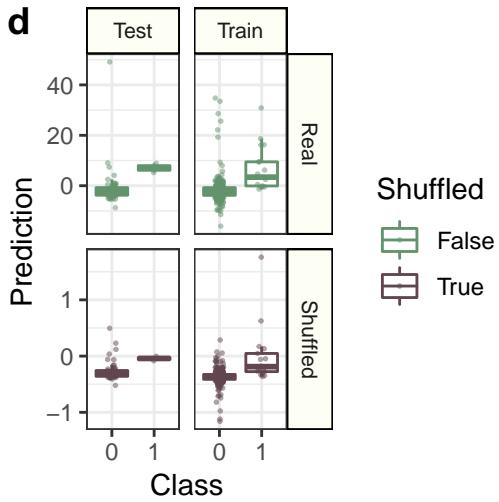
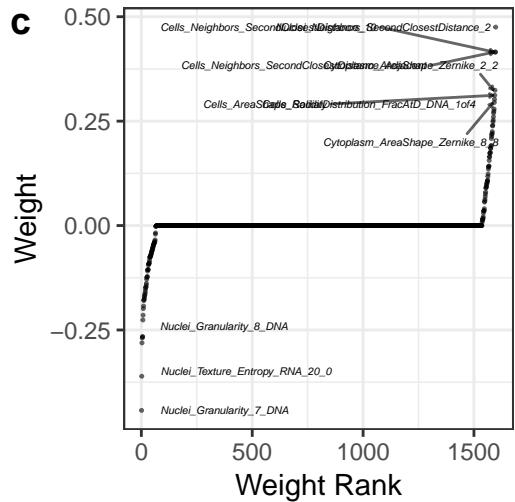
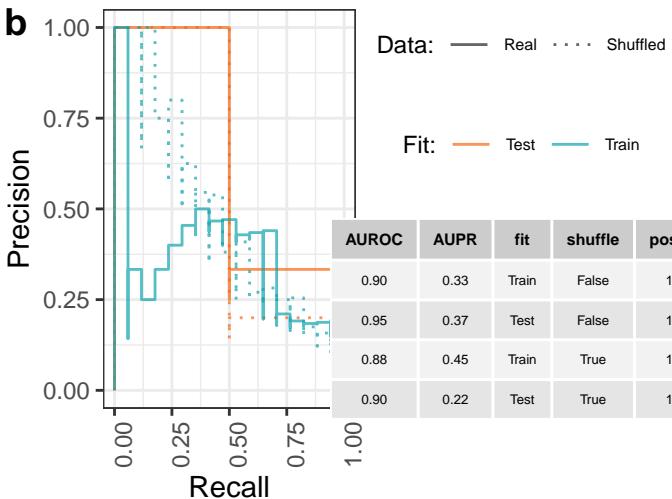
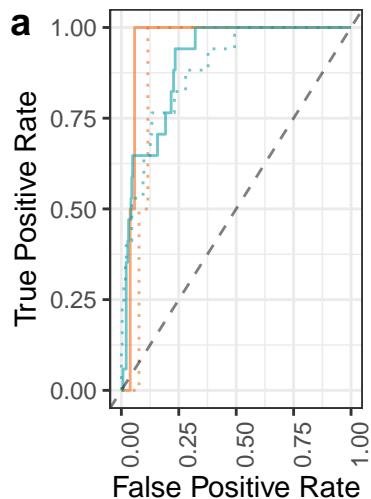
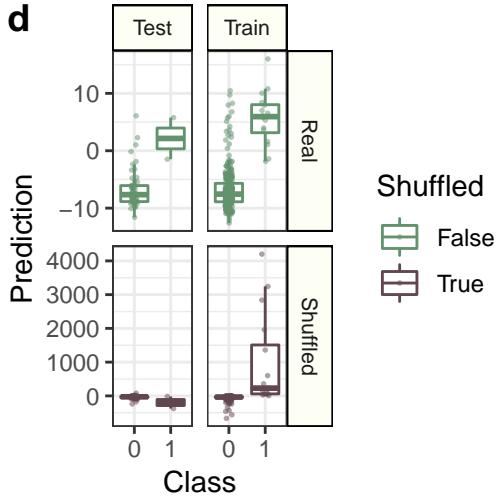
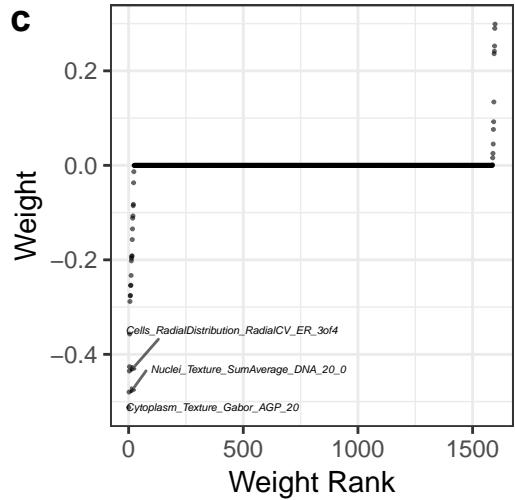
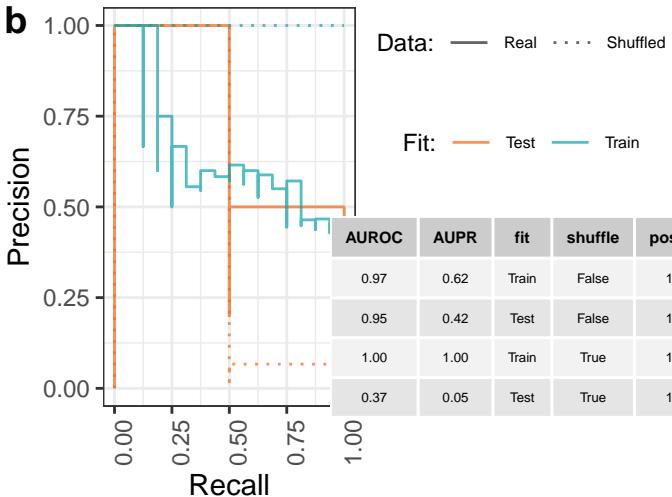
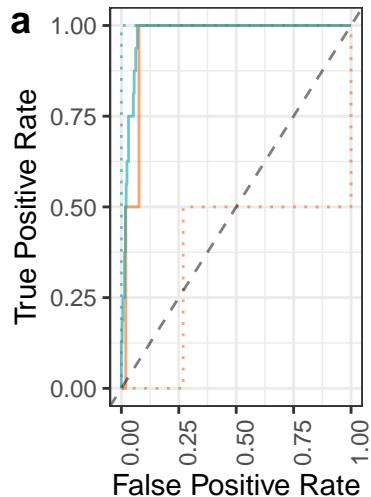


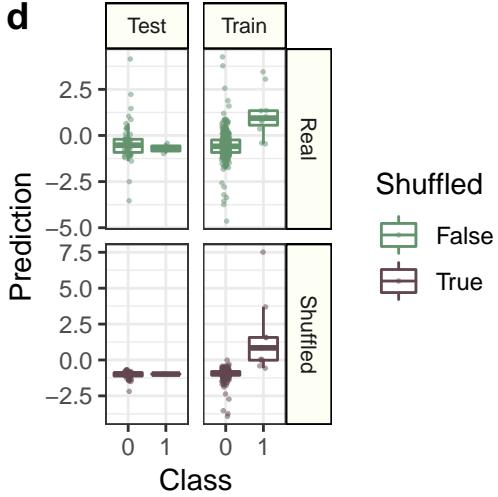
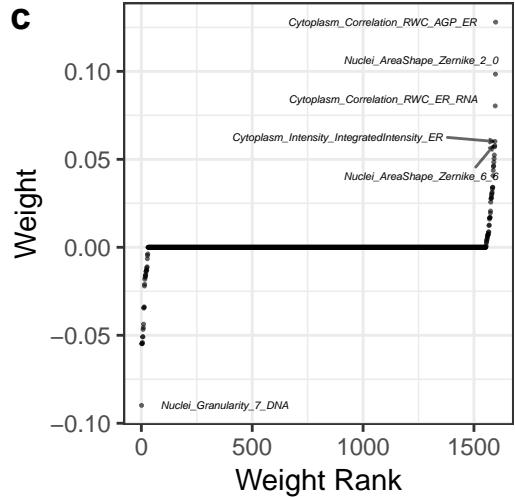
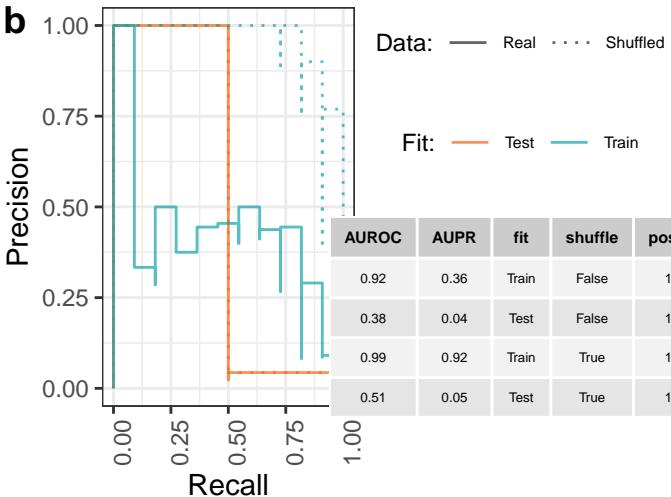
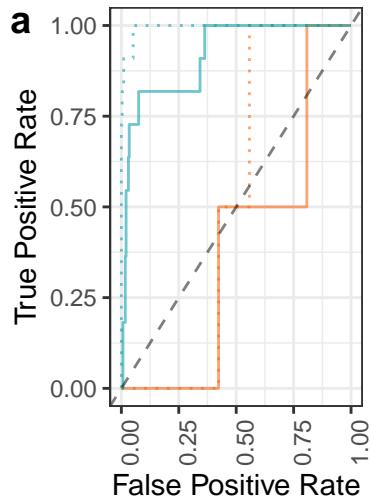
Performance: cc_g2_ph3_pos_high_n_spots_h2ax_mean



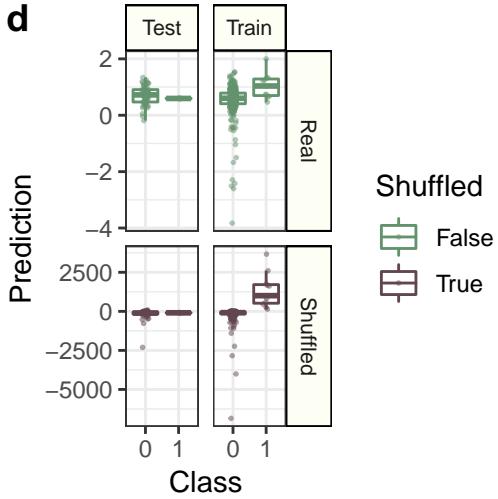
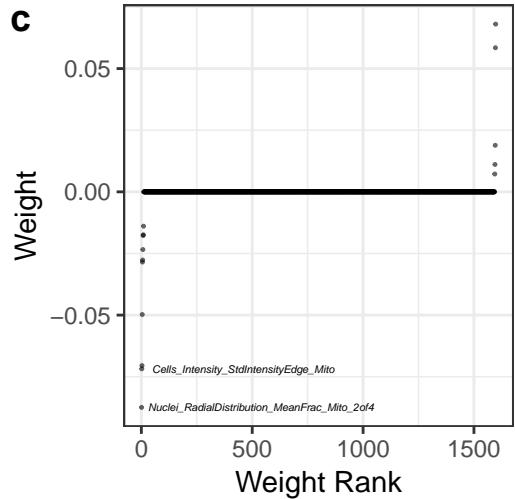
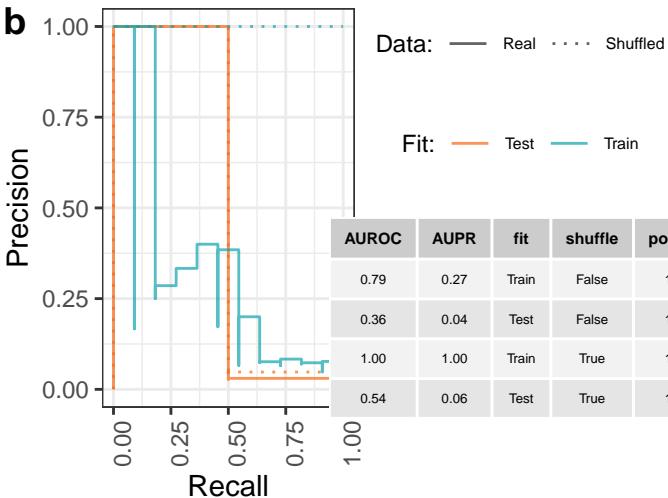
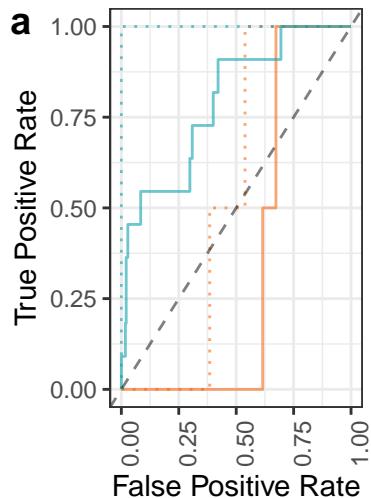
Performance: cc_g2_plus_all_m



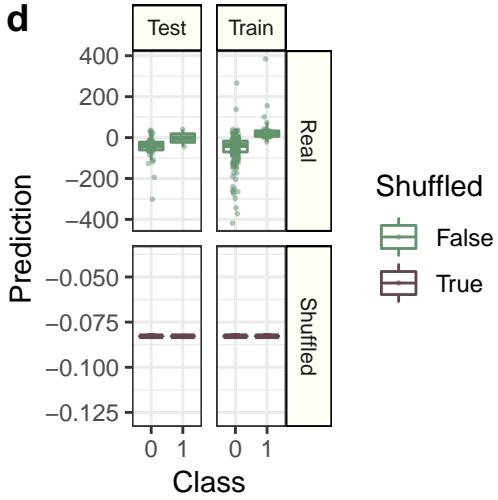
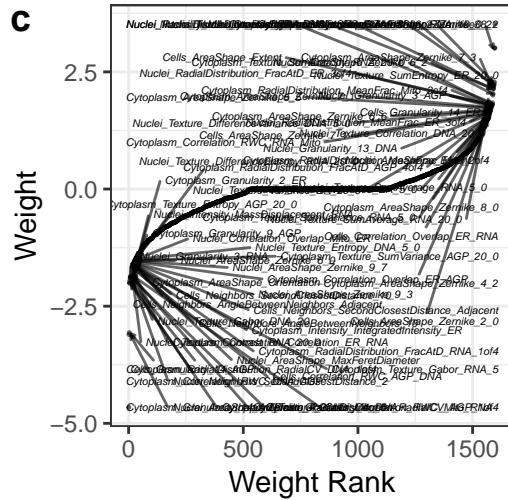
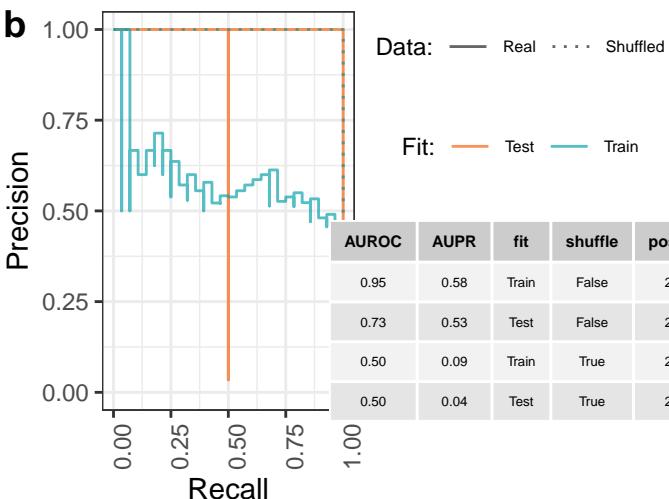
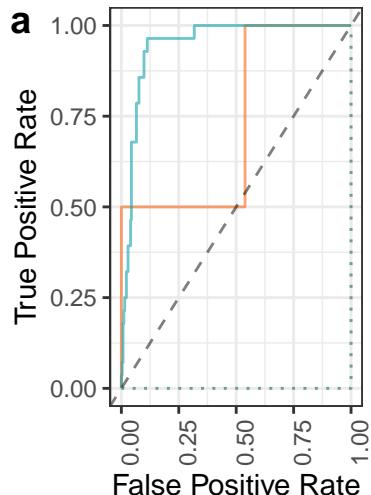
Performance: cc_infection_percentage



