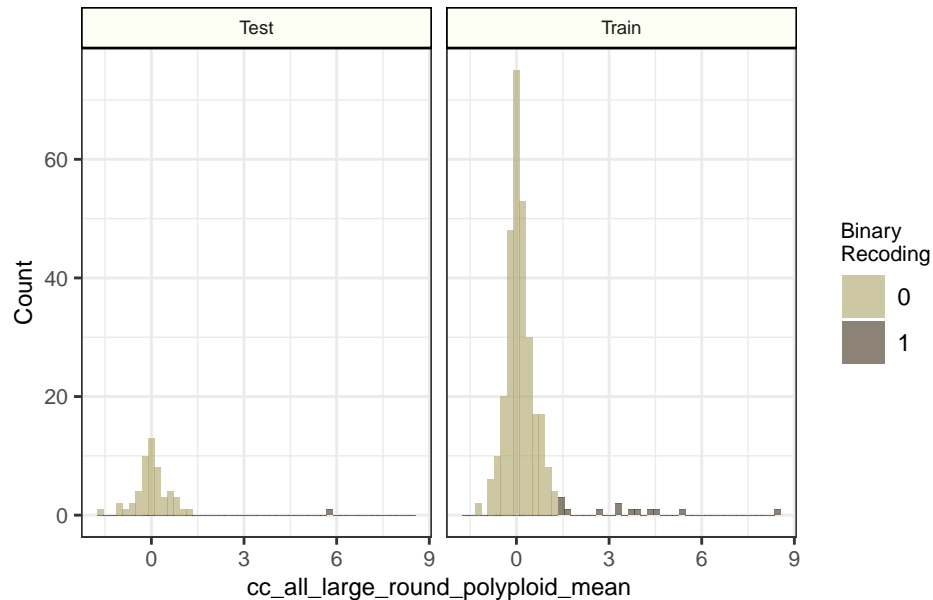
cc_all_high_h2ax



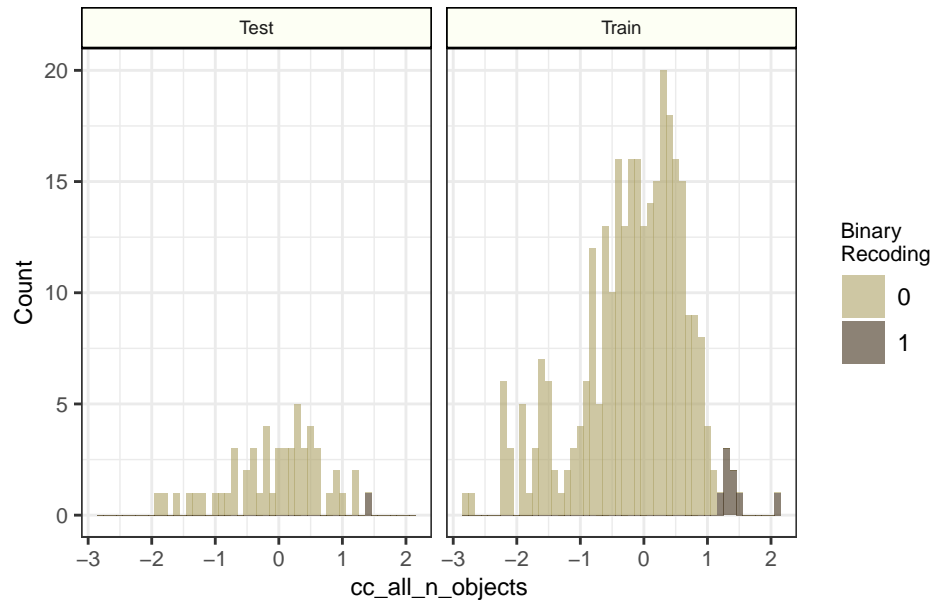
cc_all_large_notround_polynuclear_mean



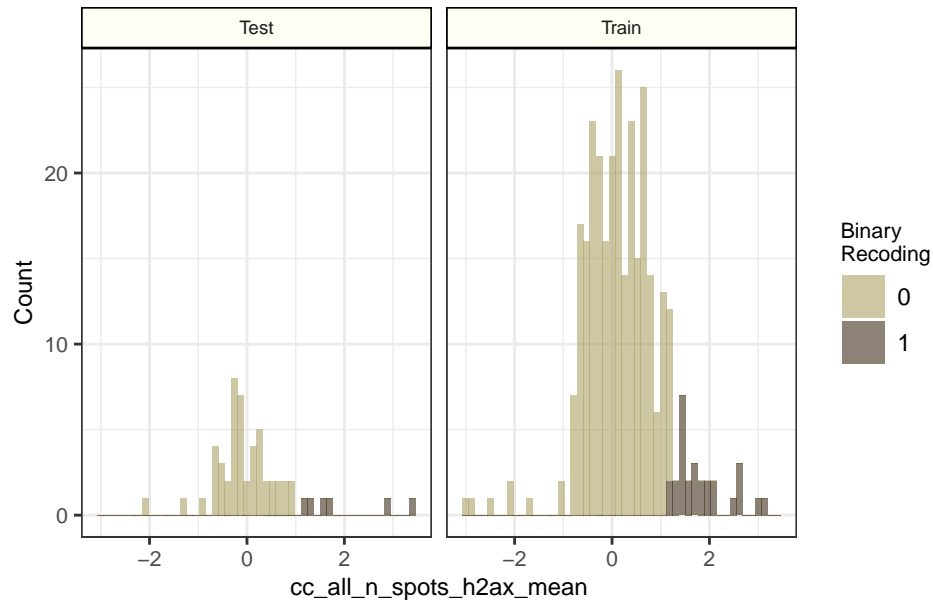
cc_all_large_round_polyploid_mean



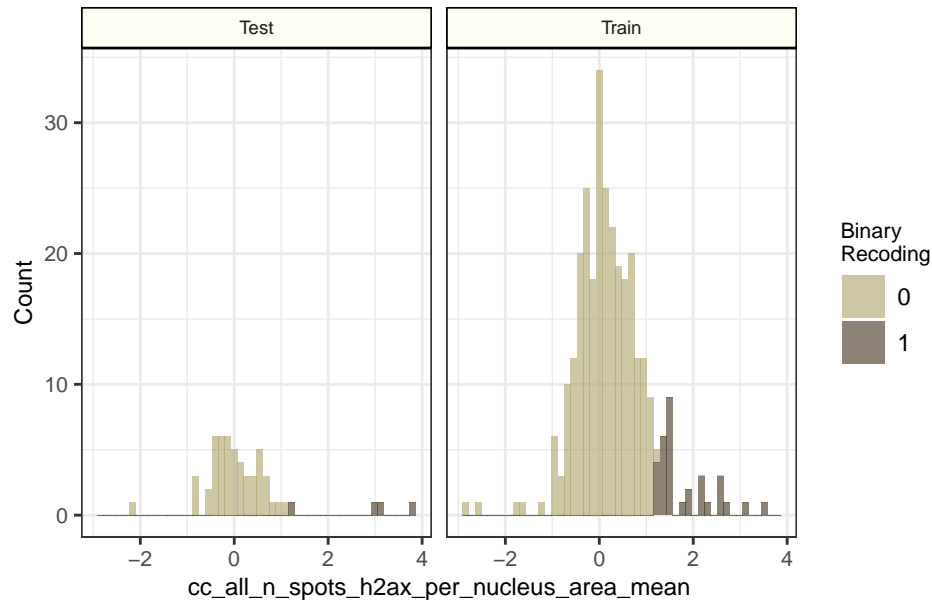
cc_all_n_objects



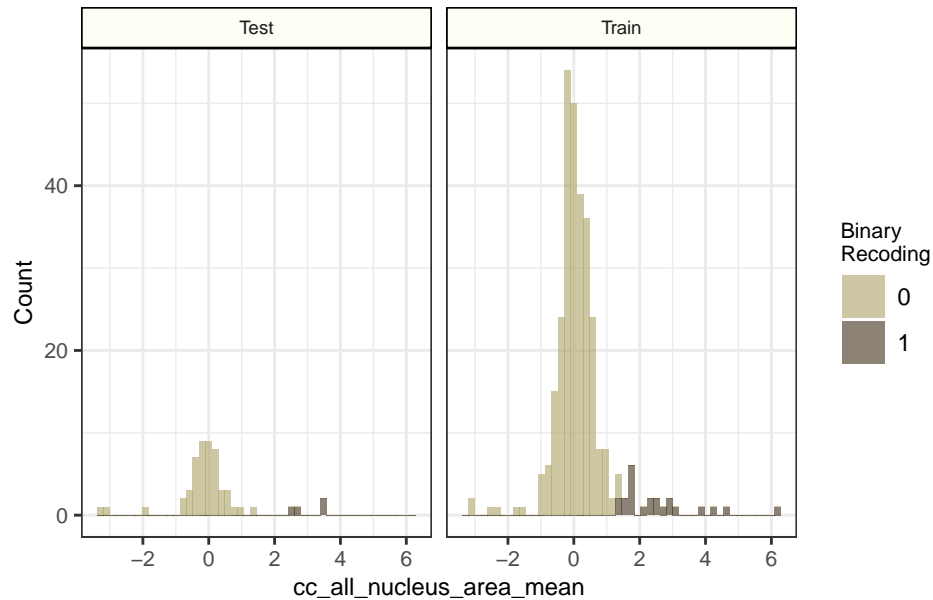
cc_all_n_spots_h2ax_mean



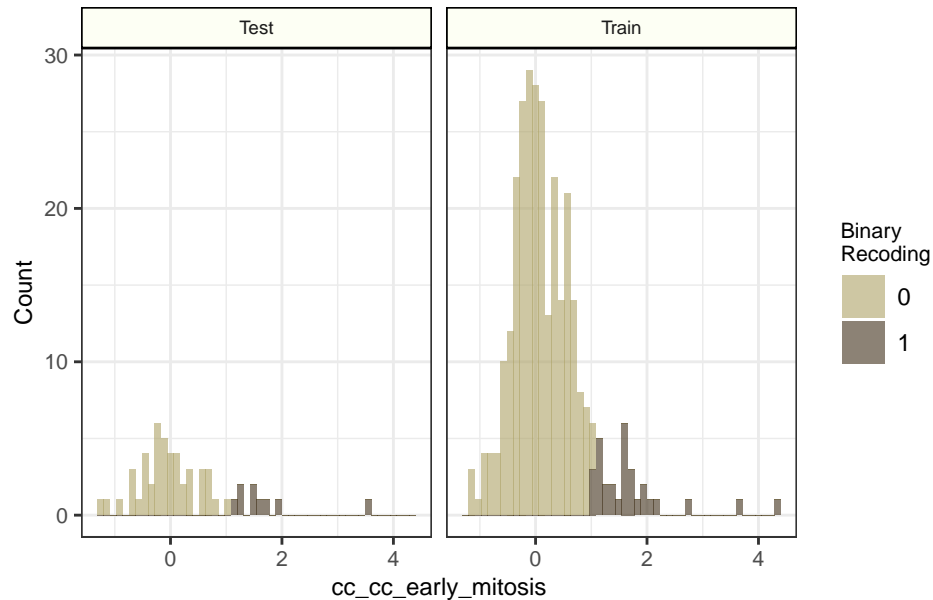
cc_all_n_spots_h2ax_per_nucleus_area_mean



