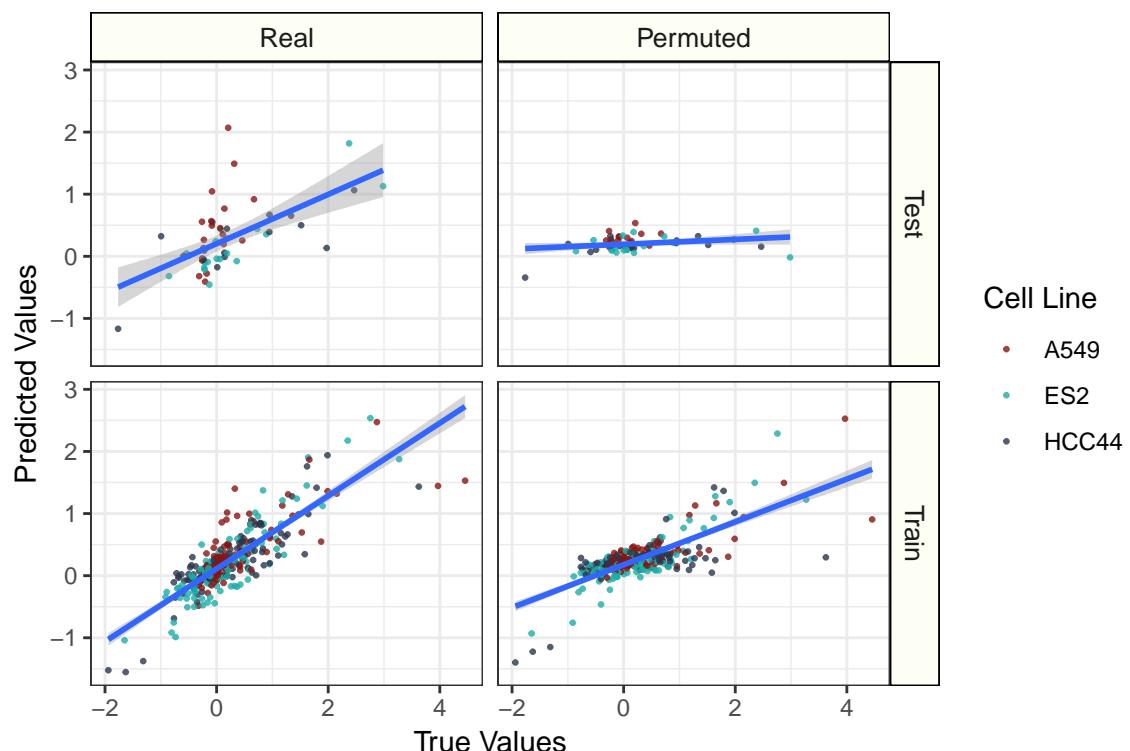
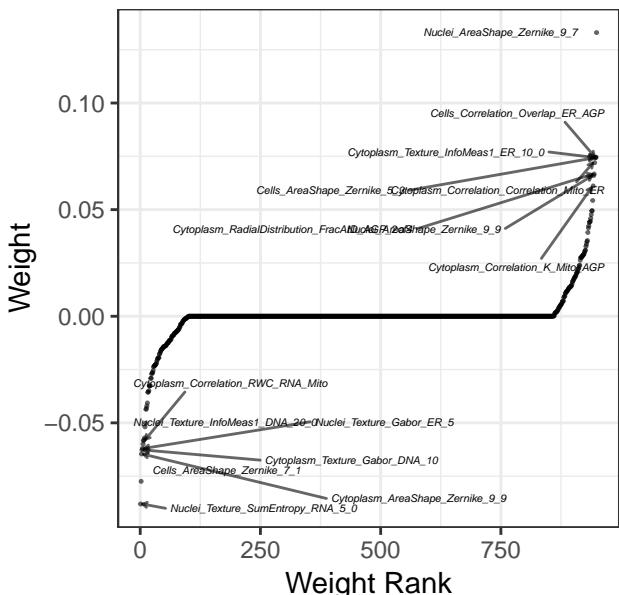


Performance: cc_all_high_h2ax

Transform: raw

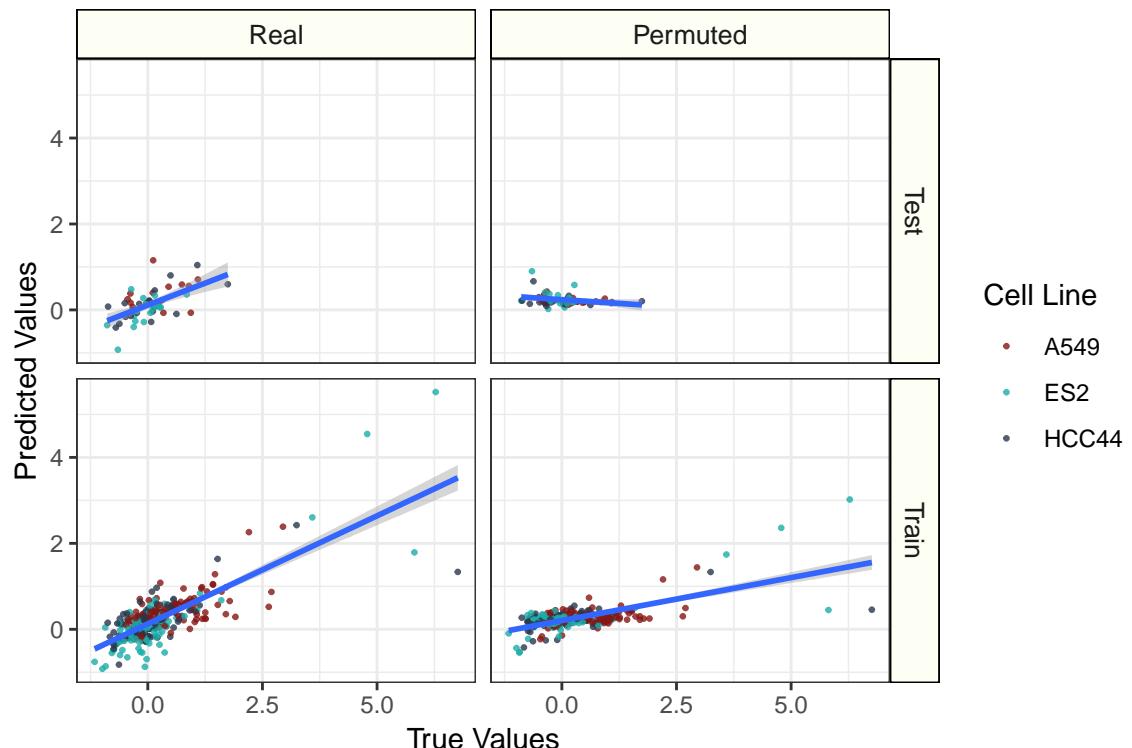


fit	shuffle	transform	r2	value
Train	Real	raw	0.68	0.19
Test	Real	raw	0.34	0.46
Train	Shuffle	raw	0.48	0.32
Test	Shuffle	raw	0.05	0.65

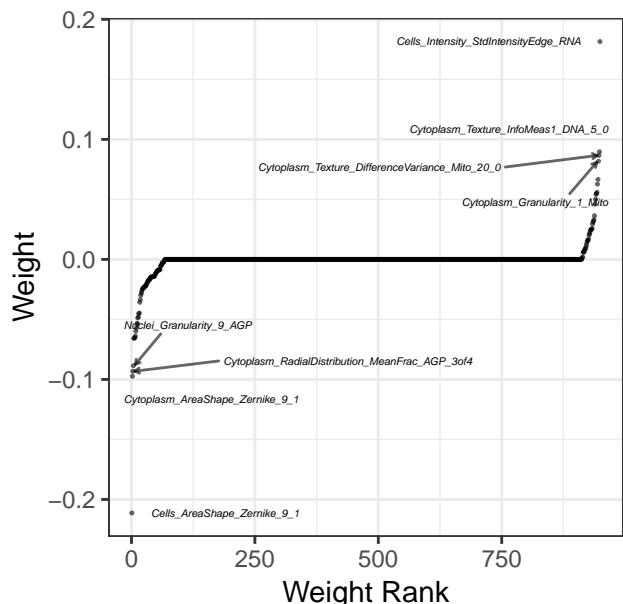


Performance: cc_all_large_notround_polinuclear_mean

Transform: raw

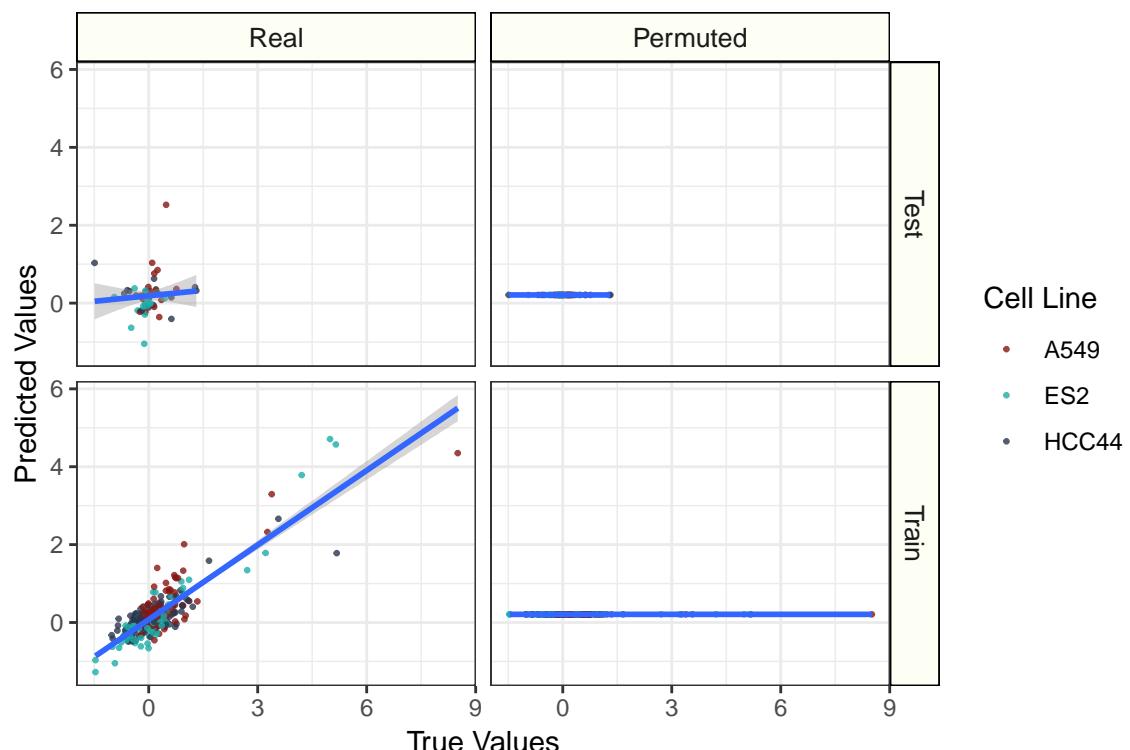


fit	shuffle	transform	r2	value
Train	Real	raw	0.59	0.36
Test	Real	raw	0.30	0.19
Train	Shuffle	raw	0.31	0.62
Test	Shuffle	raw	-0.34	0.36

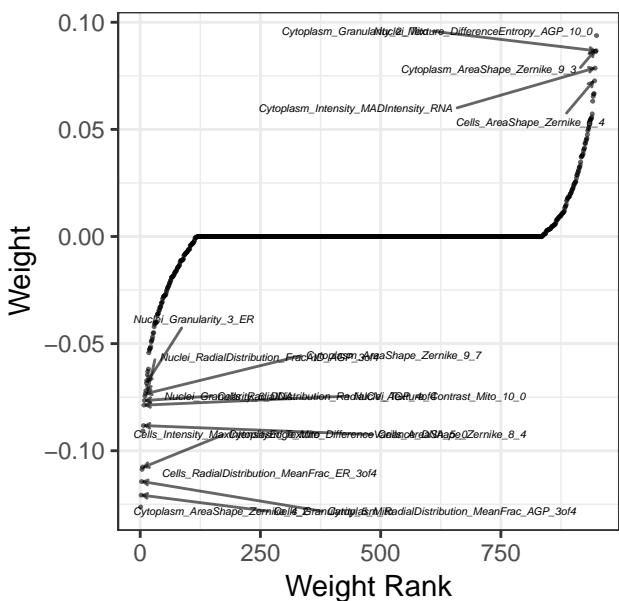


Performance: cc_all_large_round_polyplloid_mean

Transform: raw

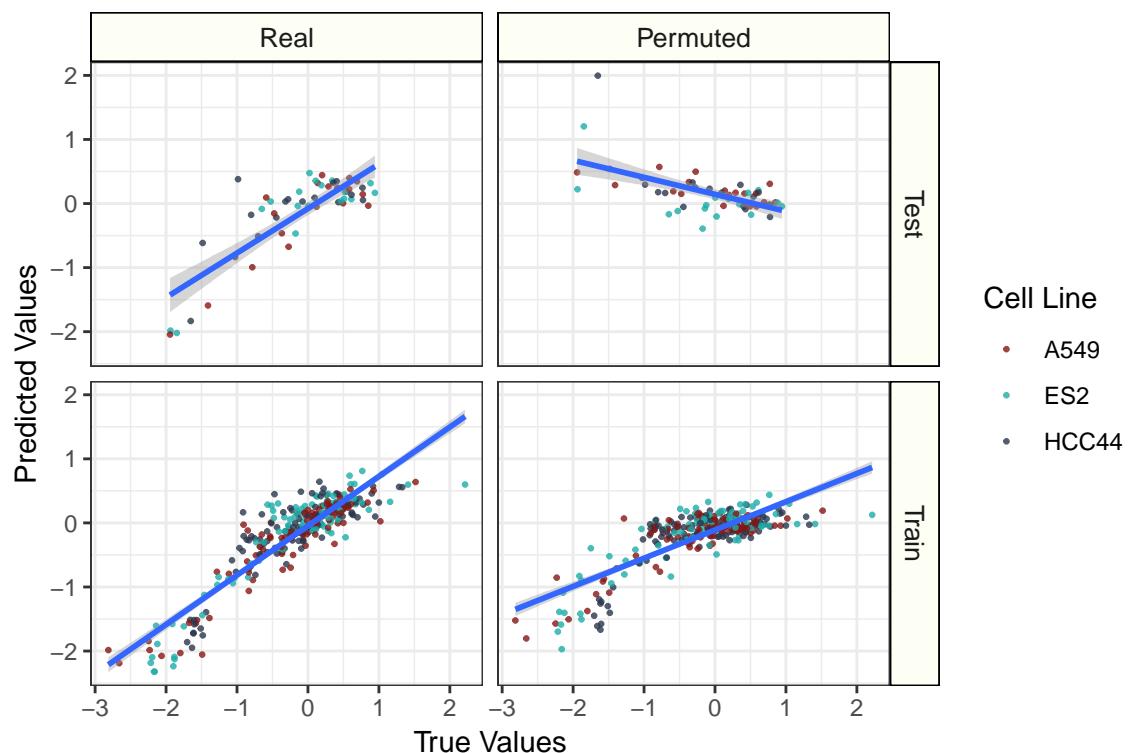


fit	shuffle	transform	r2	value
Train	Real	raw	0.74	0.23
Test	Real	raw	-1.14	0.43
Train	Shuffle	raw	0.00	0.90
Test	Shuffle	raw	-0.21	0.24

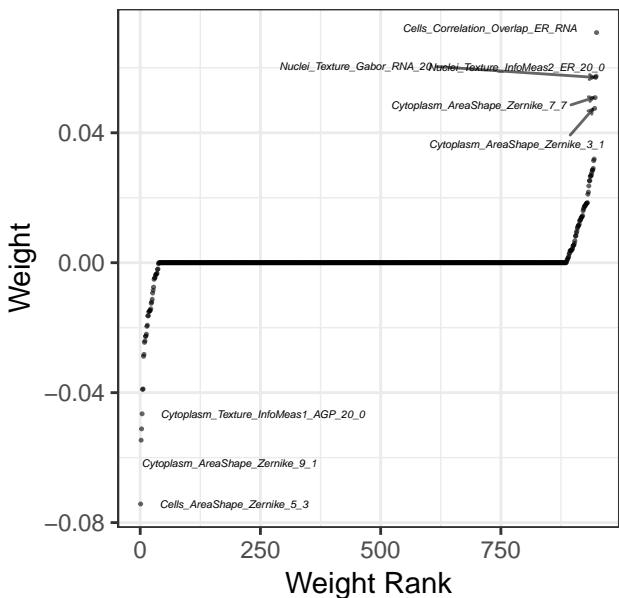


Performance: cc_all_n_objects

Transform: raw

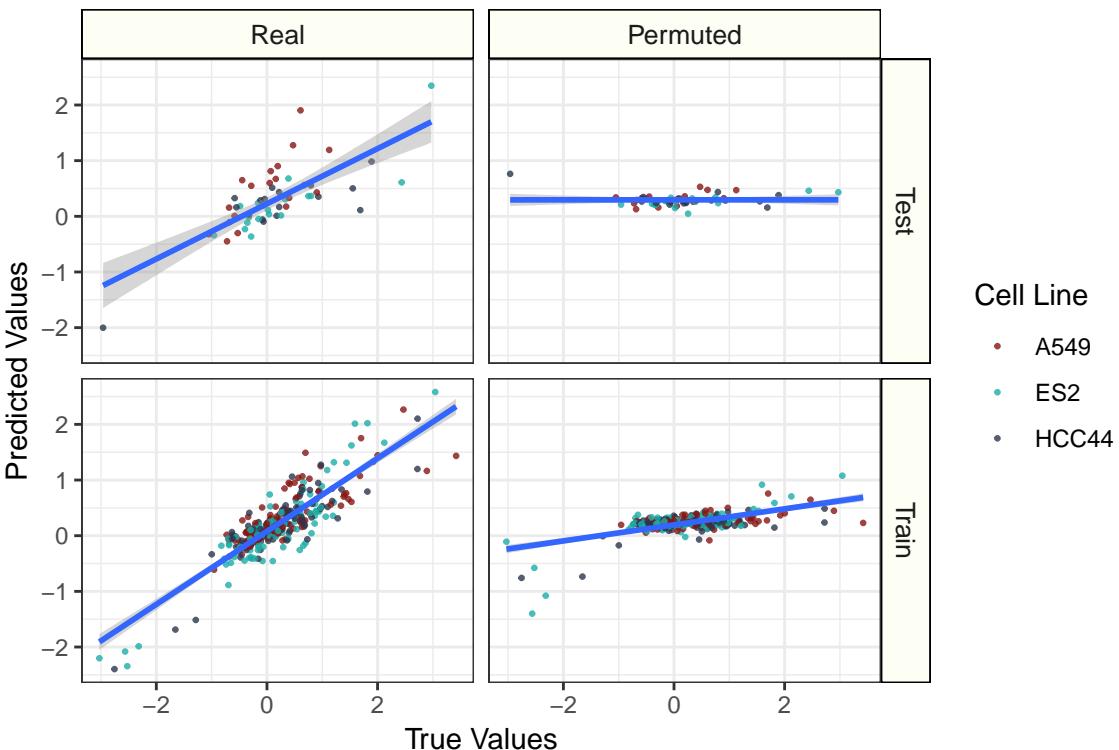


fit	shuffle	transform	r2	value
Train	Real	raw	0.81	0.12
Test	Real	raw	0.67	0.19
Train	Shuffle	raw	0.57	0.26
Test	Shuffle	raw	-0.86	1.08

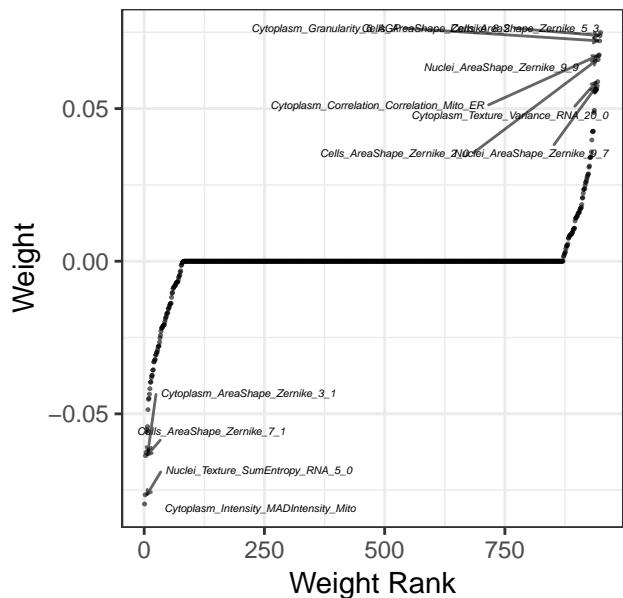


Performance: cc_all_n_spots_h2ax_mean

Transform: raw

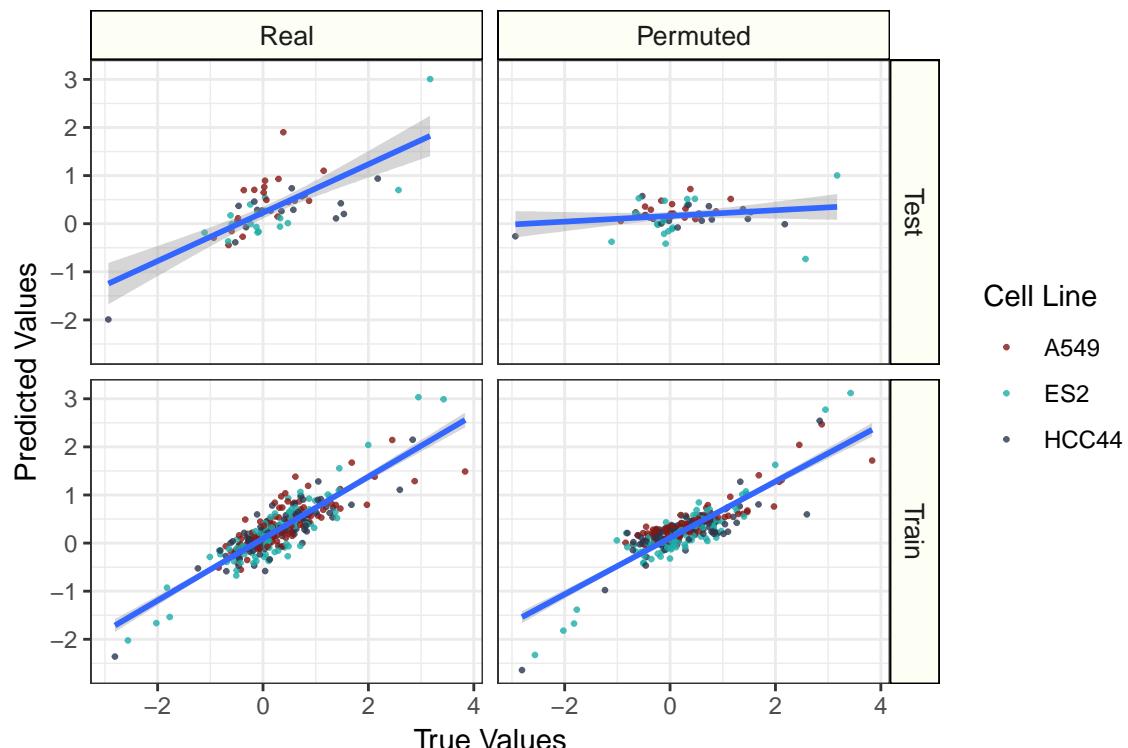


fit	shuffle	transform	r2	value
Train	Real	raw	0.74	0.17
Test	Real	raw	0.51	0.40
Train	Shuffle	raw	0.23	0.49
Test	Shuffle	raw	-0.04	0.85

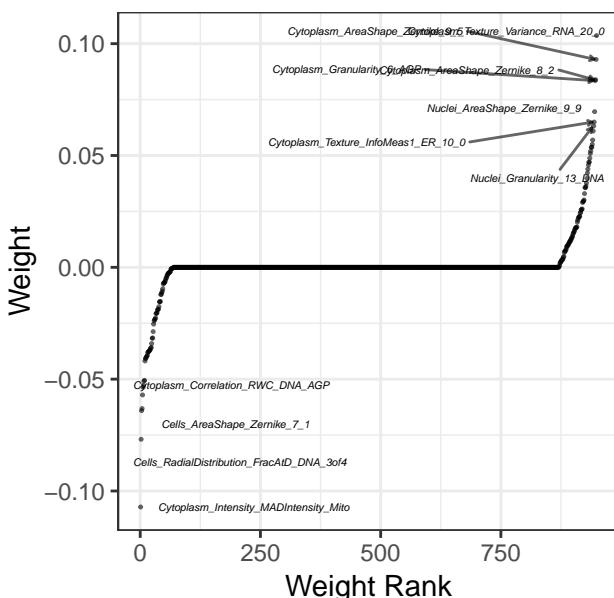


Performance: cc_all_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

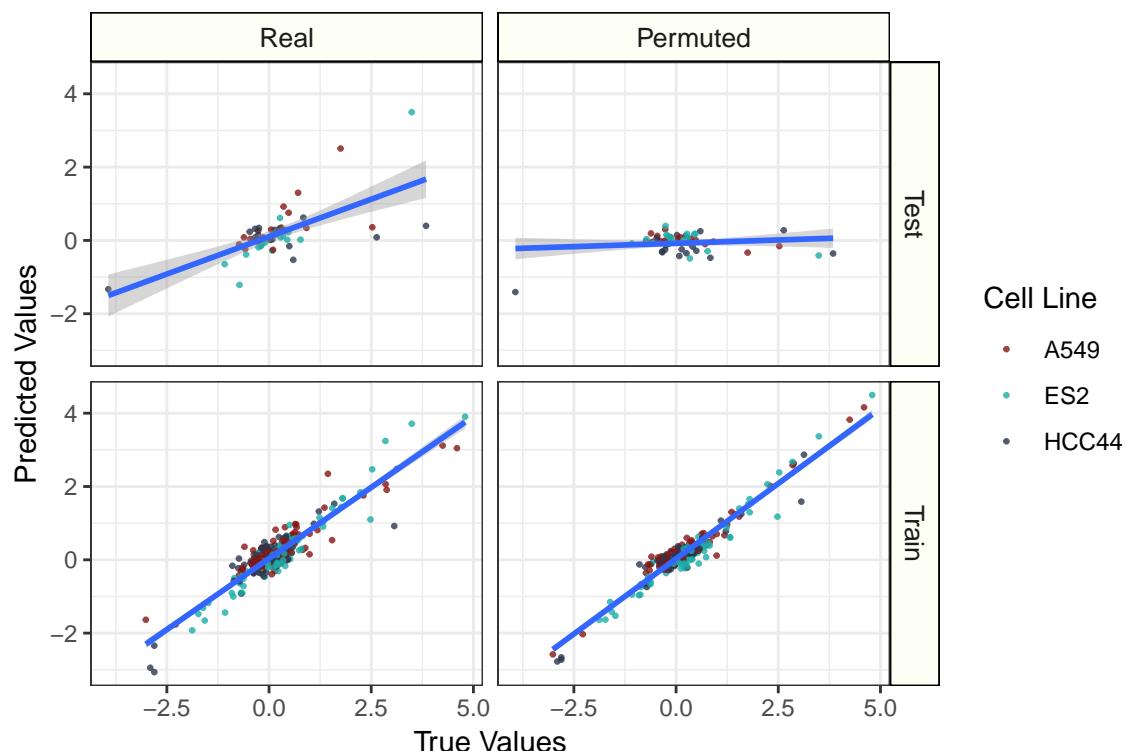


fit	shuffle	transform	r2	value
Train	Real	raw	0.73	0.16
Test	Real	raw	0.50	0.43
Train	Shuffle	raw	0.71	0.17
Test	Shuffle	raw	0.02	0.84