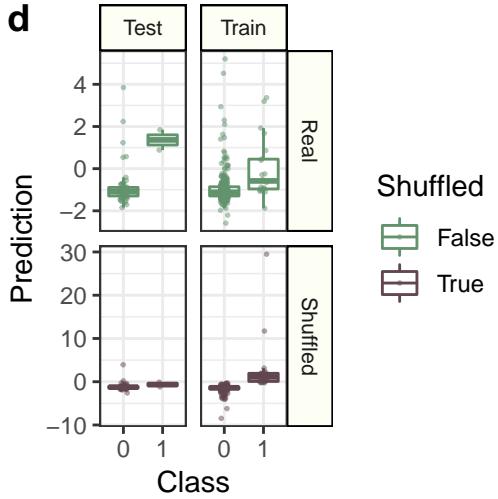
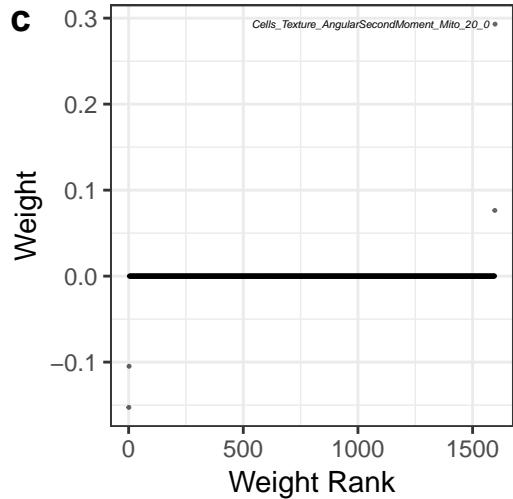
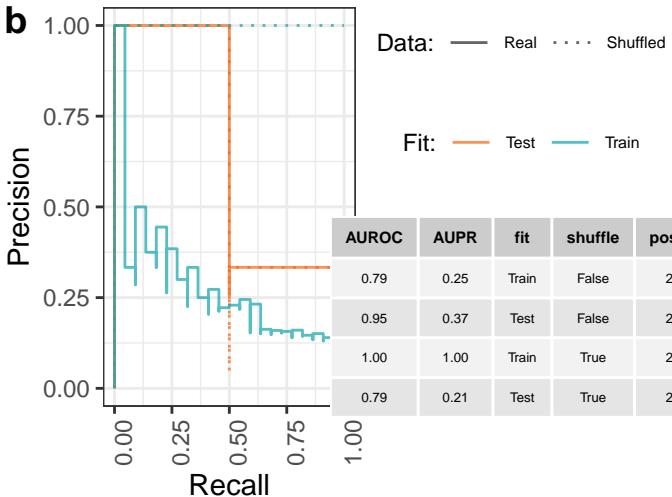
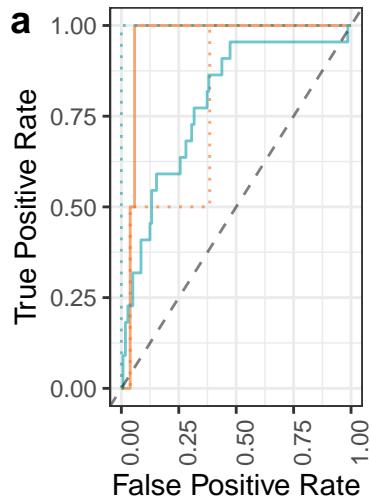
Performance: cc_mitosis_ph3_neg_n_spots_per_nucleus



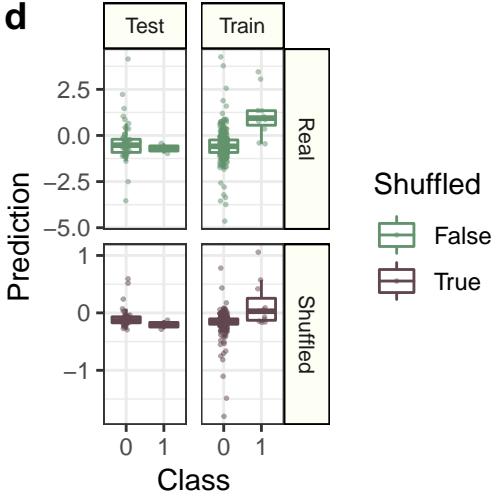
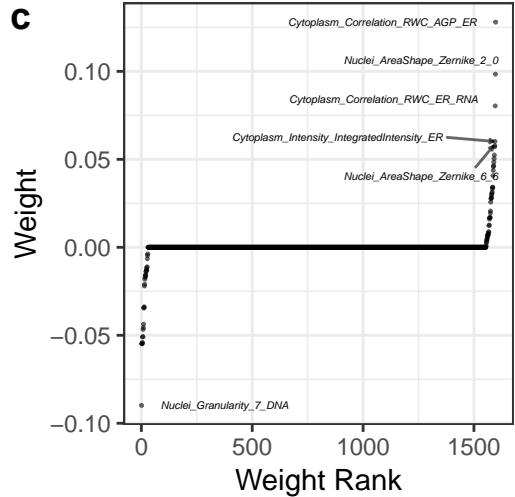
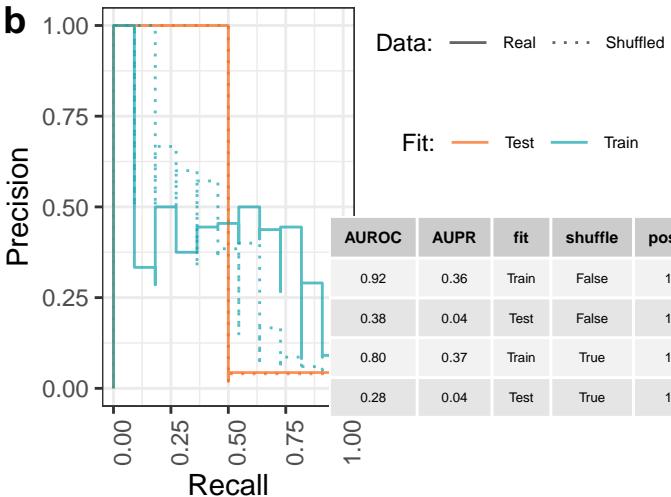
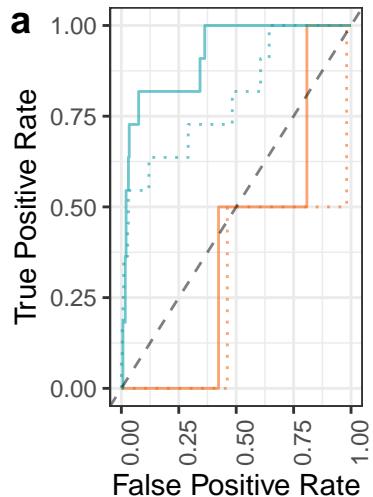
Performance: cc_mitosis_ph3_pos_n_spots_per_nucleus



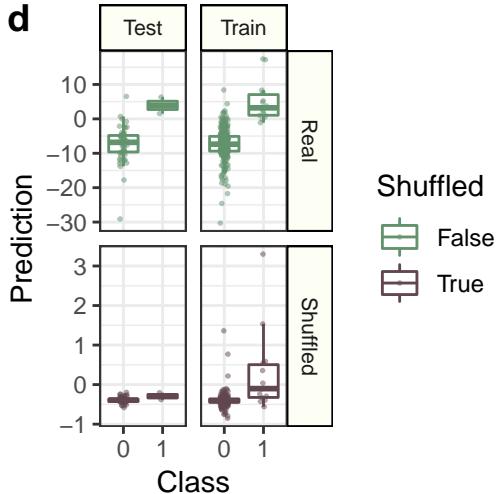
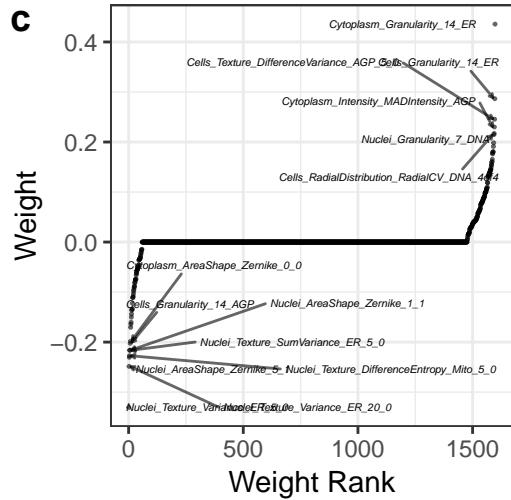
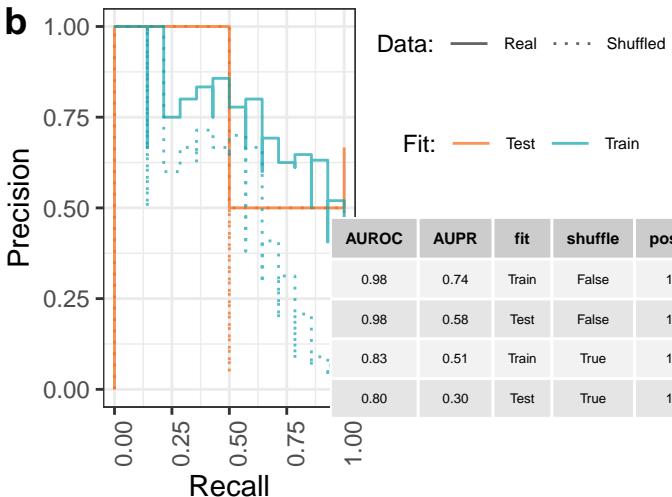
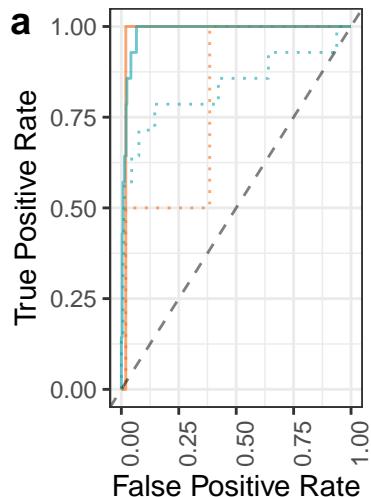
Performance: cc_polynuclear_high_n_spots_h2ax_mean



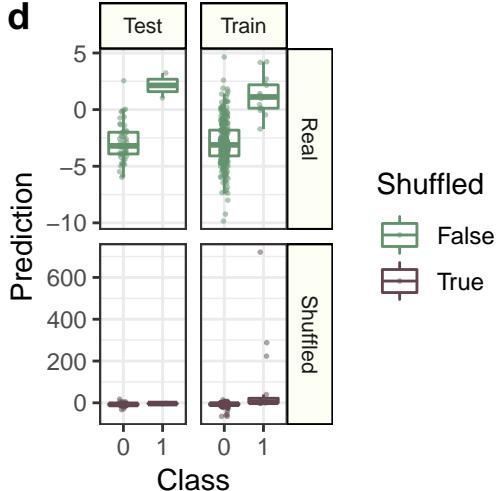
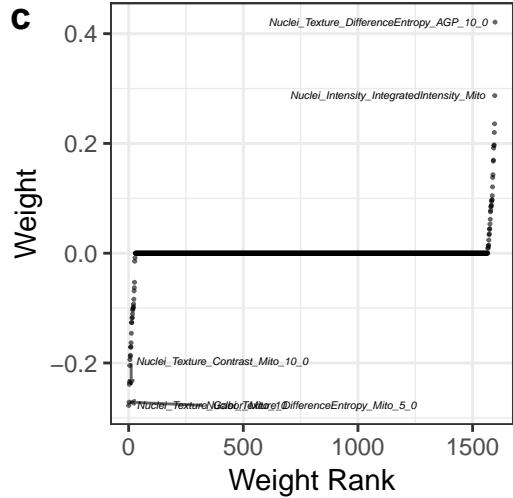
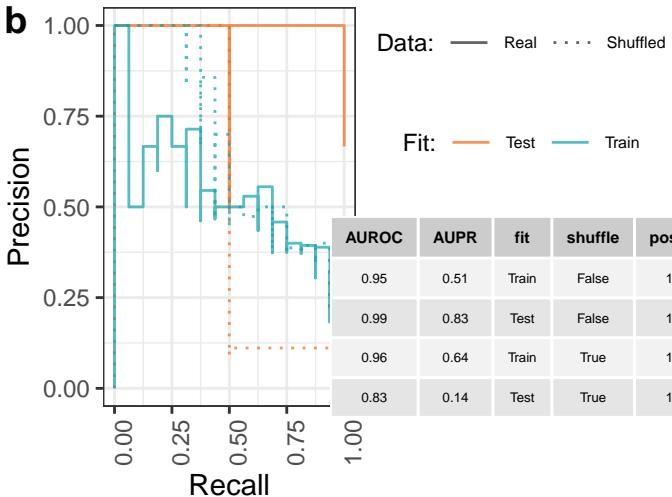
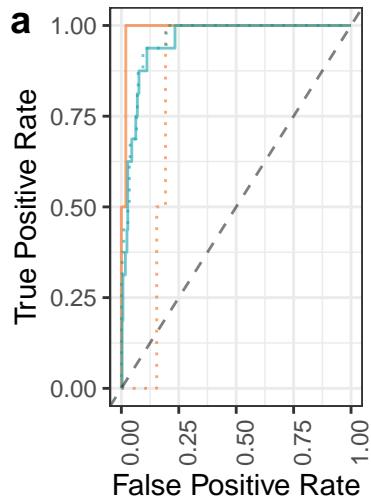
Performance: vb_infection_percentage



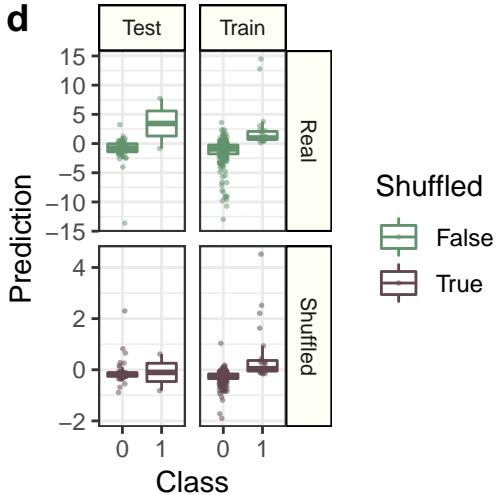
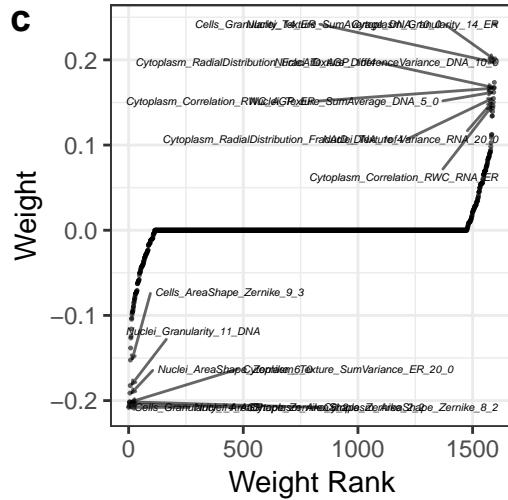
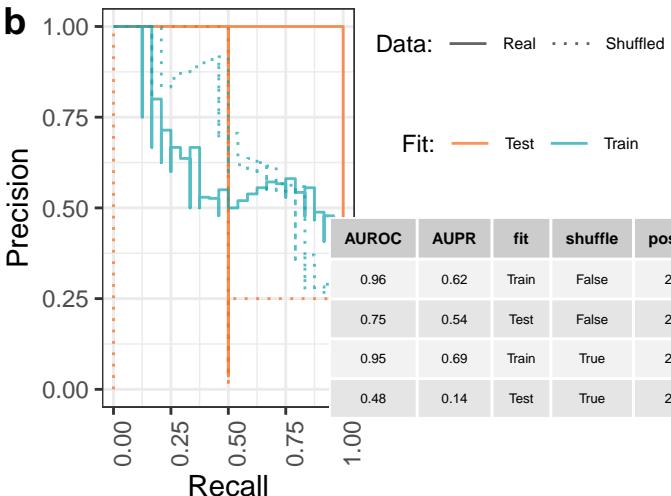
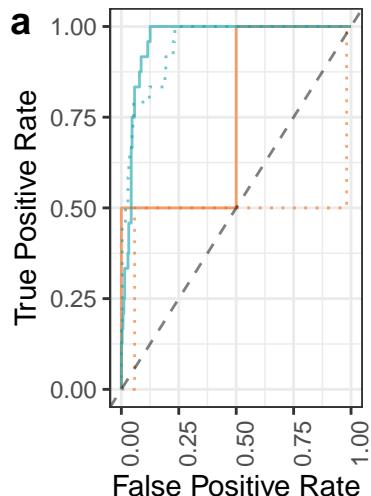
Performance: vb_percent_all_apoptosis