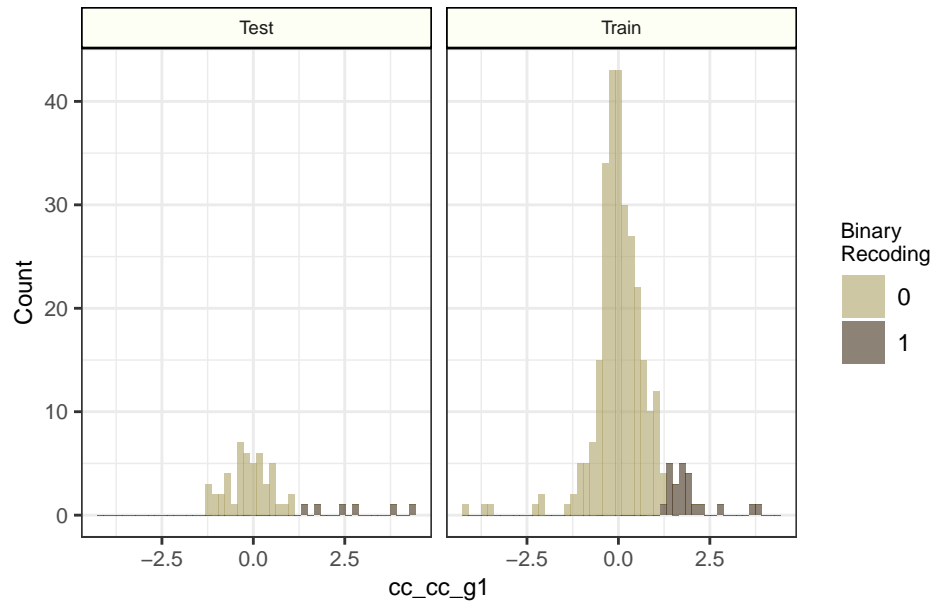
cc_all_nucleus_area_mean



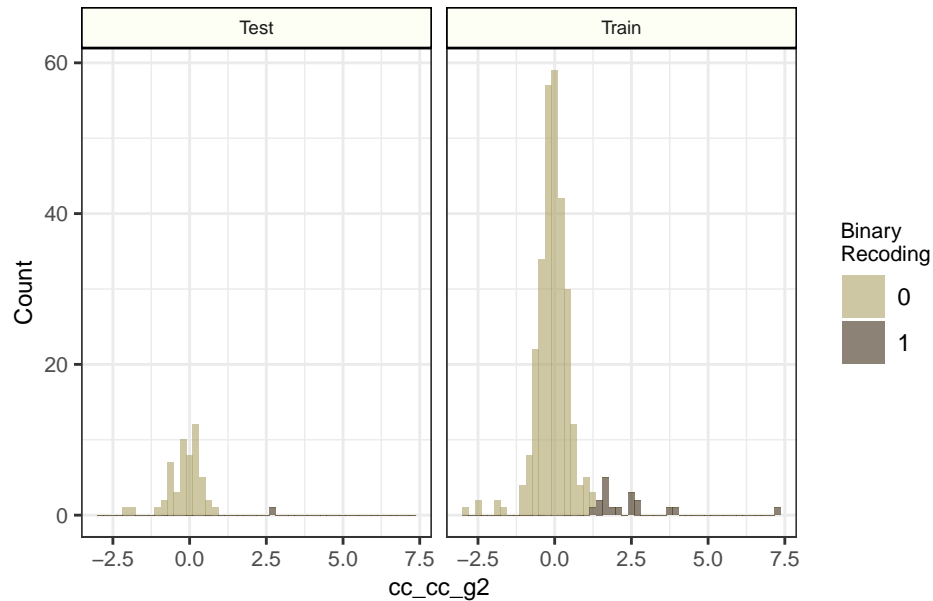
cc_cc_early_mitosis



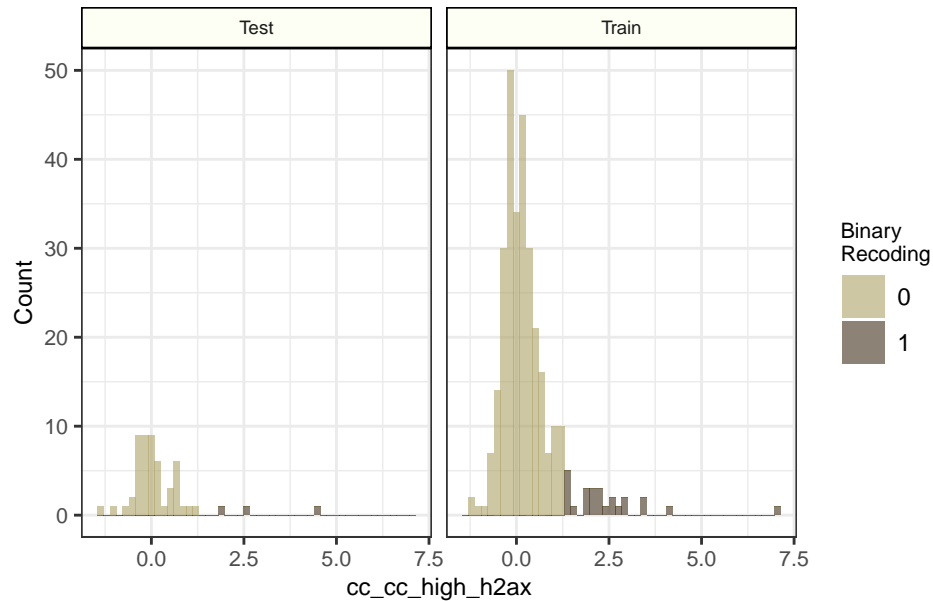
cc_cc_g1



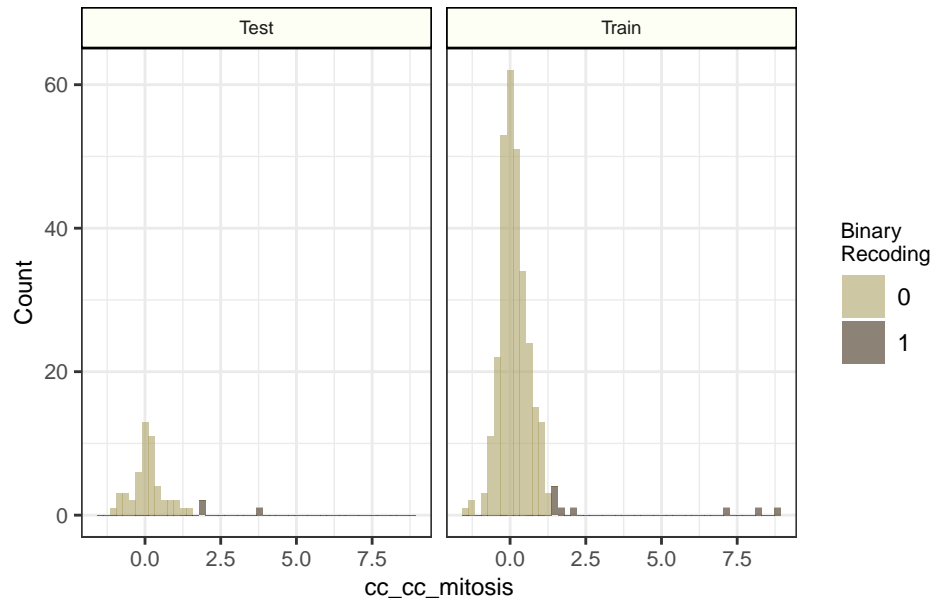
cc_cc_g2



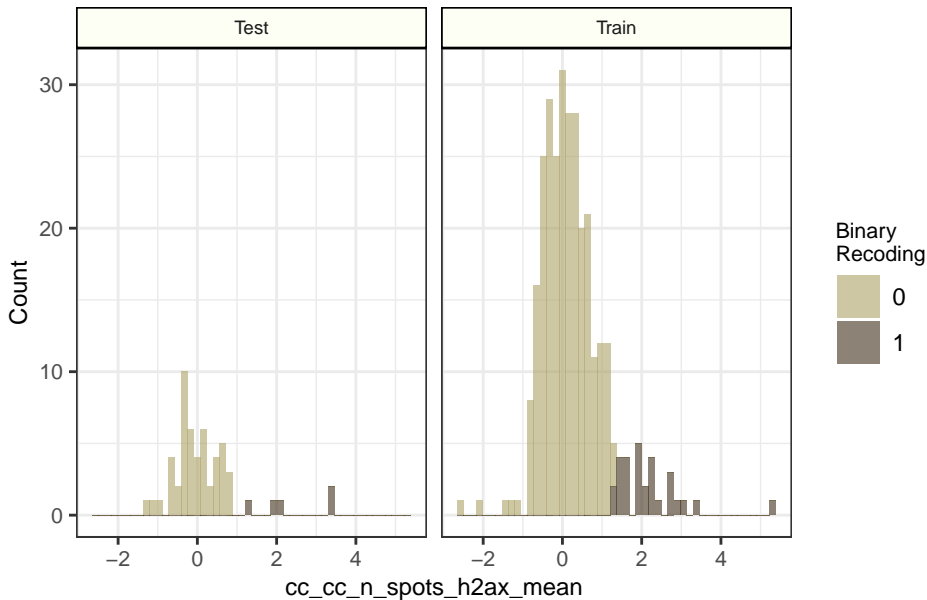
cc_cc_high_h2ax



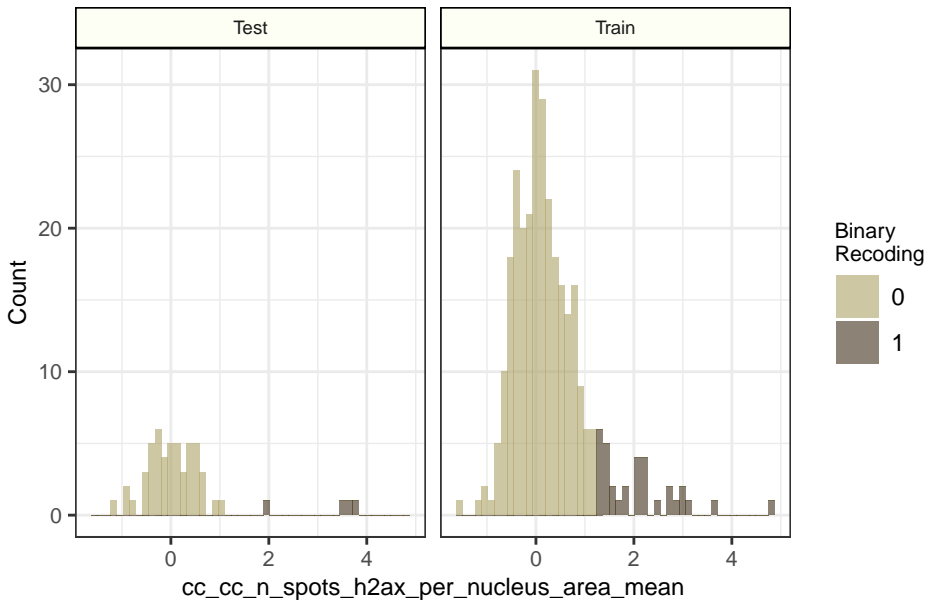
cc_cc_mitosis



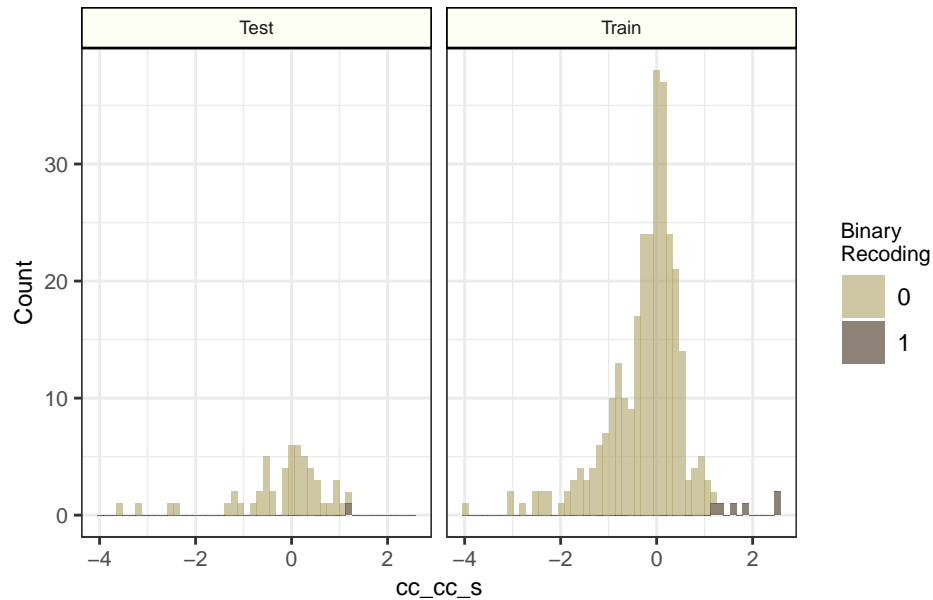
cc_cc_n_spots_h2ax_mean



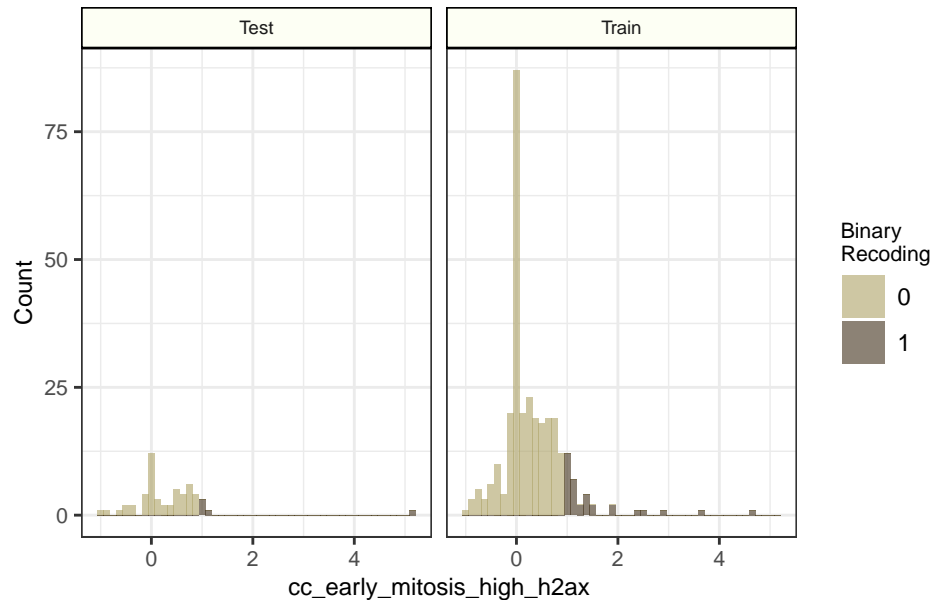
cc_cc_n_spots_h2ax_per_nucleus_area_mean