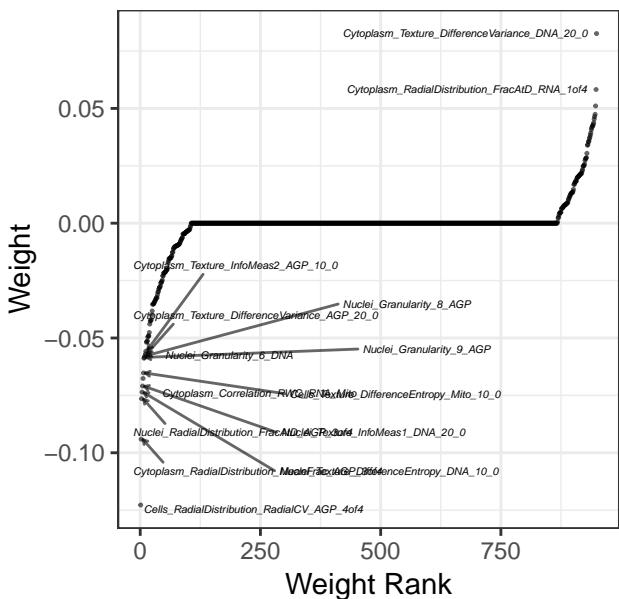


Performance: cc_all_nucleus_area_mean

Transform: raw

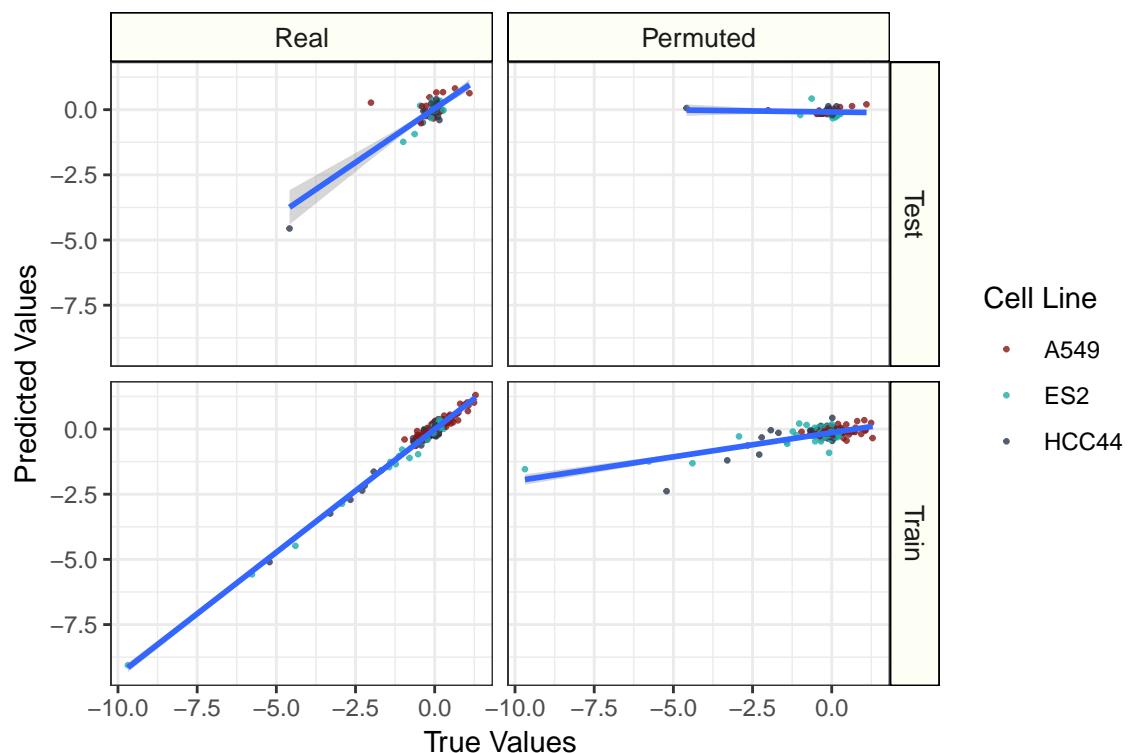


fit	shuffle	transform	r2	value
Train	Real	raw	0.85	0.13
Test	Real	raw	0.42	0.72
Train	Shuffle	raw	0.91	0.08
Test	Shuffle	raw	-0.04	1.28

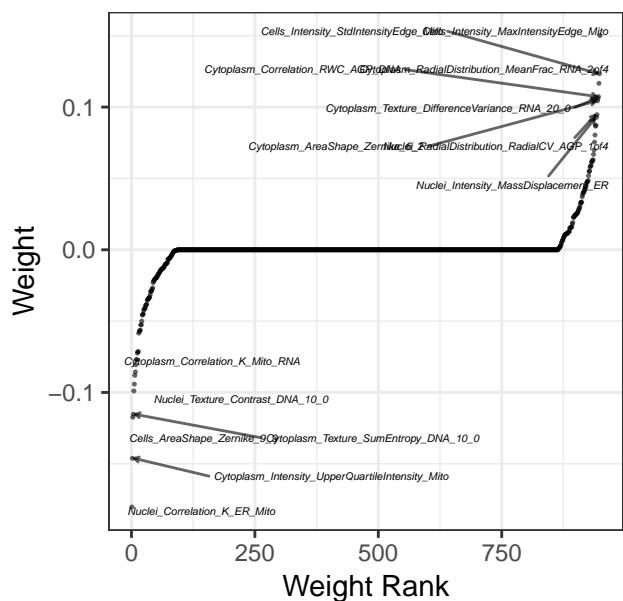


Performance: cc_all_nucleus_roundness_mean

Transform: raw

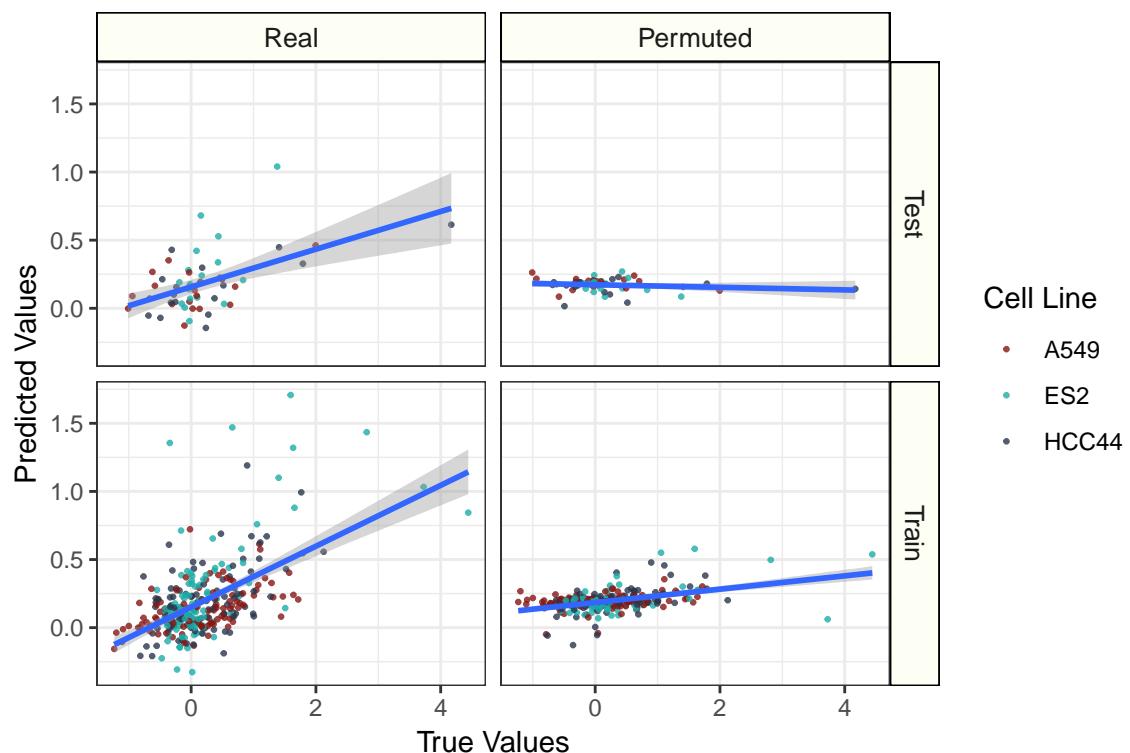


fit	shuffle	transform	r2	value
Train	Real	raw	0.97	0.02
Test	Real	raw	0.68	0.17
Train	Shuffle	raw	0.30	0.56
Test	Shuffle	raw	-0.07	0.57

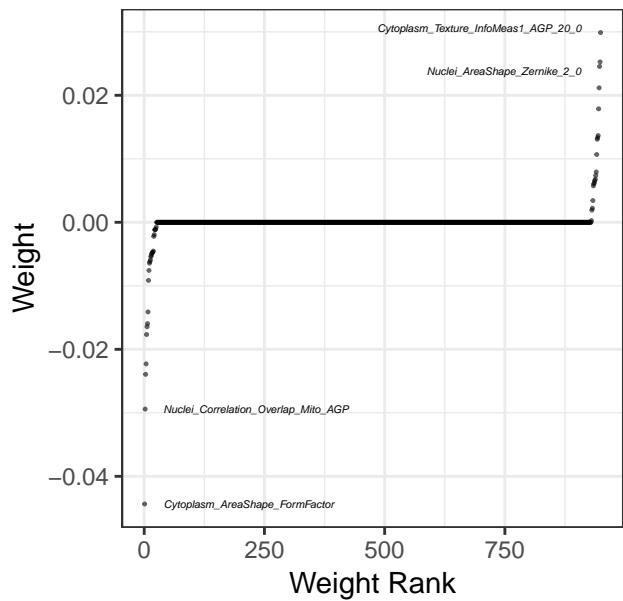


Performance: cc_cc_early_mitosis

Transform: raw

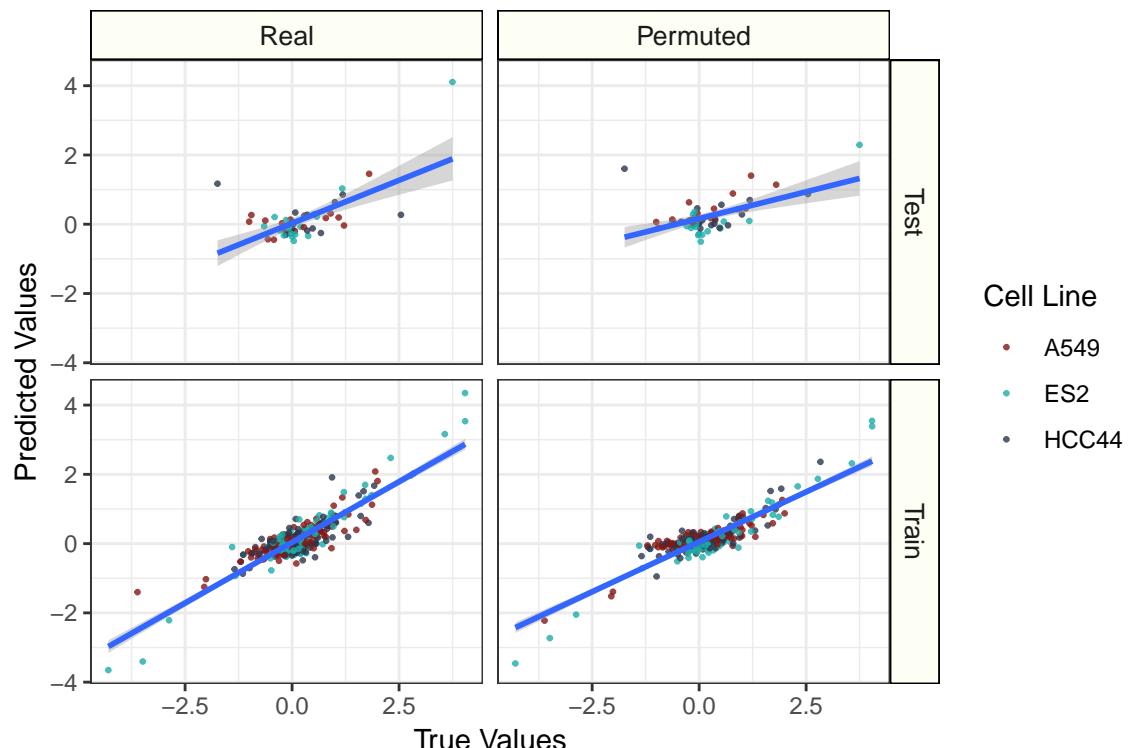


fit	shuffle	transform	r2	value
Train	Real	raw	0.28	0.33
Test	Real	raw	0.21	0.51
Train	Shuffle	raw	0.09	0.42
Test	Shuffle	raw	-0.02	0.66

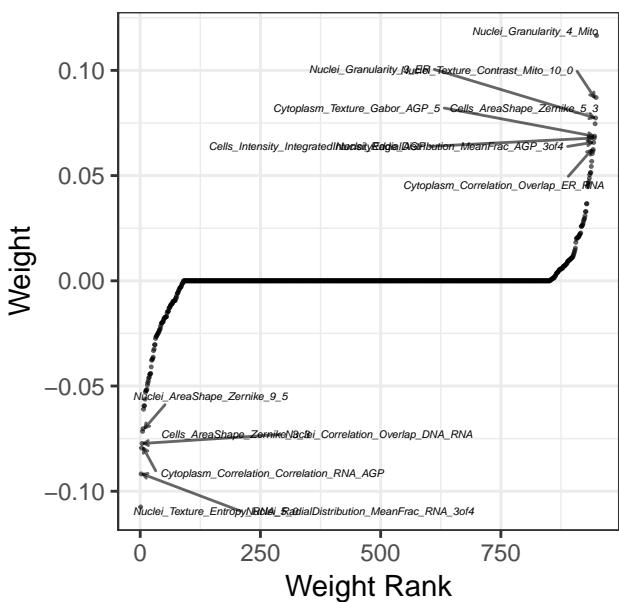


Performance: cc_cc_g1

Transform: raw

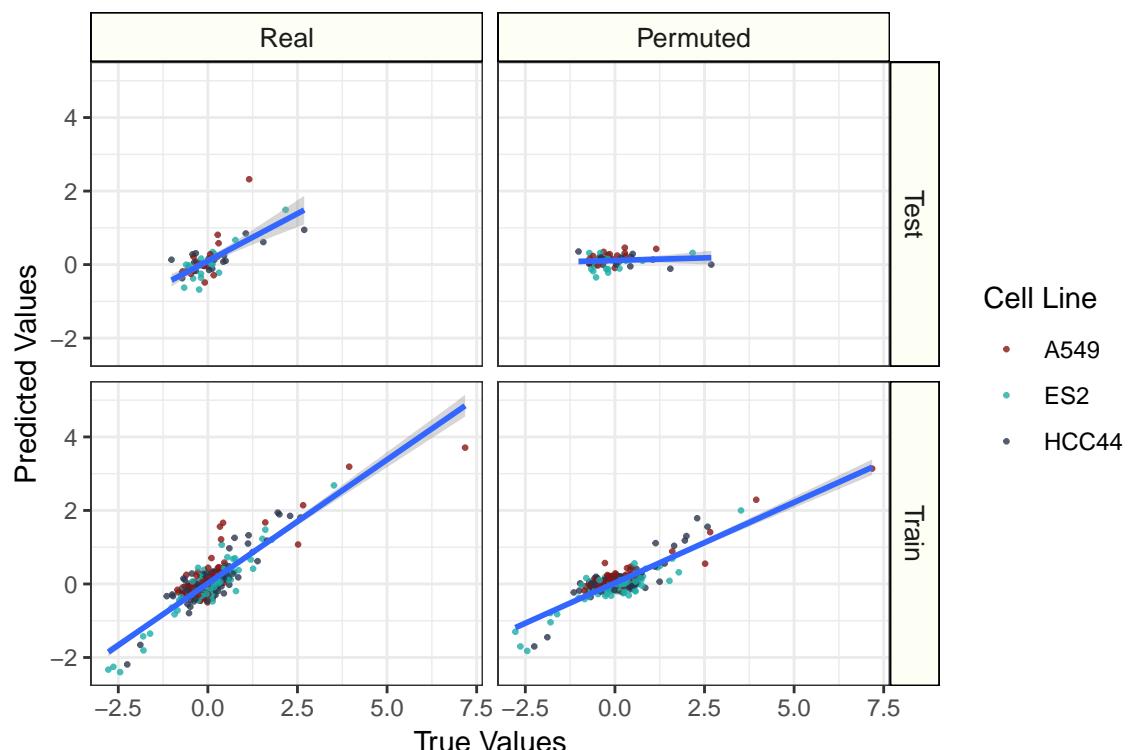


fit	shuffle	transform	r2	value
Train	Real	raw	0.78	0.17
Test	Real	raw	0.35	0.46
Train	Shuffle	raw	0.73	0.22
Test	Shuffle	raw	0.28	0.51

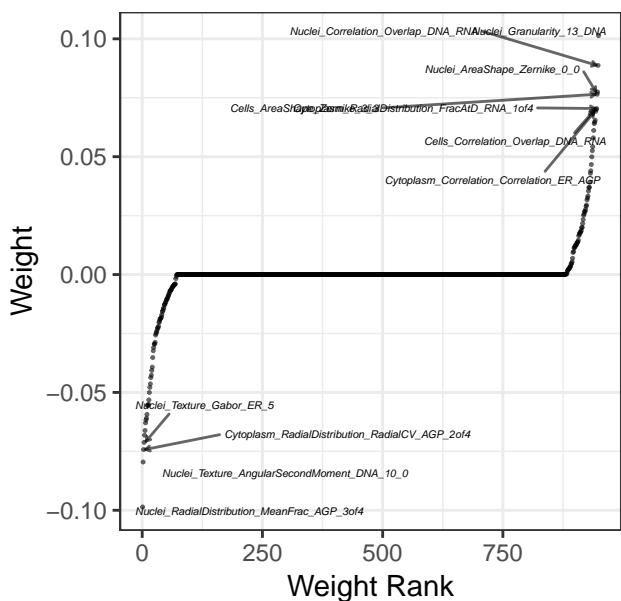


Performance: cc_cc_g2

Transform: raw

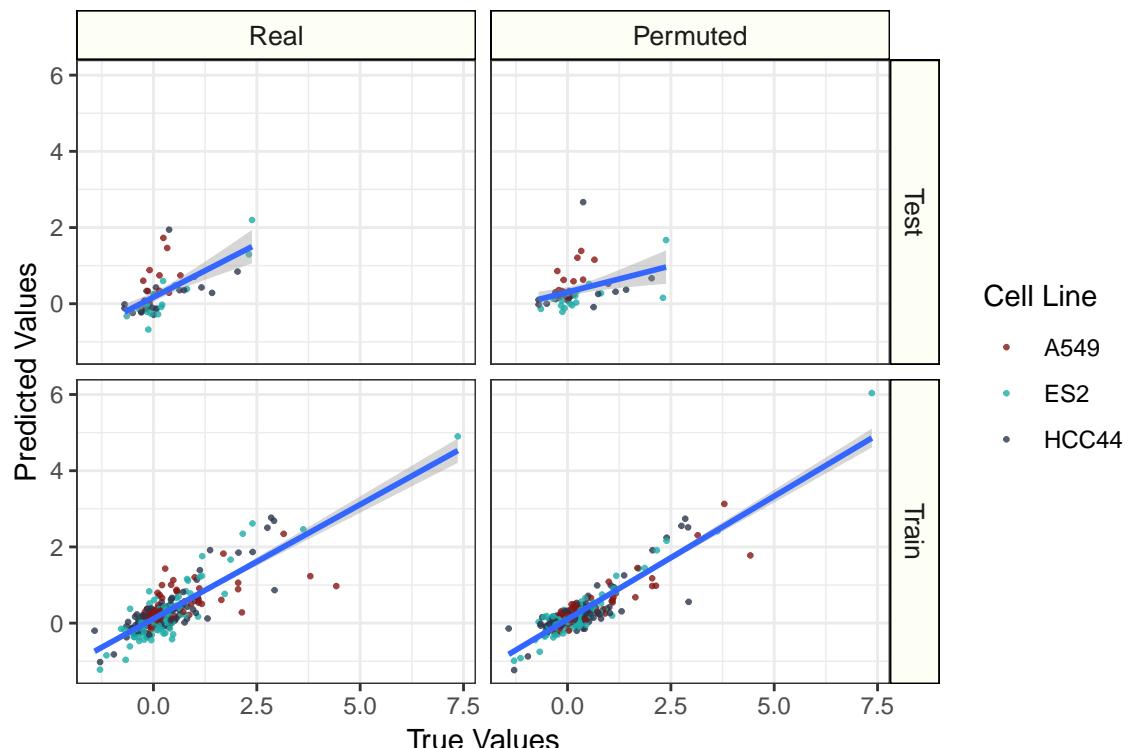


fit	shuffle	transform	r2	value
Train	Real	raw	0.76	0.17
Test	Real	raw	0.49	0.24
Train	Shuffle	raw	0.62	0.27
Test	Shuffle	raw	-0.03	0.48

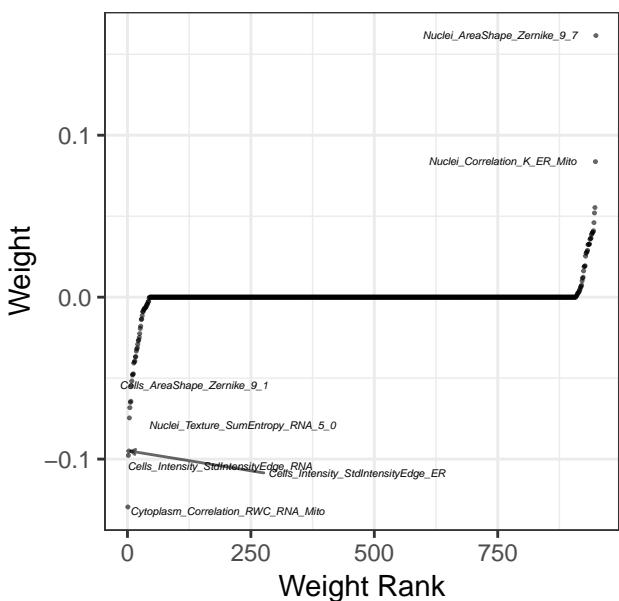


Performance: cc_cc_high_h2ax

Transform: raw

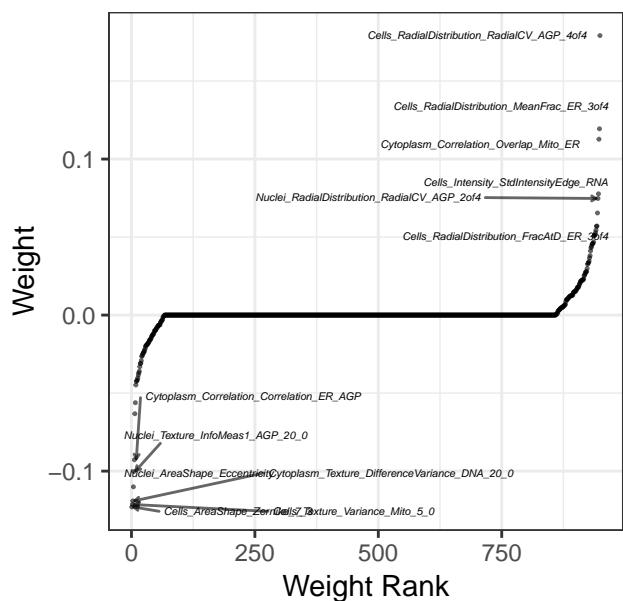
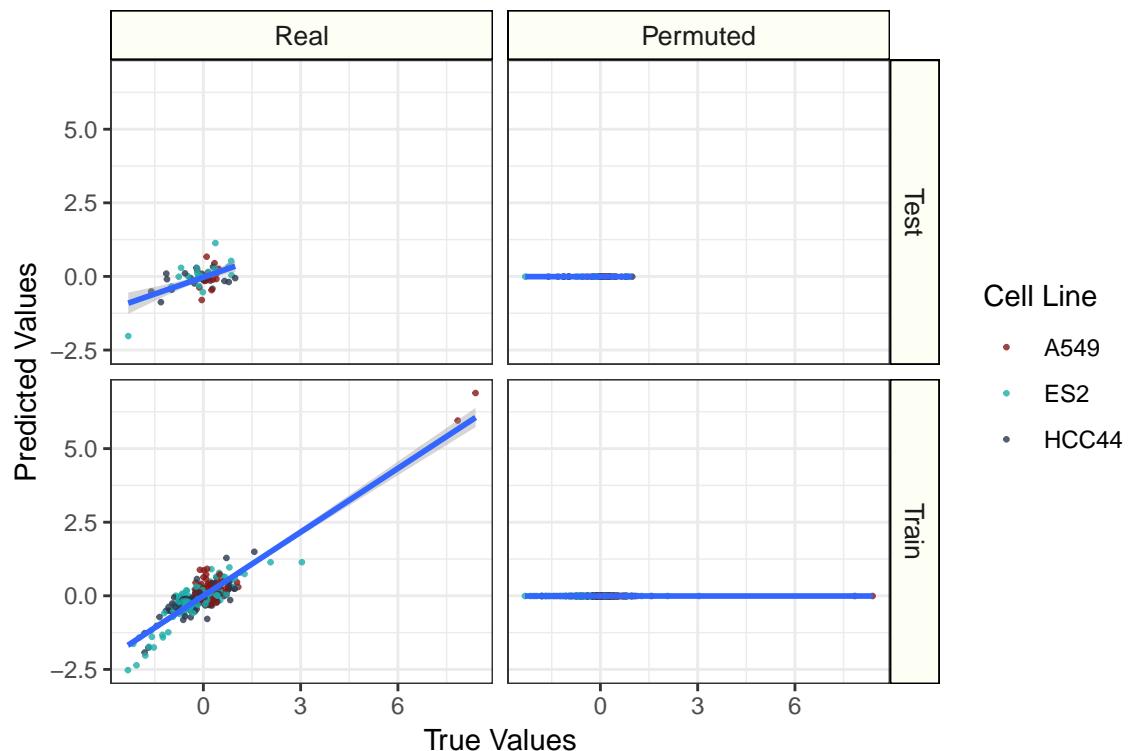


fit	shuffle	transform	r2	value
Train	Real	raw	0.68	0.24
Test	Real	raw	0.31	0.29
Train	Shuffle	raw	0.78	0.16
Test	Shuffle	raw	-0.06	0.46