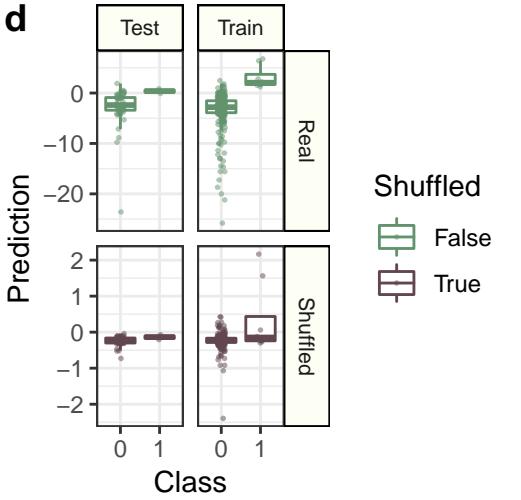
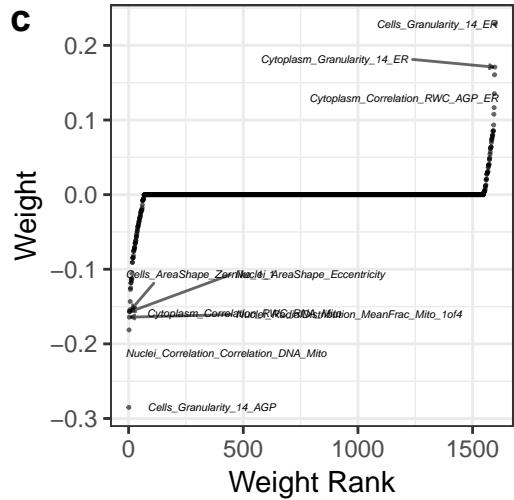
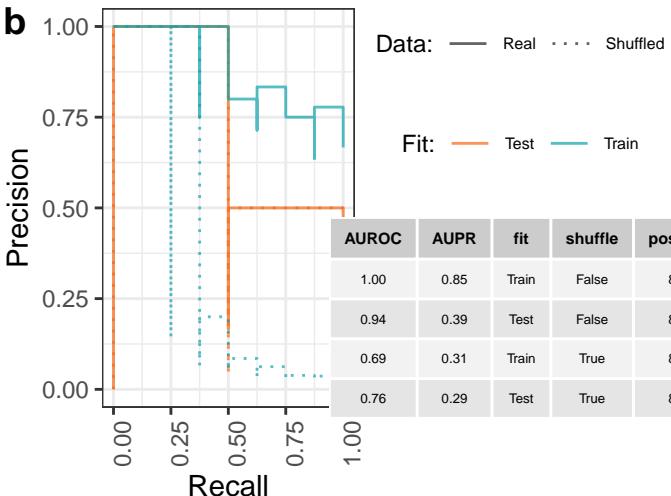
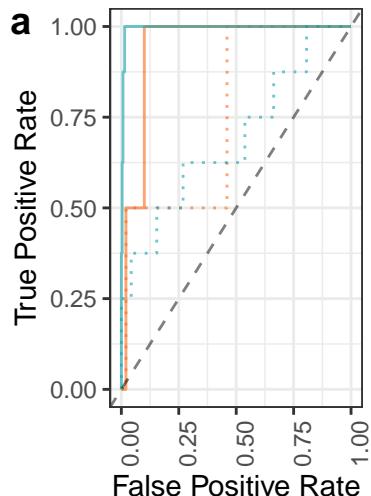
Performance: vb_percent_all_early_apoptosis



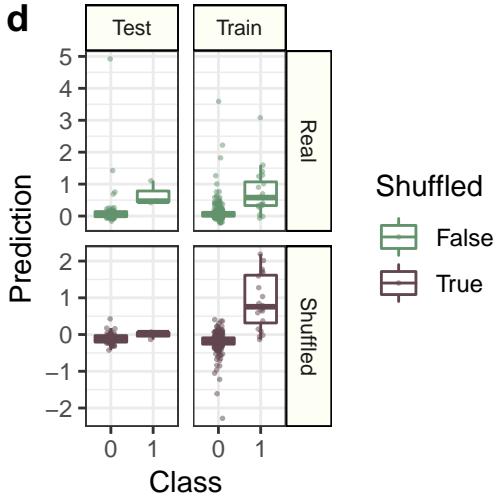
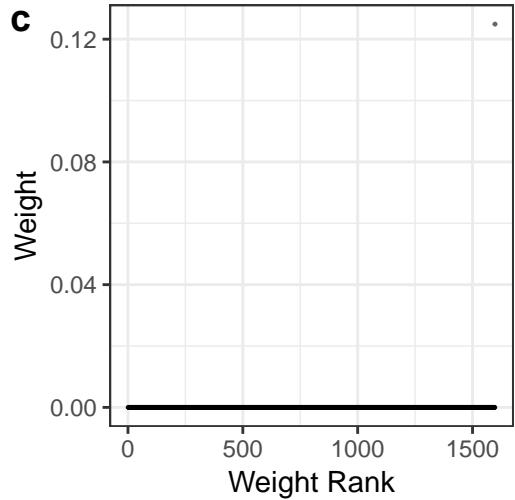
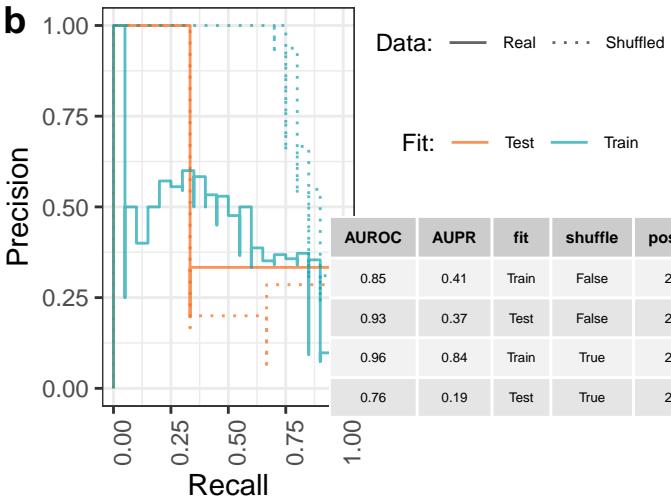
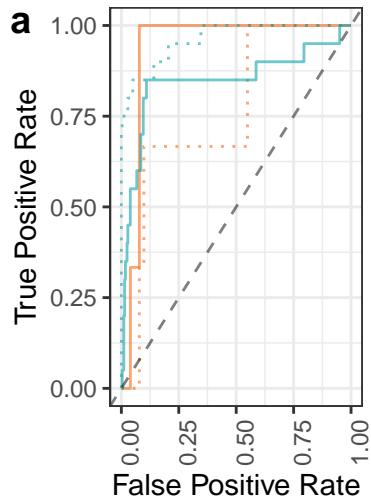
Performance: vb_percent_caspase_dead_only



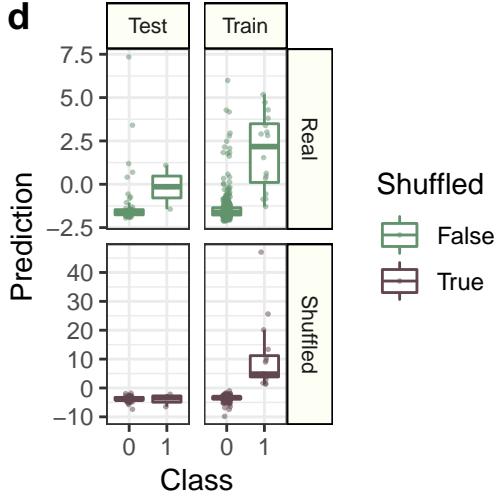
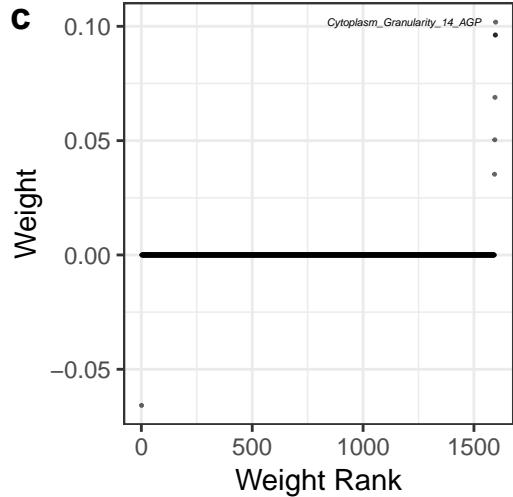
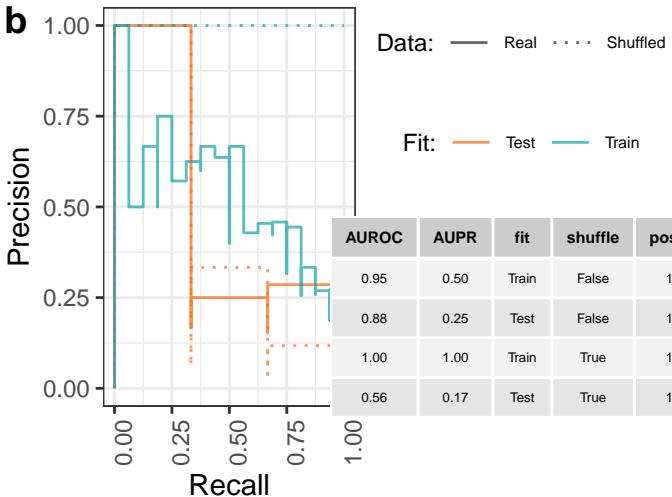
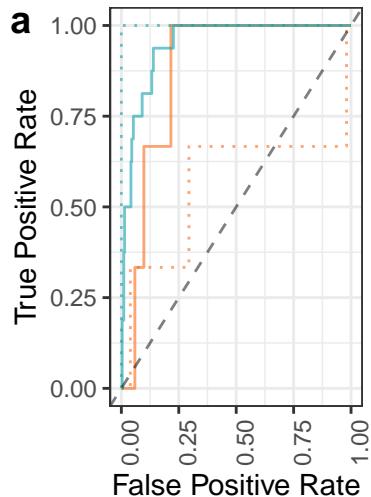
Performance: vb_ros_mean



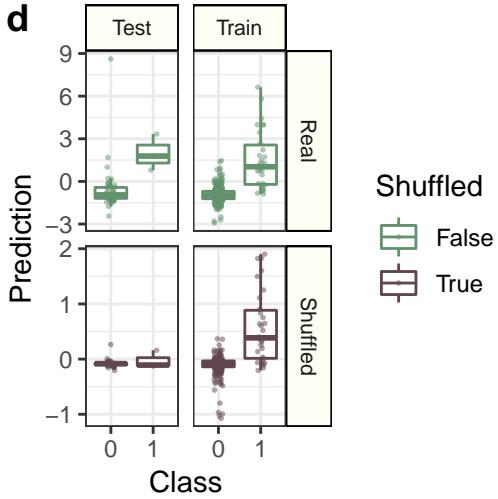
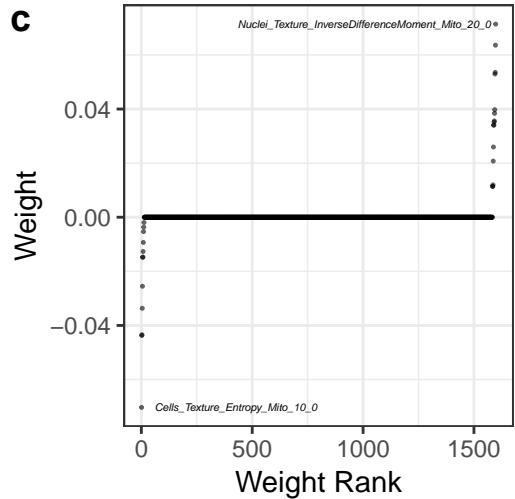
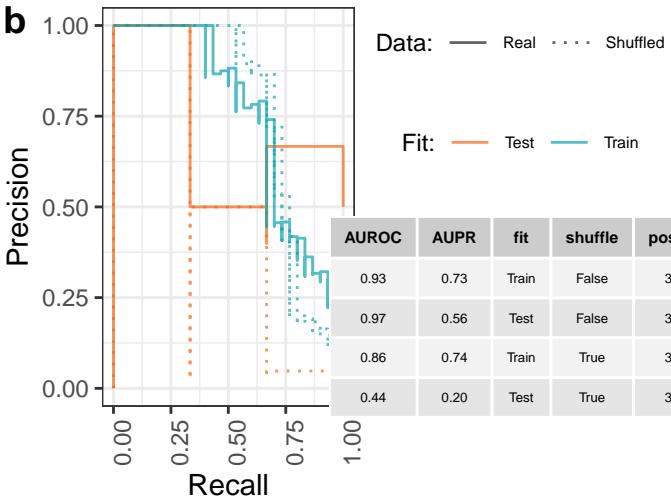
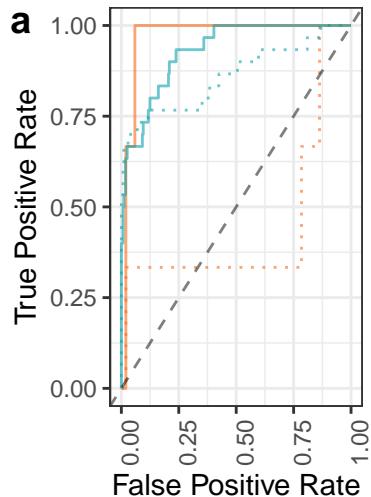
Performance: cc_cc_g1_mean



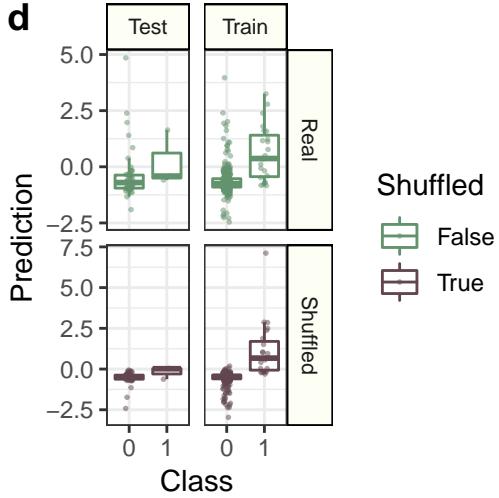
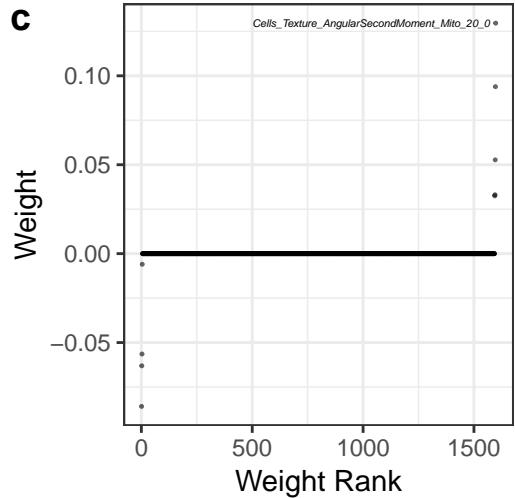
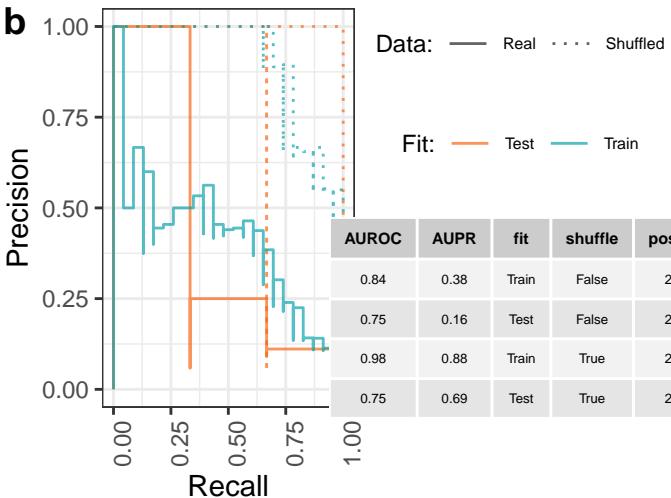
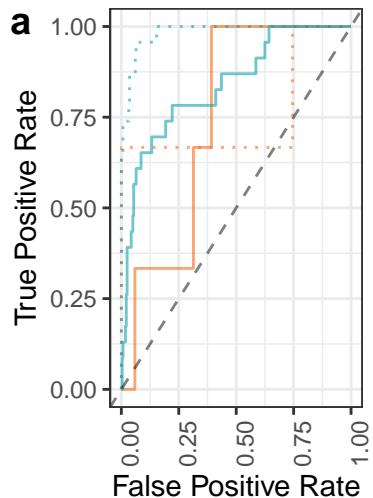
Performance: cc_cc_g2_ph3_neg_mean