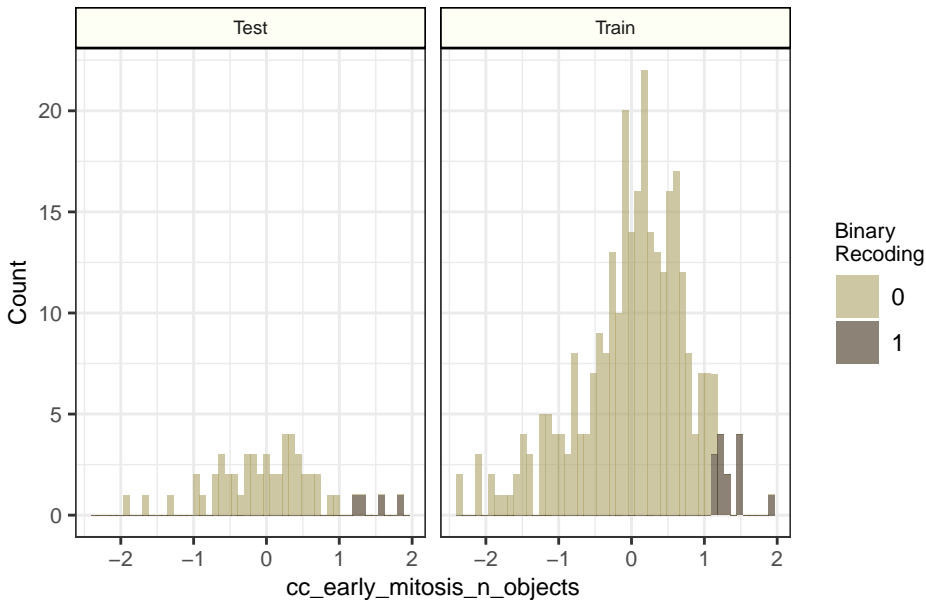
CC_CC_S



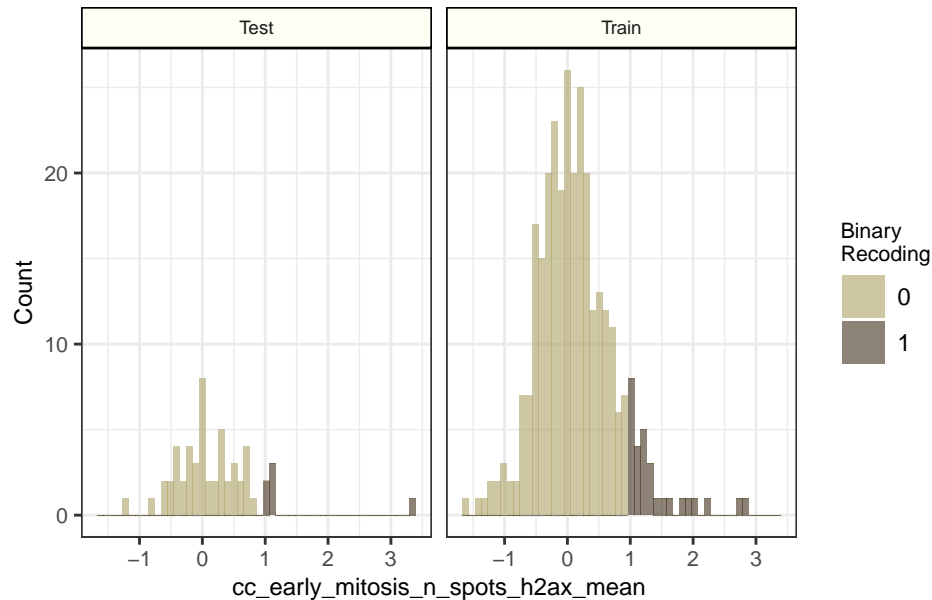
cc_early_mitosis_high_h2ax



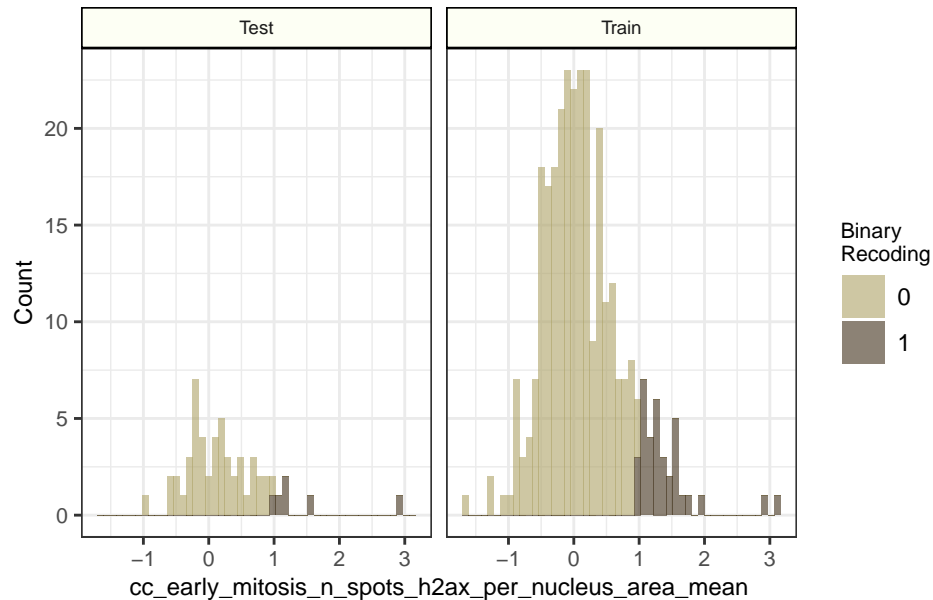
cc_early_mitosis_n_objects



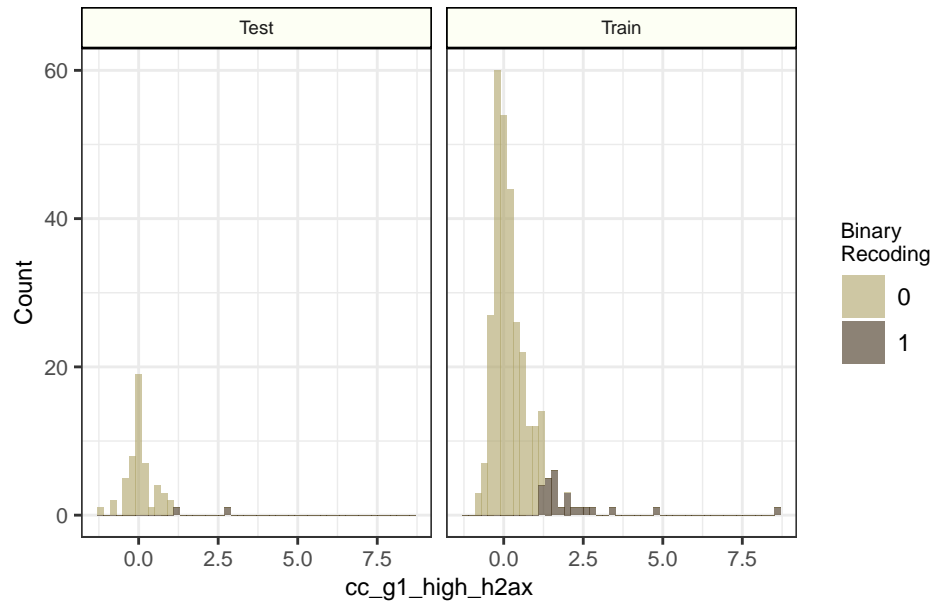
cc_early_mitosis_n_spots_h2ax_mean



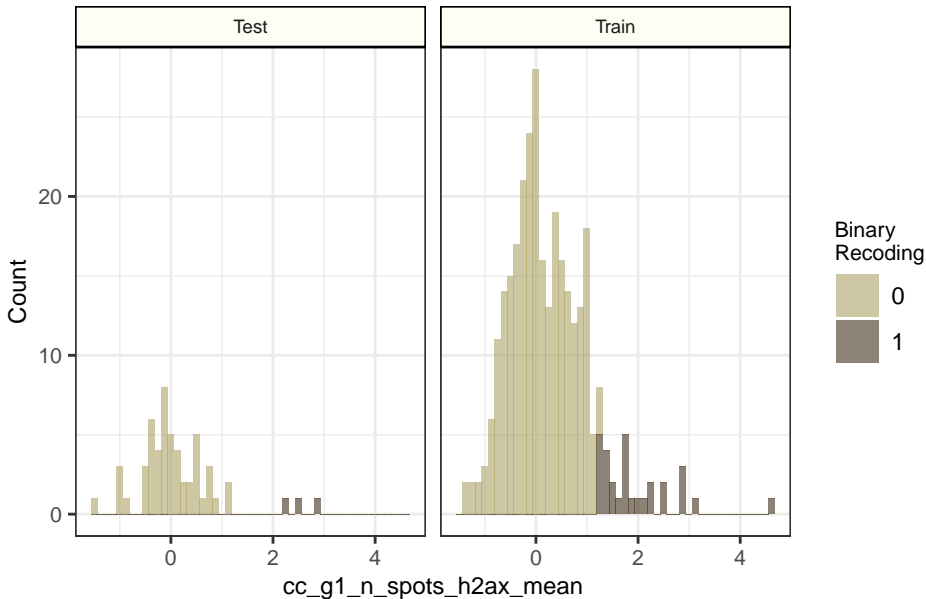
cc_early_mitosis_n_spots_h2ax_per_nucleus_area_mean



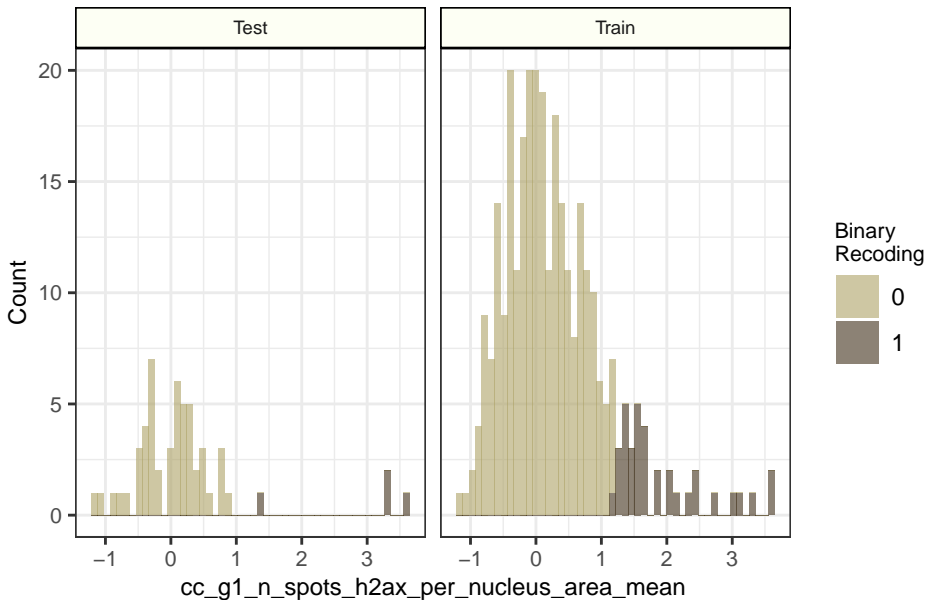
cc_g1_high_h2ax



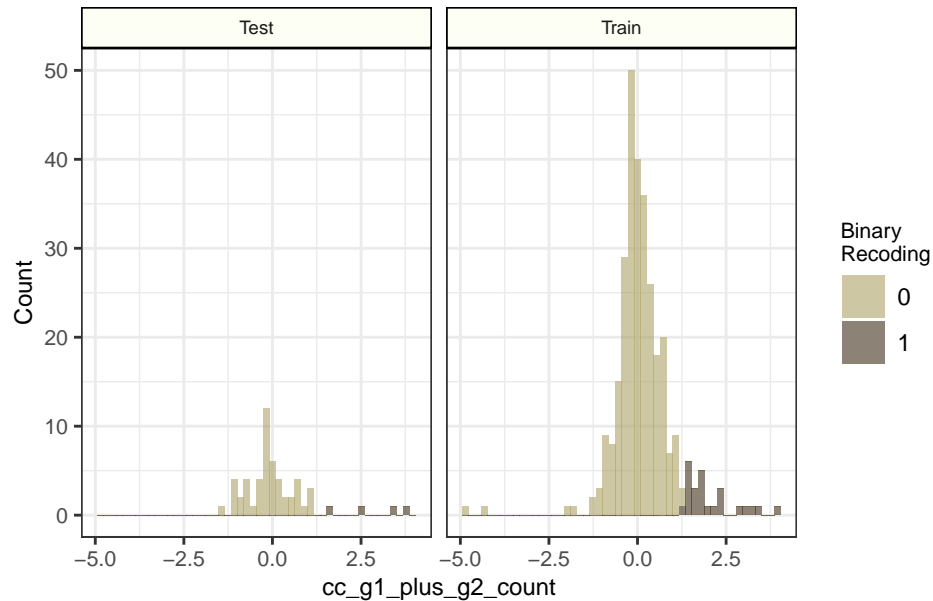
cc_g1_n_spots_h2ax_mean



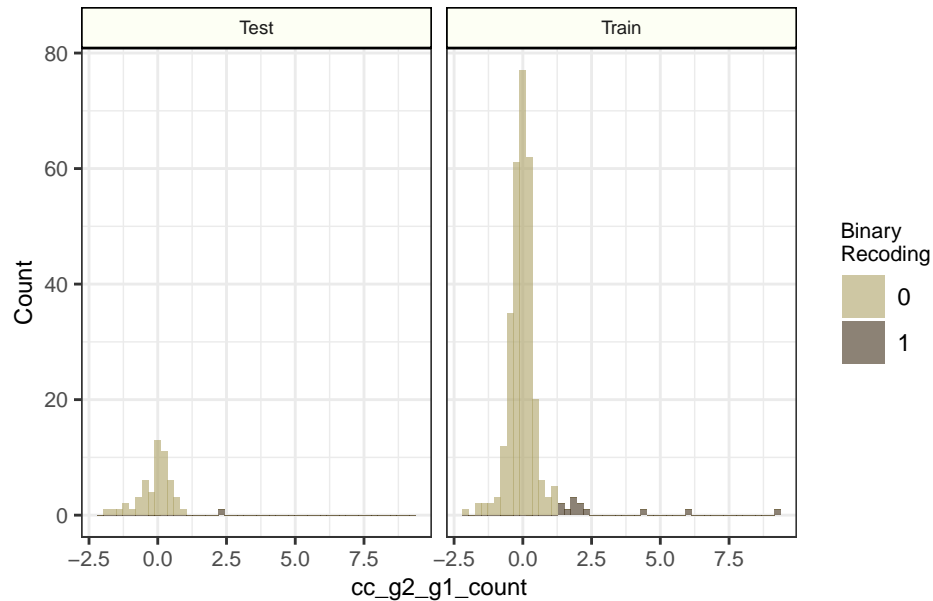
cc_g1_n_spots_h2ax_per_nucleus_area_mean