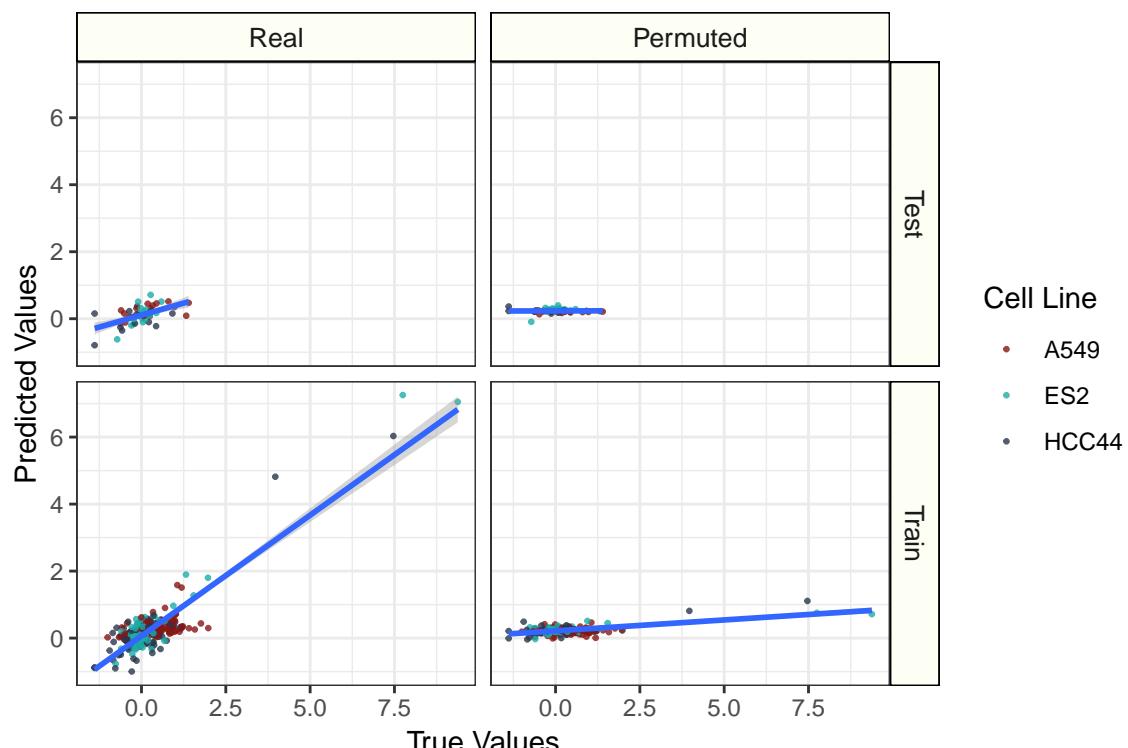
Performance: cc_cc_late_mitosis

Transform: raw

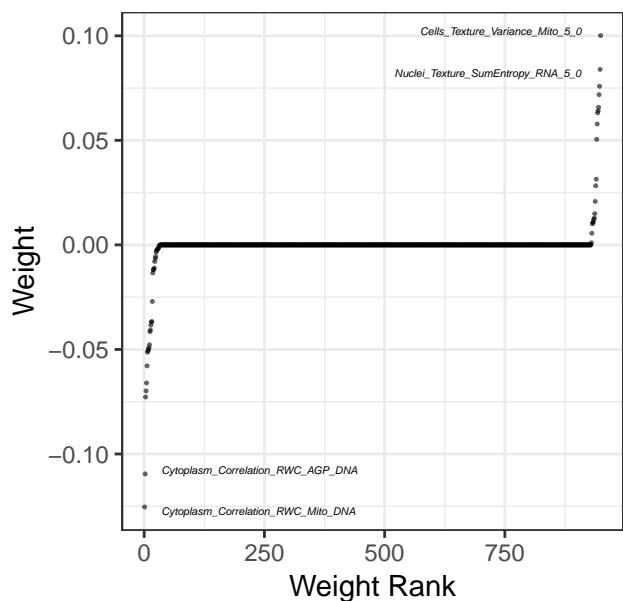


Performance: cc_cc_mitosis

Transform: raw

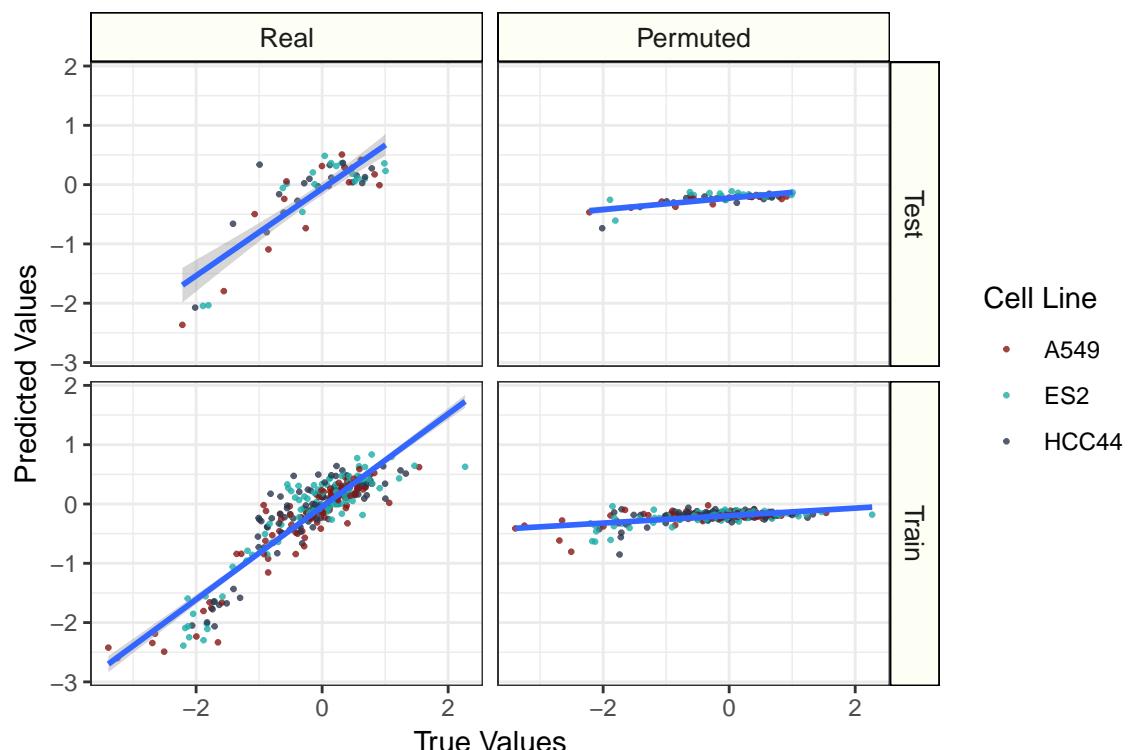


fit	shuffle	transform	r2	value
Train	Real	raw	0.79	0.20
Test	Real	raw	0.28	0.20
Train	Shuffle	raw	0.12	0.84
Test	Shuffle	raw	-0.17	0.32

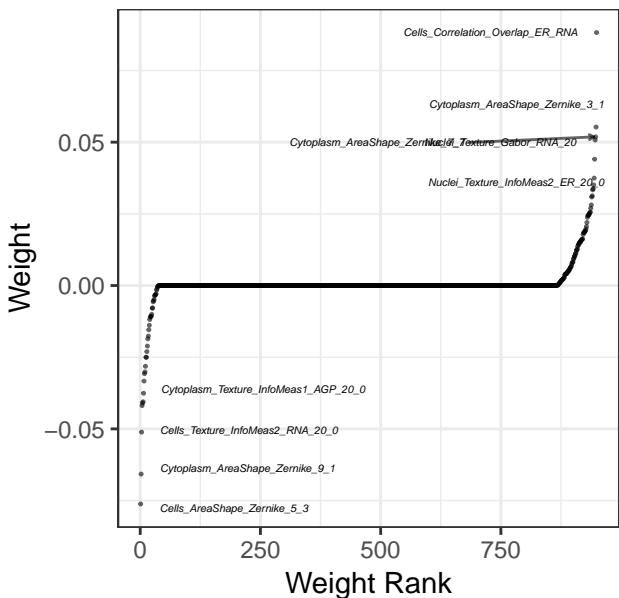


Performance: cc_cc_n_objects

Transform: raw

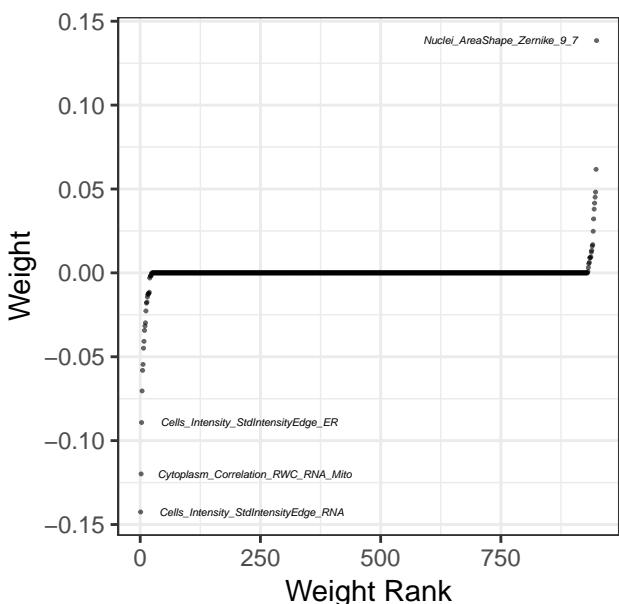
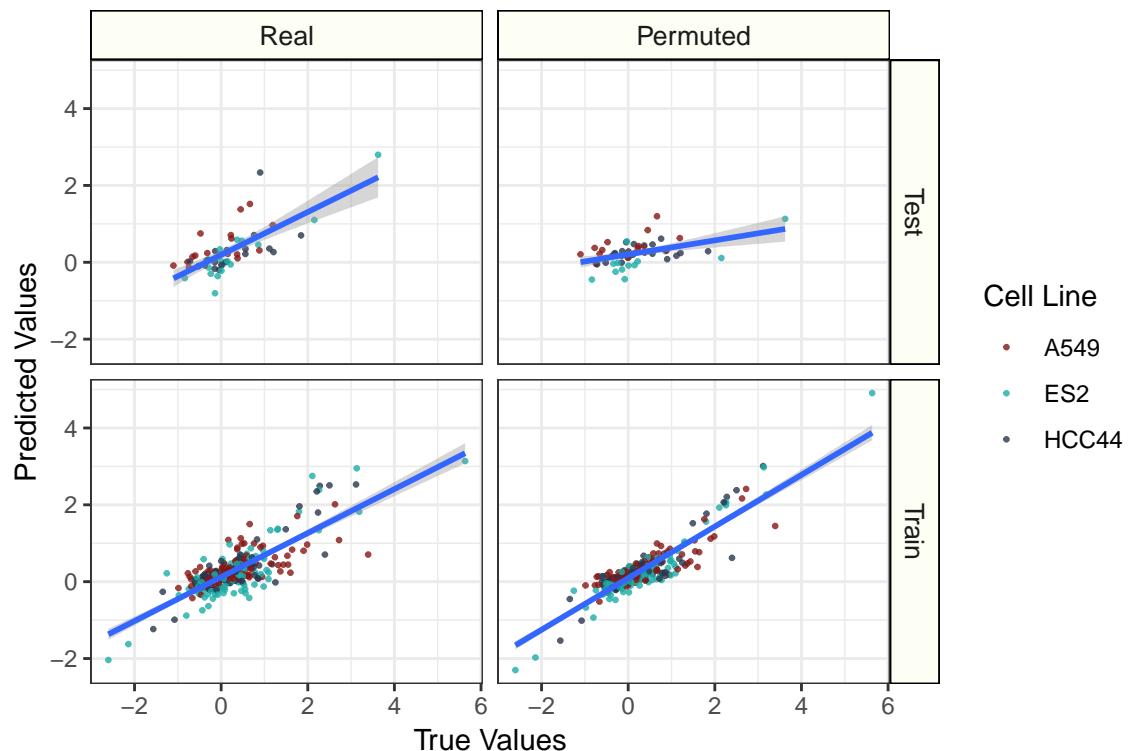


fit	shuffle	transform	r2	value
Train	Real	raw	0.82	0.12
Test	Real	raw	0.71	0.19
Train	Shuffle	raw	0.11	0.61
Test	Shuffle	raw	0.15	0.54



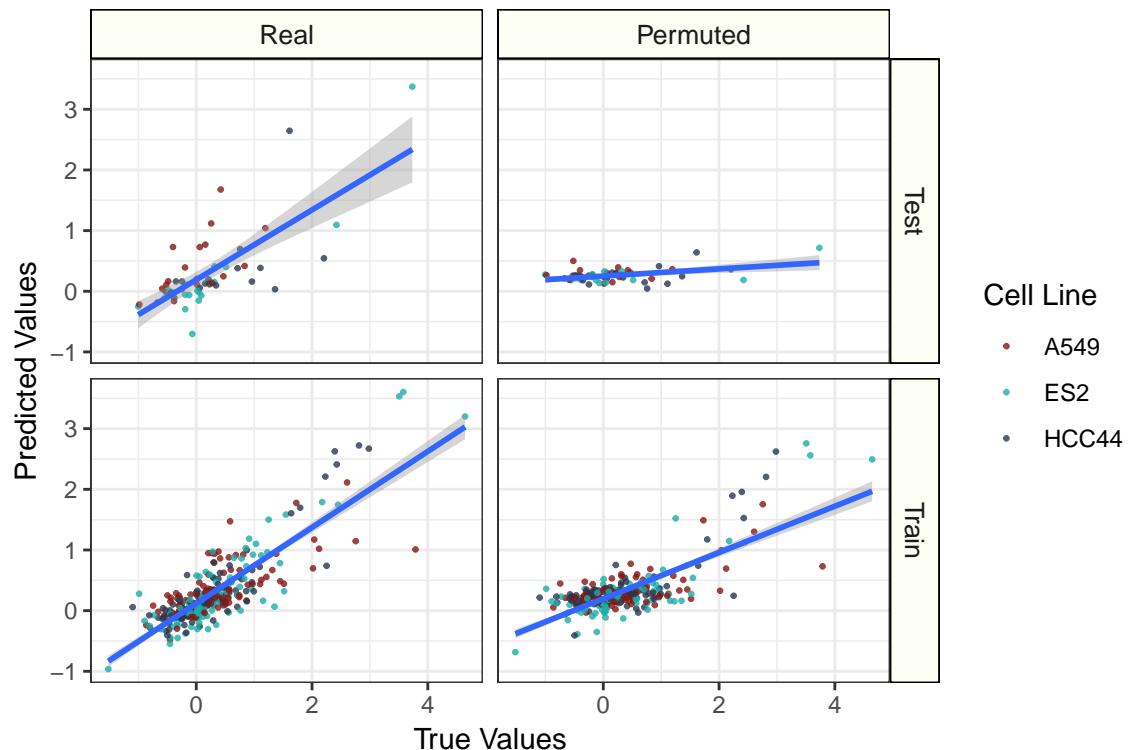
Performance: cc_cc_n_spots_h2ax_mean

Transform: raw

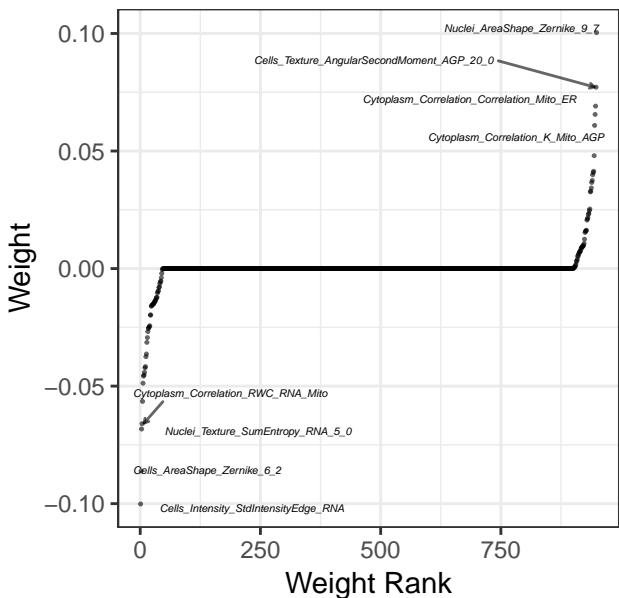


Performance: cc_cc_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

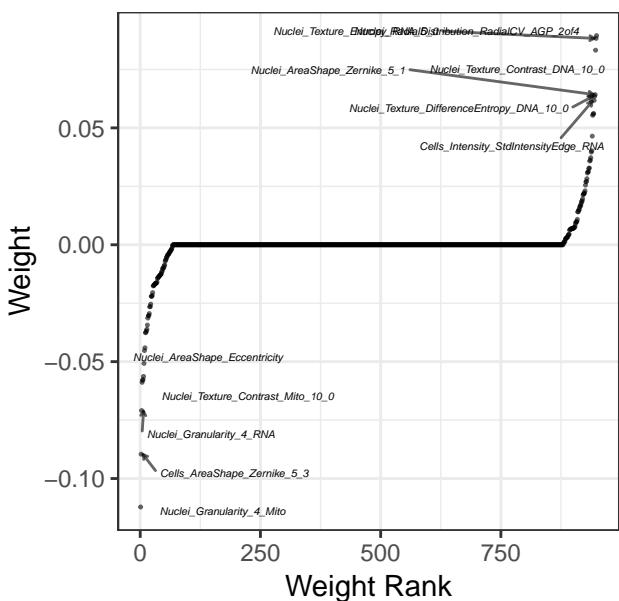
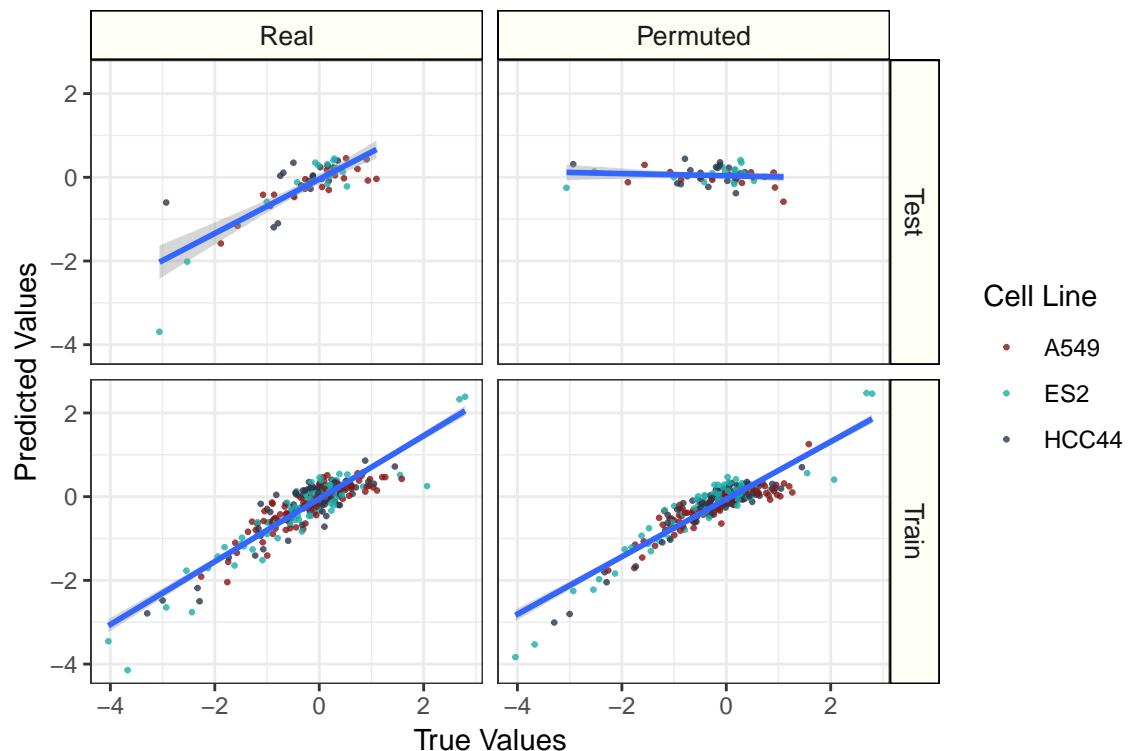


fit	shuffle	transform	r2	value
Train	Real	raw	0.70	0.20
Test	Real	raw	0.51	0.35
Train	Shuffle	raw	0.51	0.32
Test	Shuffle	raw	0.10	0.65



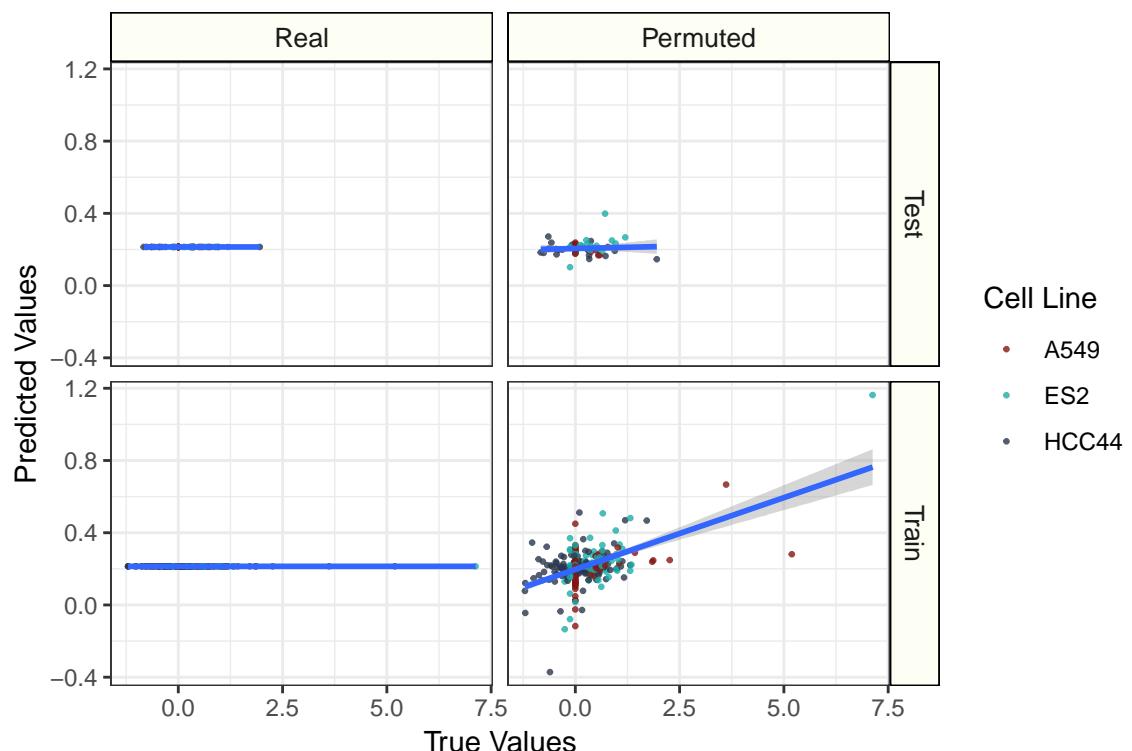
Performance: cc_cc_s

Transform: raw

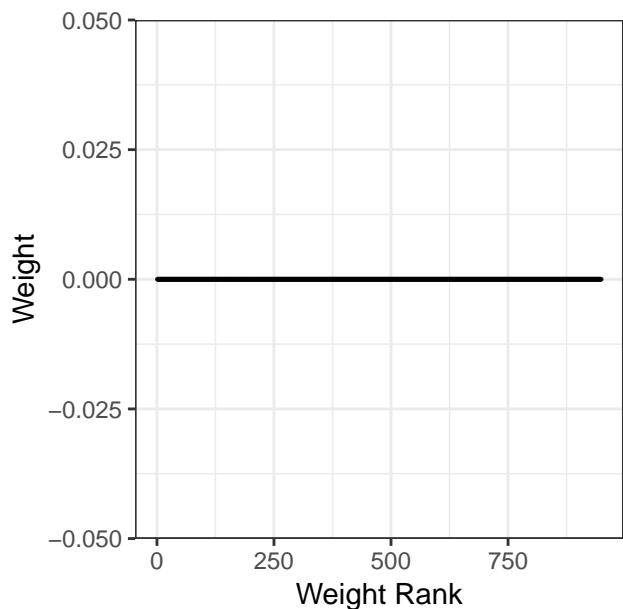


Performance: cc_early_mitosis_high_h2ax

Transform: raw

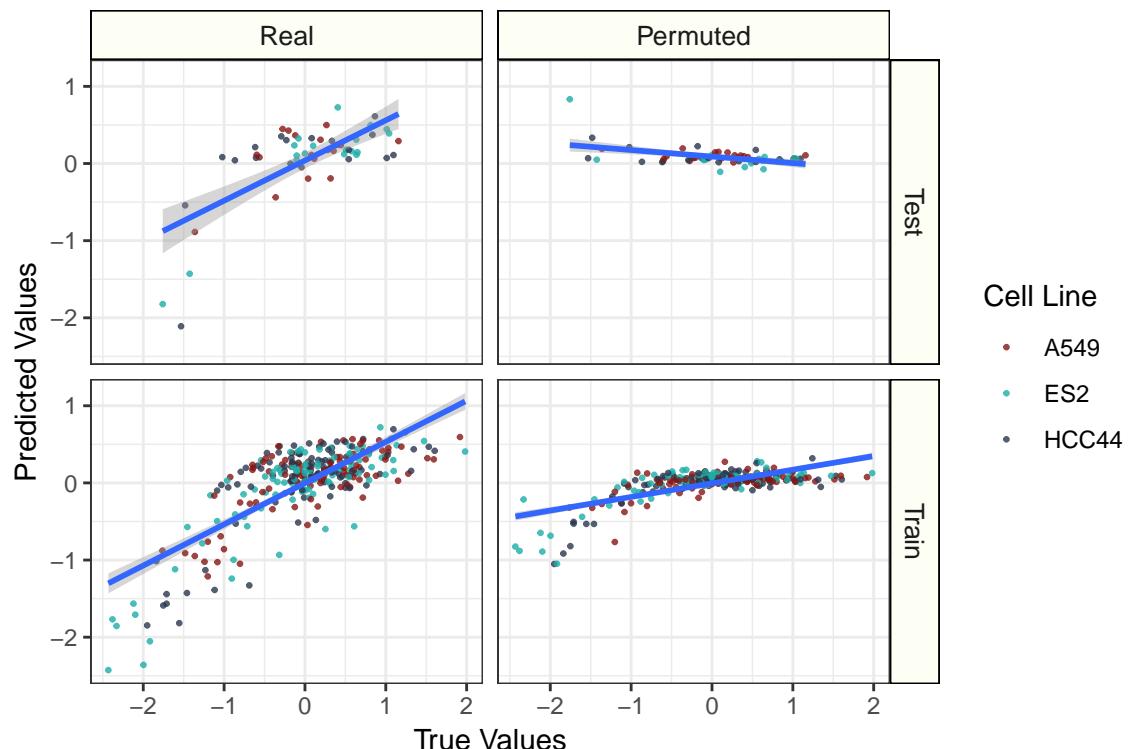


fit	shuffle	transform	r2	value
Train	Real	raw	0.00	0.52
Test	Real	raw	0.00	0.27
Train	Shuffle	raw	0.14	0.45
Test	Shuffle	raw	0.00	0.27