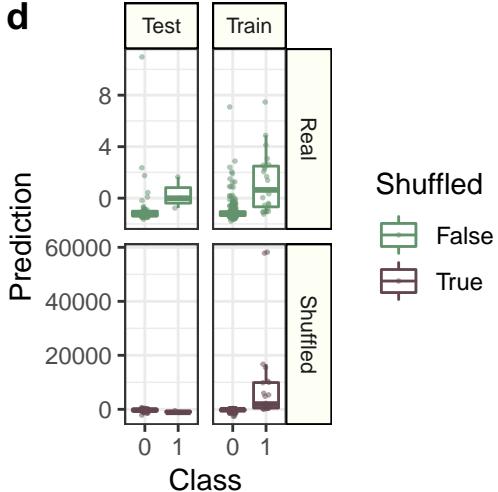
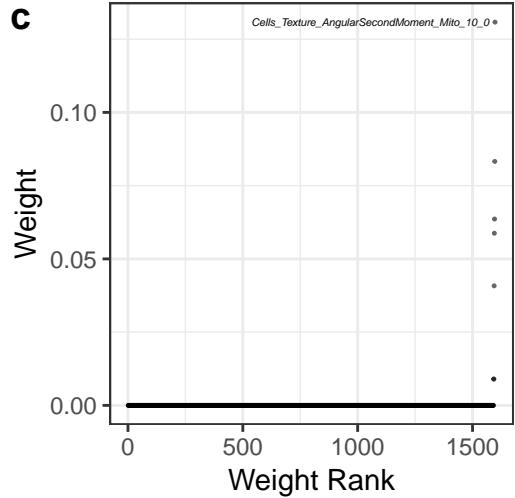
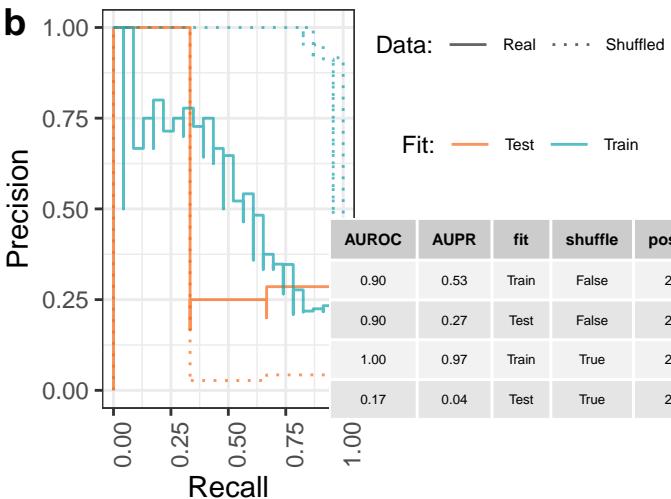
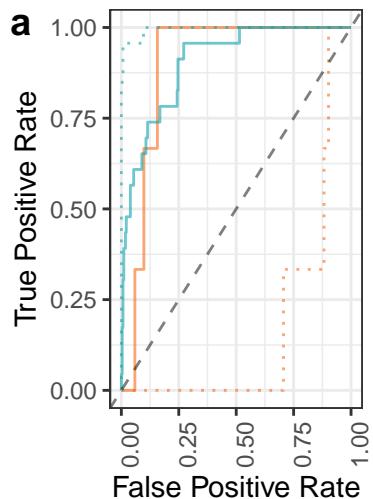
Performance: cc_cc_n_spots_mean



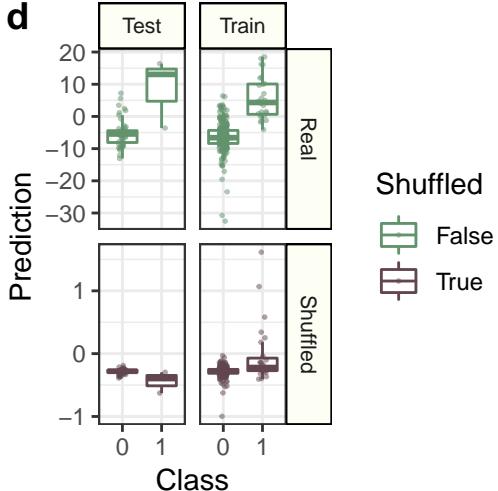
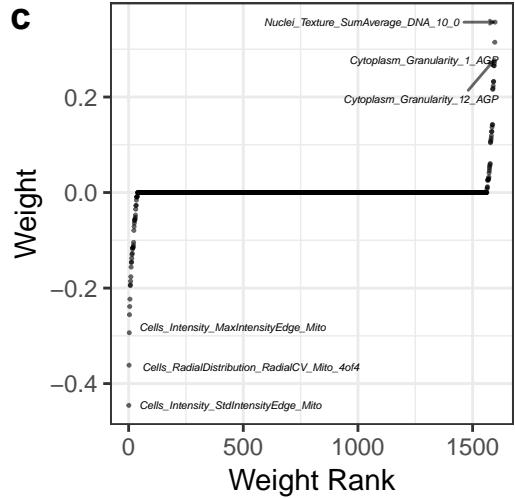
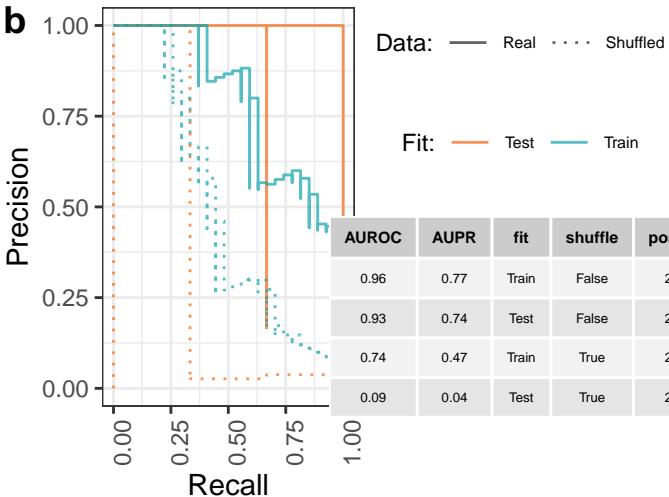
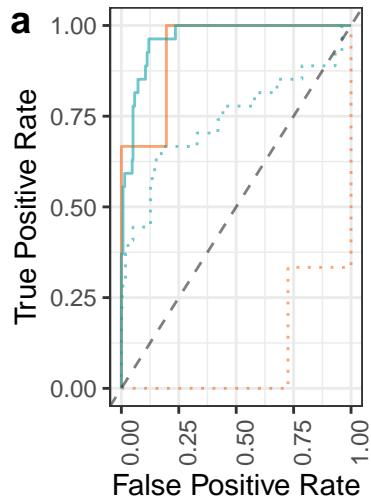
Performance: cc_edu_pos_n_spots_mean



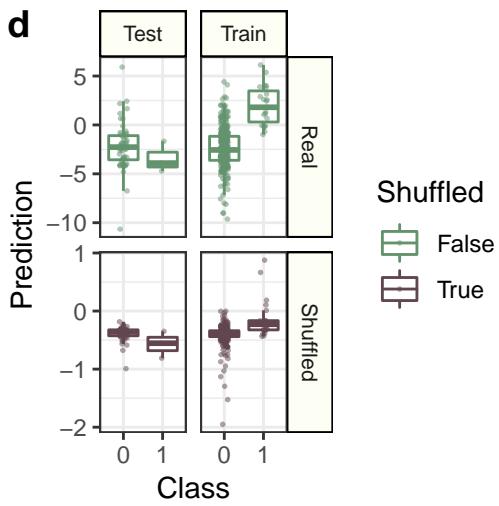
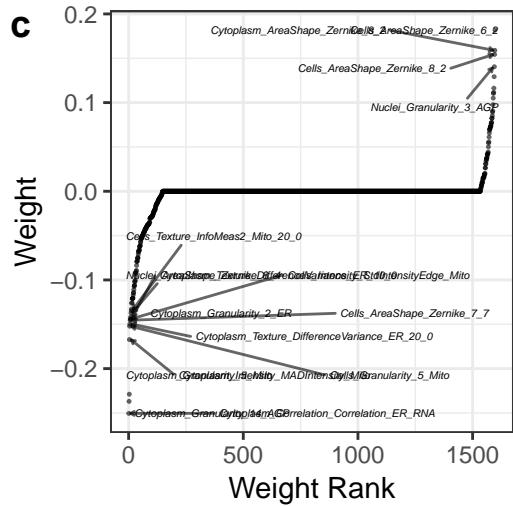
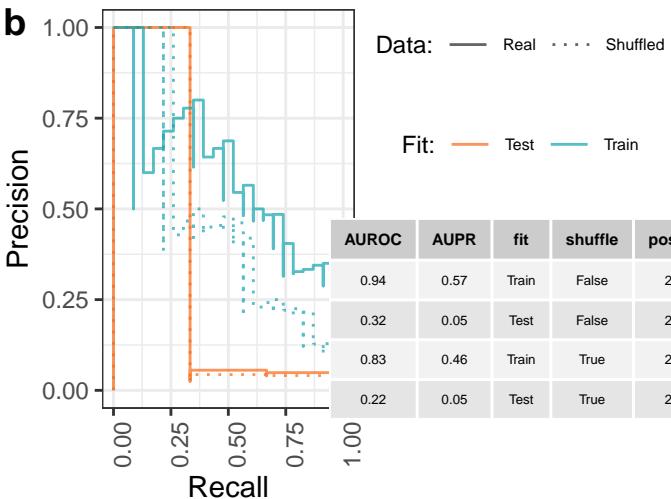
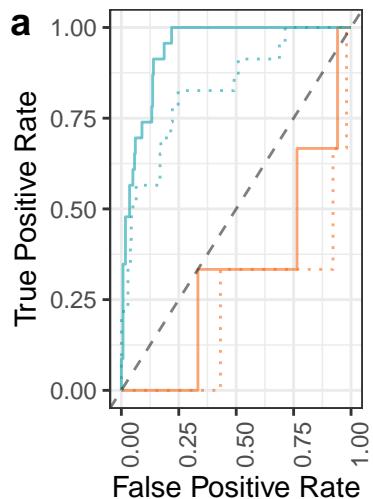
Performance: cc_g2_ph3_neg_high_n_spots_h2ax_mean



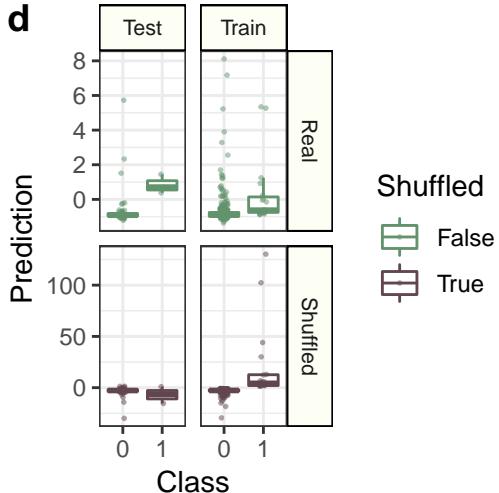
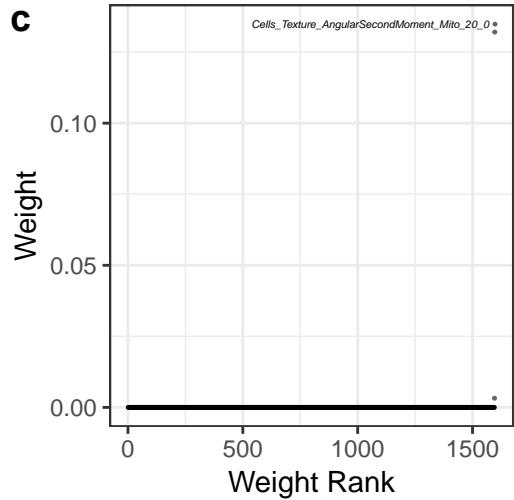
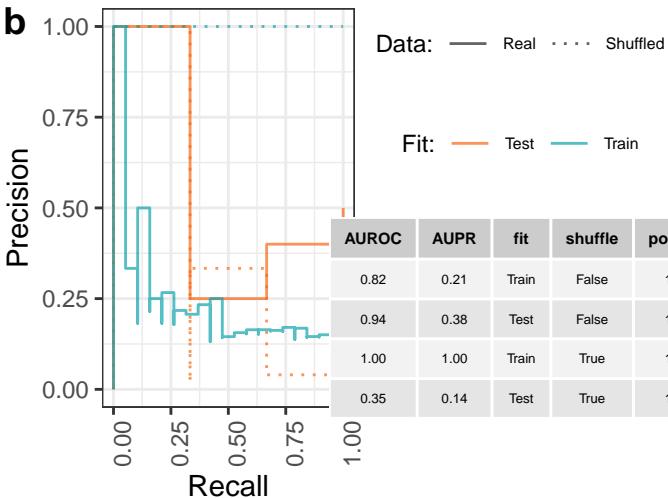
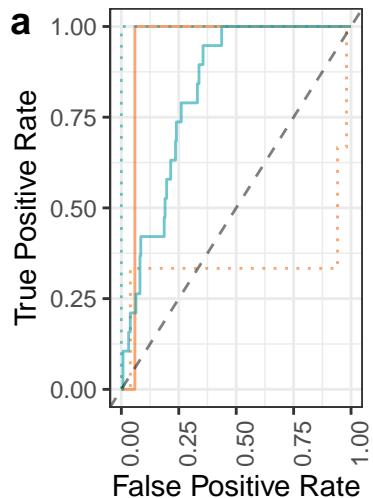
Performance: cc_g2_ph3_neg_n_spots_per_nucleus_area



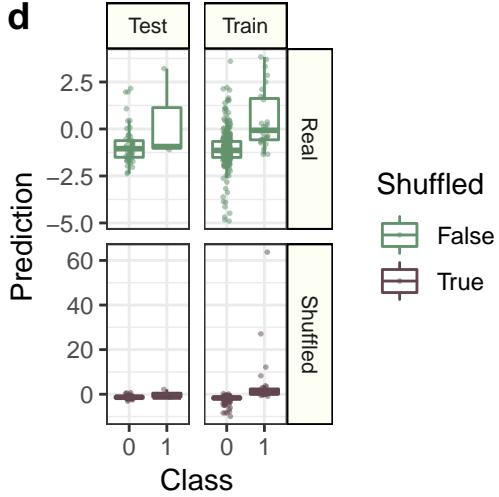
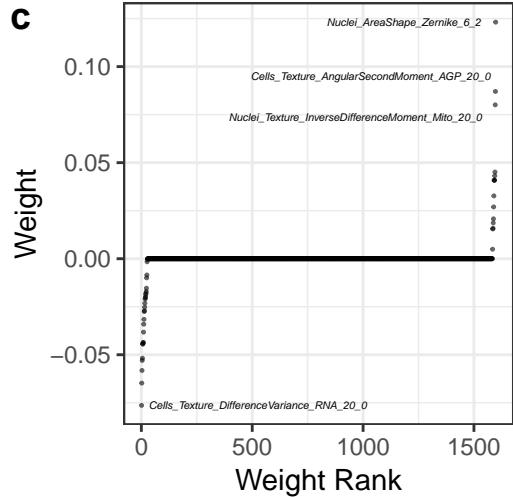
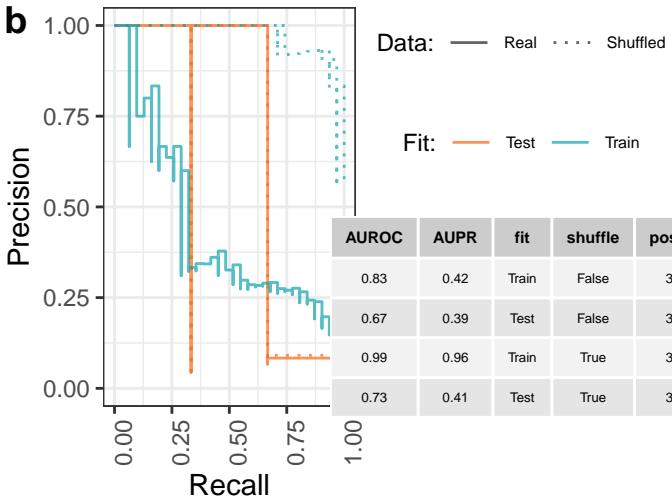
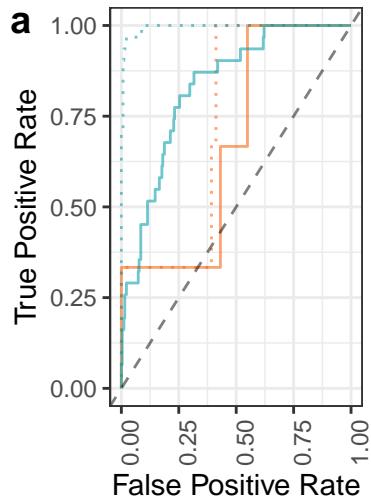
Performance: cc_mitosis_ph3_pos_n_spots_mean