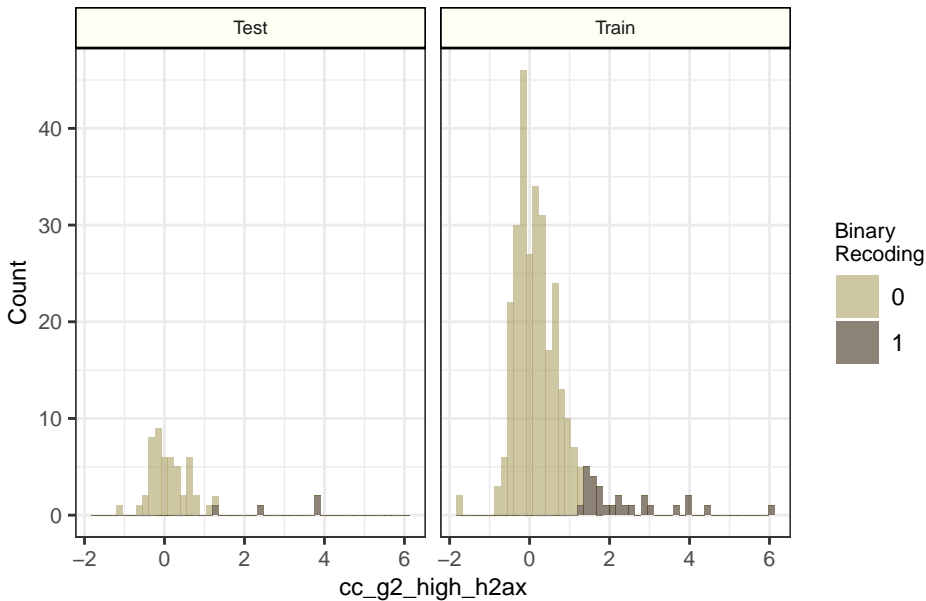
cc_g1_plus_g2_count



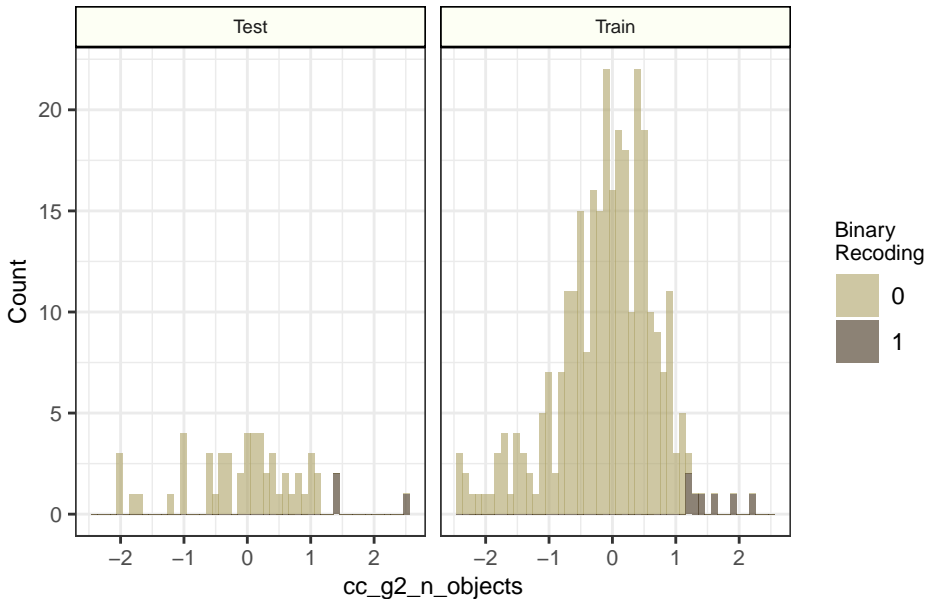
cc_g2_g1_count



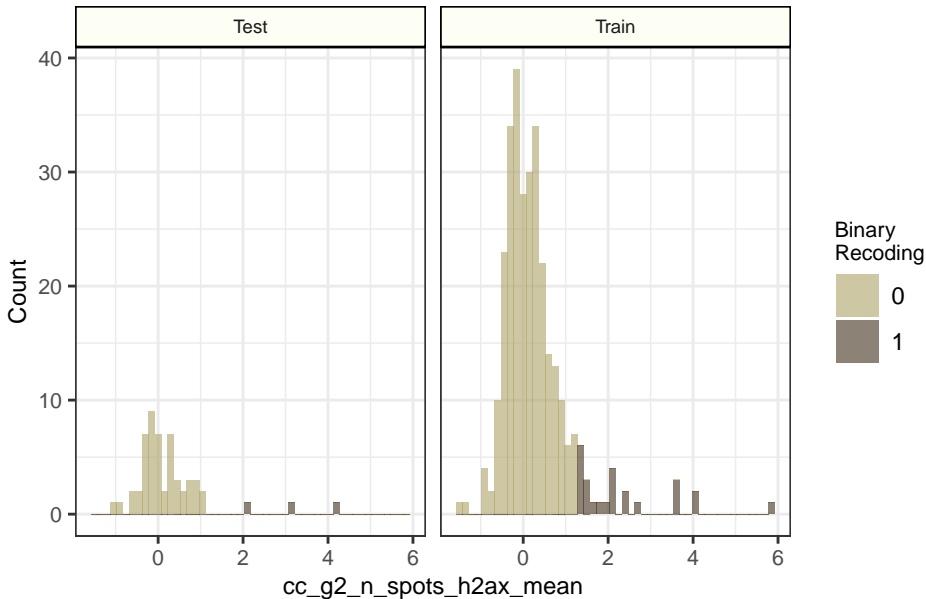
cc_g2_high_h2ax



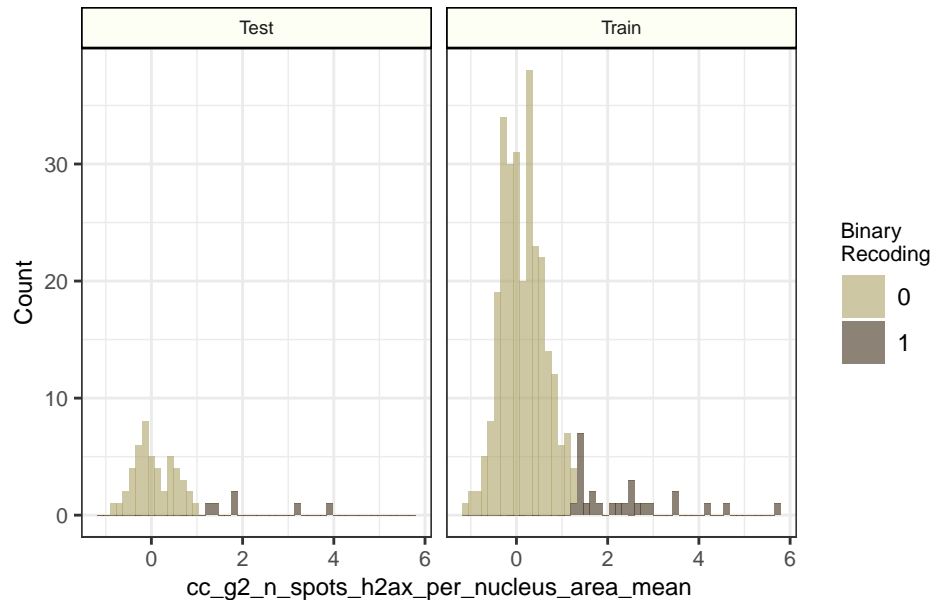
cc_g2_n_objects



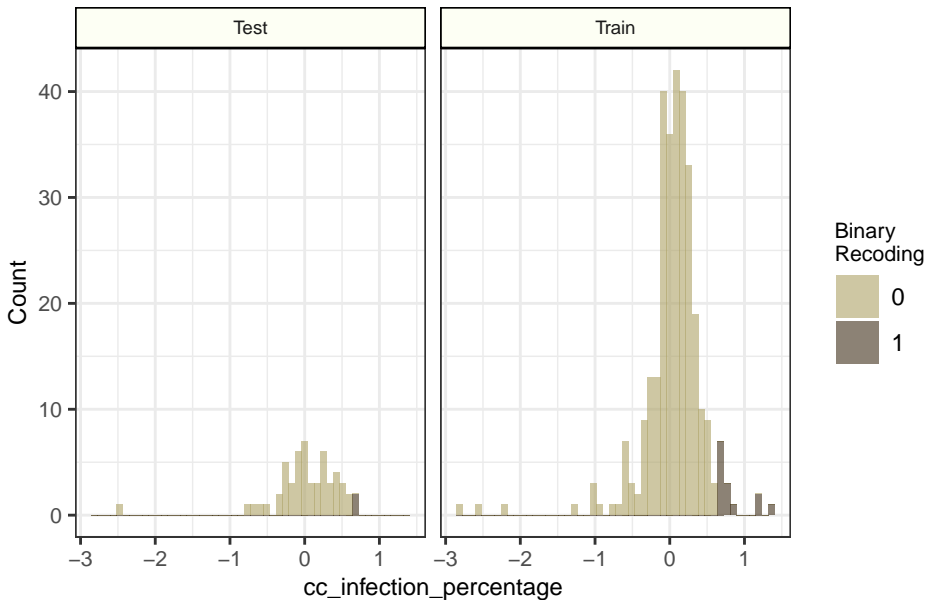
cc_g2_n_spots_h2ax_mean



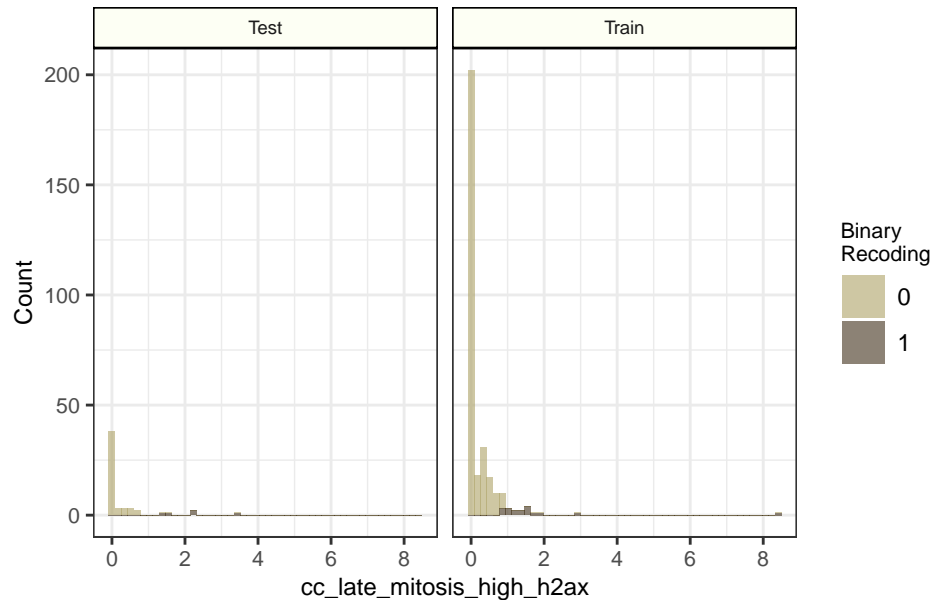
cc_g2_n_spots_h2ax_per_nucleus_area_mean