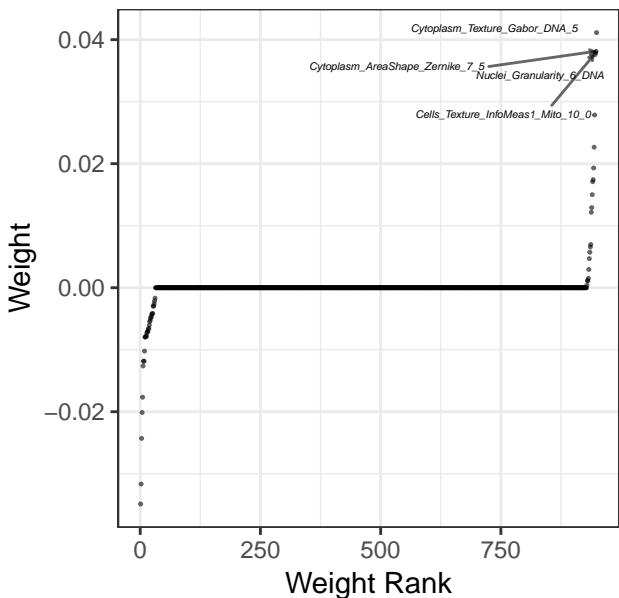


Performance: cc_early_mitosis_n_objects

Transform: raw

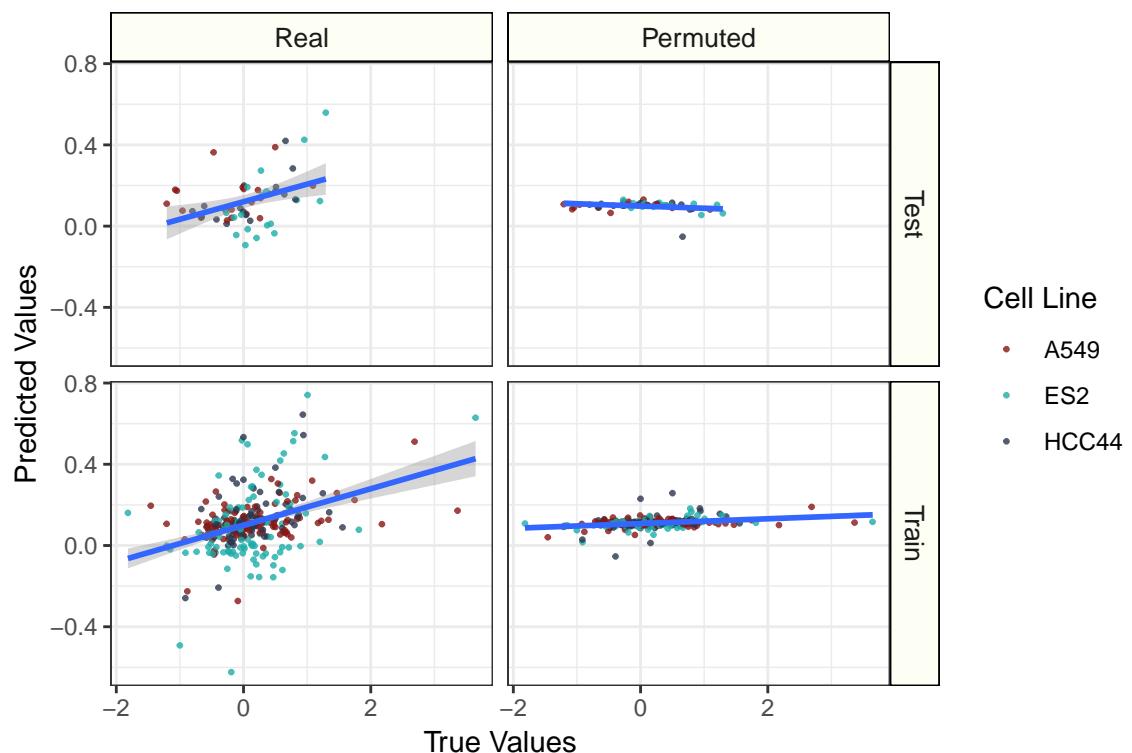


fit	shuffle	transform	r2	value
Train	Real	raw	0.58	0.25
Test	Real	raw	0.49	0.25
Train	Shuffle	raw	0.29	0.42
Test	Shuffle	raw	-0.21	0.60

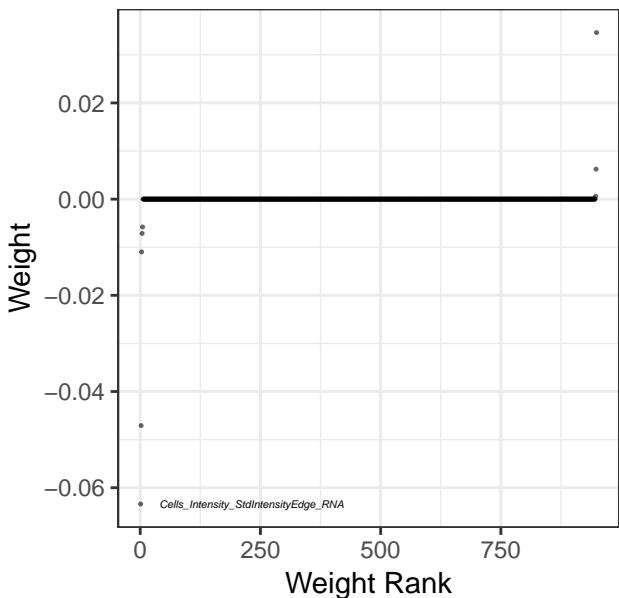


Performance: cc_early_mitosis_n_spots_h2ax_mean

Transform: raw

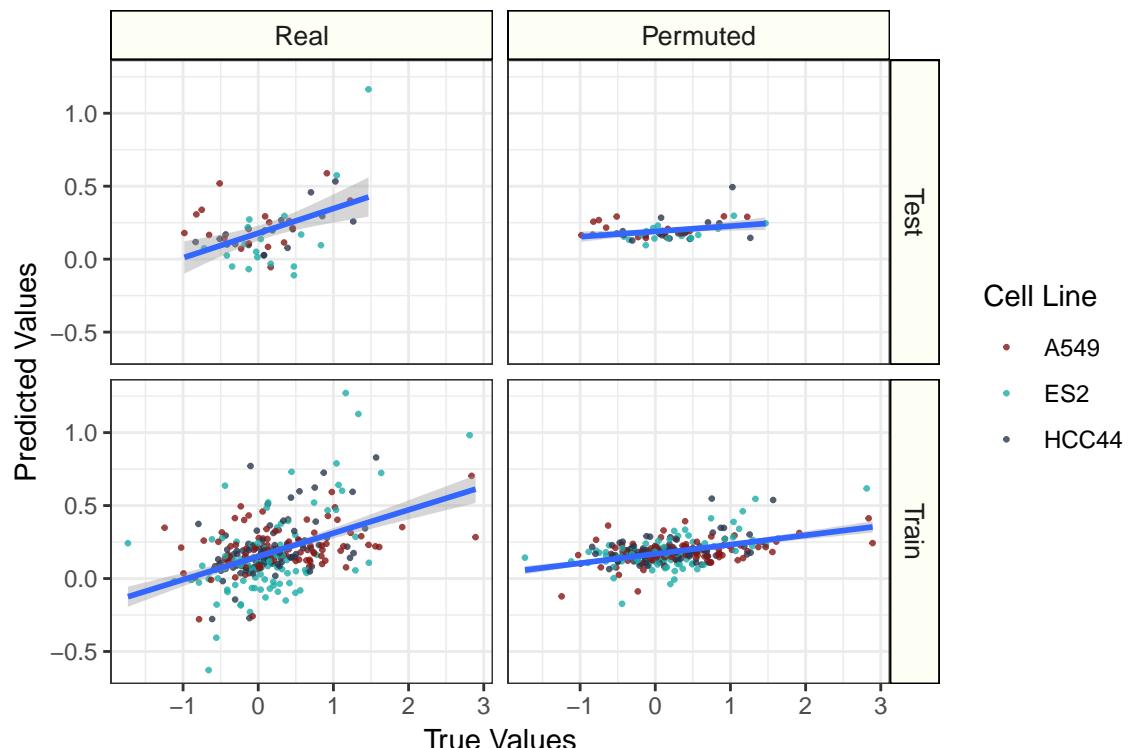


fit	shuffle	transform	r2	value
Train	Real	raw	0.13	0.34
Test	Real	raw	0.11	0.28
Train	Shuffle	raw	0.02	0.38
Test	Shuffle	raw	-0.03	0.32

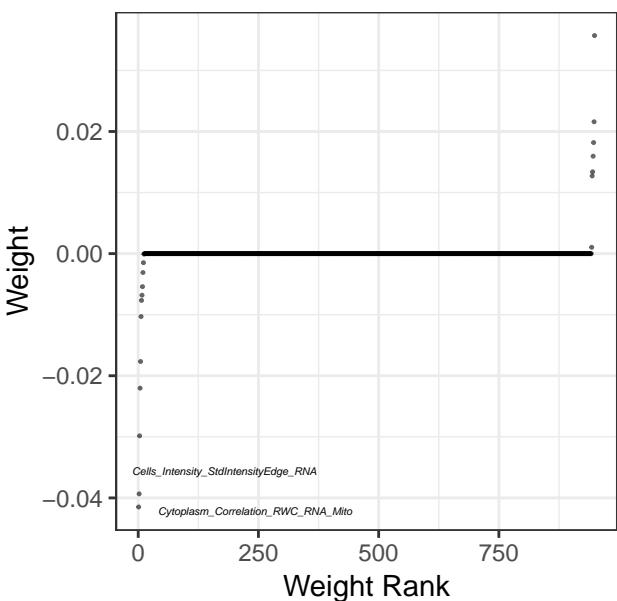


Performance: cc_early_mitosis_n_spots_h2ax_per_nucleus_are

Transform: raw

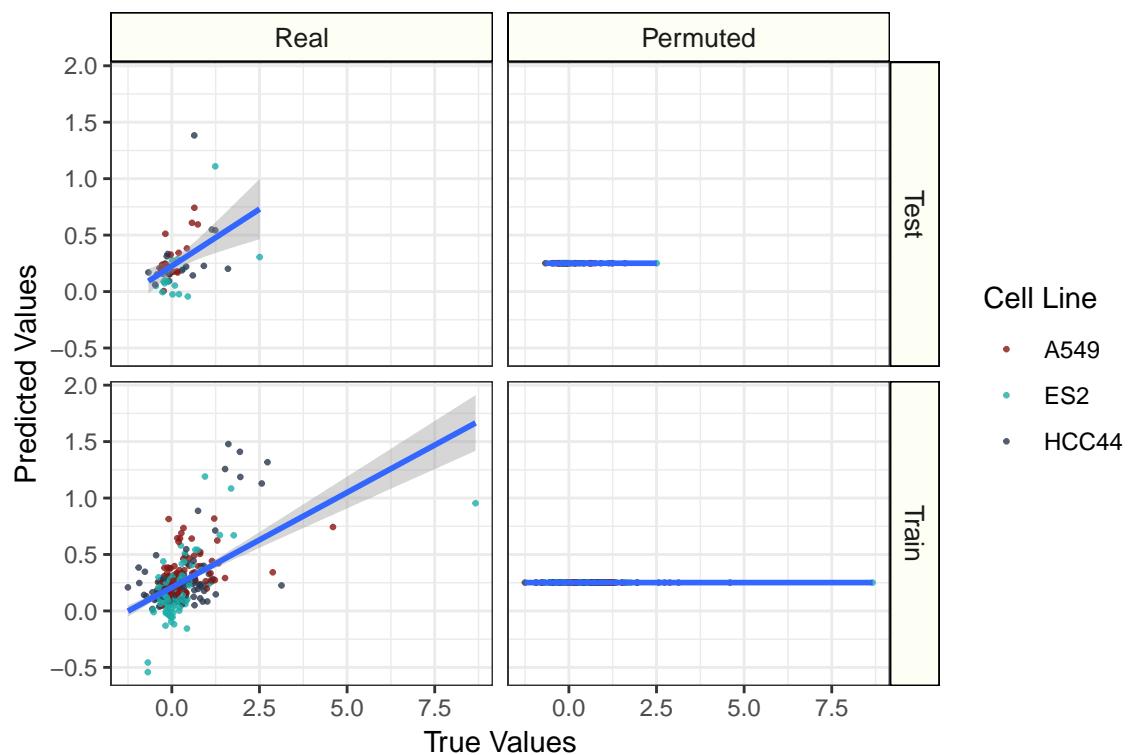


fit	shuffle	transform	r2	value
Train	Real	raw	0.20	0.30
Test	Real	raw	0.17	0.26
Train	Shuffle	raw	0.11	0.34
Test	Shuffle	raw	0.03	0.31

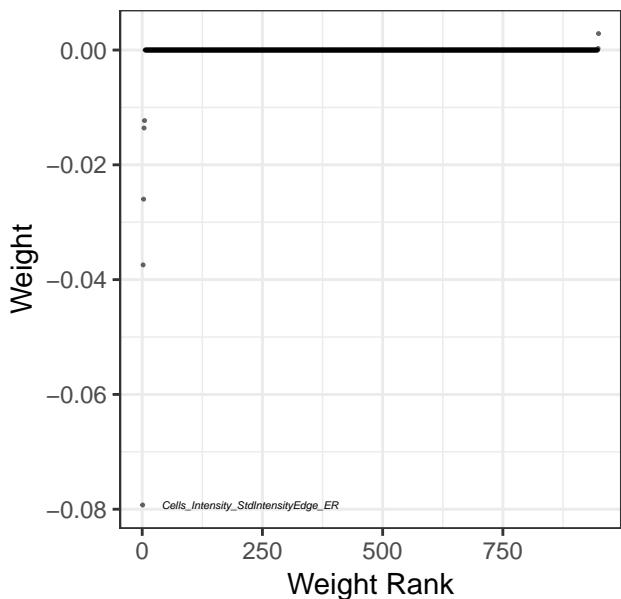


Performance: cc_g1_high_h2ax

Transform: raw

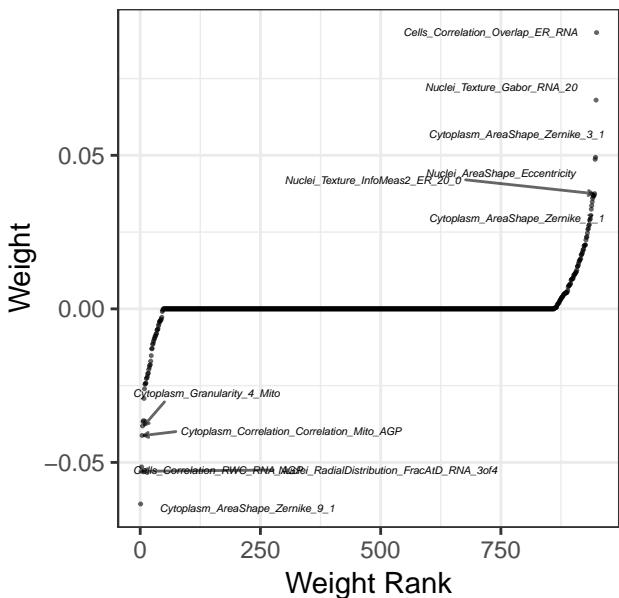
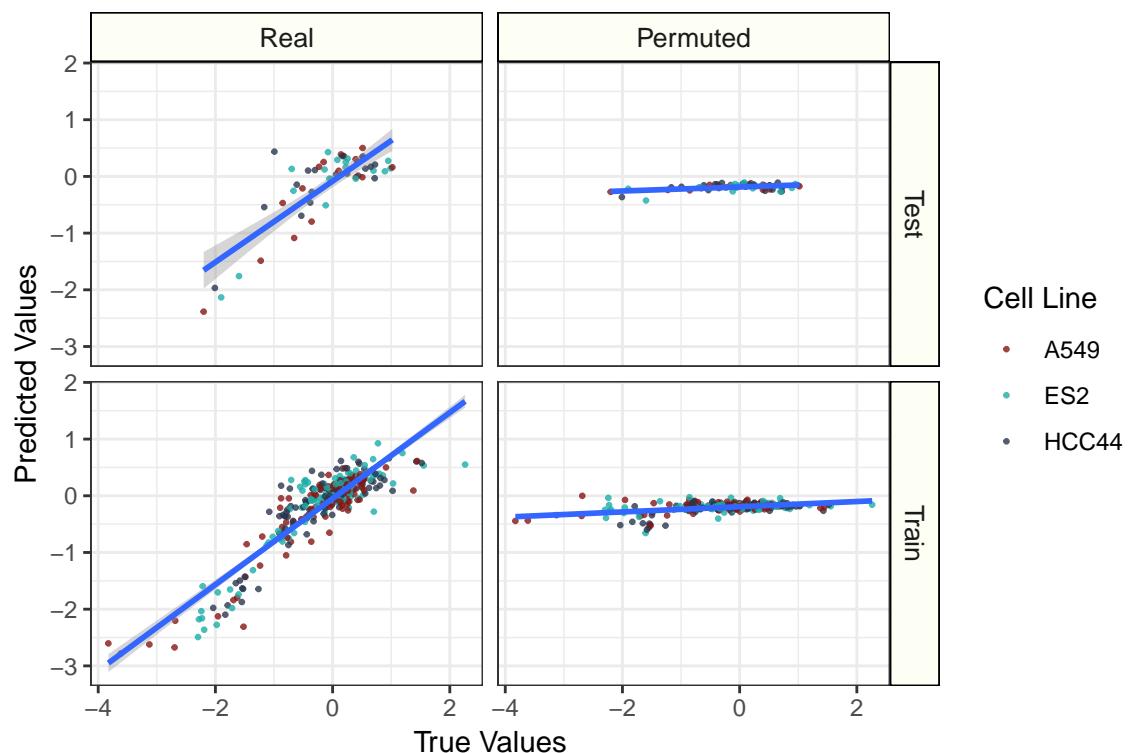


fit	shuffle	transform	r2	value
Train	Real	raw	0.24	0.48
Test	Real	raw	0.17	0.27
Train	Shuffle	raw	0.00	0.64
Test	Shuffle	raw	-0.03	0.34



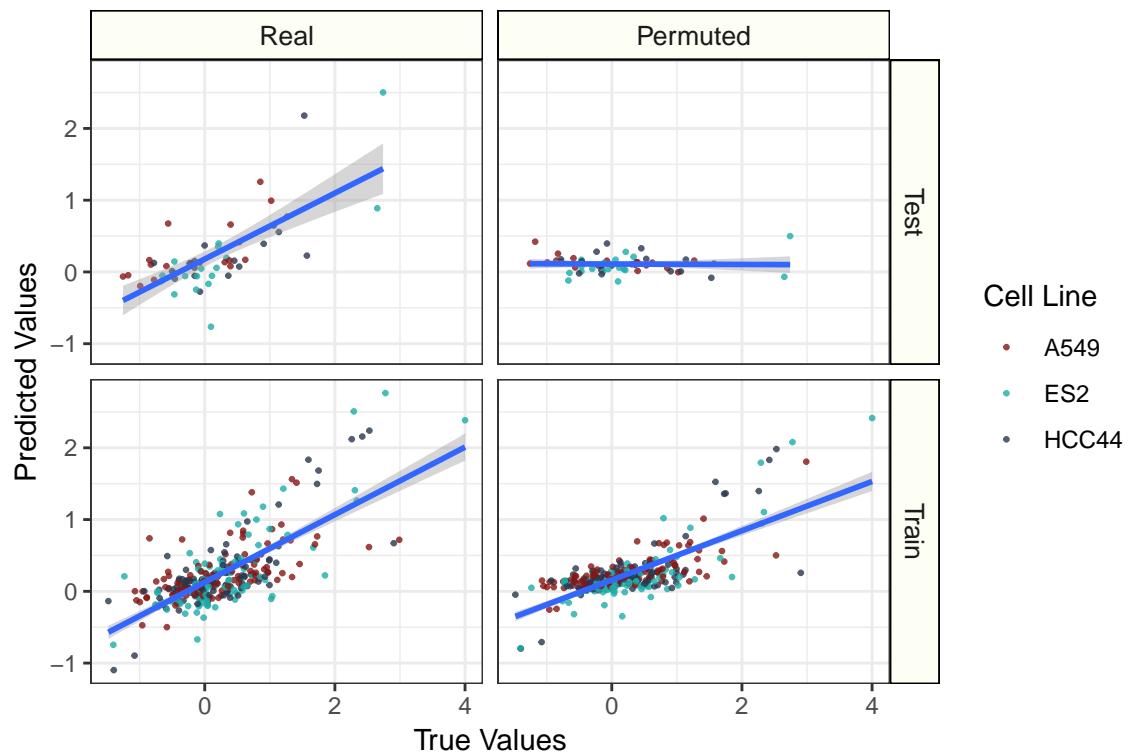
Performance: cc_g1_n_objects

Transform: raw

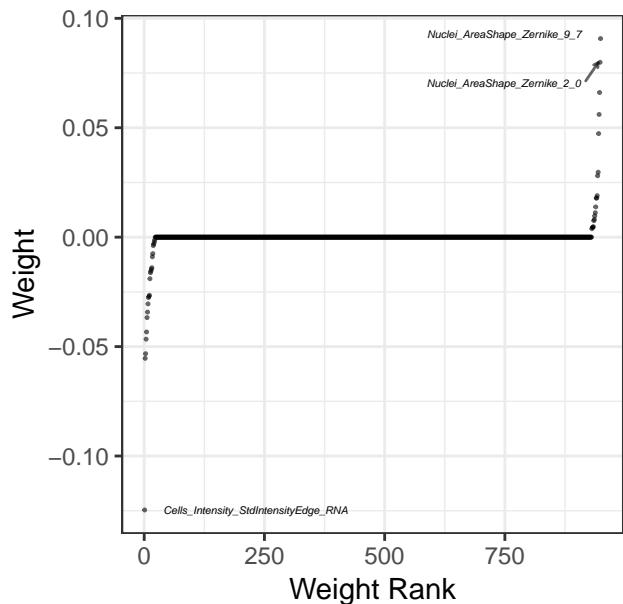


Performance: cc_g1_n_spots_h2ax_mean

Transform: raw

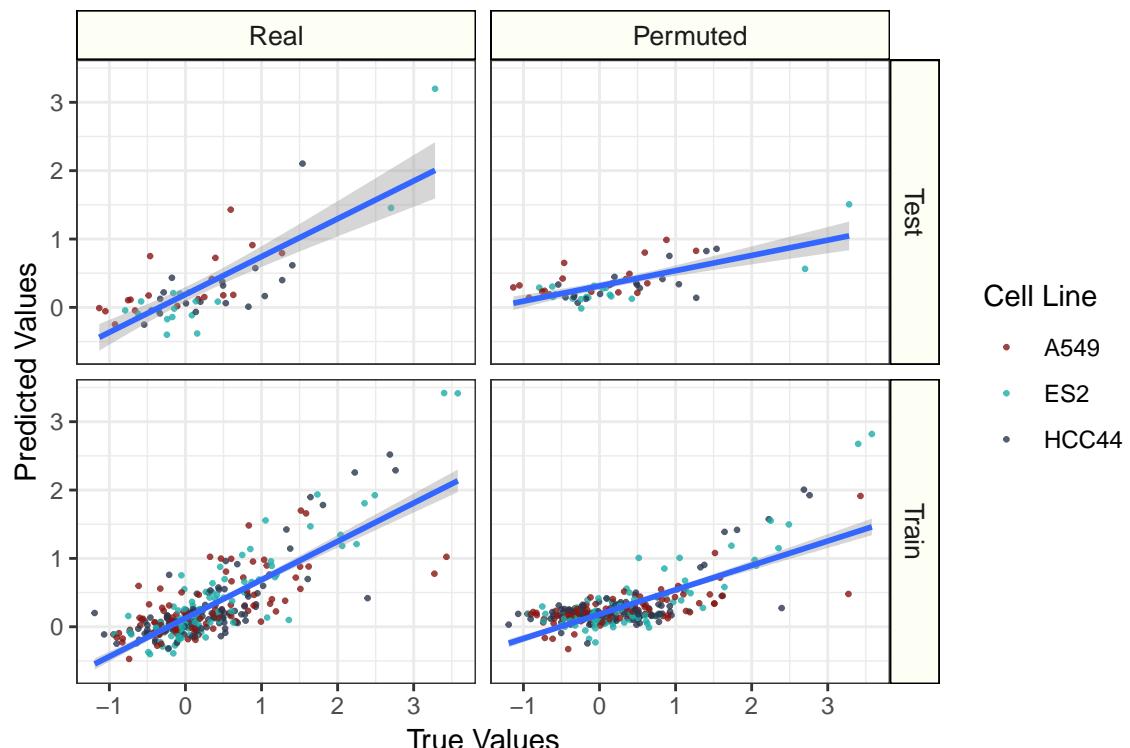


fit	shuffle	transform	r2	value
Train	Real	raw	0.54	0.28
Test	Real	raw	0.48	0.36
Train	Shuffle	raw	0.47	0.31
Test	Shuffle	raw	-0.03	0.70

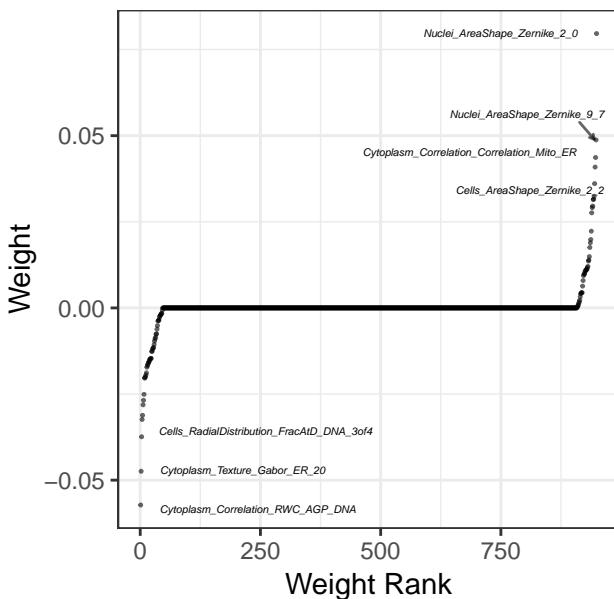


Performance: cc_g1_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

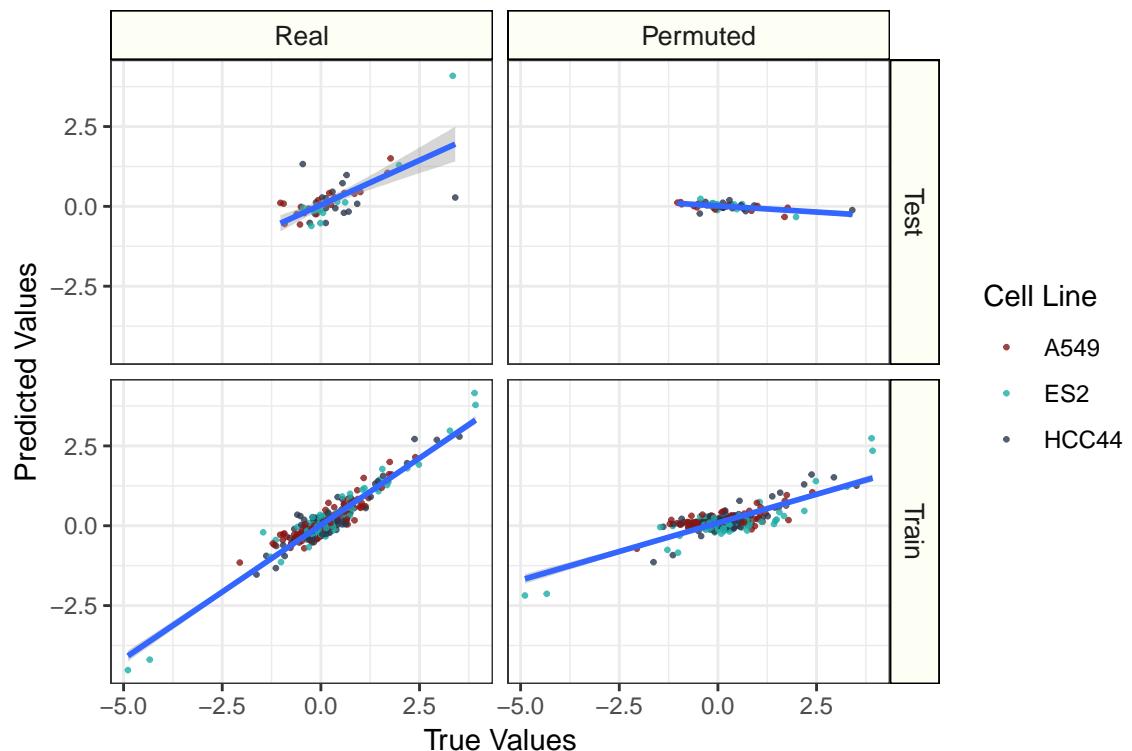


fit	shuffle	transform	r2	value
Train	Real	raw	0.62	0.22
Test	Real	raw	0.57	0.30
Train	Shuffle	raw	0.49	0.30
Test	Shuffle	raw	0.29	0.50