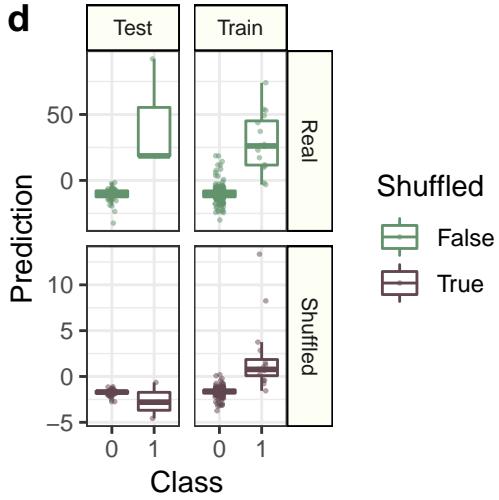
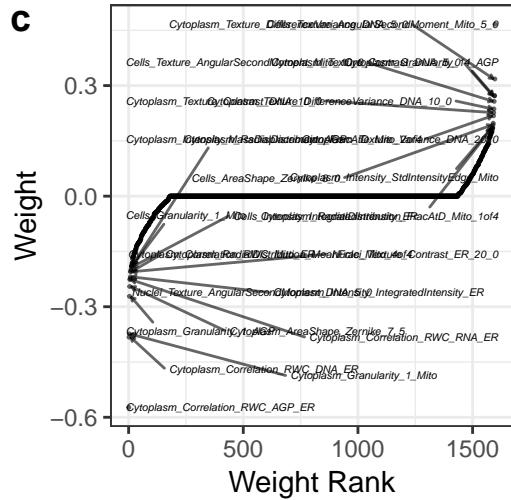
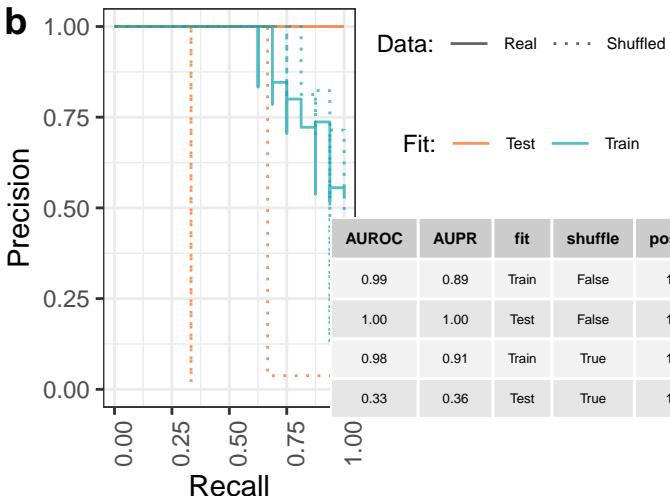
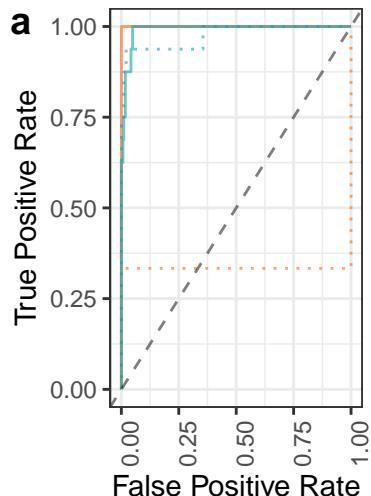
Performance: cc_polynuclear_n_spots_per_nucleus_area



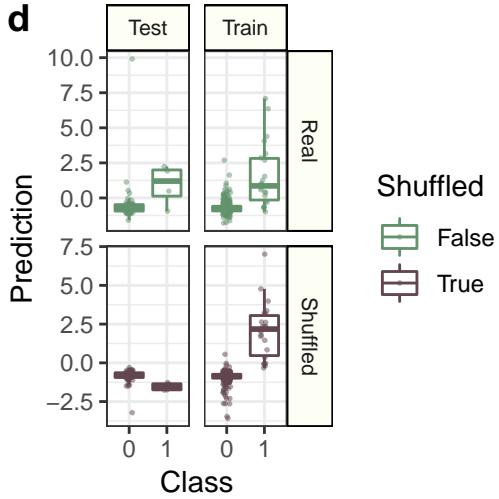
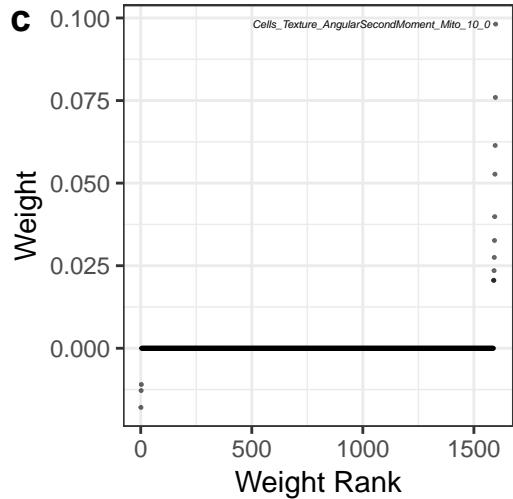
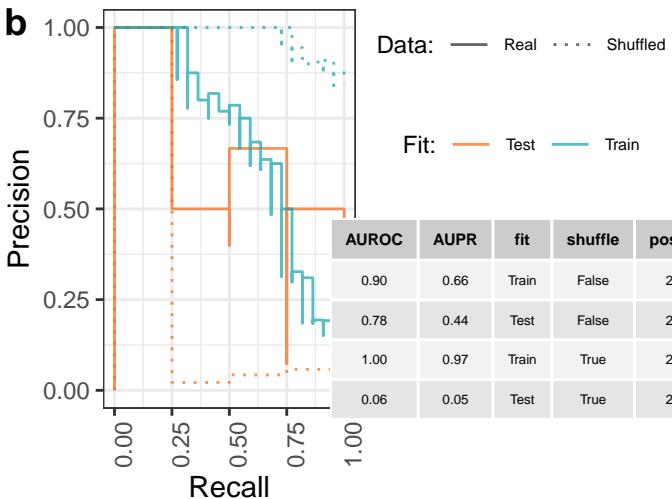
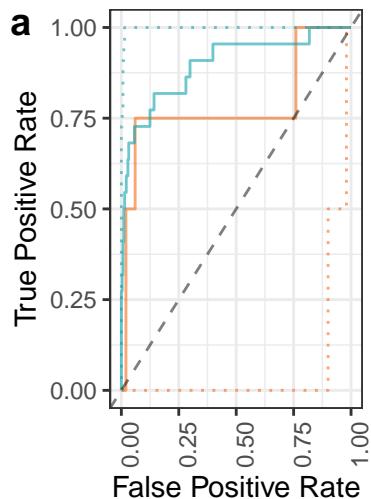
Performance: cc_polyploid_n_spots_mean



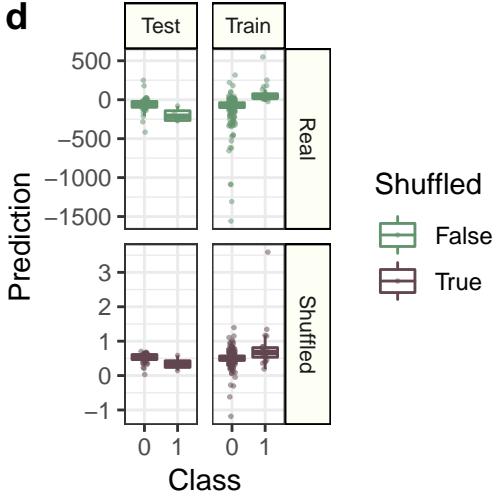
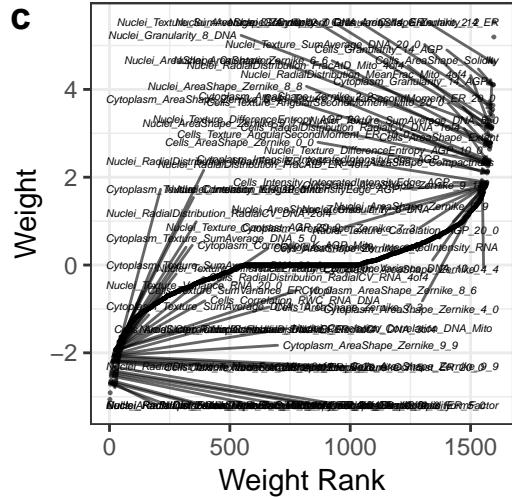
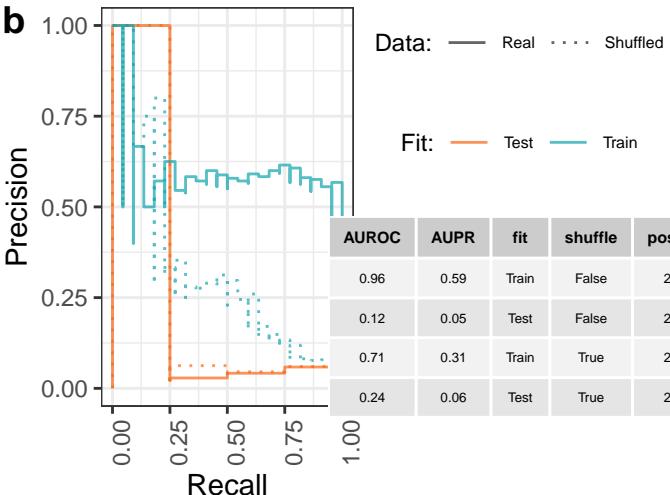
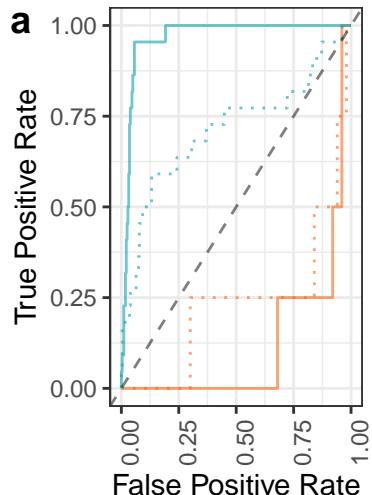
Performance: vb_percent_dead



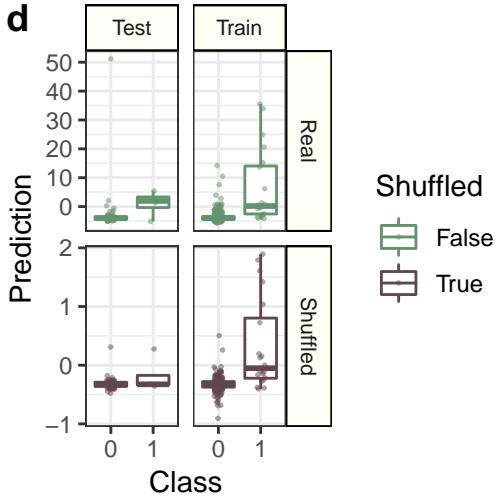
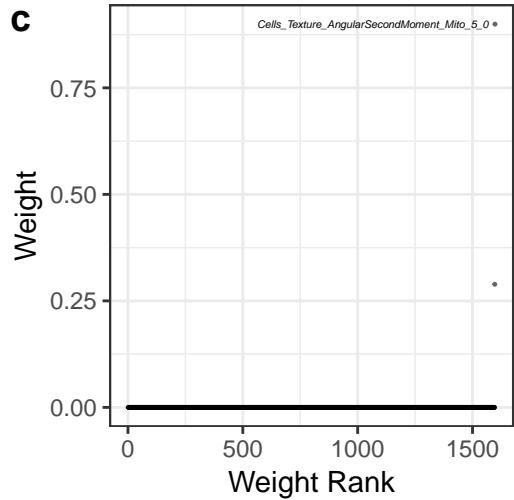
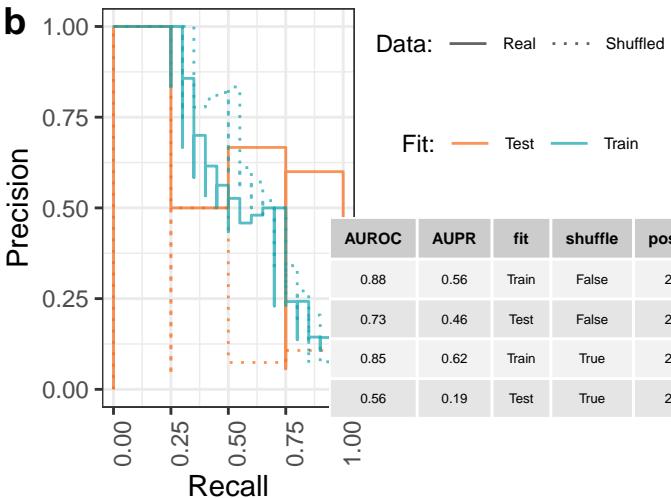
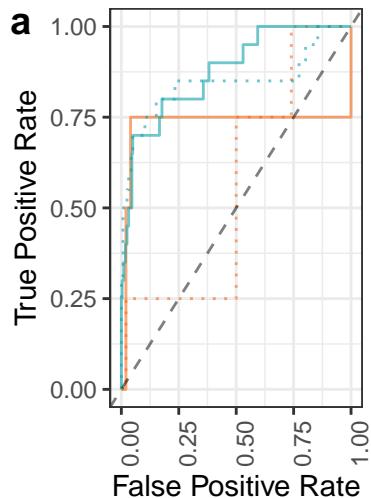
Performance: cc_cc_high_n_spots_h2ax_mean



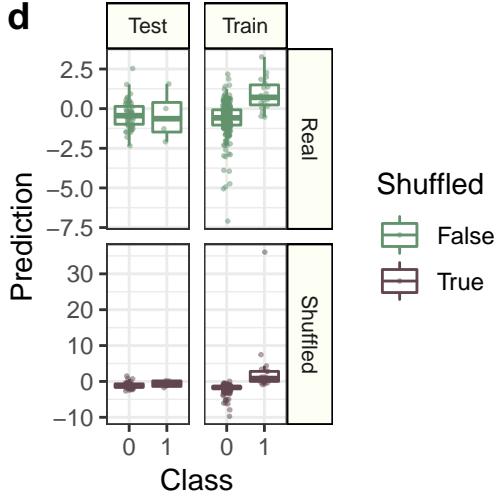
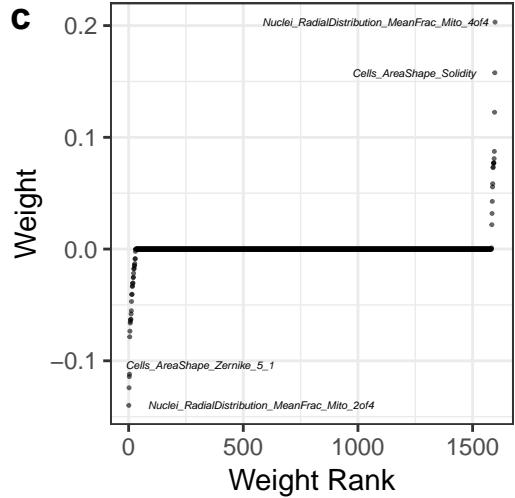
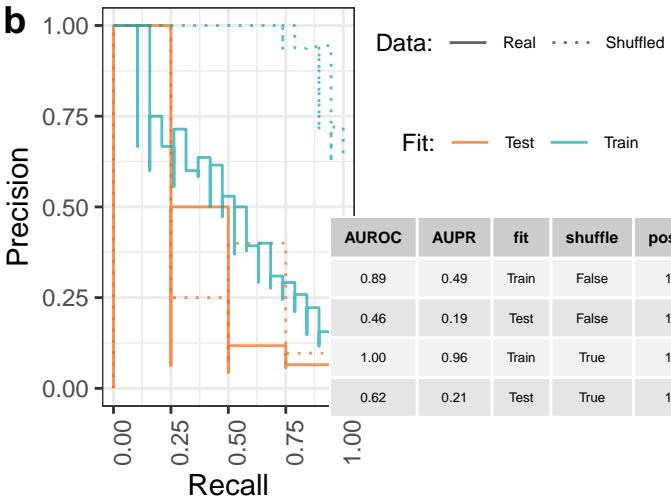
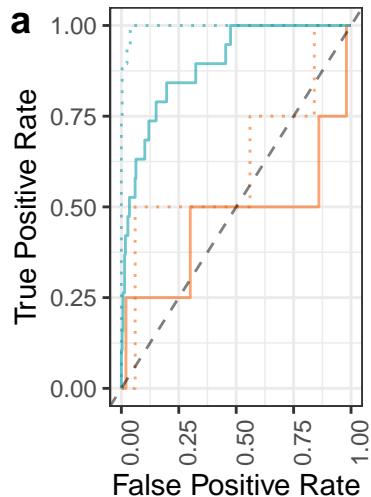
Performance: cc_edu_pos_high_n_spots_h2ax_mean