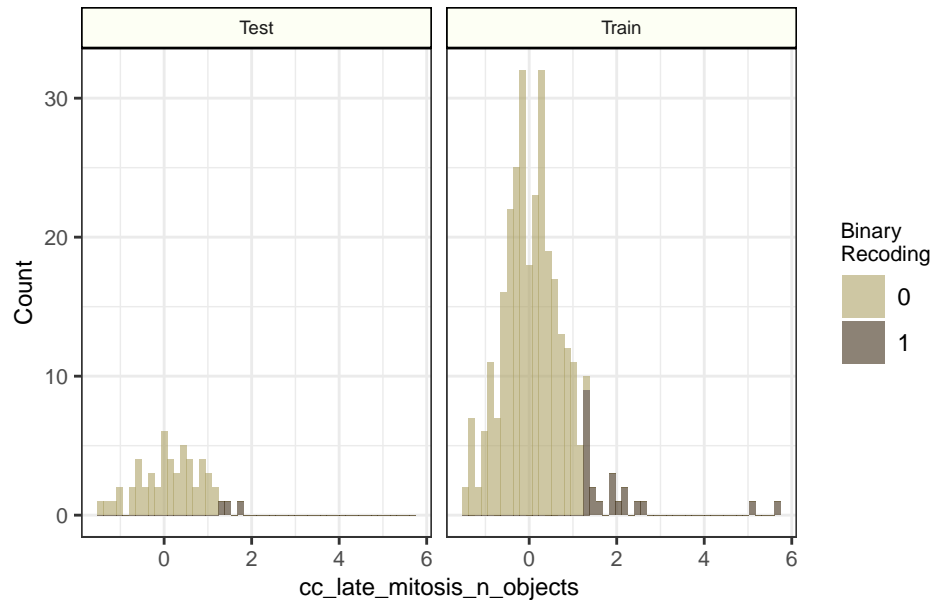
cc_infection_percentage



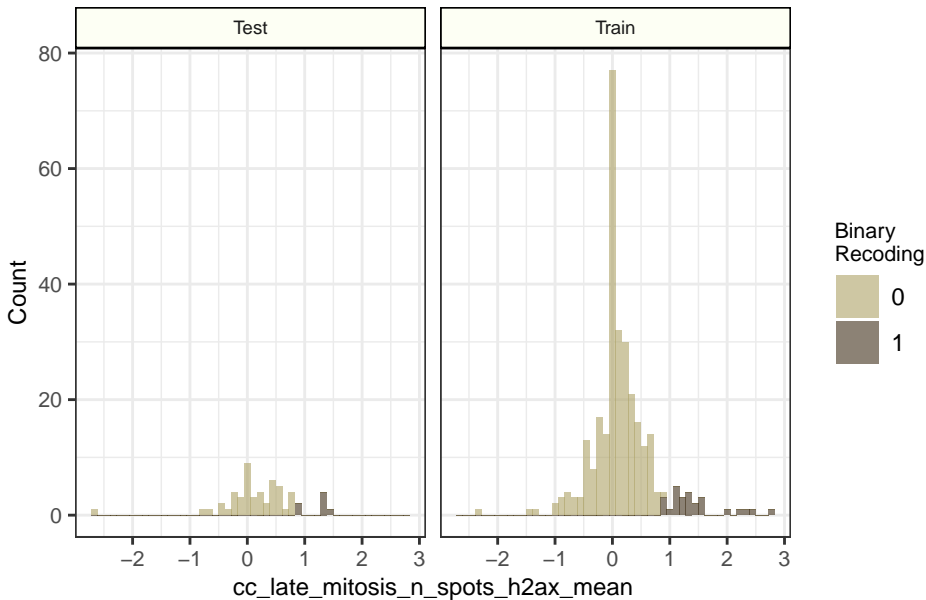
cc_late_mitosis_high_h2ax



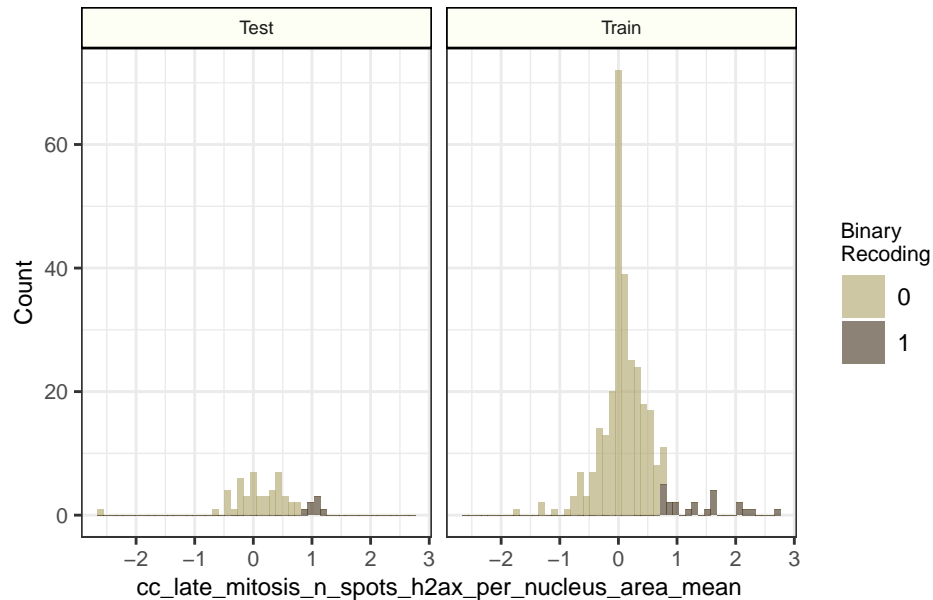
cc_late_mitosis_n_objects



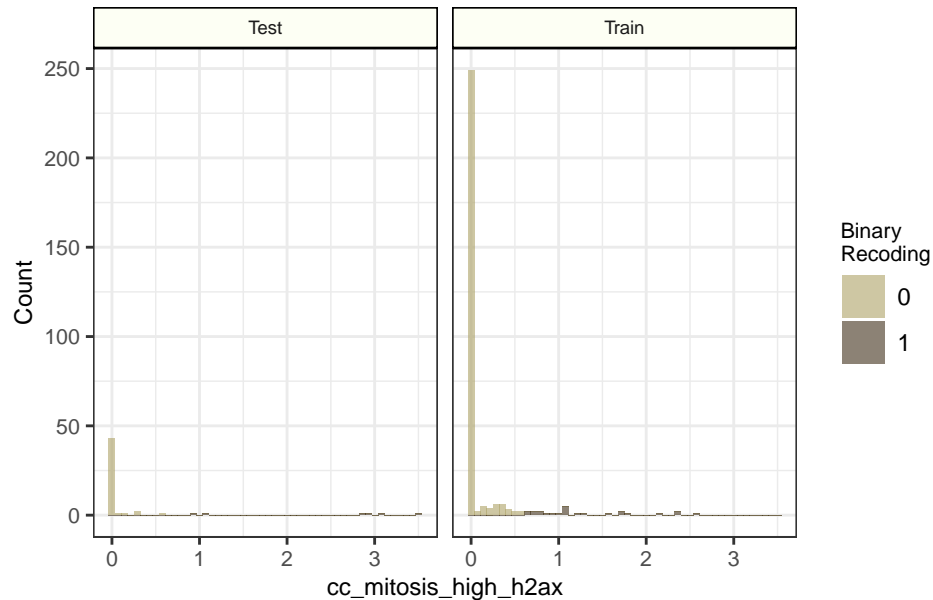
cc_late_mitosis_n_spots_h2ax_mean



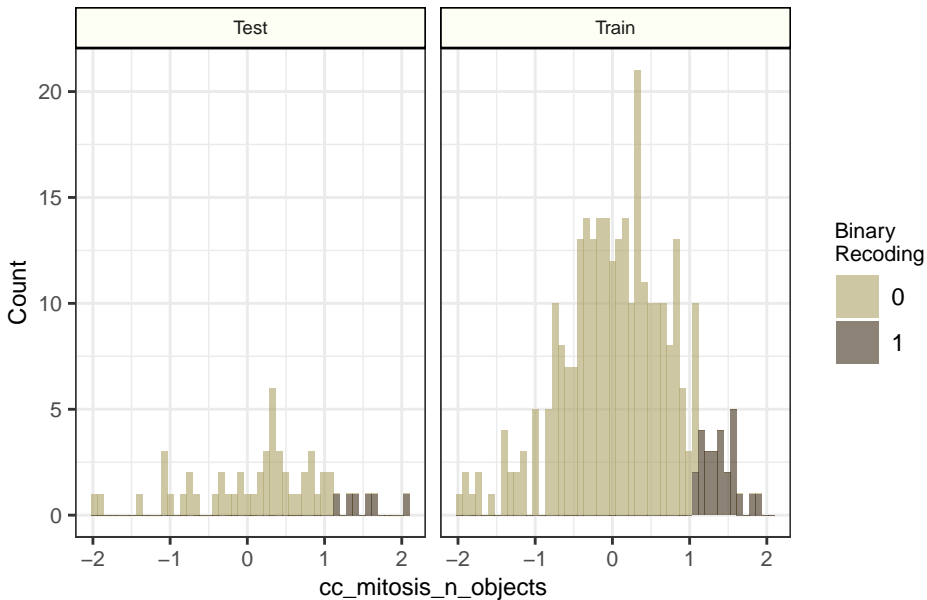
cc_late_mitosis_n_spots_h2ax_per_nucleus_area_m