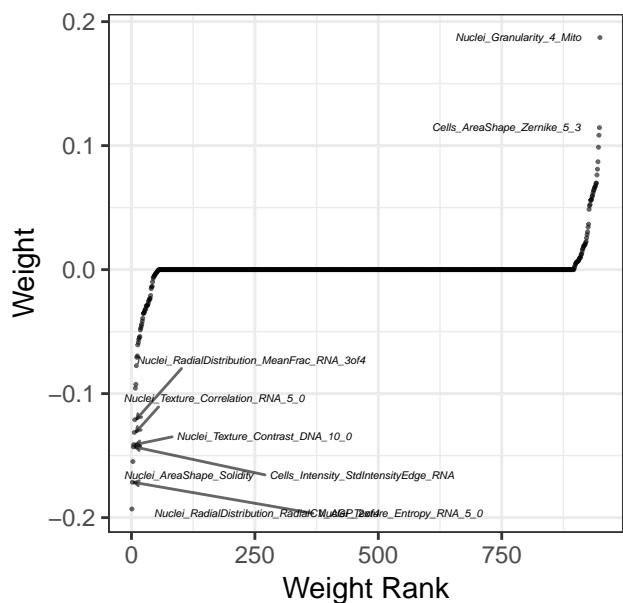


Performance: cc_g1_plus_g2_count

Transform: raw

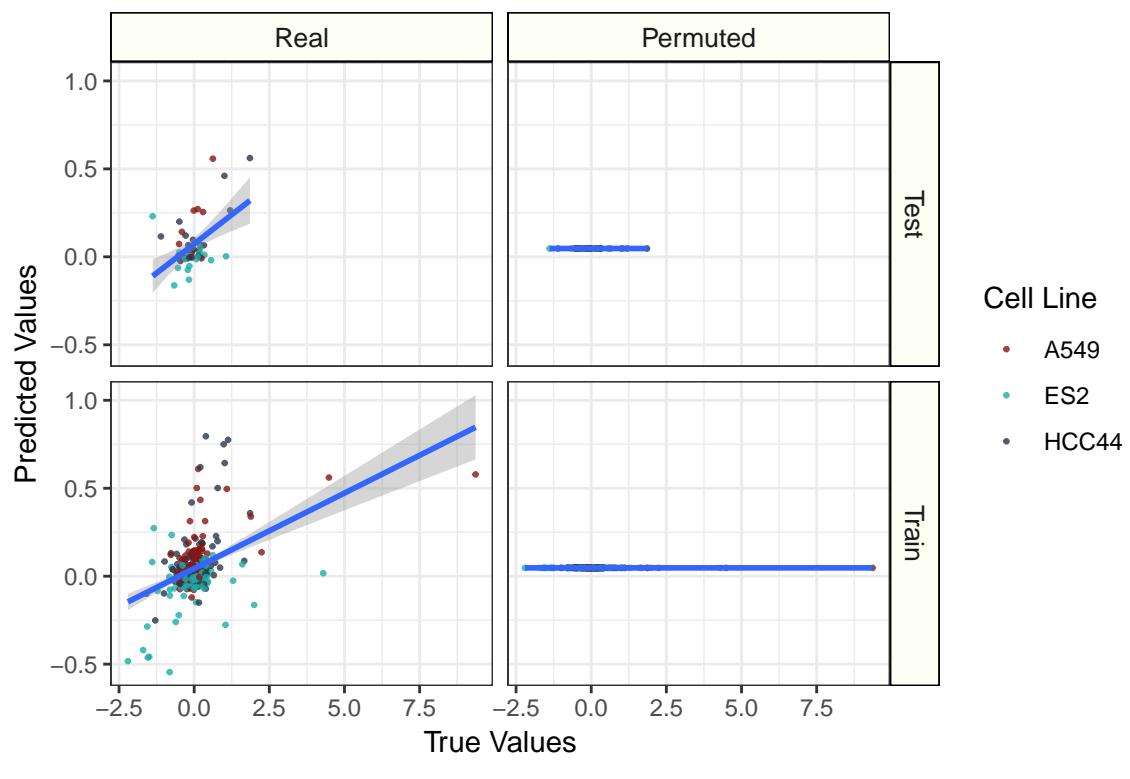


fit	shuffle	transform	r2	value
Train	Real	raw	0.89	0.09
Test	Real	raw	0.45	0.42
Train	Shuffle	raw	0.52	0.38
Test	Shuffle	raw	-0.25	0.94

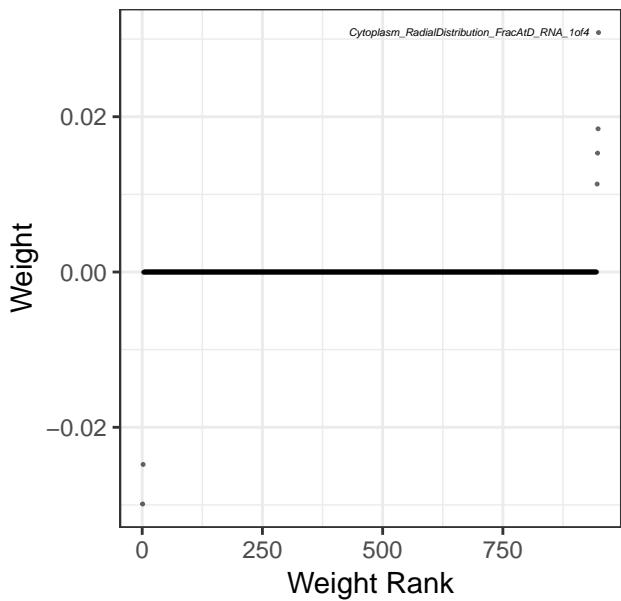


Performance: cc_g2_g1_count

Transform: raw

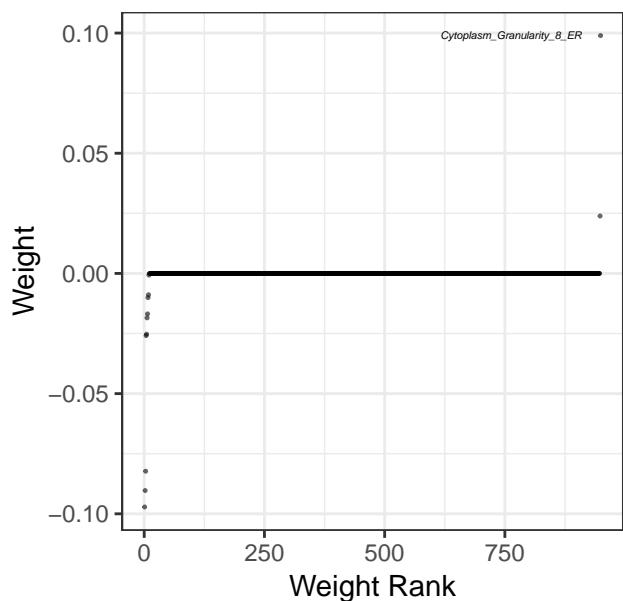
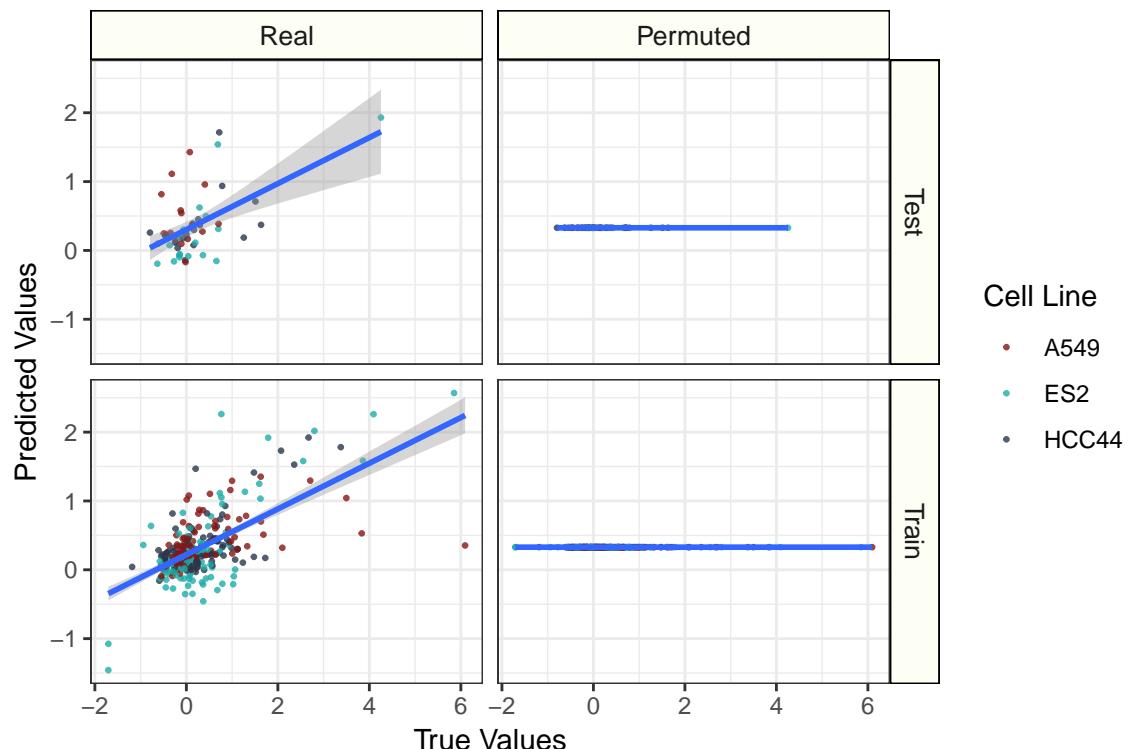


fit	shuffle	transform	r2	value
Train	Real	raw	0.13	0.59
Test	Real	raw	0.14	0.23
Train	Shuffle	raw	0.00	0.69
Test	Shuffle	raw	-0.03	0.28



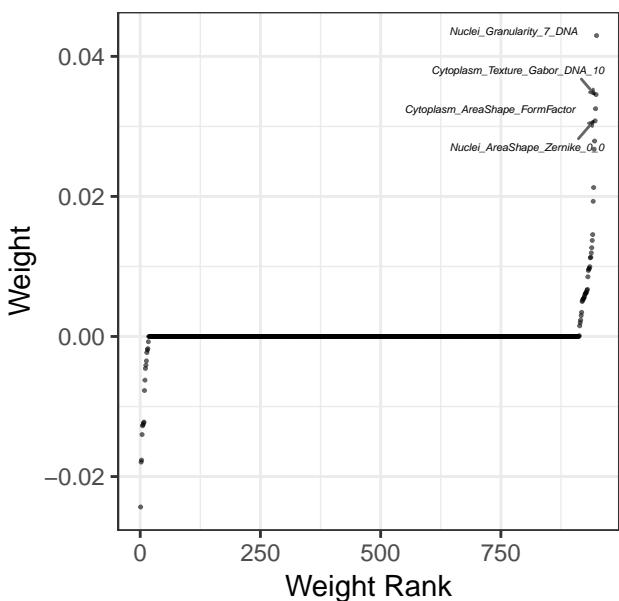
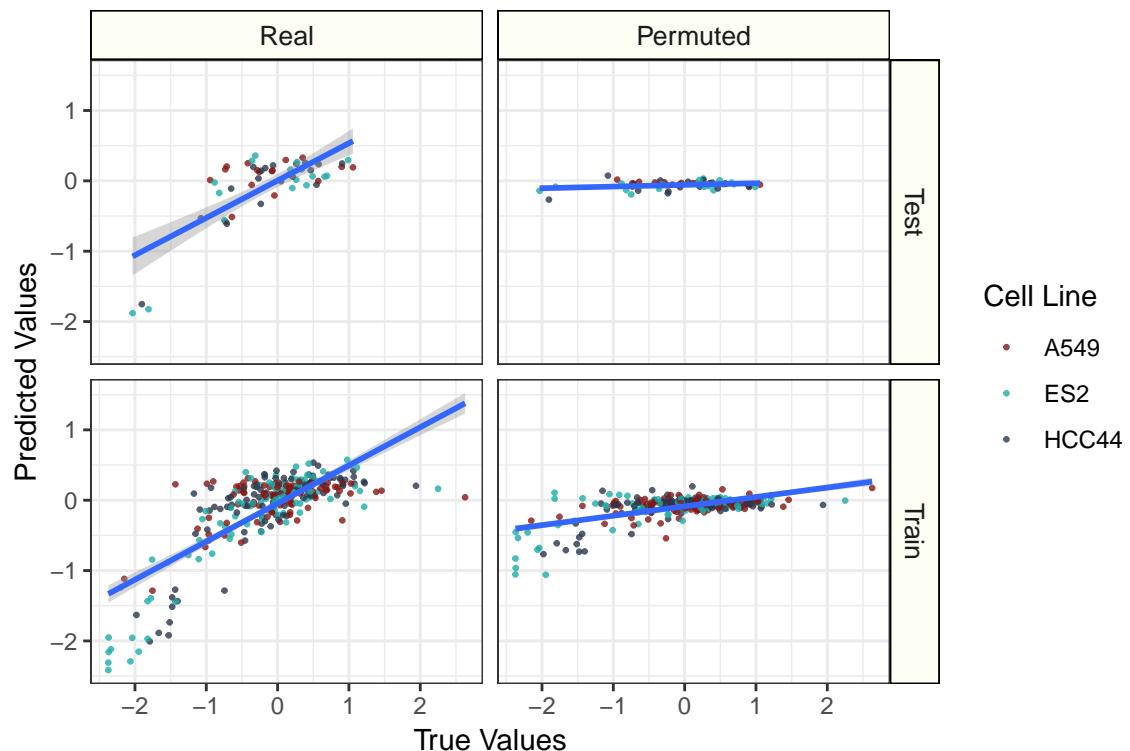
Performance: cc_g2_high_h2ax

Transform: raw



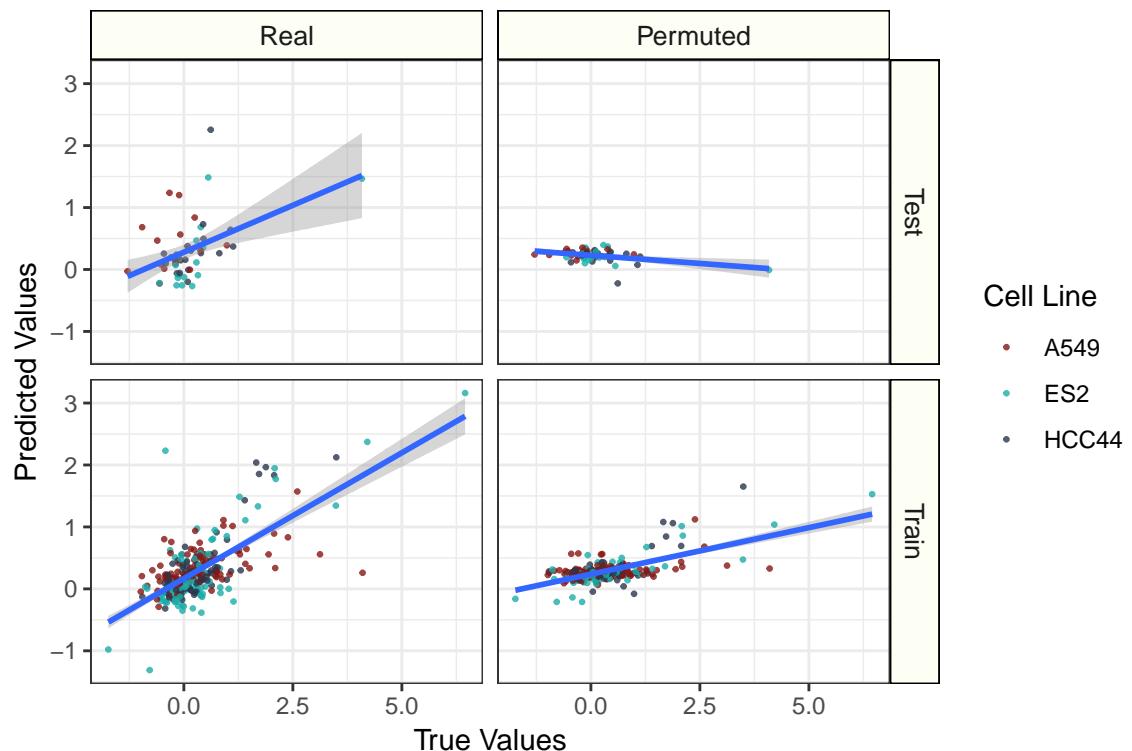
Performance: cc_g2_n_objects

Transform: raw

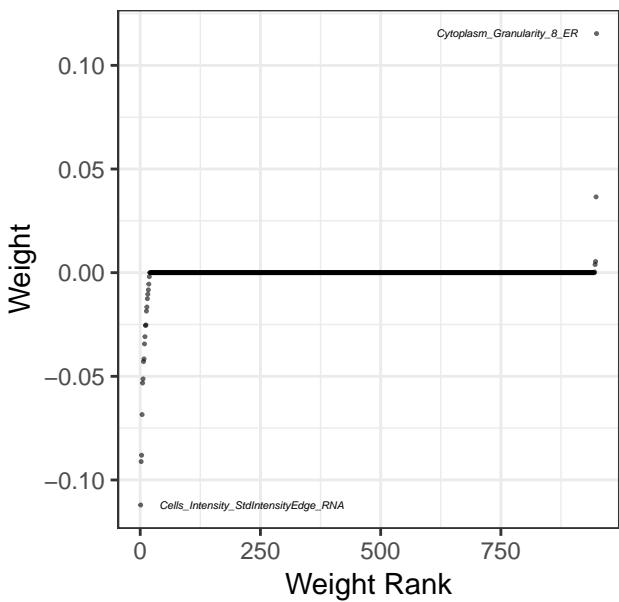


Performance: cc_g2_n_spots_h2ax_mean

Transform: raw

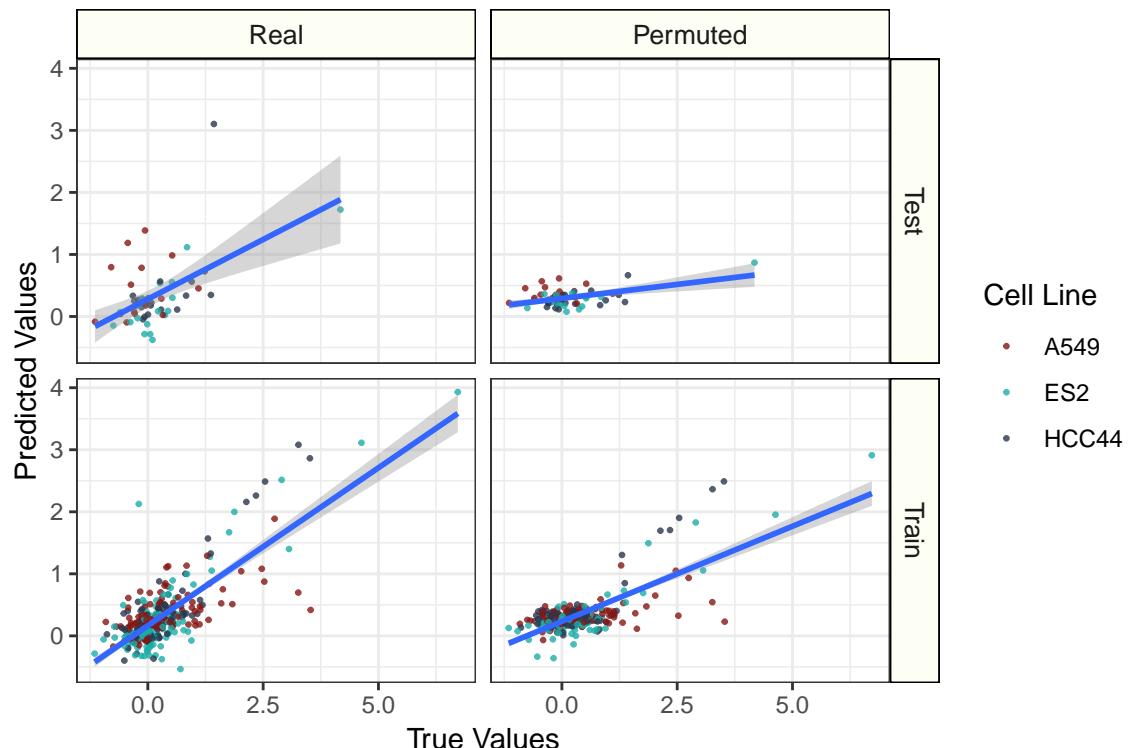


fit	shuffle	transform	r2	value
Train	Real	raw	0.48	0.36
Test	Real	raw	0.05	0.48
Train	Shuffle	raw	0.25	0.52
Test	Shuffle	raw	-0.16	0.59

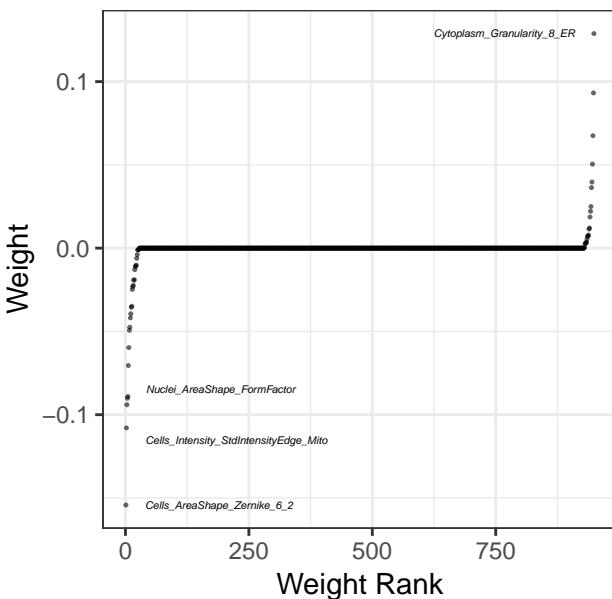


Performance: cc_g2_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

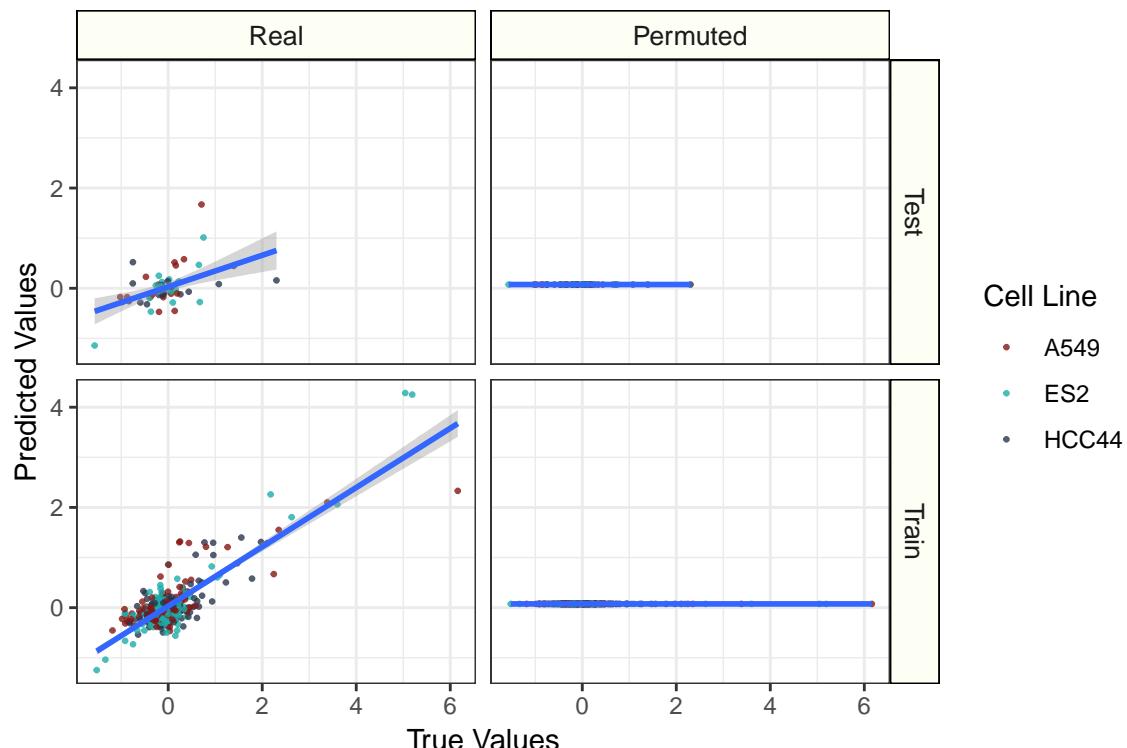


fit	shuffle	transform	r2	value
Train	Real	raw	0.59	0.30
Test	Real	raw	0.18	0.48
Train	Shuffle	raw	0.44	0.41
Test	Shuffle	raw	0.11	0.52