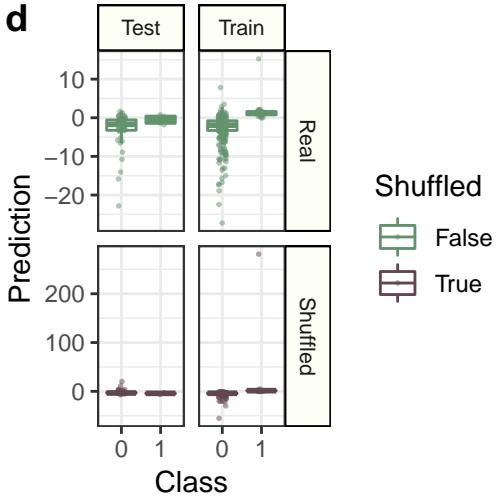
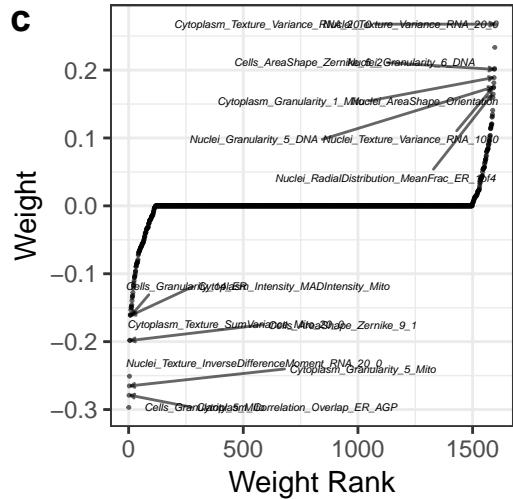
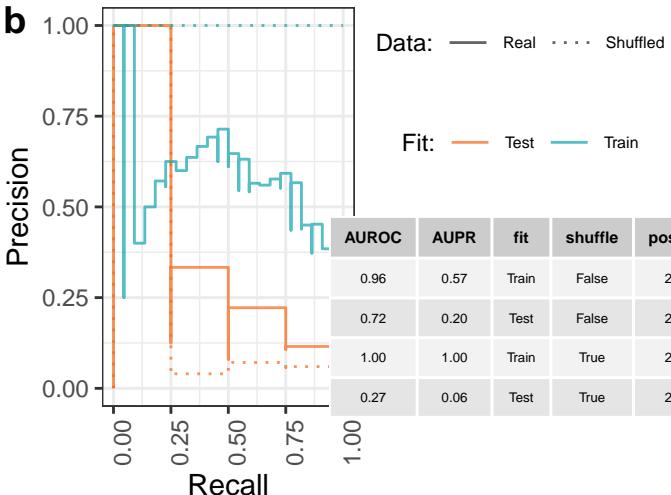
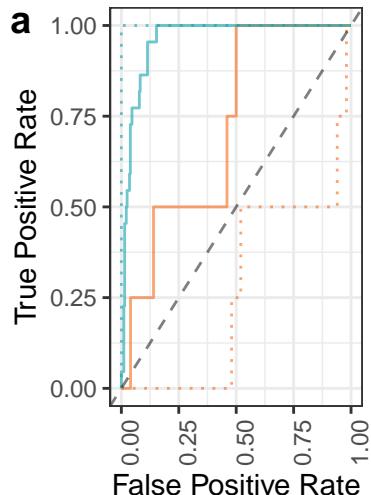
Performance: cc_g1_high_n_spots_h2ax_mean



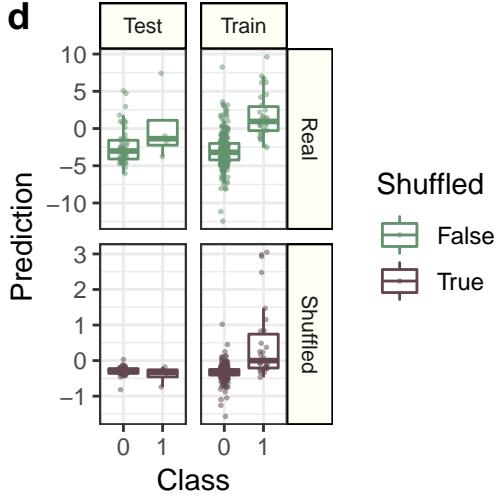
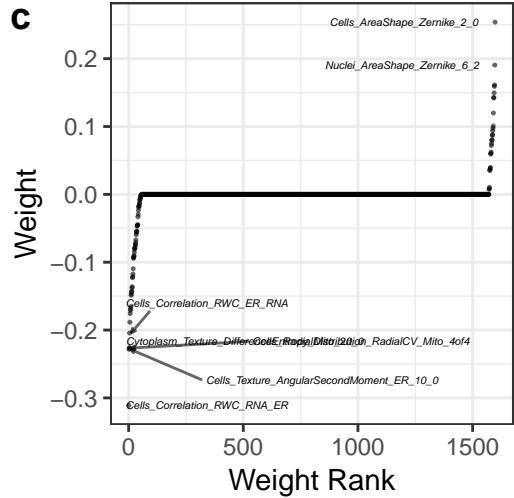
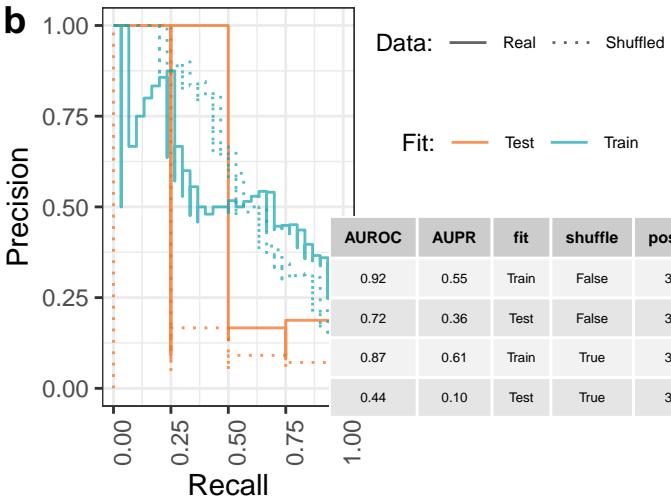
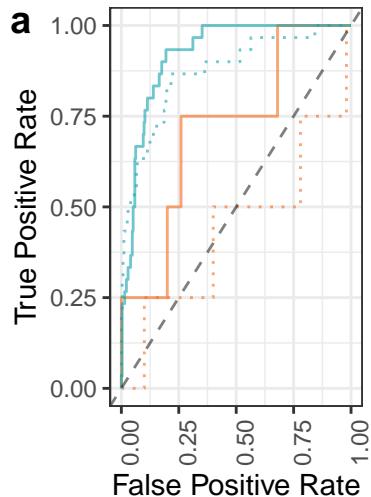
Performance: cc_g2_ph3_pos_n_spots_mean



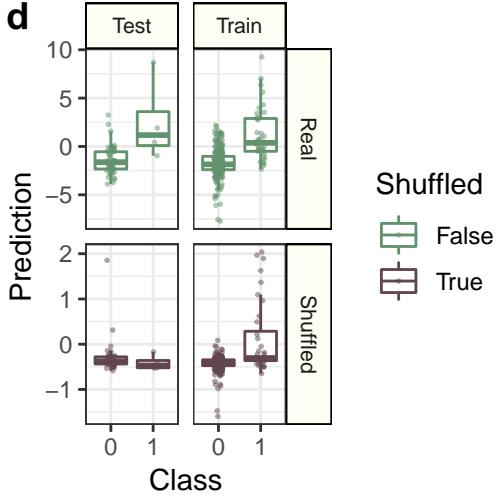
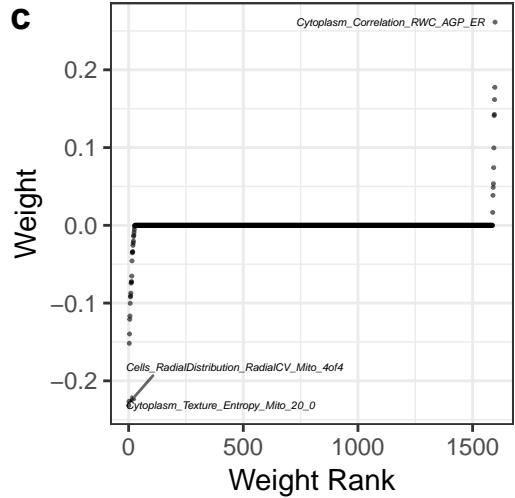
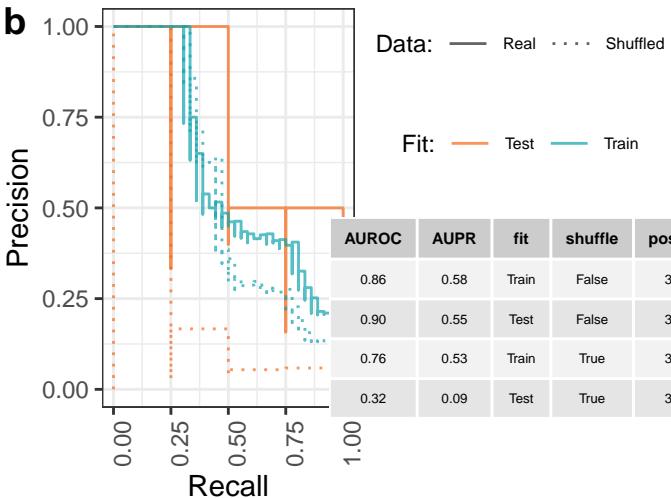
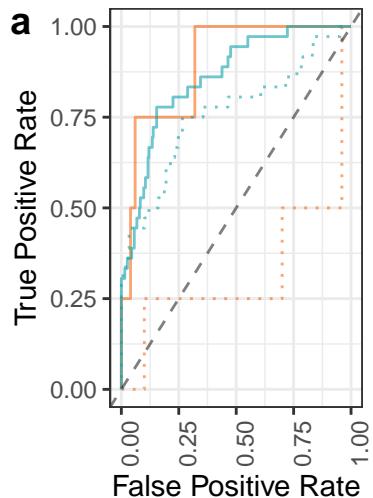
Performance: cc_mitosis_ph3_pos_n_objects



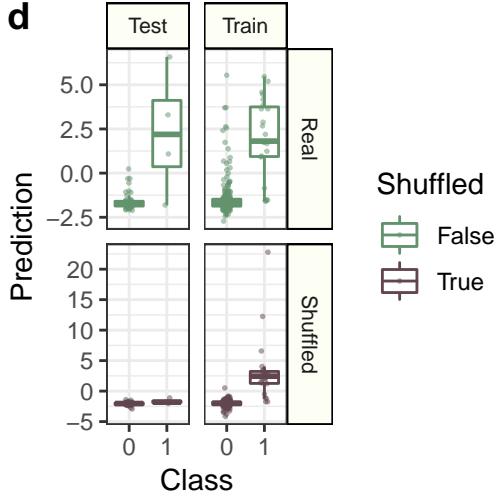
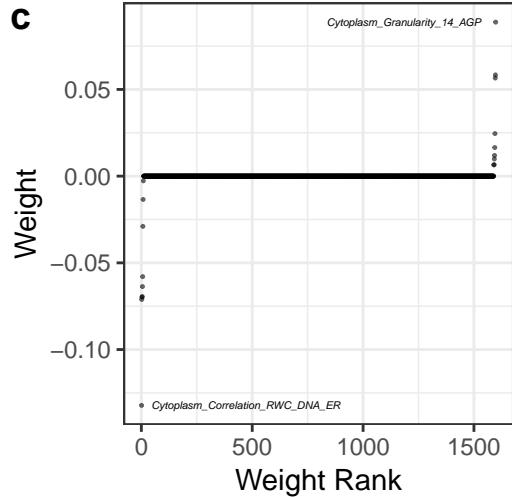
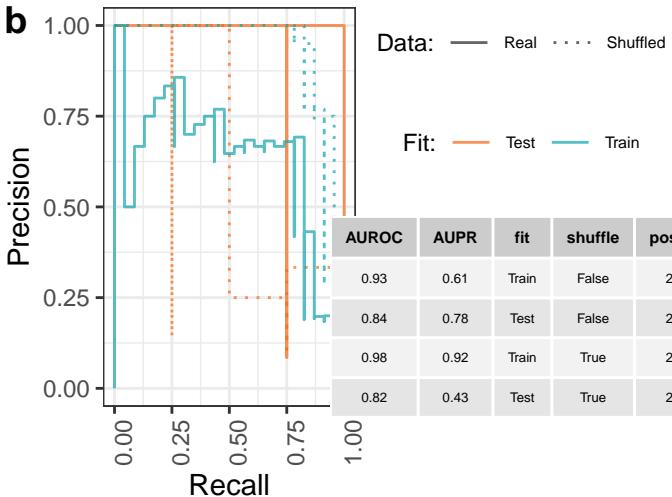
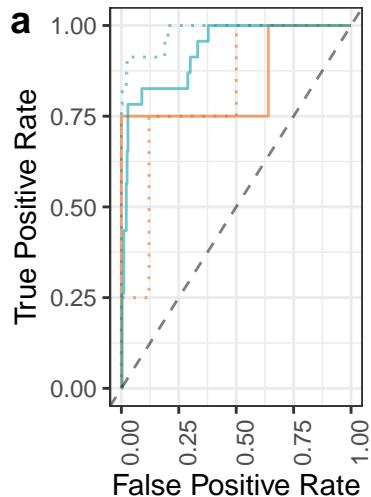
Performance: cc_polyploid_high_n_spots_h2ax_mean



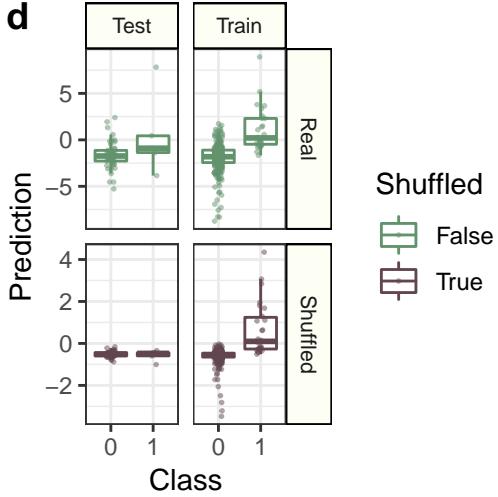
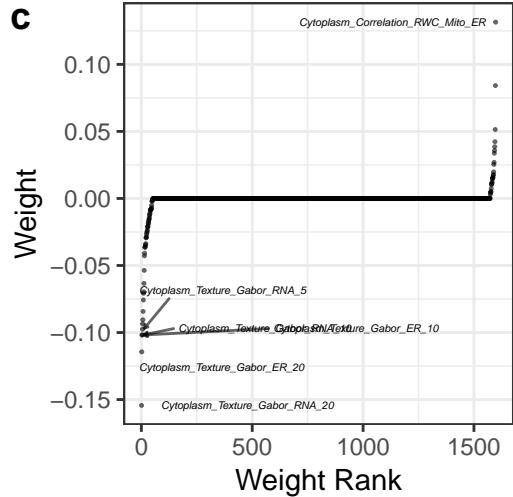
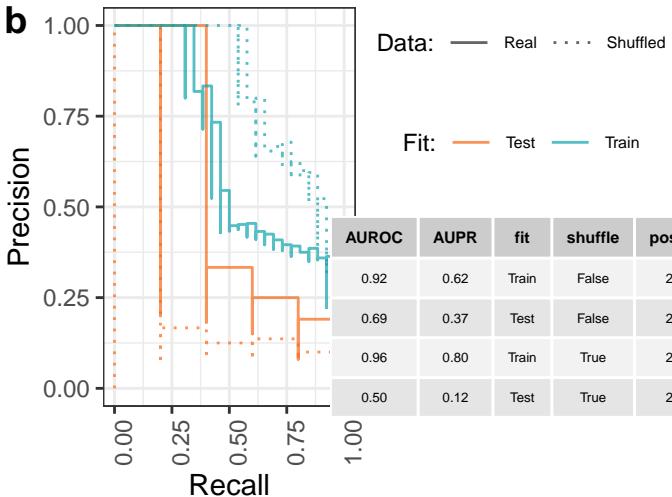
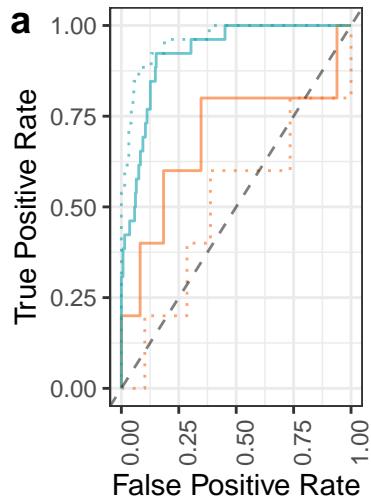
Performance: cc_polyploid_n_spots_per_nucleus_area_mean