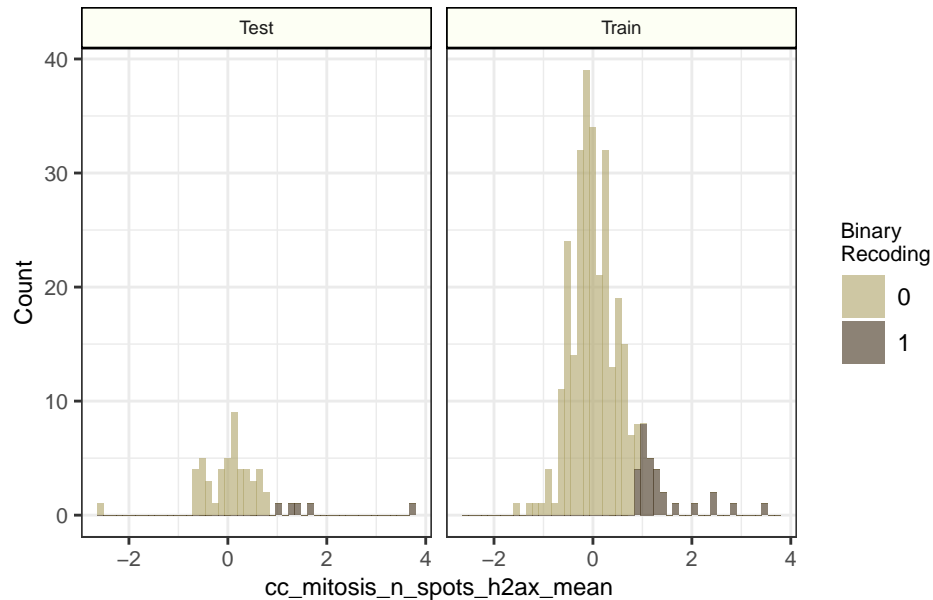
cc_mitosis_high_h2ax



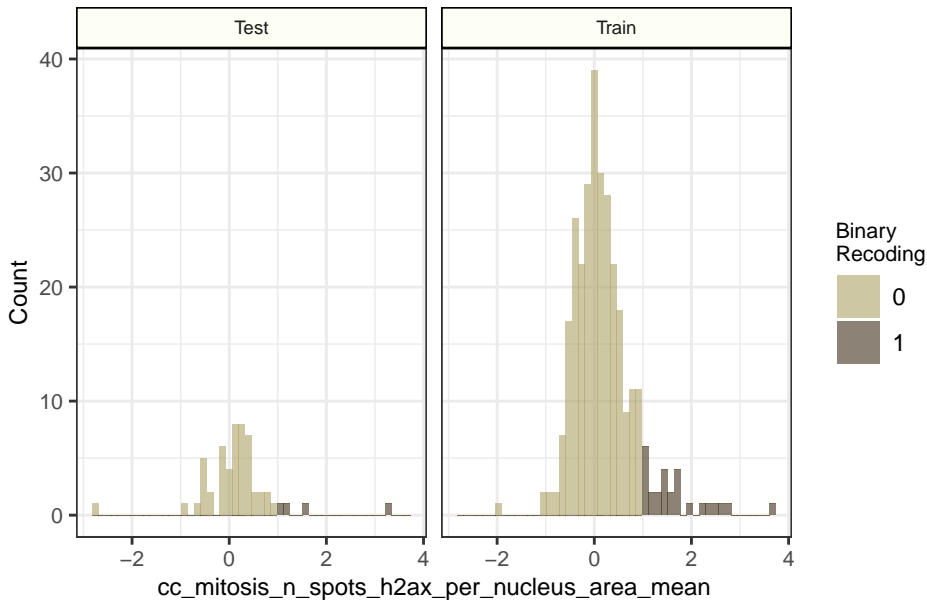
cc_mitosis_n_objects



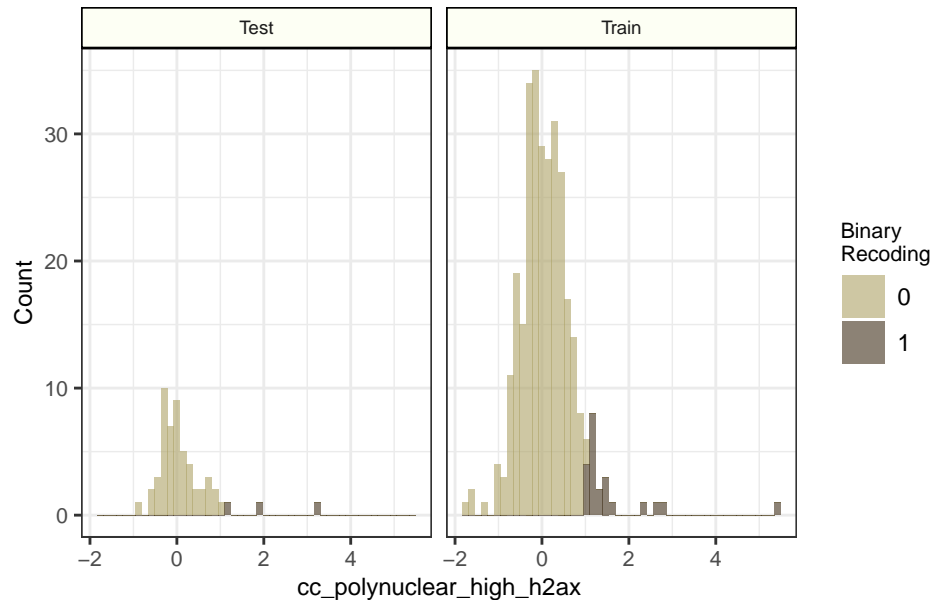
cc_mitosis_n_spots_h2ax_mean



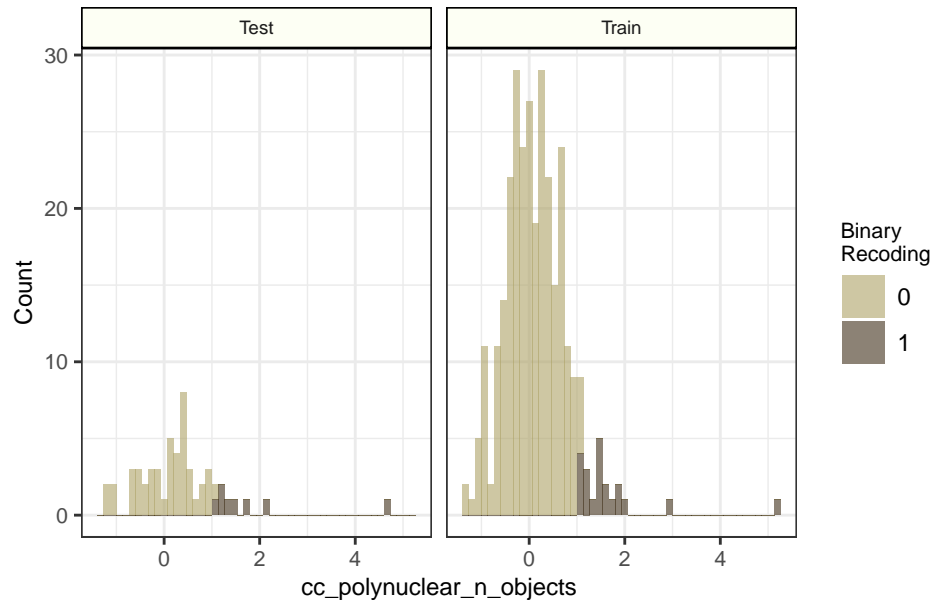
cc_mitosis_n_spots_h2ax_per_nucleus_area_mean



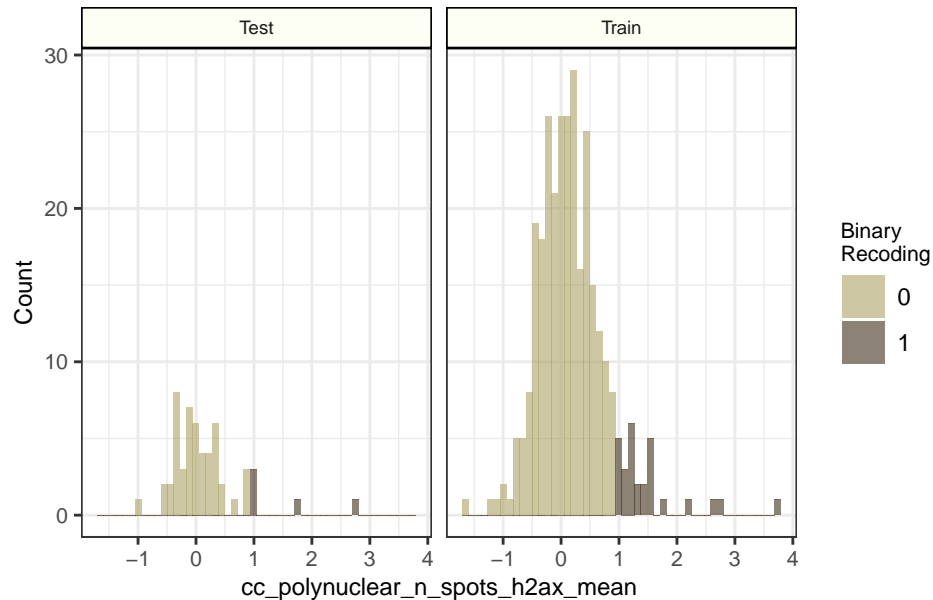
cc_polynuclear_high_h2ax



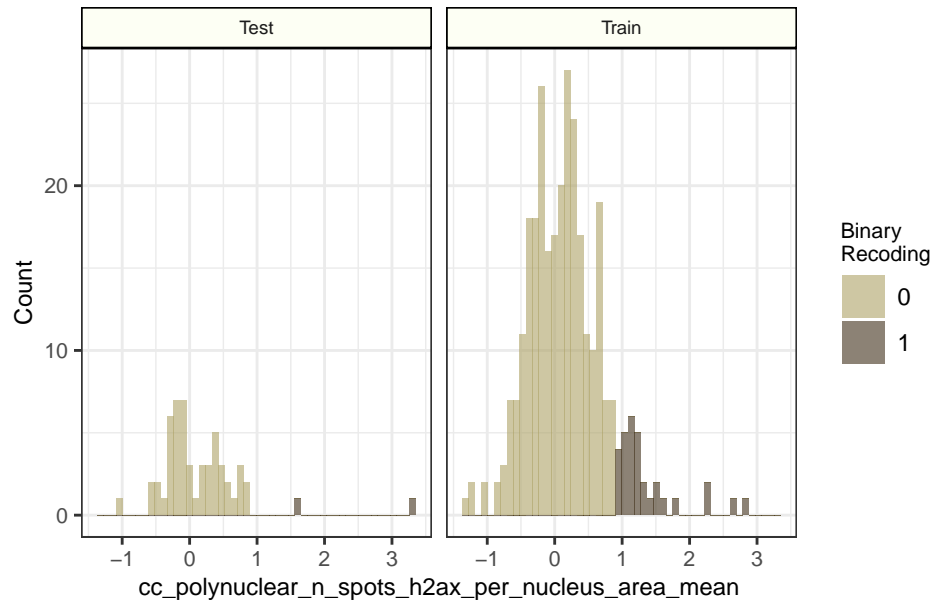
cc_polynuclear_n_objects



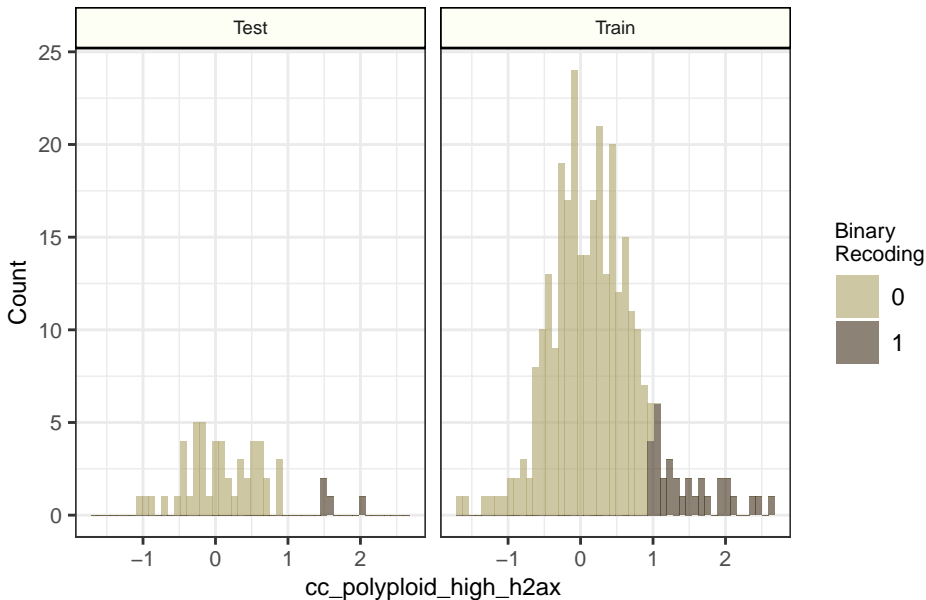
cc_polynuclear_n_spots_h2ax_mean



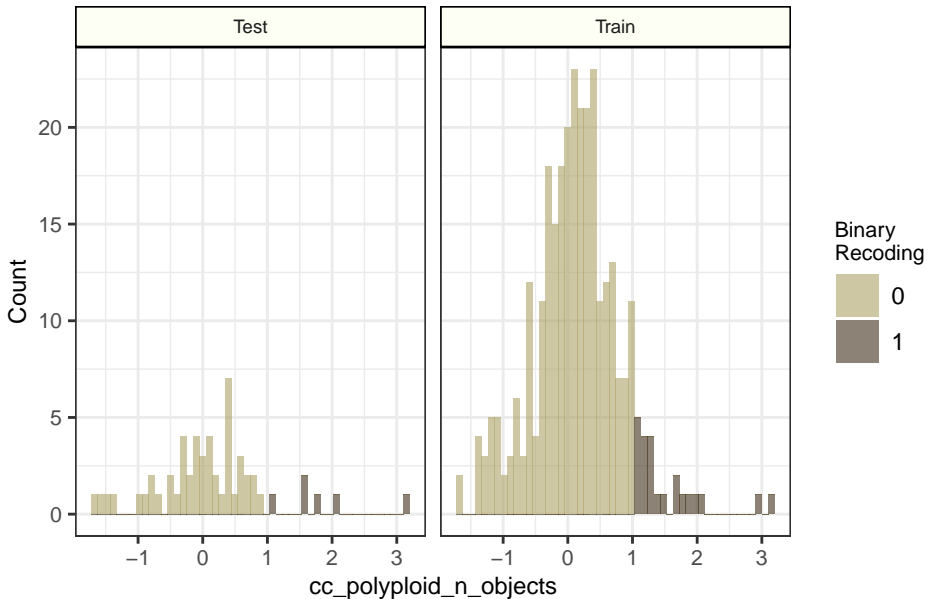
cc_polynuclear_n_spots_h2ax_per_nucleus_area_mean