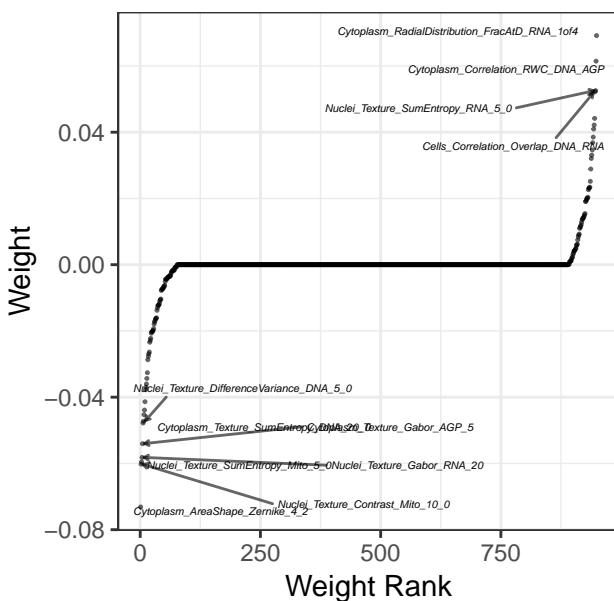


Performance: cc_g2_plus_all_m_count

Transform: raw

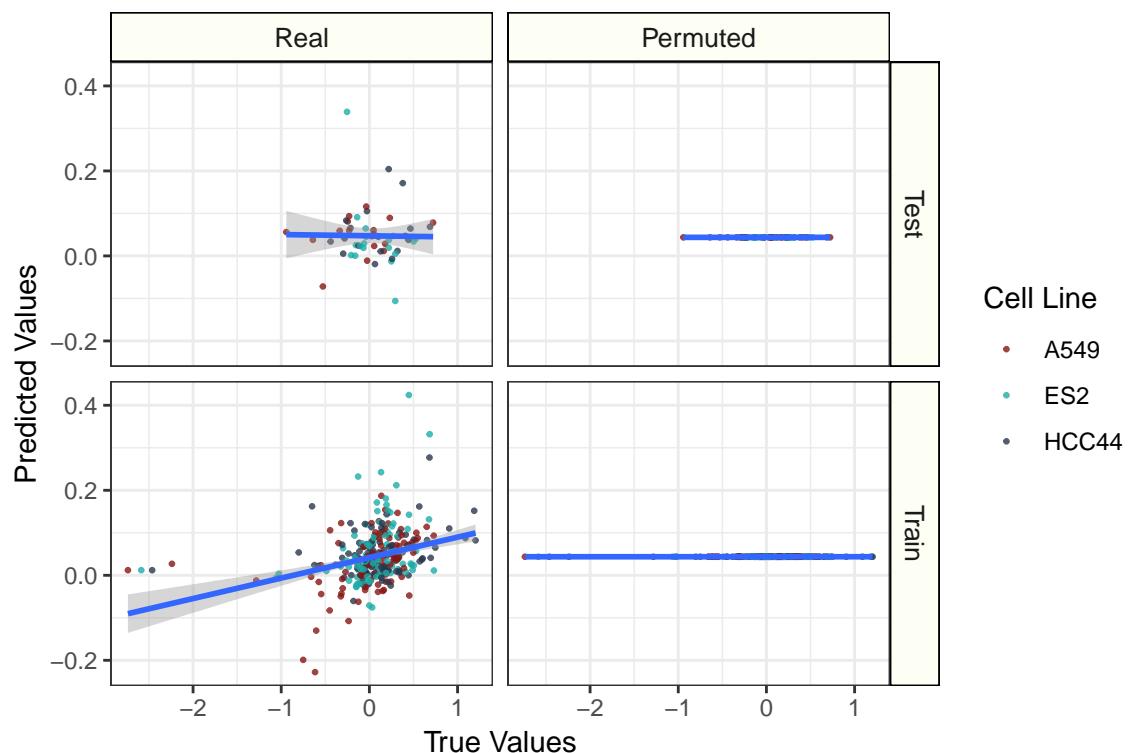


fit	shuffle	transform	r2	value
Train	Real	raw	0.68	0.21
Test	Real	raw	0.20	0.29
Train	Shuffle	raw	0.00	0.67
Test	Shuffle	raw	-0.03	0.38

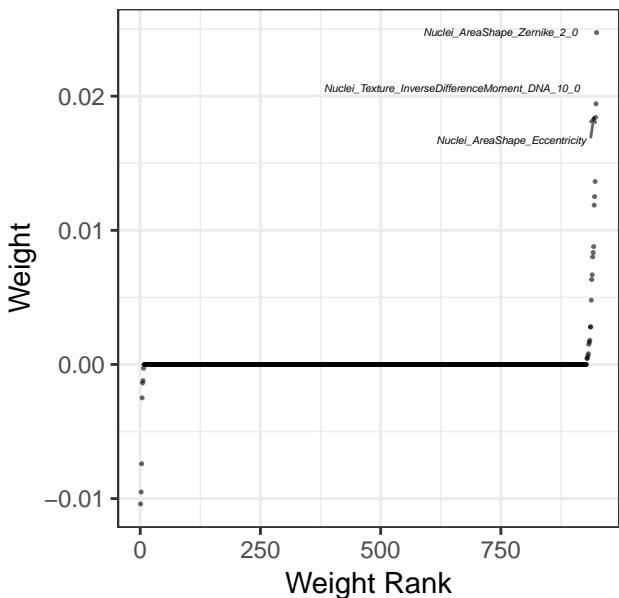


Performance: cc_infection_percentage

Transform: raw

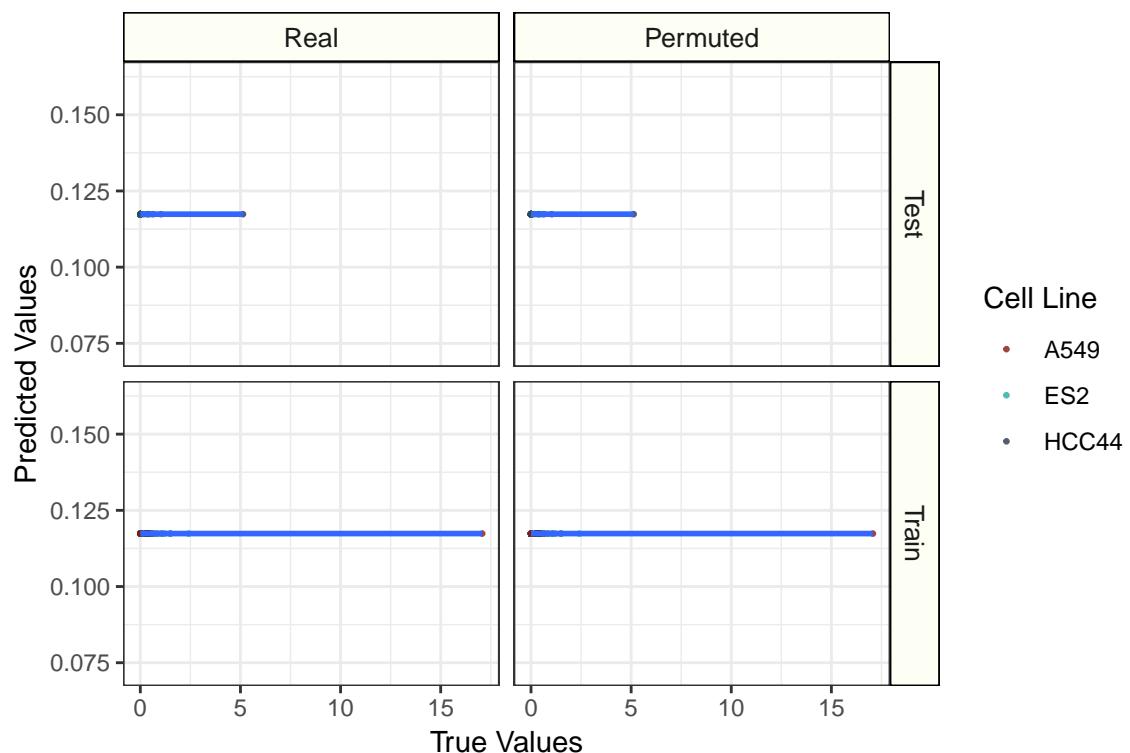


fit	shuffle	transform	r2	value
Train	Real	raw	0.07	0.18
Test	Real	raw	-0.05	0.11
Train	Shuffle	raw	0.00	0.19
Test	Shuffle	raw	0.00	0.10

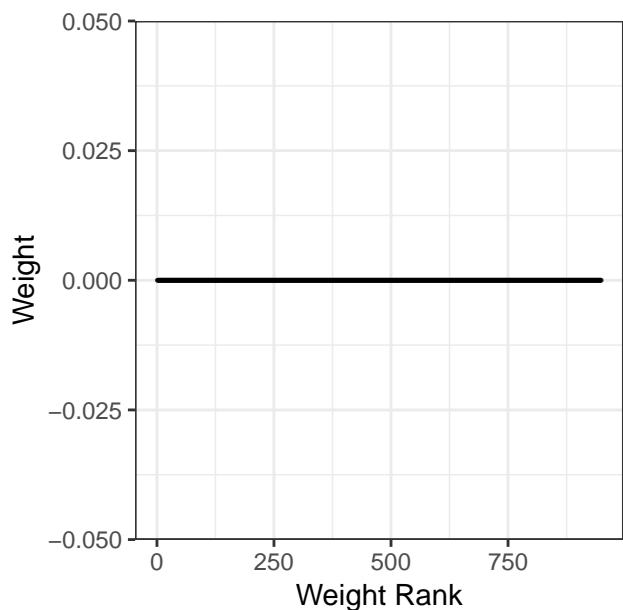


Performance: cc_late_mitosis_high_h2ax

Transform: raw

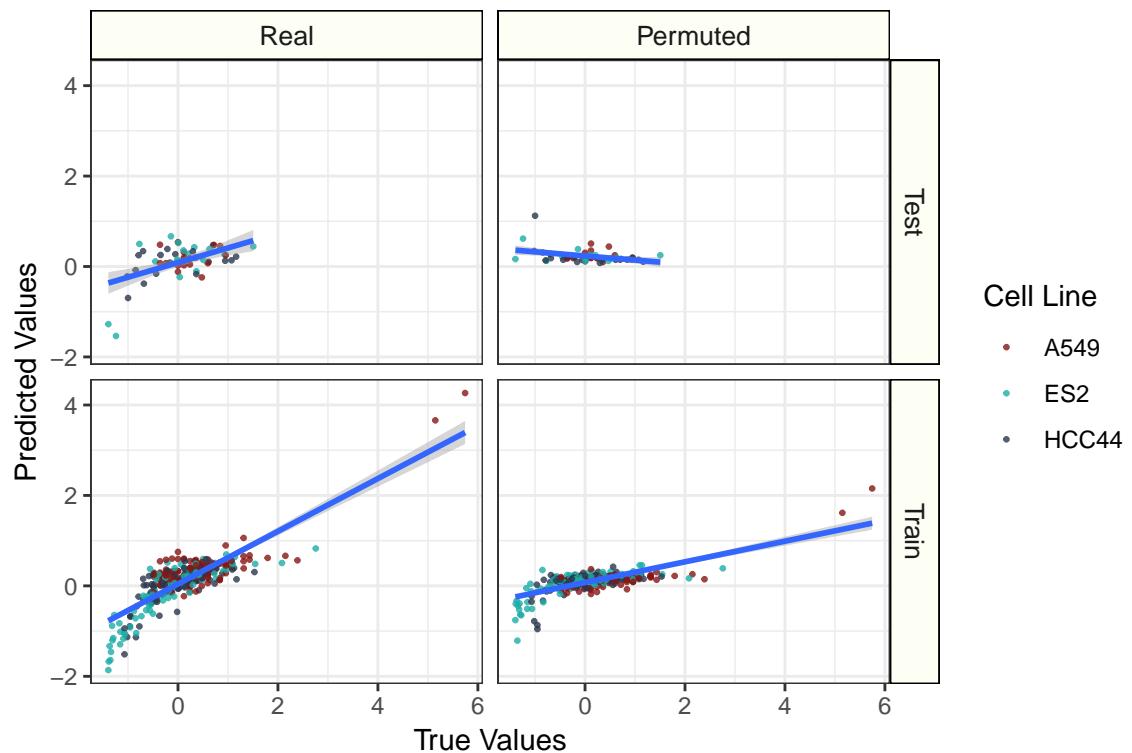


fit	shuffle	transform	r2	value
Train	Real	raw	0	1.03
Test	Real	raw	0	0.50
Train	Shuffle	raw	0	1.03
Test	Shuffle	raw	0	0.50

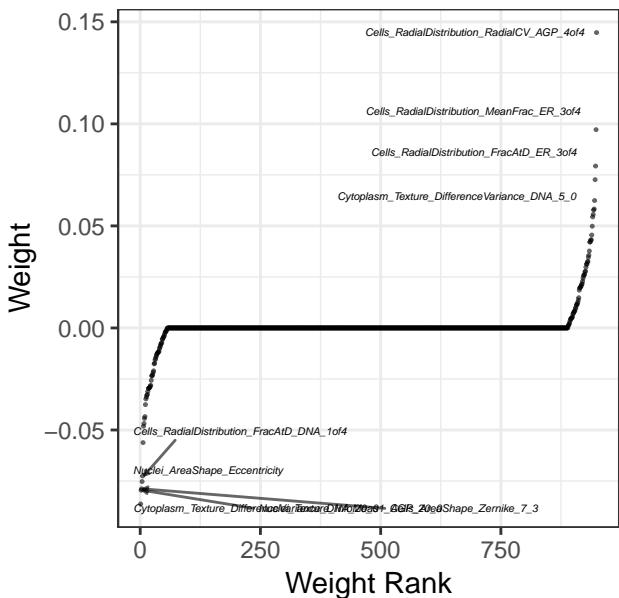


Performance: cc_late_mitosis_n_objects

Transform: raw

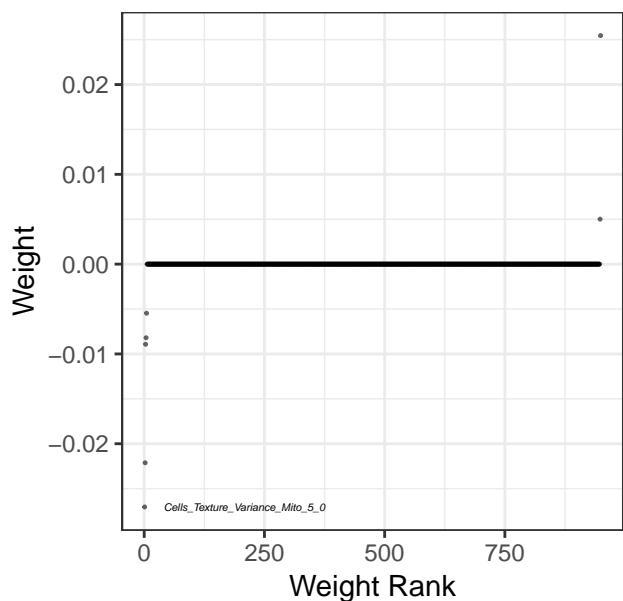
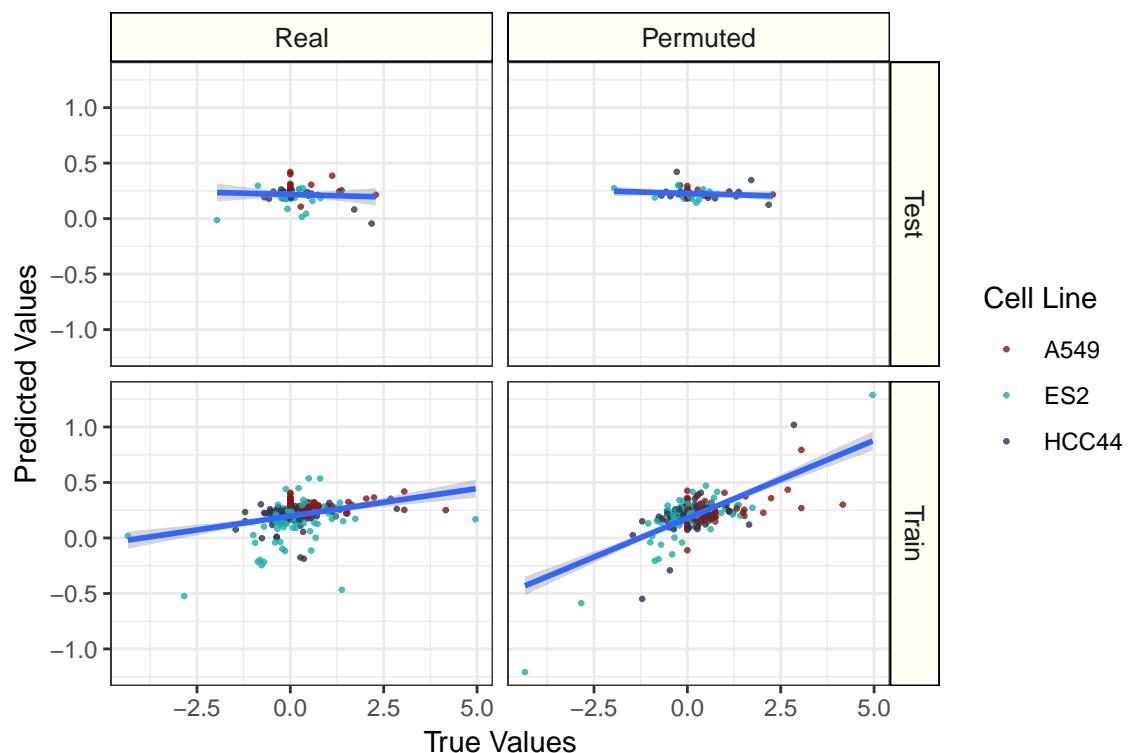


fit	shuffle	transform	r2	value
Train	Real	raw	0.67	0.21
Test	Real	raw	0.25	0.29
Train	Shuffle	raw	0.36	0.41
Test	Shuffle	raw	-0.31	0.51



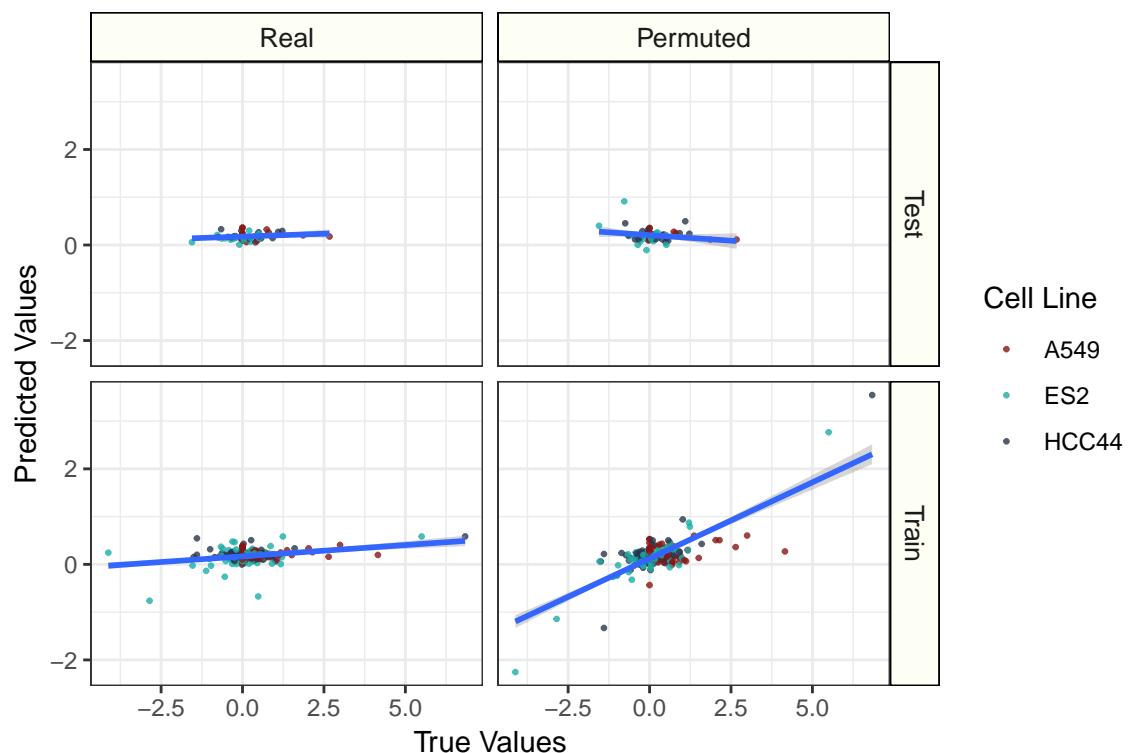
Performance: cc_late_mitosis_n_spots_h2ax_mean

Transform: raw

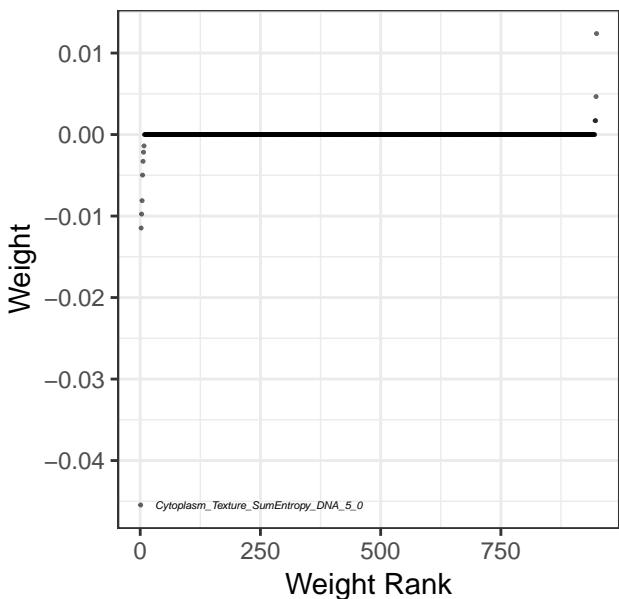


Performance: cc_late_mitosis_n_spots_h2ax_per_nucleus_area

Transform: raw

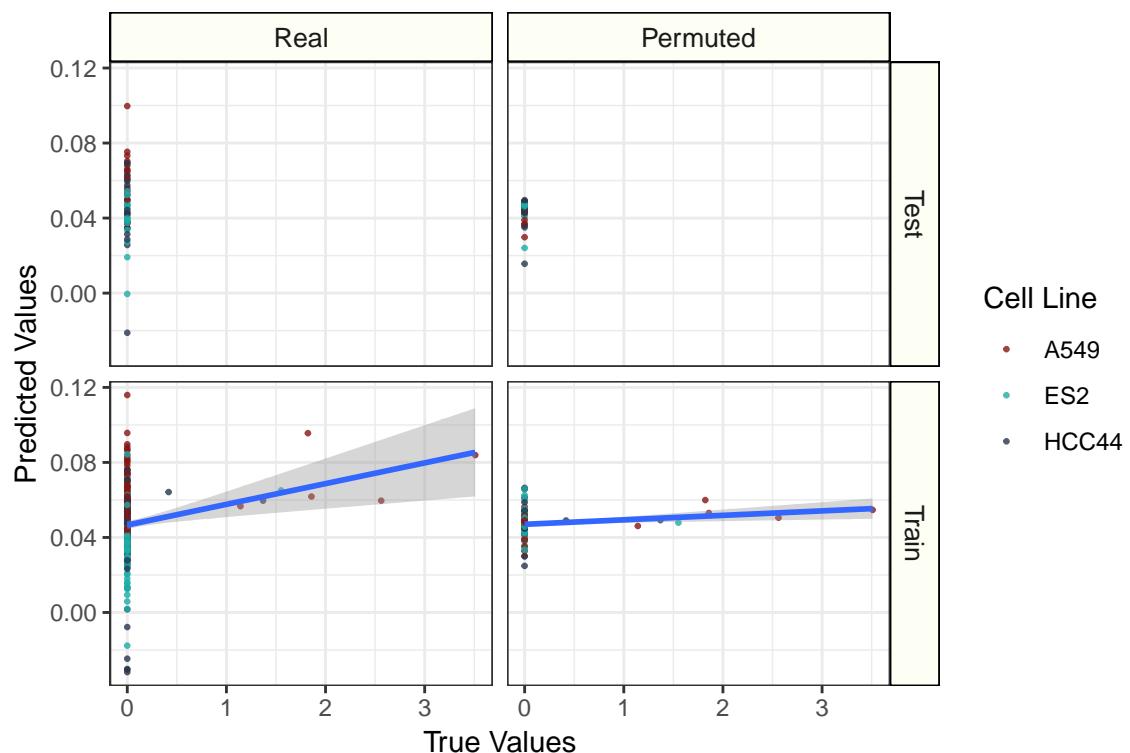


fit	shuffle	transform	r2	value
Train	Real	raw	0.07	0.62
Test	Real	raw	0.03	0.40
Train	Shuffle	raw	0.47	0.36
Test	Shuffle	raw	-0.16	0.47

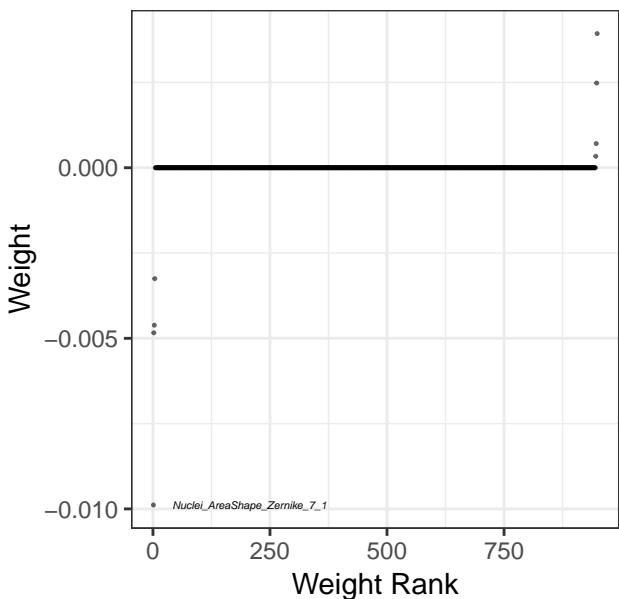


Performance: cc_mitosis_high_h2ax

Transform: raw

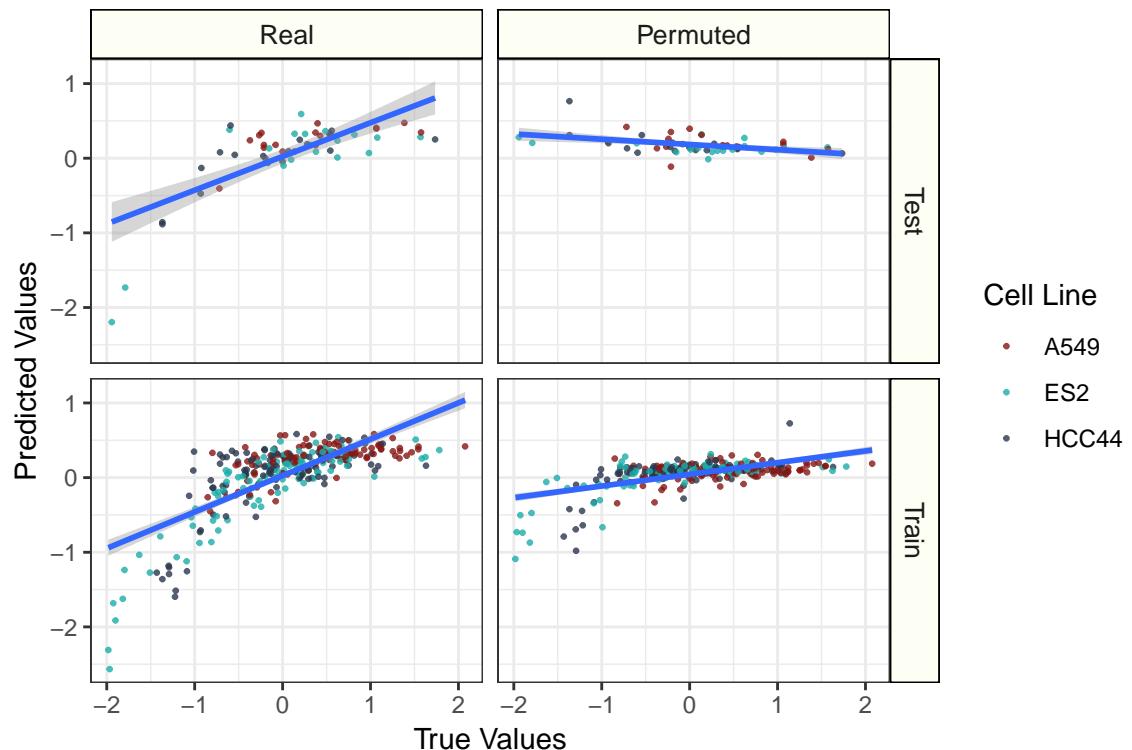


fit	shuffle	transform	r2	value
Train	Real	raw	0.02	0.1
Test	Real	raw	0.00	0.0
Train	Shuffle	raw	0.00	0.1
Test	Shuffle	raw	0.00	0.0