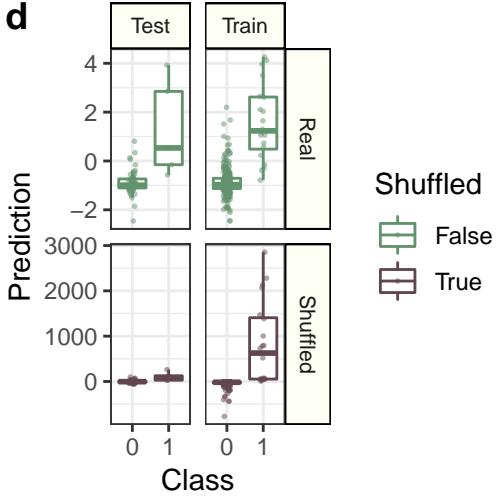
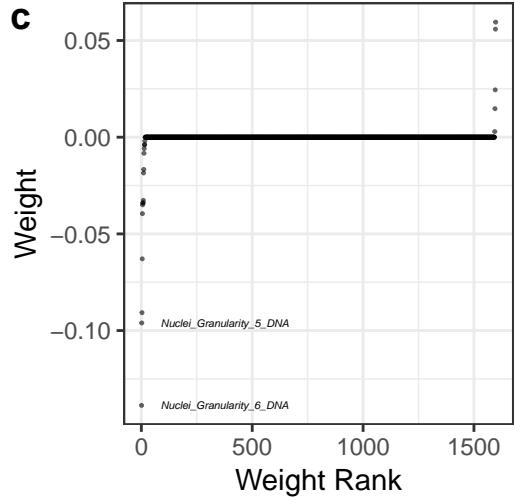
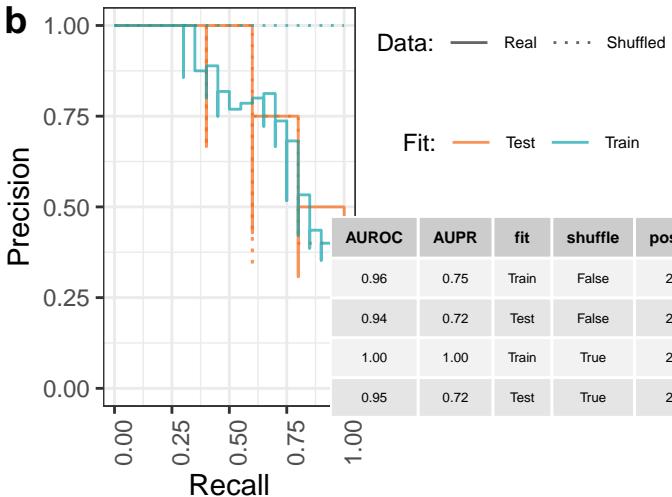
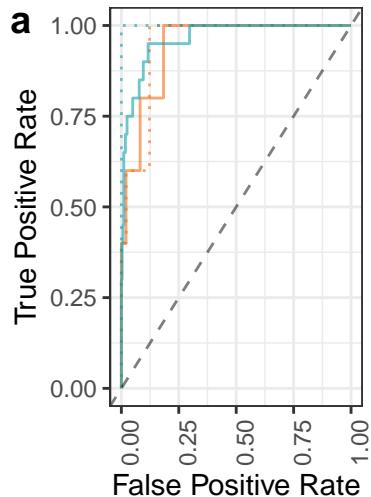
Performance: vb_percent_dead_only



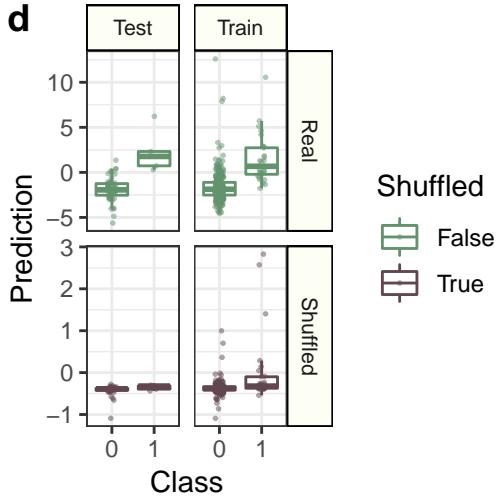
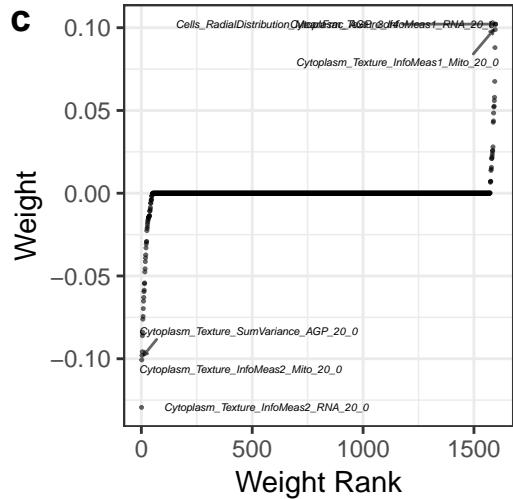
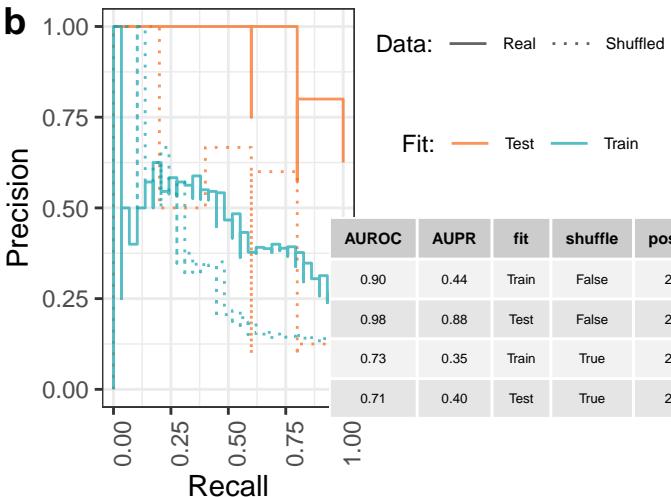
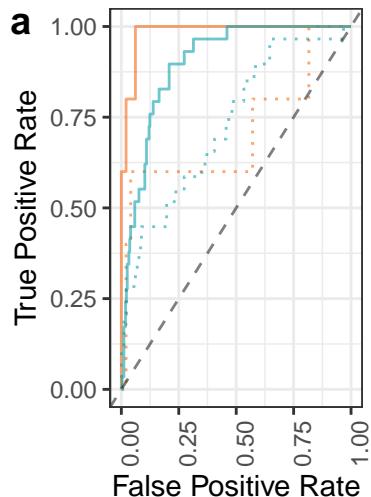
Performance: cc_all_n_spots_mean



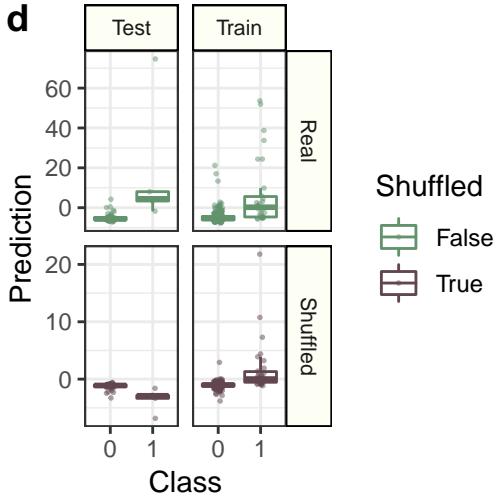
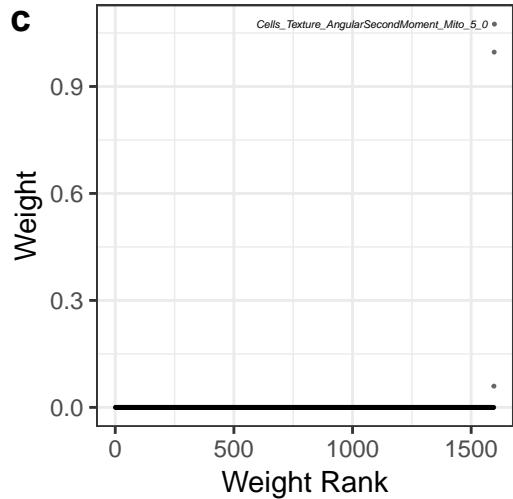
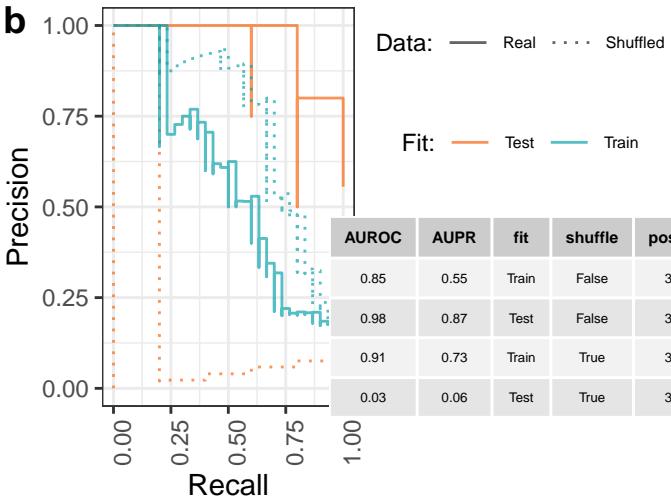
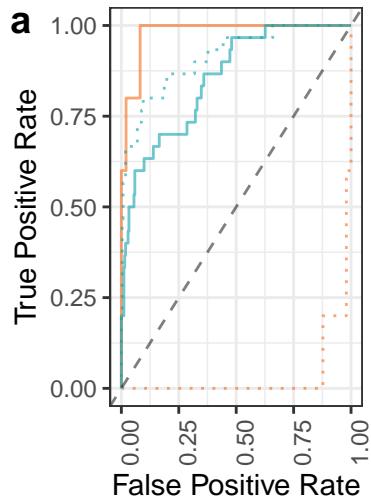
Performance: cc_all_nucleus_area_mean



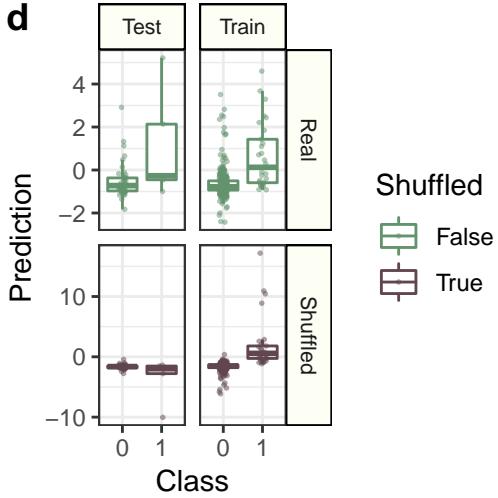
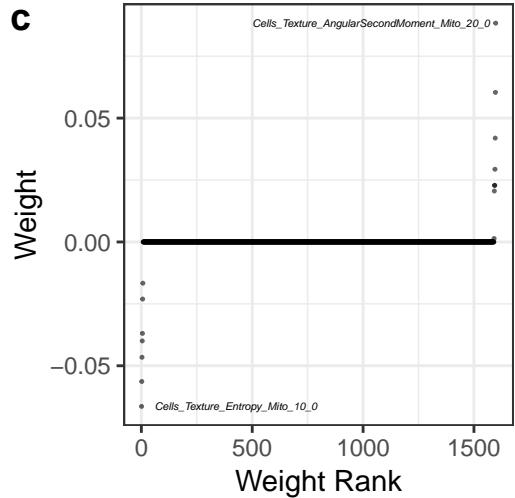
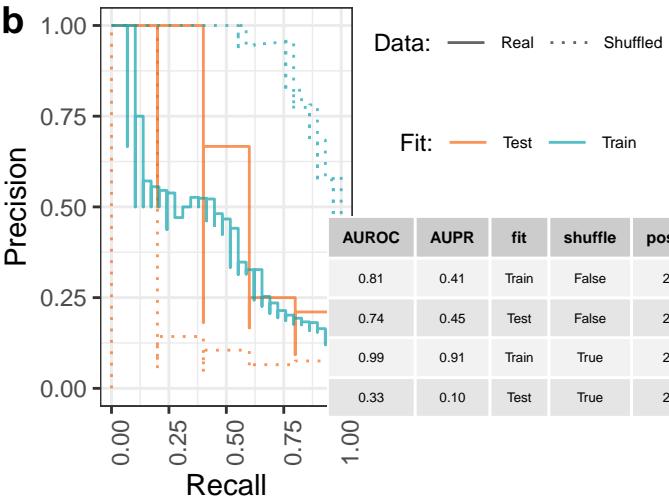
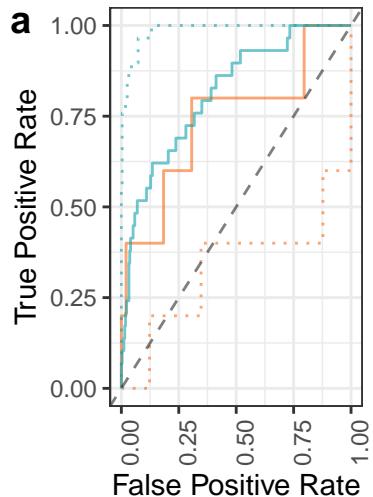
Performance: cc_cc_g2_ph3_pos_early_mitosis_mean



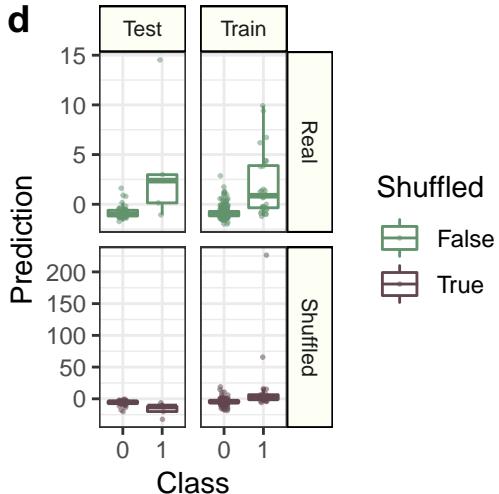
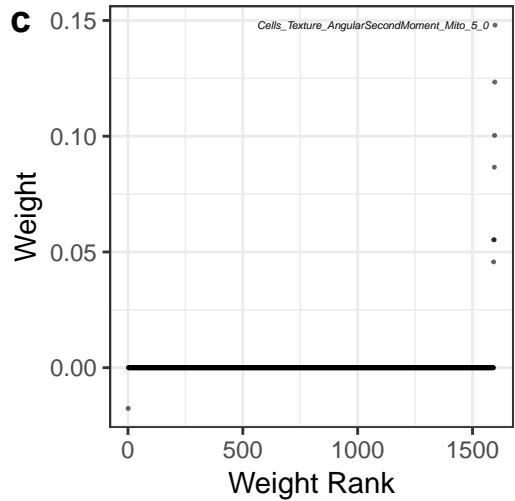
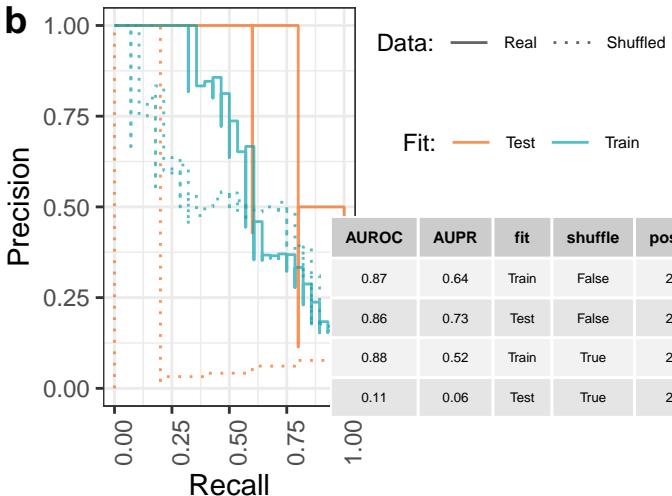
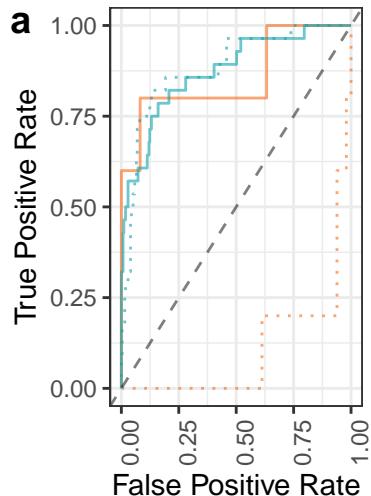
Performance: cc_cc_n_spots_per_nucleus_area_mean