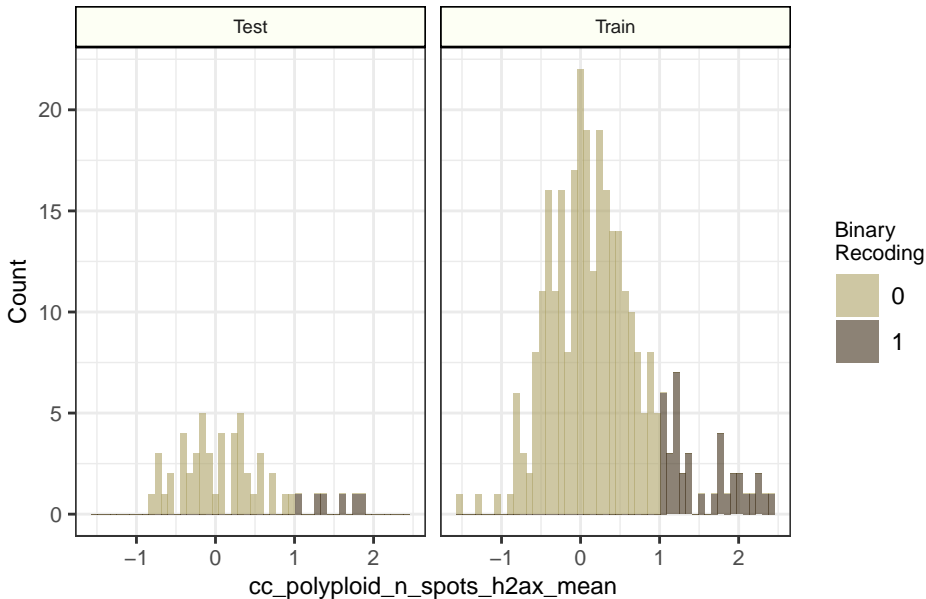
cc_polyploid_high_h2ax



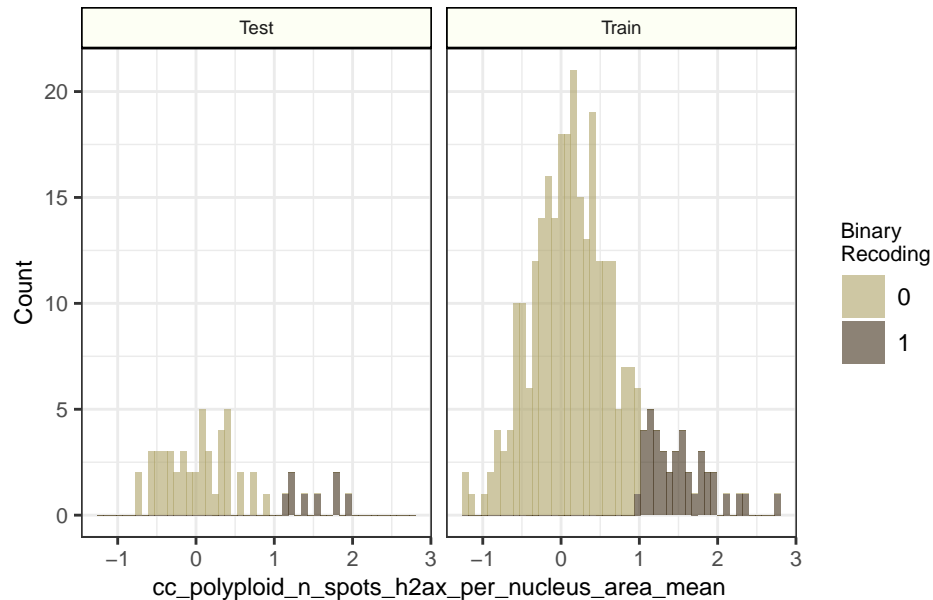
cc_polyplloid_n_objects



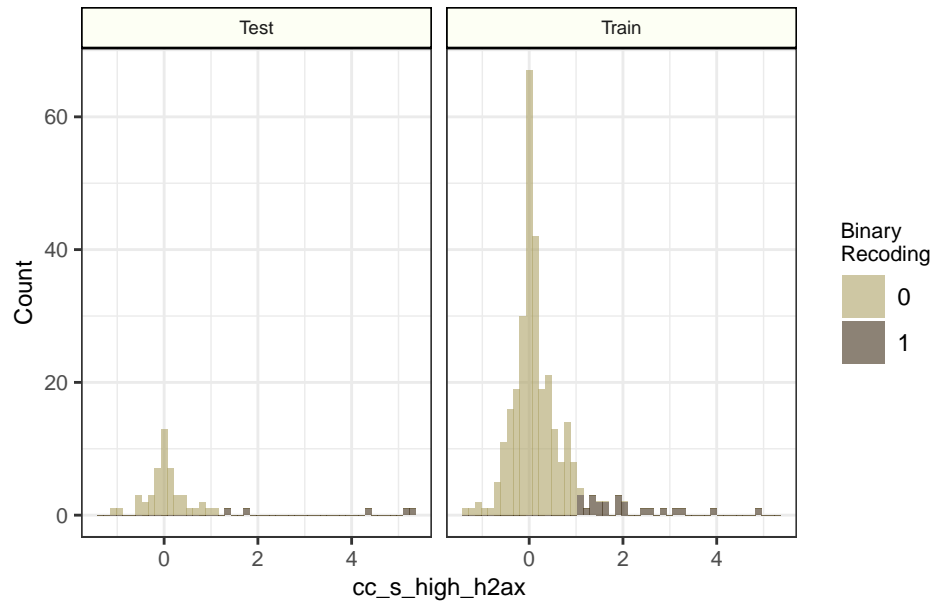
cc_polyploid_n_spots_h2ax_mean



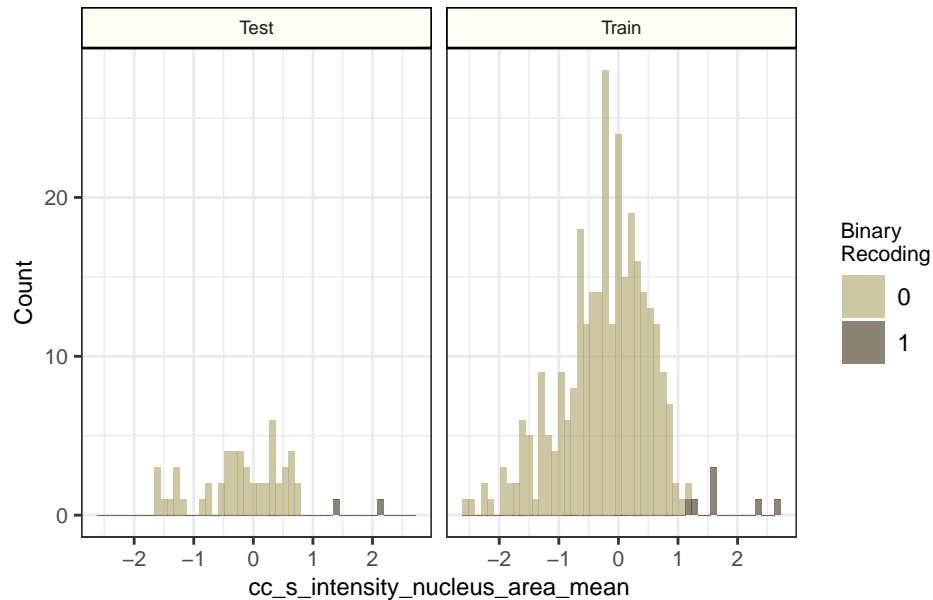
cc_polyploid_n_spots_h2ax_per_nucleus_area_mean