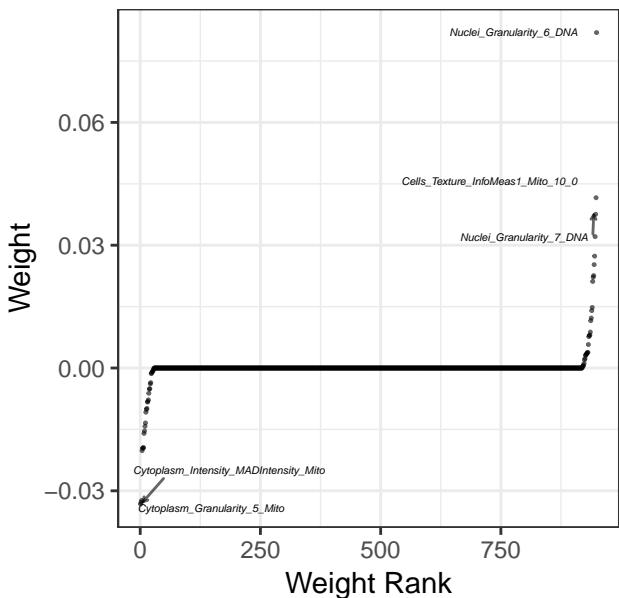


Performance: cc_mitosis_n_objects

Transform: raw

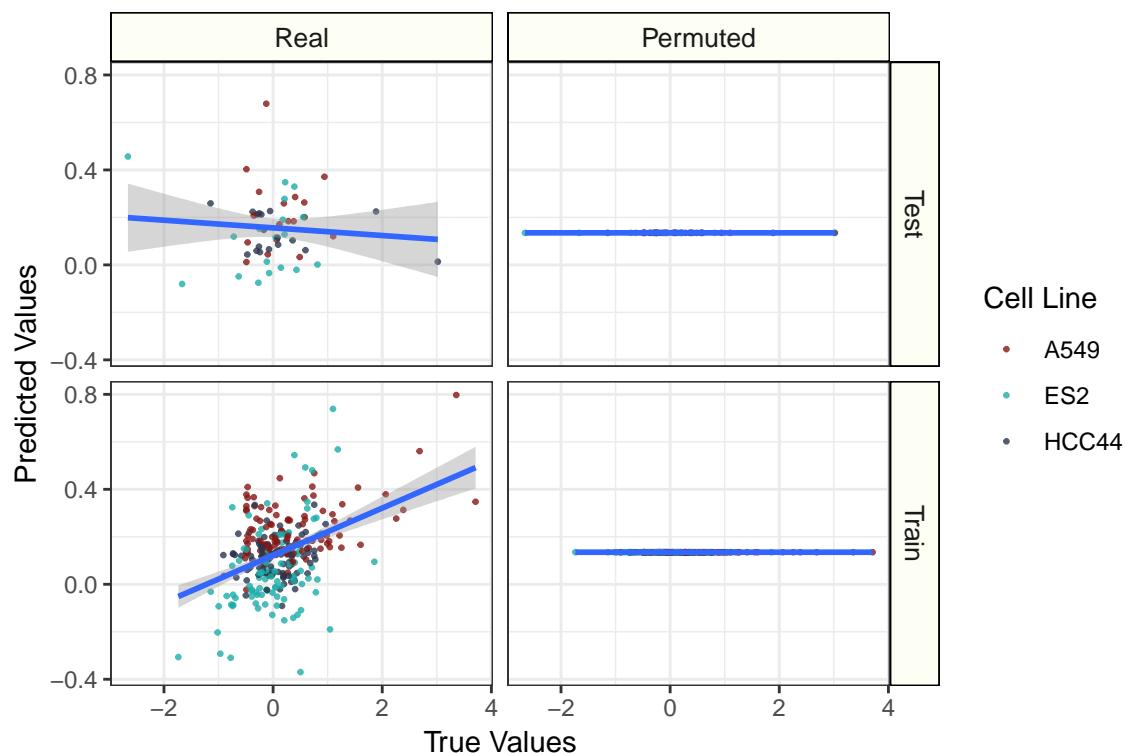


fit	shuffle	transform	r2	value
Train	Real	raw	0.54	0.24
Test	Real	raw	0.51	0.30
Train	Shuffle	raw	0.24	0.40
Test	Shuffle	raw	-0.18	0.71

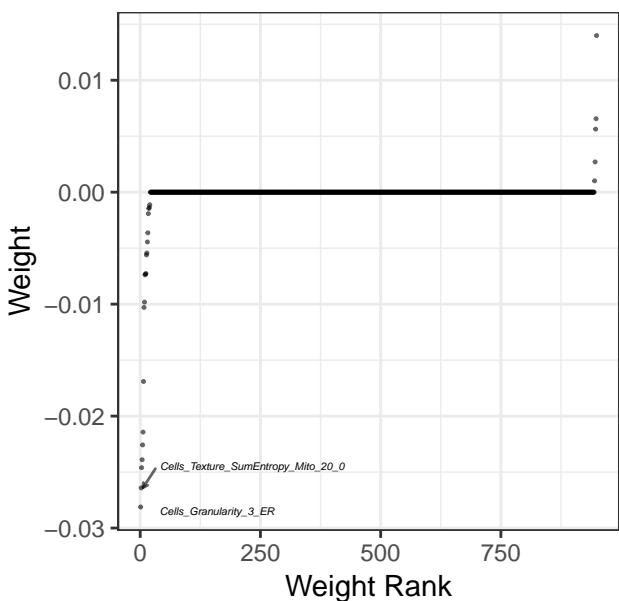


Performance: cc_mitosis_n_spots_h2ax_mean

Transform: raw

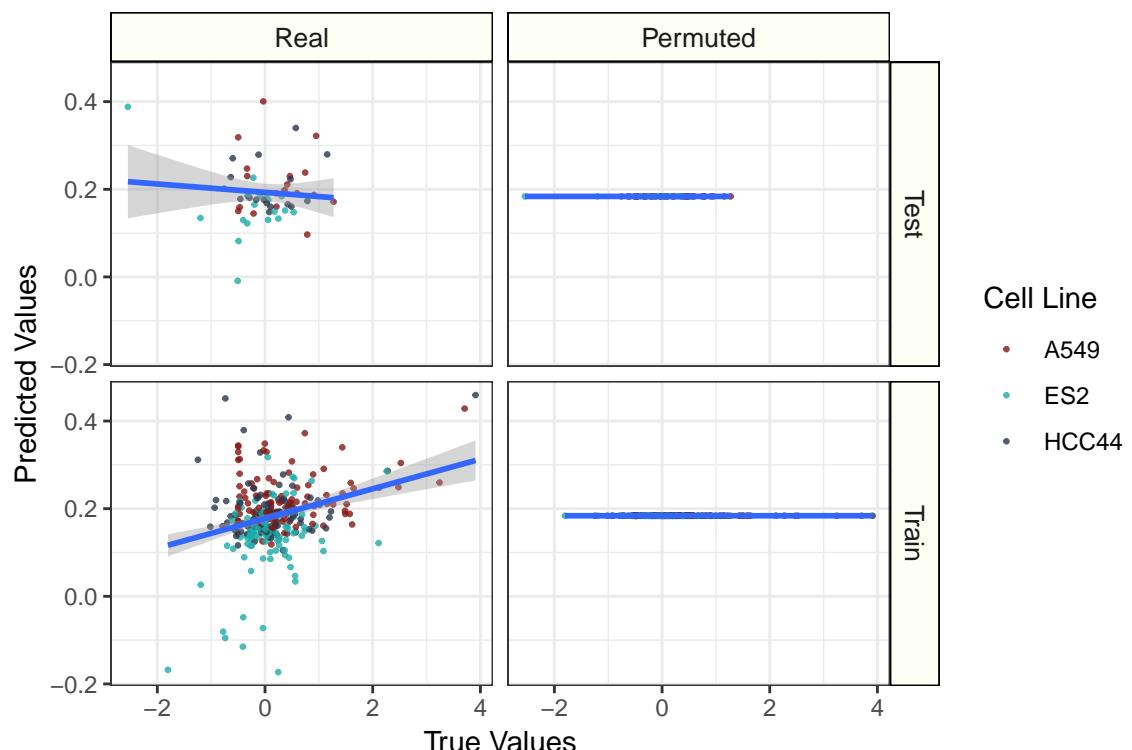


fit	shuffle	transform	r2	value
Train	Real	raw	0.14	0.35
Test	Real	raw	-0.09	0.65
Train	Shuffle	raw	0.00	0.41
Test	Shuffle	raw	-0.02	0.60

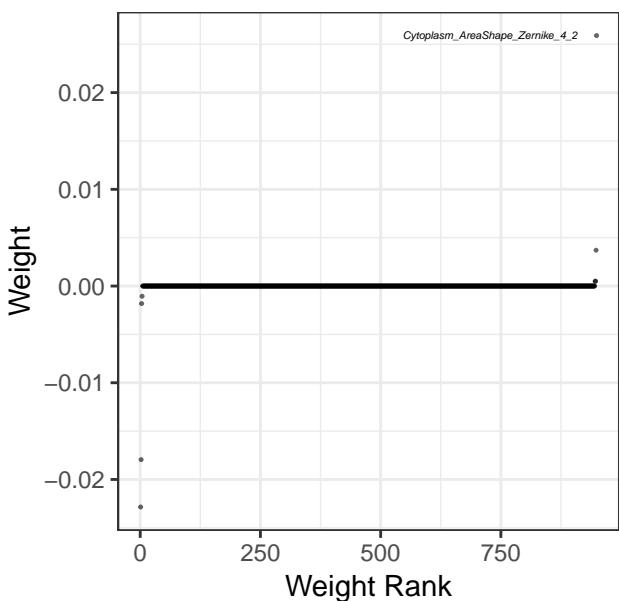


Performance: cc_mitosis_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

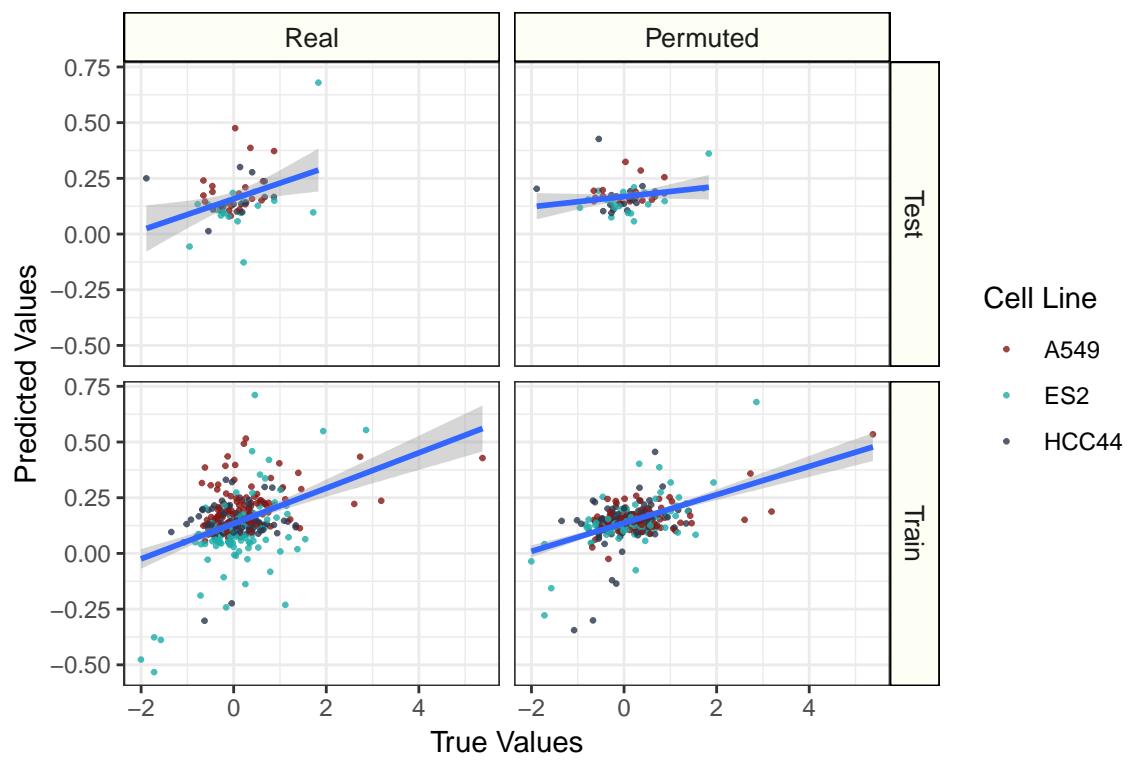


fit	shuffle	transform	r2	value
Train	Real	raw	0.06	0.46
Test	Real	raw	-0.11	0.43
Train	Shuffle	raw	0.00	0.49
Test	Shuffle	raw	-0.07	0.41

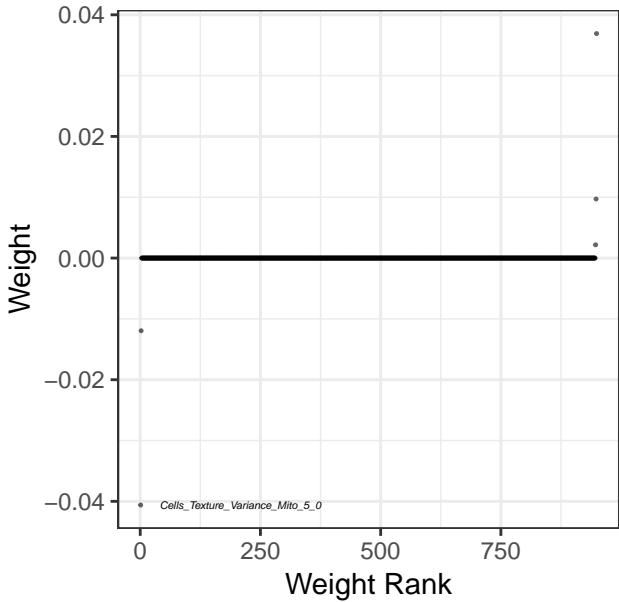


Performance: cc_polynuclear_high_h2ax

Transform: raw

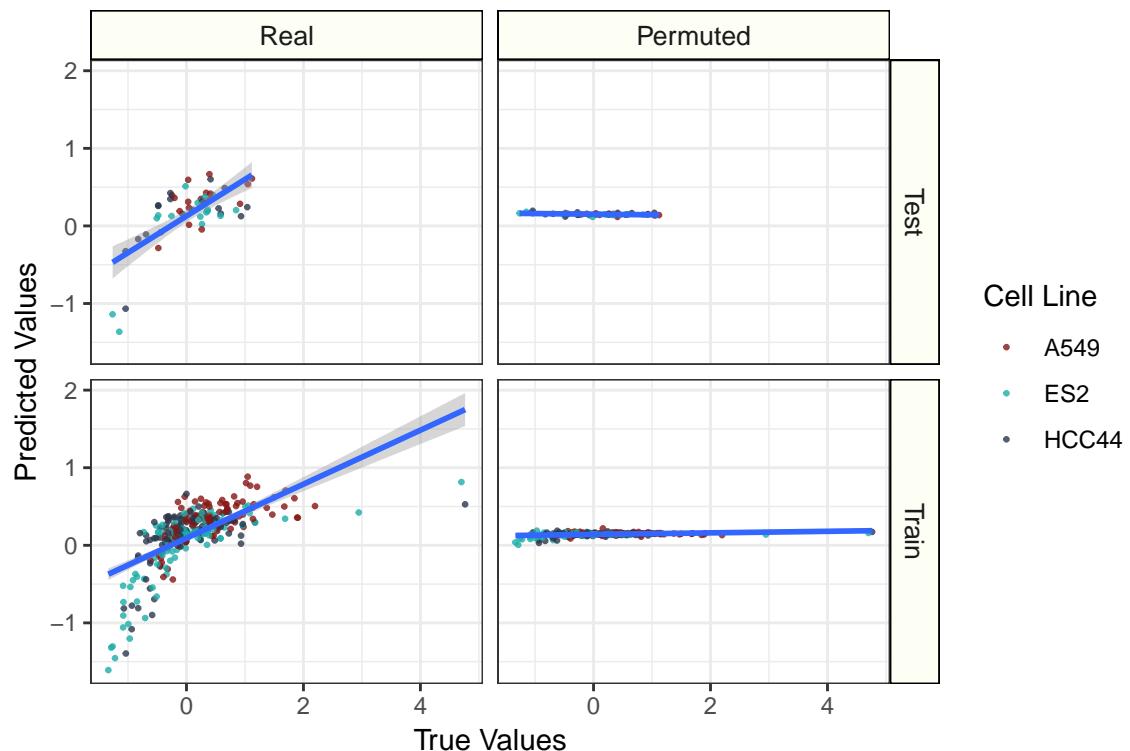


fit	shuffle	transform	r2	value
Train	Real	raw	0.12	0.44
Test	Real	raw	0.07	0.34
Train	Shuffle	raw	0.11	0.44
Test	Shuffle	raw	-0.01	0.37

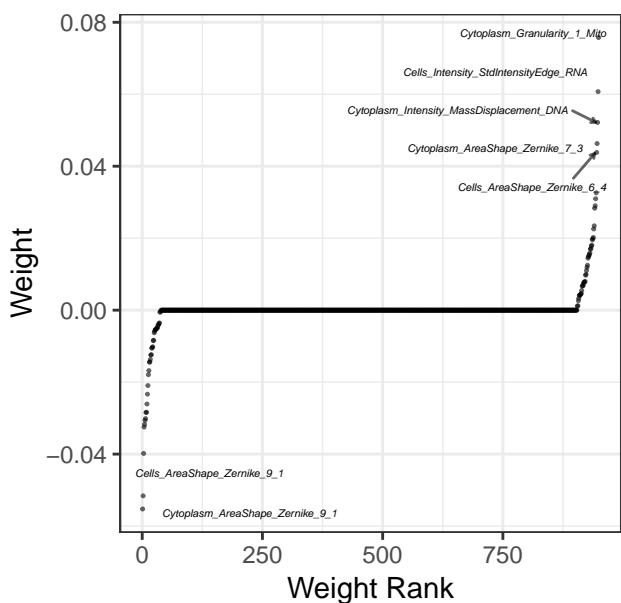


Performance: cc_polynuclear_n_objects

Transform: raw

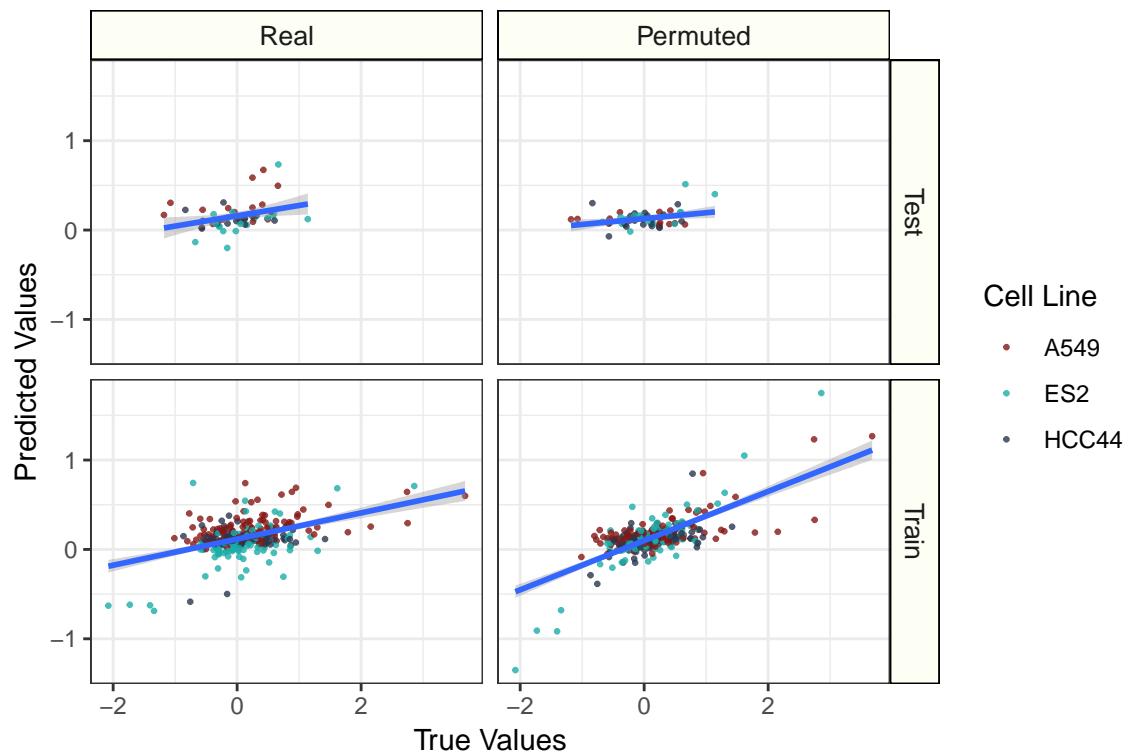


fit	shuffle	transform	r2	value
Train	Real	raw	0.41	0.32
Test	Real	raw	0.44	0.18
Train	Shuffle	raw	0.02	0.53
Test	Shuffle	raw	-0.03	0.32

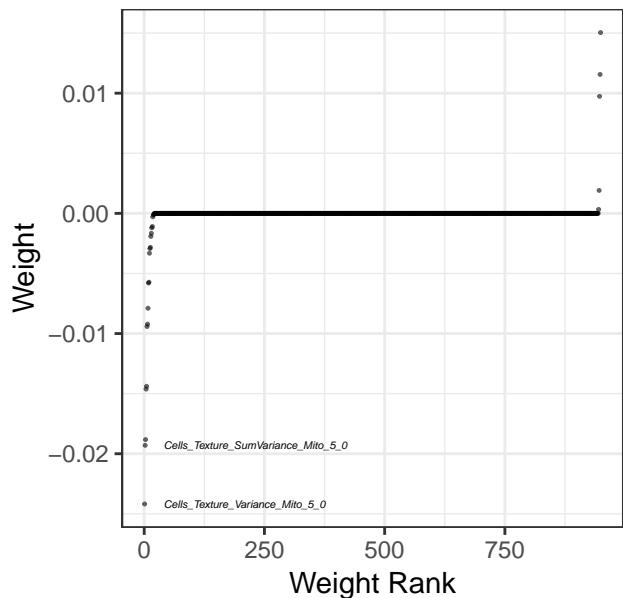


Performance: cc_polynuclear_n_spots_h2ax_mean

Transform: raw

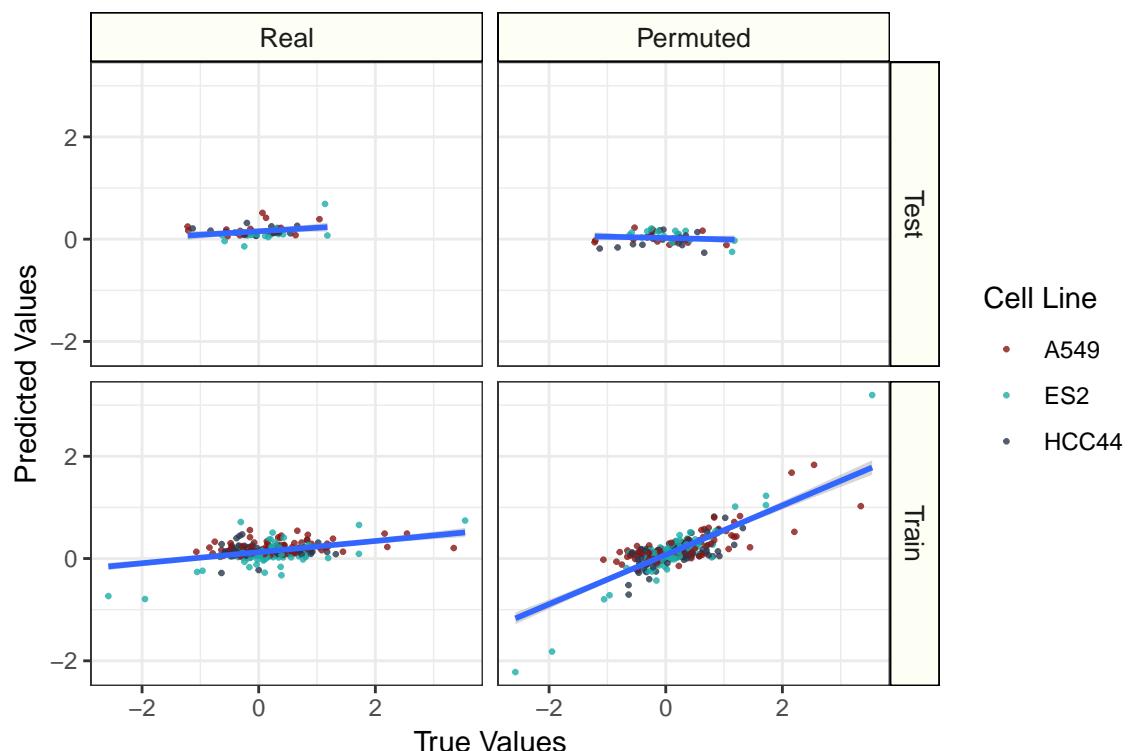


fit	shuffle	transform	r2	value
Train	Real	raw	0.20	0.30
Test	Real	raw	-0.06	0.22
Train	Shuffle	raw	0.40	0.22
Test	Shuffle	raw	-0.02	0.21