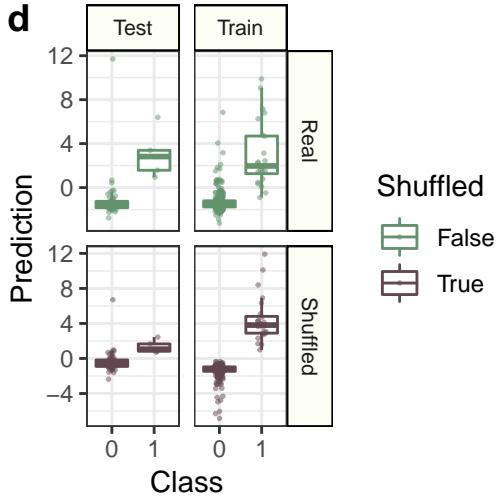
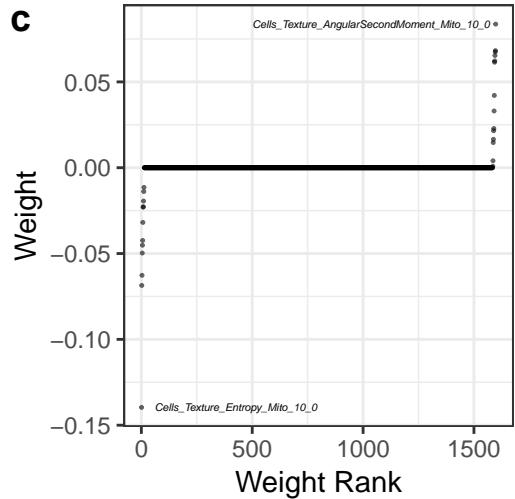
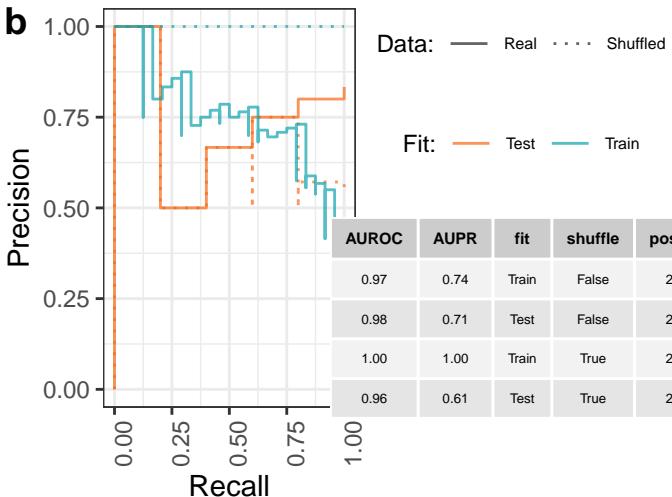
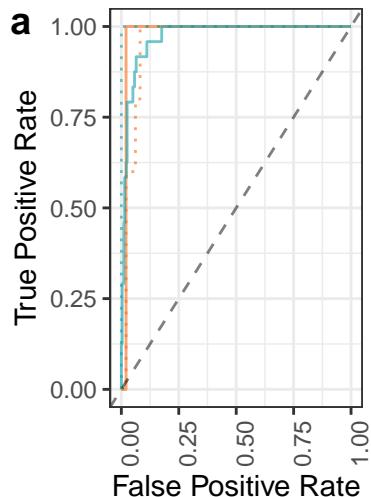
Performance: cc_edu_pos_n_spots_per_nucleus_area_mean



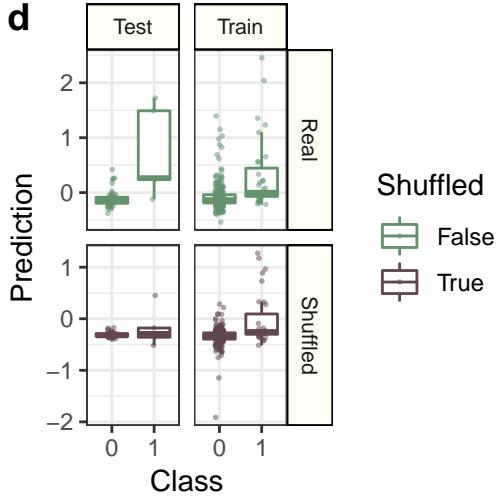
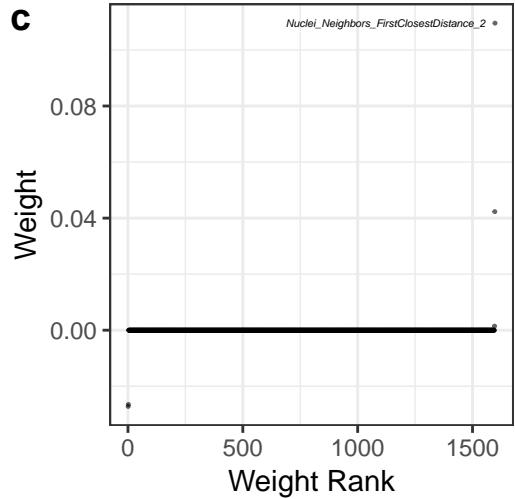
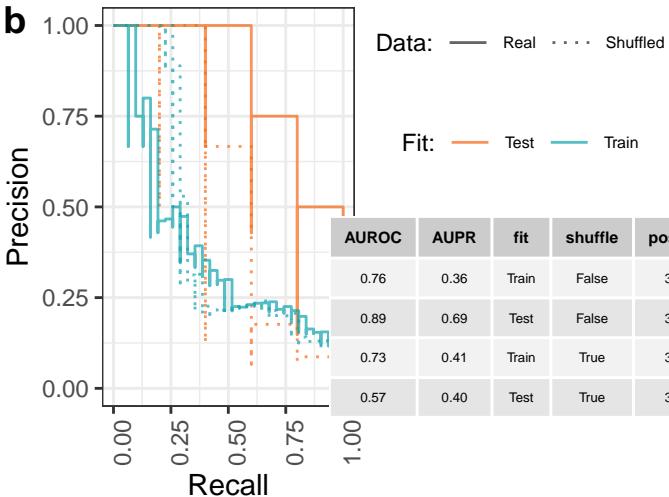
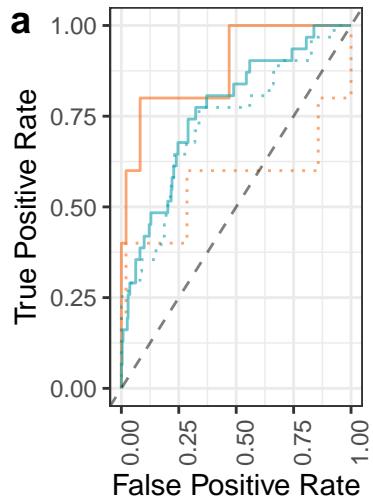
Performance: cc_g1_n_spots_mean



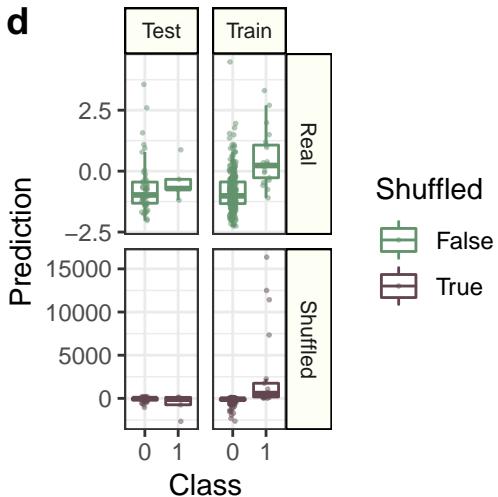
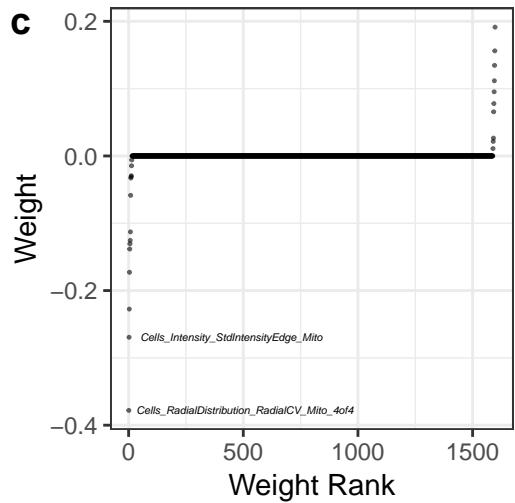
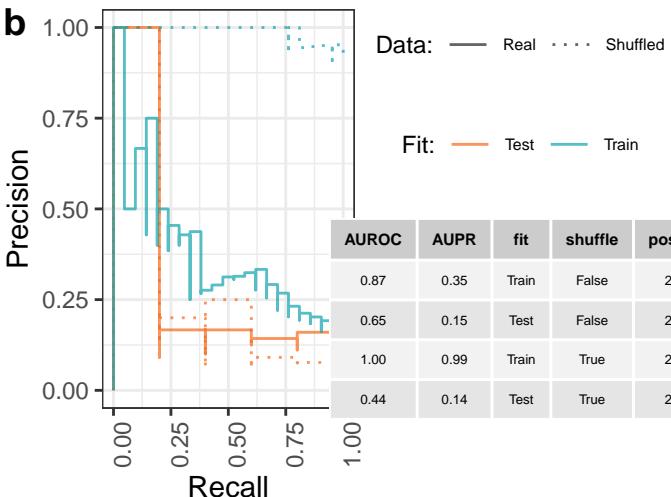
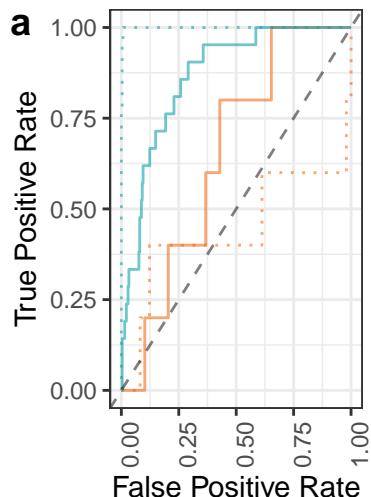
Performance: cc_g1_plus_g2



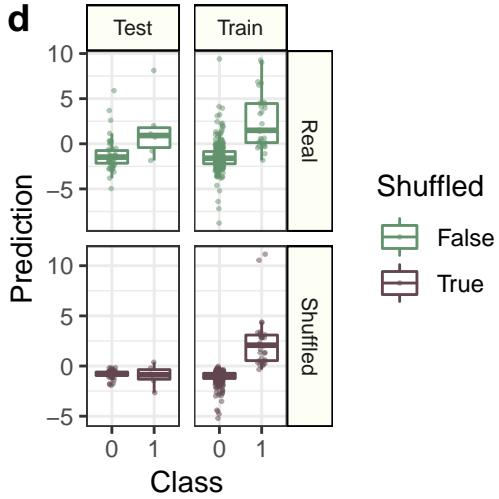
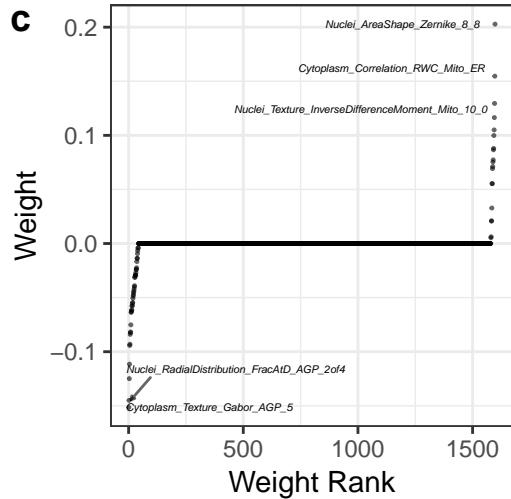
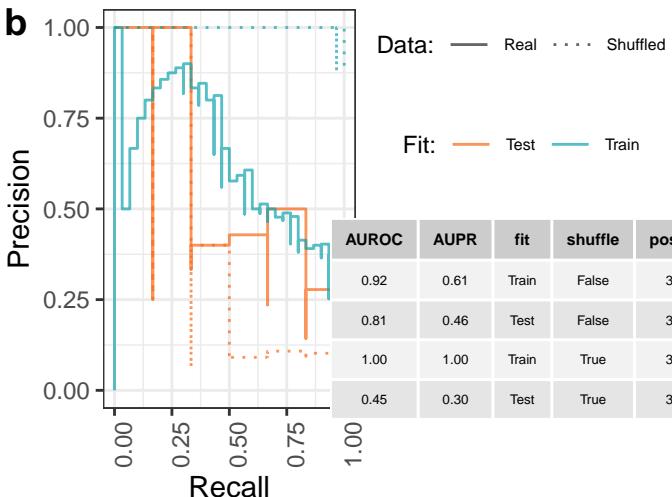
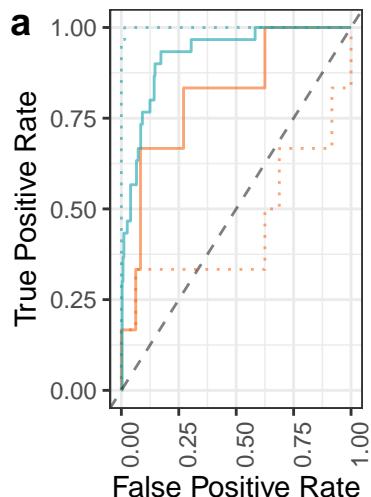
Performance: cc_g2_ph3_pos_n_spots_per_nucleus_area



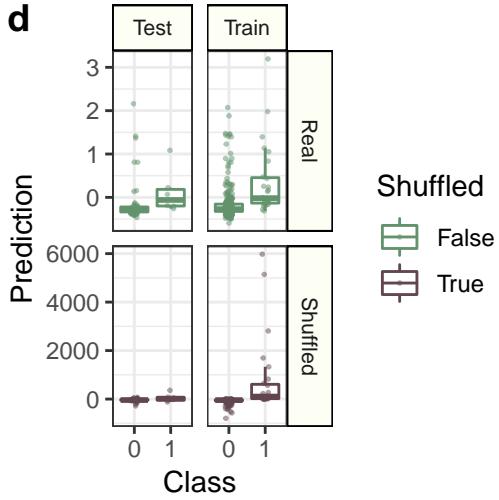
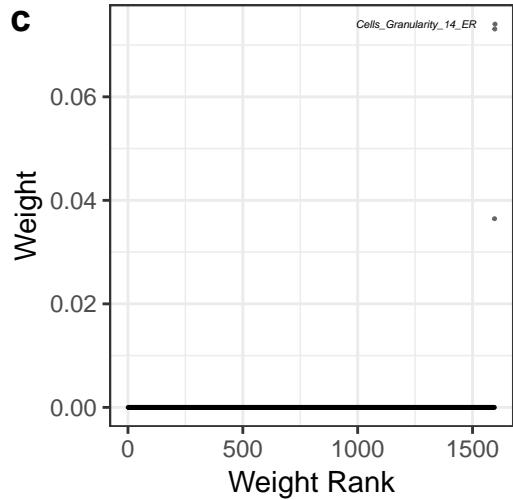
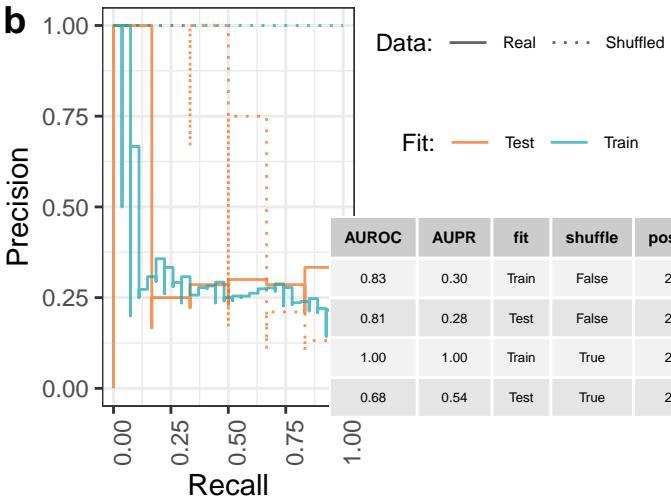
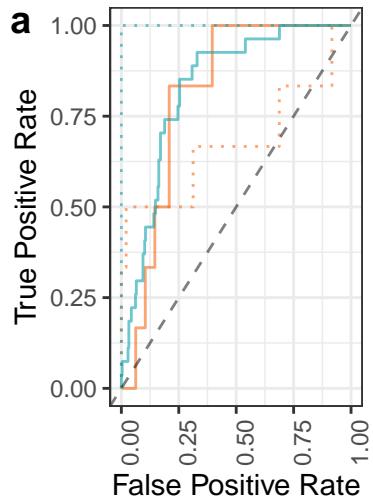
Performance: cc_mitosis_ph3_neg_n_spots_mean



Performance: cc_all_high_n_spots_h2ax_mean



Performance: cc_all_n_spots_per_nucleus_area_mean



Performance: cc_g1_n_spots_per_nucleus_area_mean