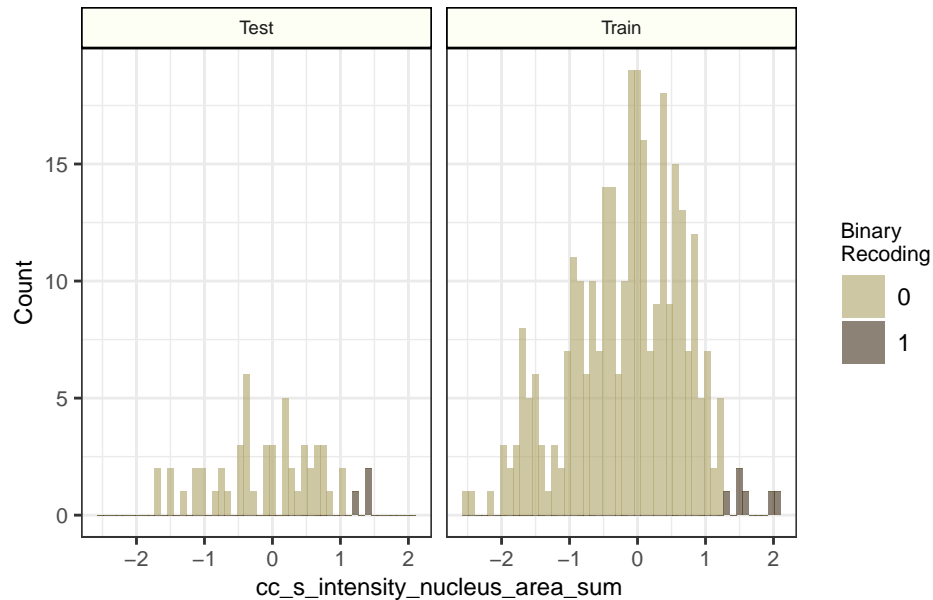
cc_s_high_h2ax



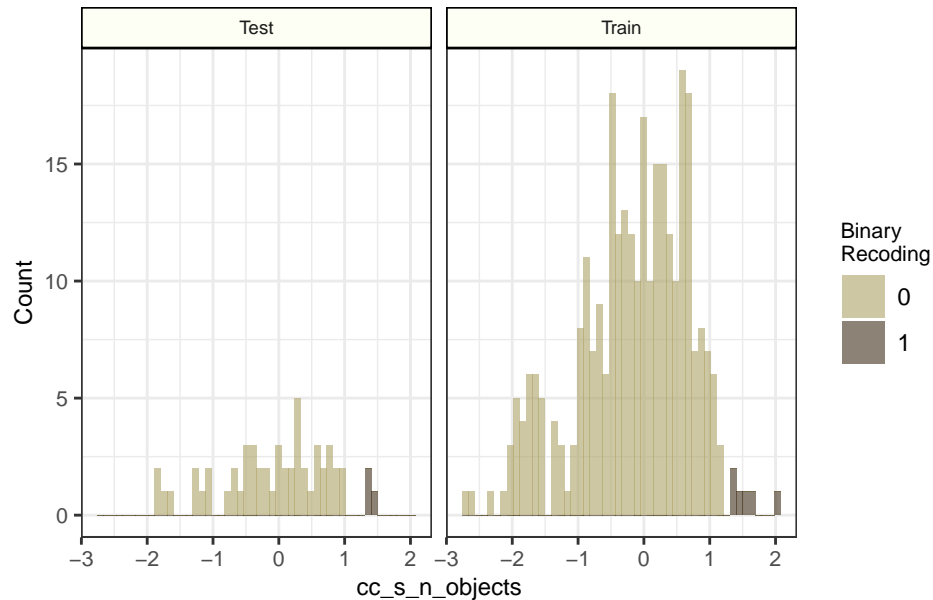
cc_s_intensity_nucleus_area_mean



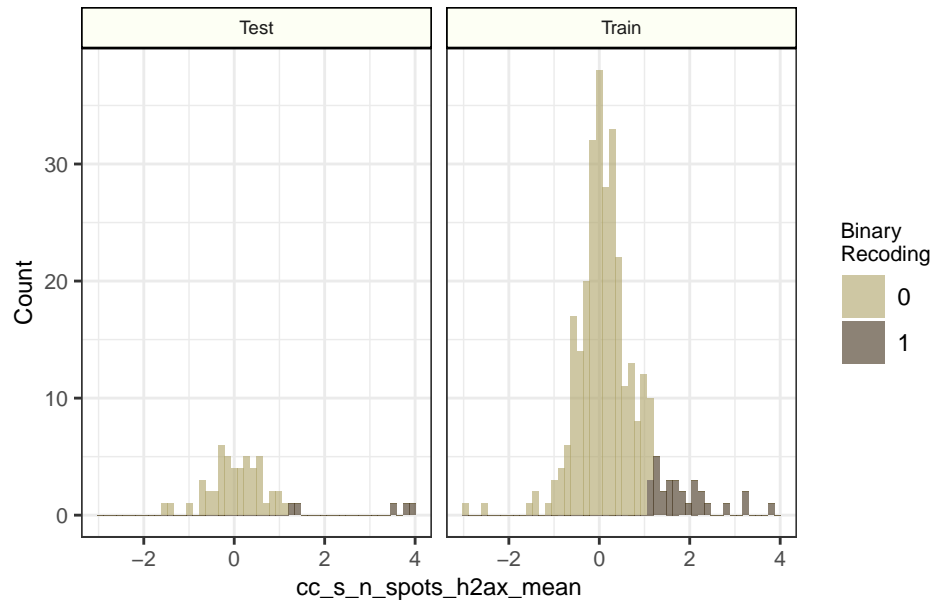
cc_s_intensity_nucleus_area_sum



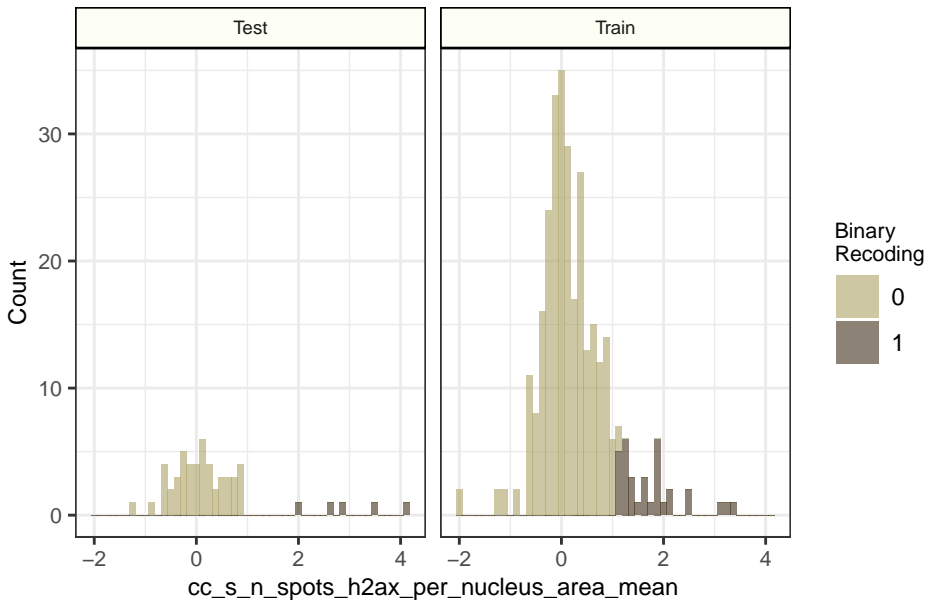
cc_s_n_objects



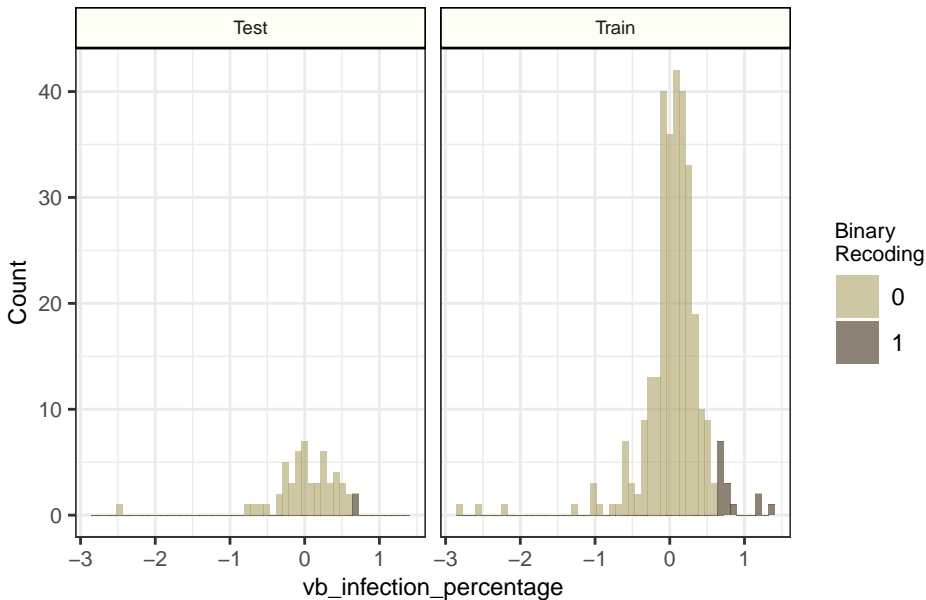
cc_s_n_spots_h2ax_mean



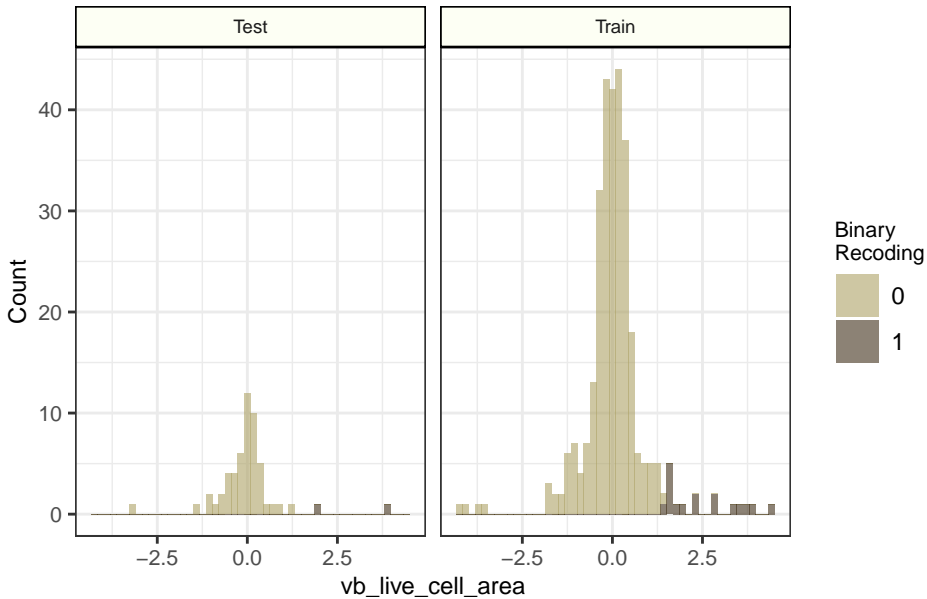
cc_s_n_spots_h2ax_per_nucleus_area_mean



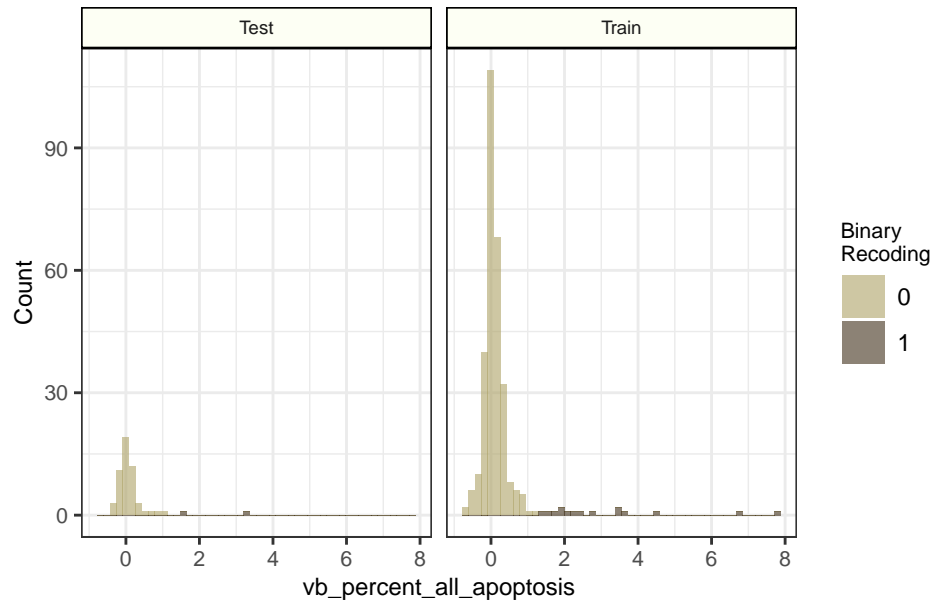
vb_infection_percentage



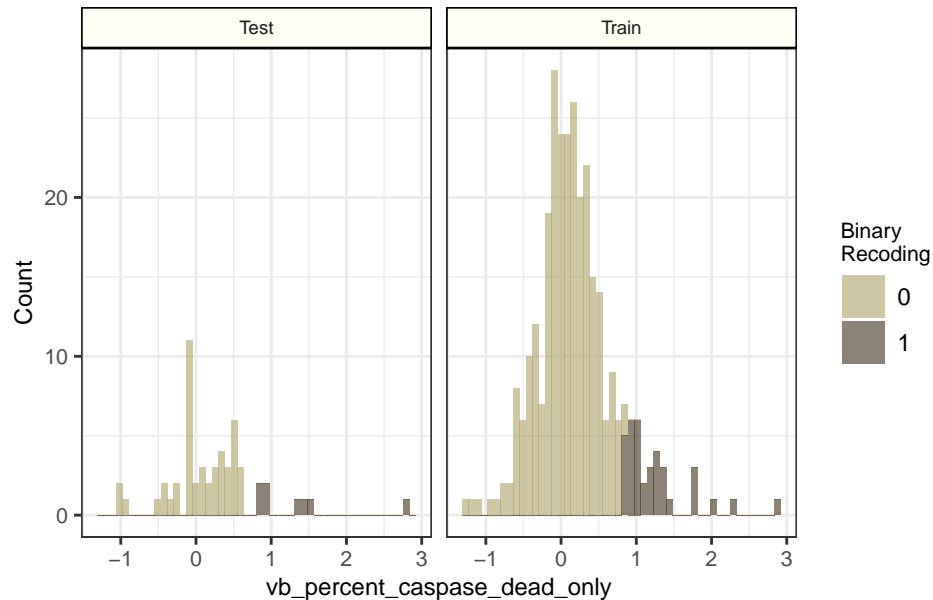
vb_live_cell_area



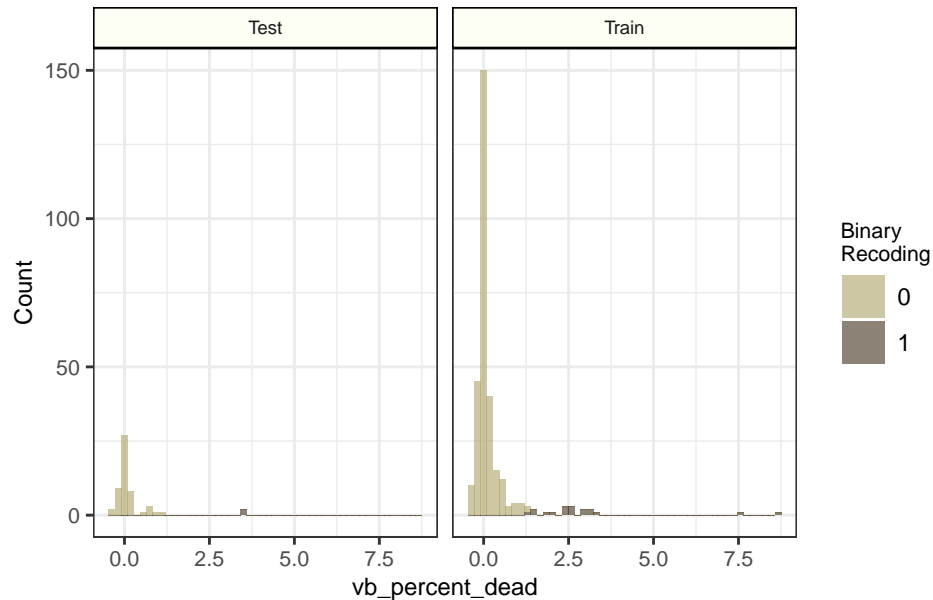
vb_percent_all_apoptosis



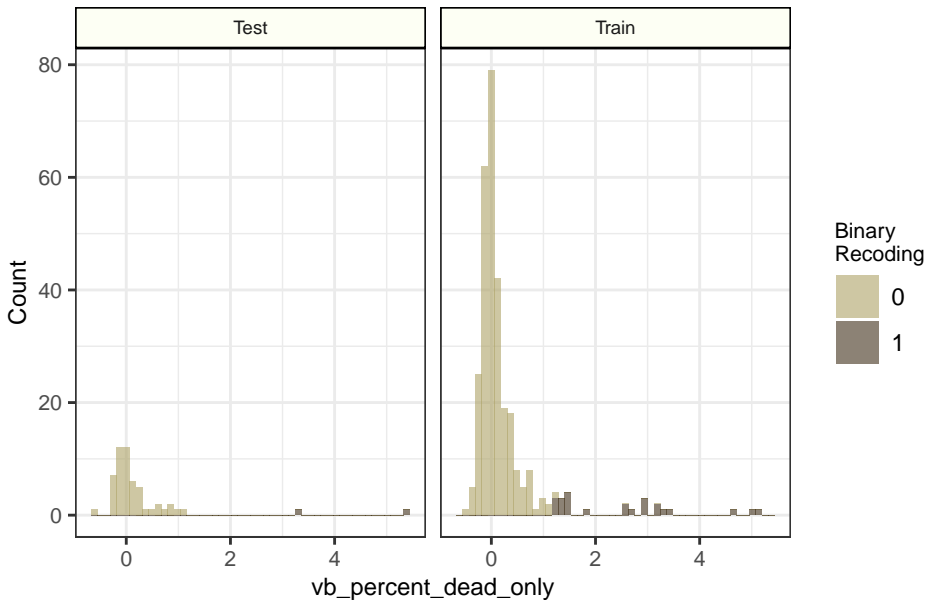
vb_percent_caspase_dead_only



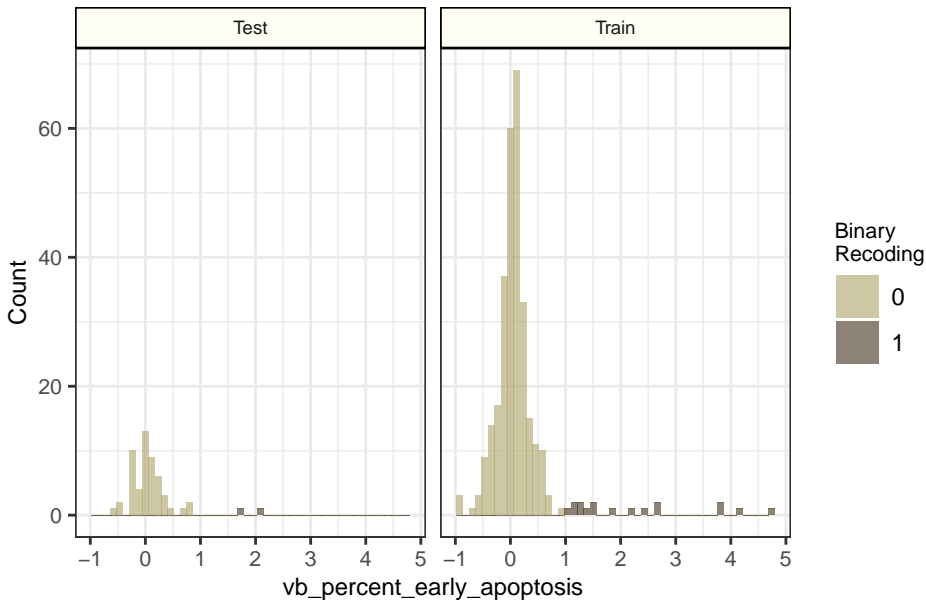
vb_percent_dead



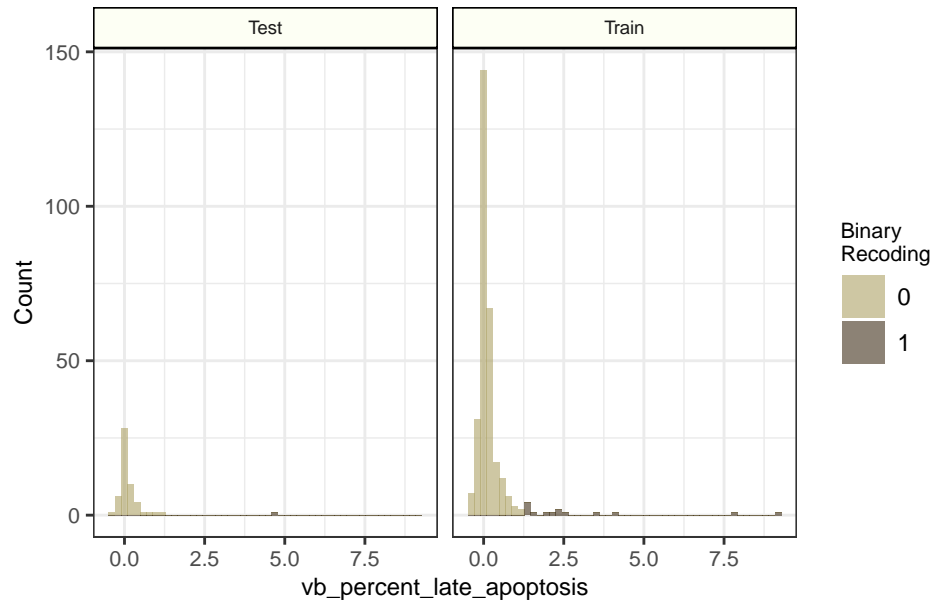
vb_percent_dead_only



vb_percent_early_apoptosis



vb_percent_late_apoptosis



vb_ros_mean