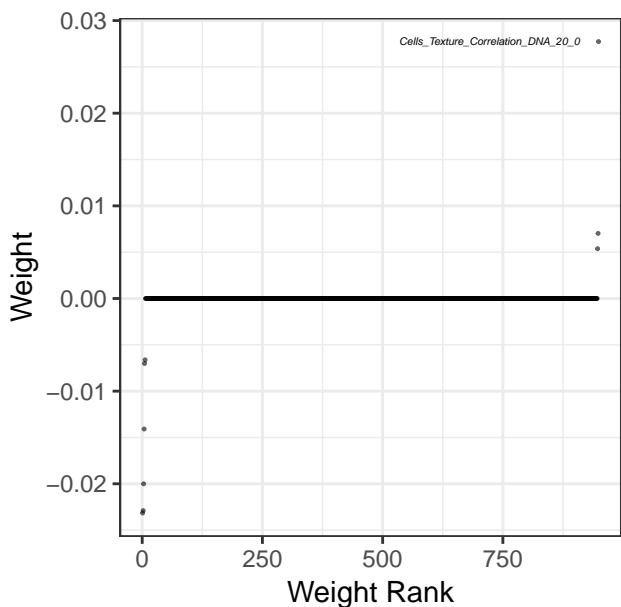


Performance: cc_polynuclear_n_spots_h2ax_per_nucleus_area

Transform: raw

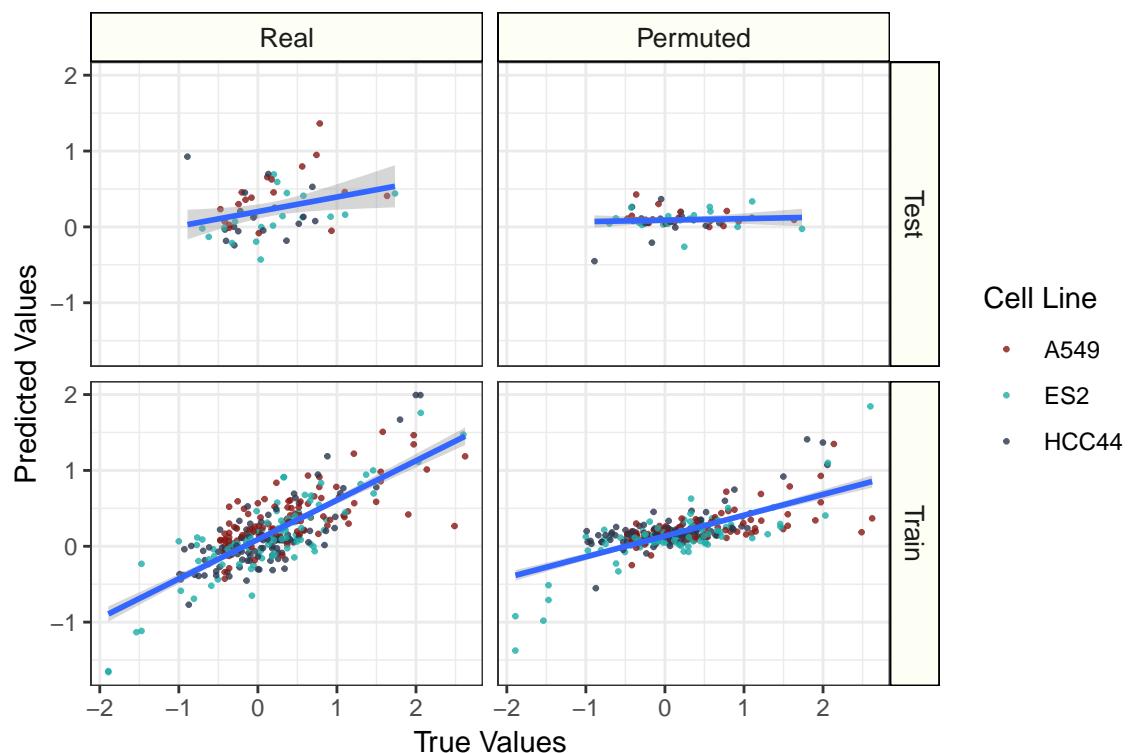


fit	shuffle	transform	r2	value
Train	Real	raw	0.15	0.32
Test	Real	raw	-0.05	0.27
Train	Shuffle	raw	0.60	0.15
Test	Shuffle	raw	-0.11	0.29

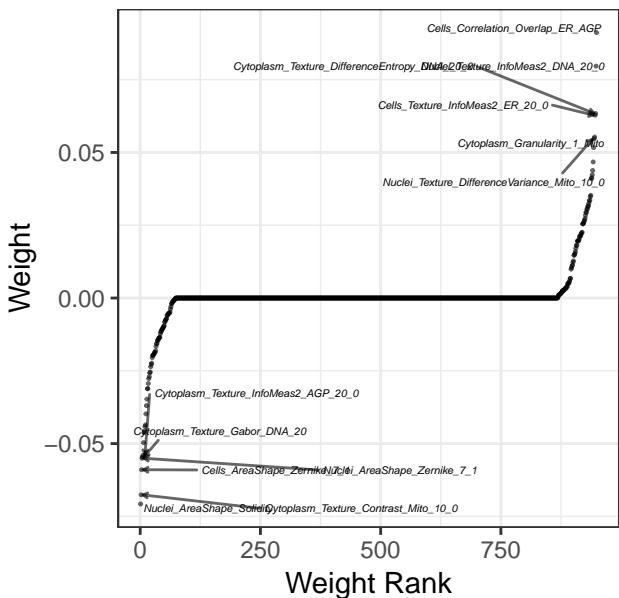


Performance: cc_polyplloid_high_h2ax

Transform: raw

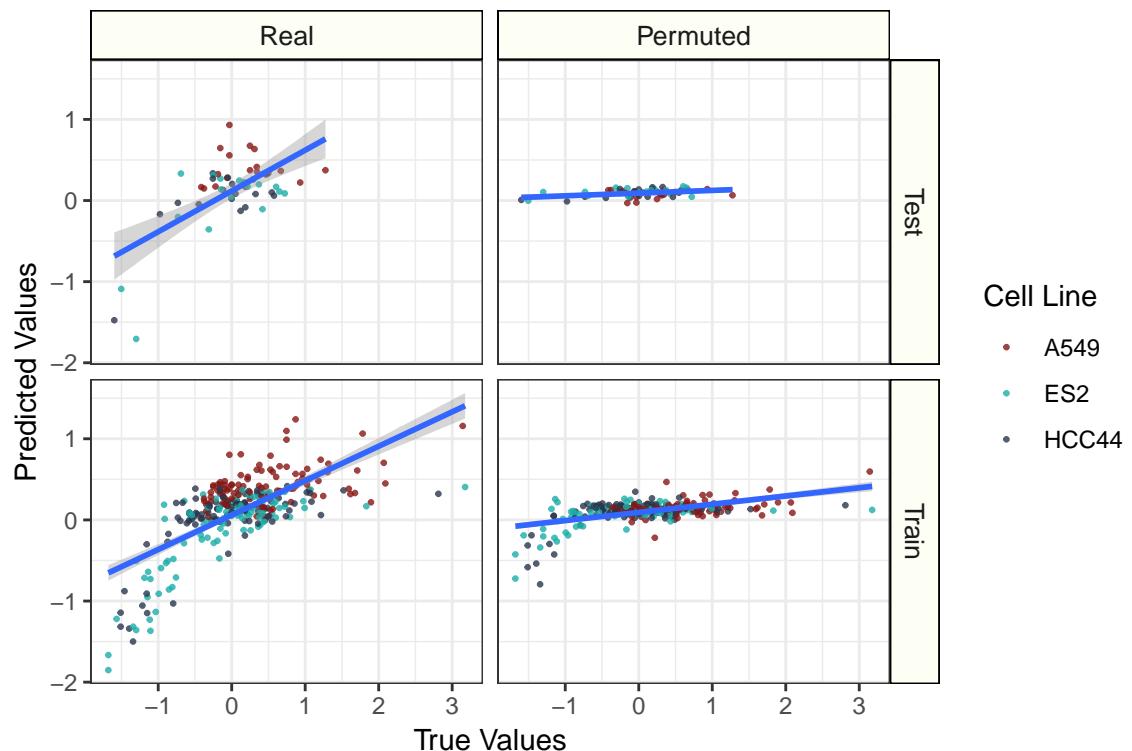


fit	shuffle	transform	r2	value
Train	Real	raw	0.61	0.19
Test	Real	raw	-0.03	0.31
Train	Shuffle	raw	0.39	0.29
Test	Shuffle	raw	-0.03	0.32

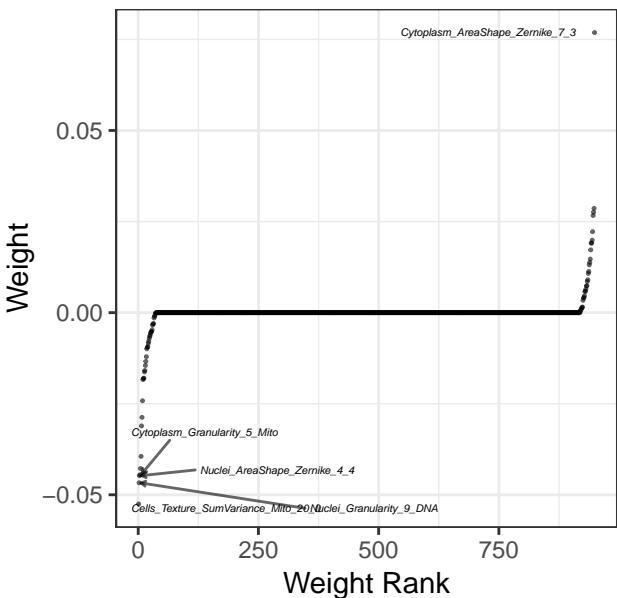


Performance: cc_polyplloid_n_objects

Transform: raw

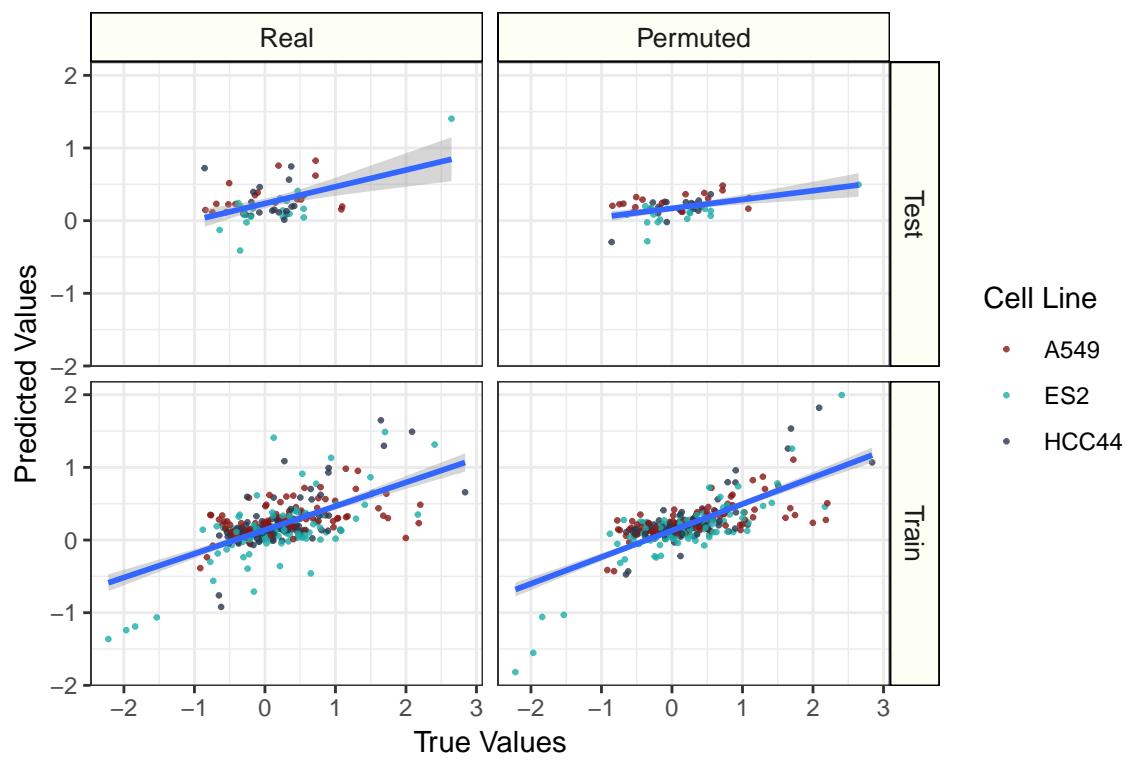


fit	shuffle	transform	r2	value
Train	Real	raw	0.48	0.29
Test	Real	raw	0.33	0.21
Train	Shuffle	raw	0.17	0.46
Test	Shuffle	raw	0.03	0.30

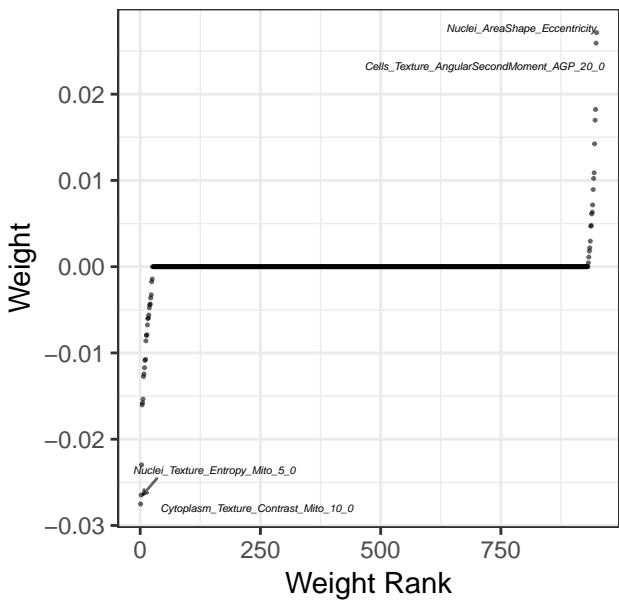


Performance: cc_polyplloid_n_spots_h2ax_mean

Transform: raw

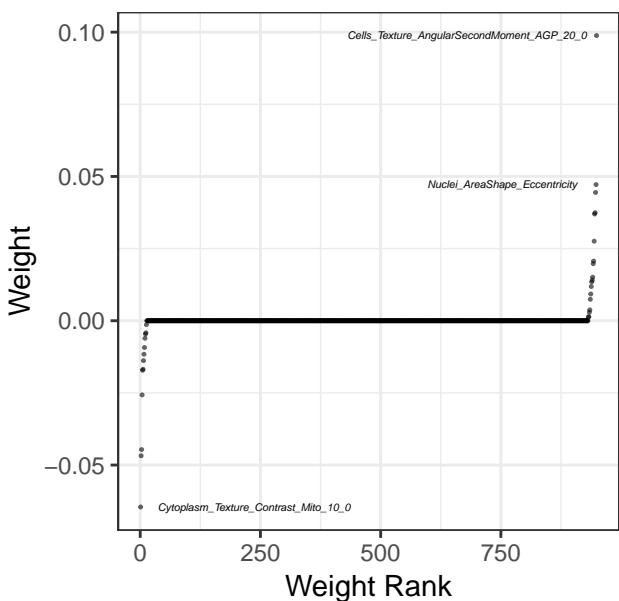
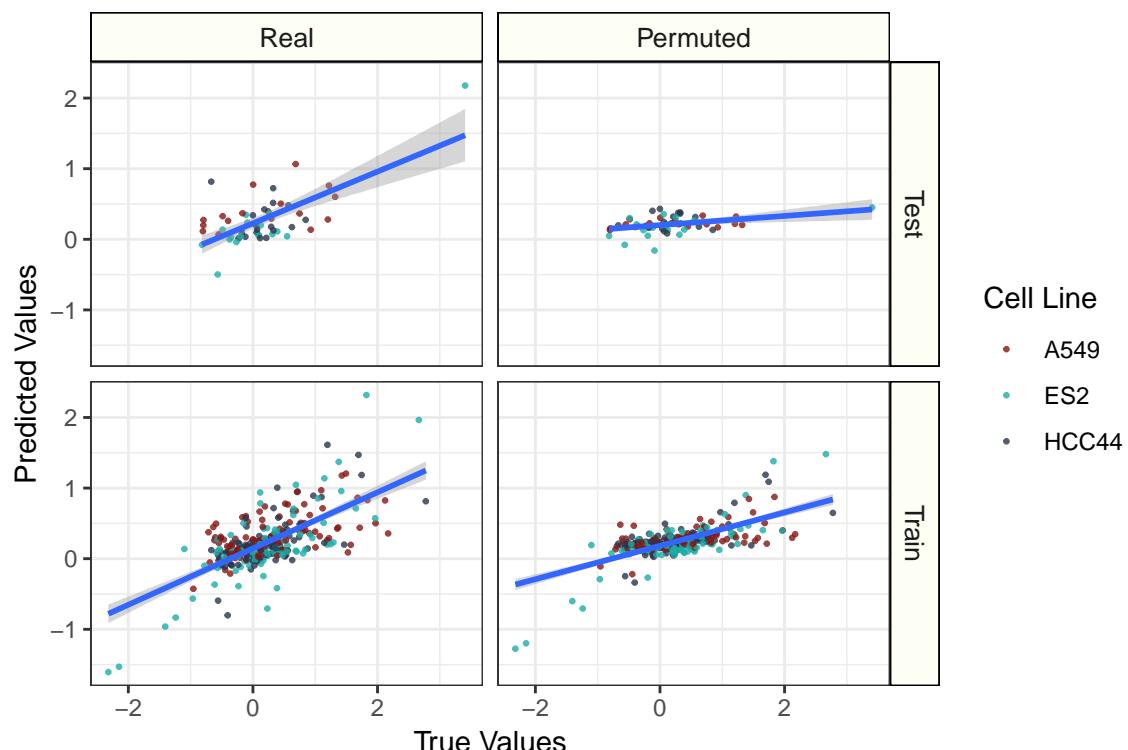


fit	shuffle	transform	r2	value
Train	Real	raw	0.38	0.28
Test	Real	raw	0.14	0.28
Train	Shuffle	raw	0.49	0.23
Test	Shuffle	raw	0.15	0.28



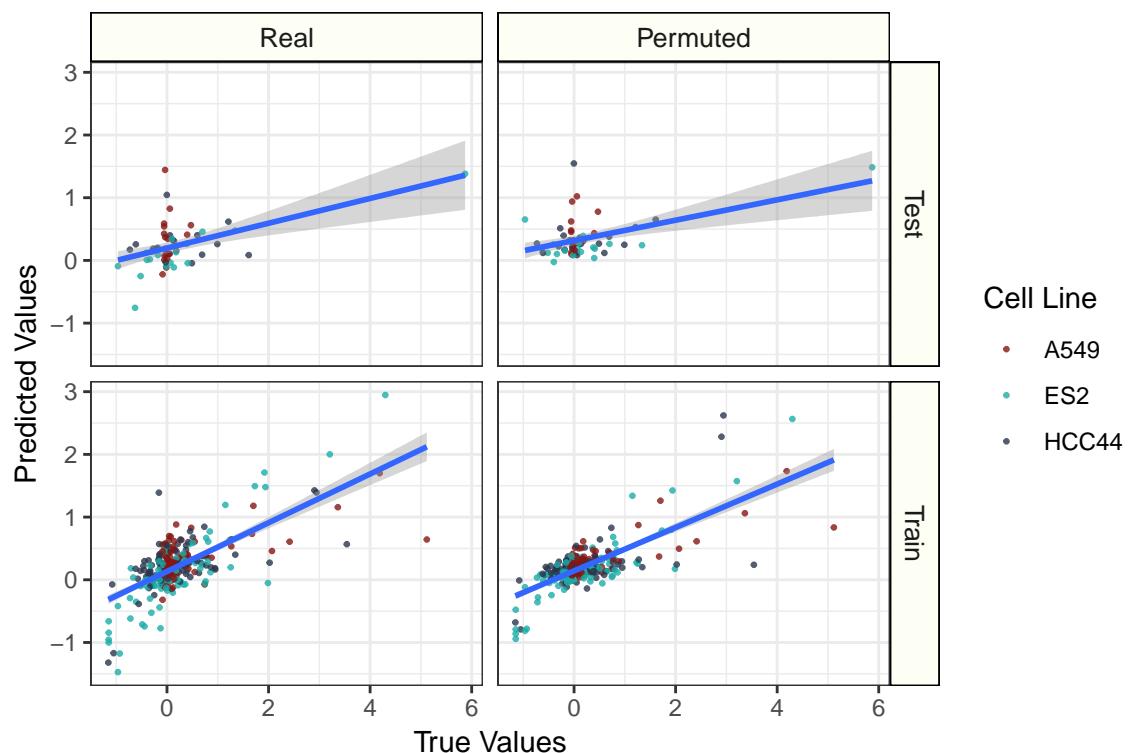
Performance: cc_polyplloid_n_spots_h2ax_per_nucleus_area_m

Transform: raw

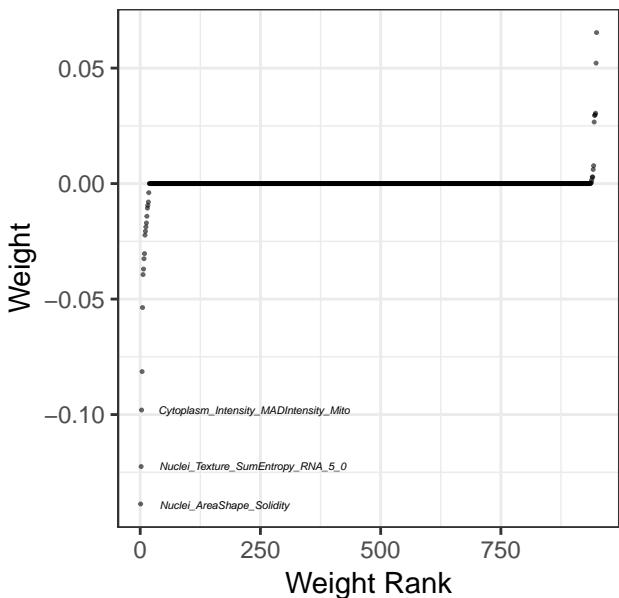


Performance: cc_s_high_h2ax

Transform: raw

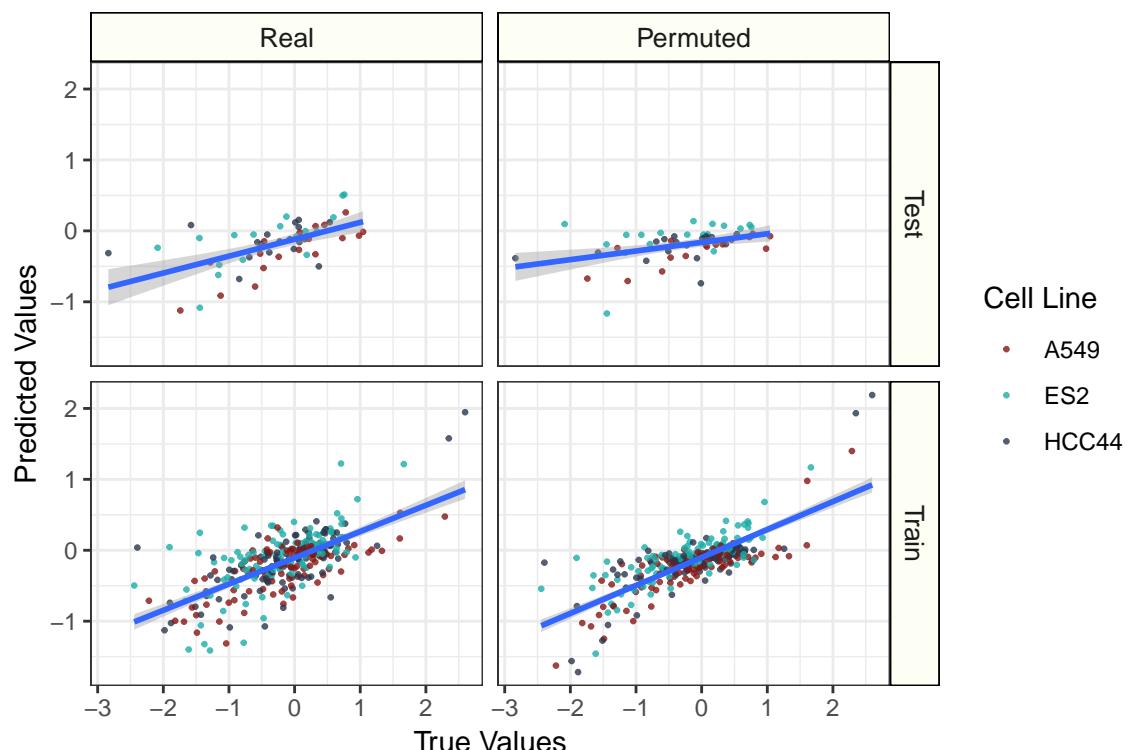


fit	shuffle	transform	r2	value
Train	Real	raw	0.46	0.33
Test	Real	raw	0.24	0.64
Train	Shuffle	raw	0.48	0.32
Test	Shuffle	raw	0.18	0.68

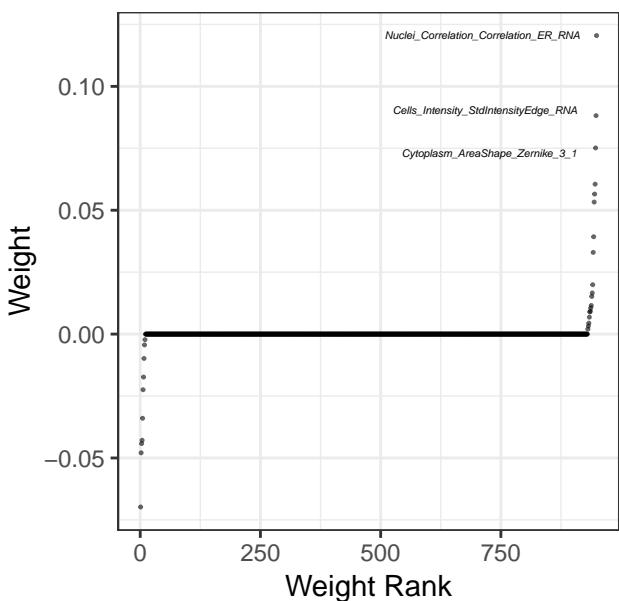


Performance: cc_s_intensity_nucleus_area_mean

Transform: raw

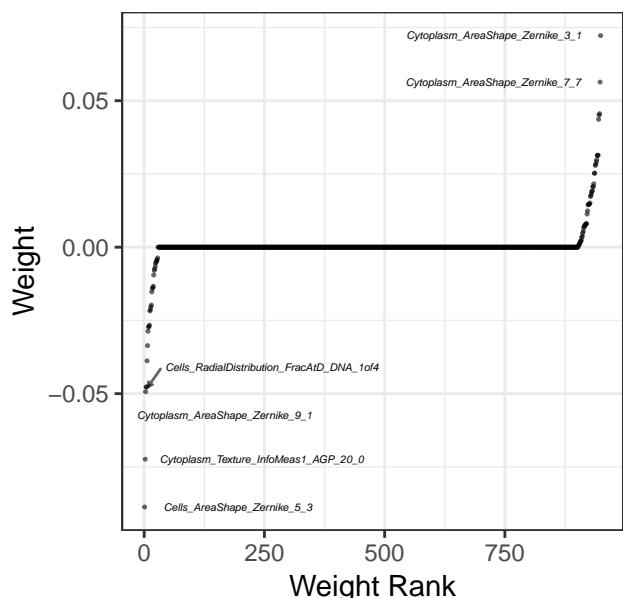
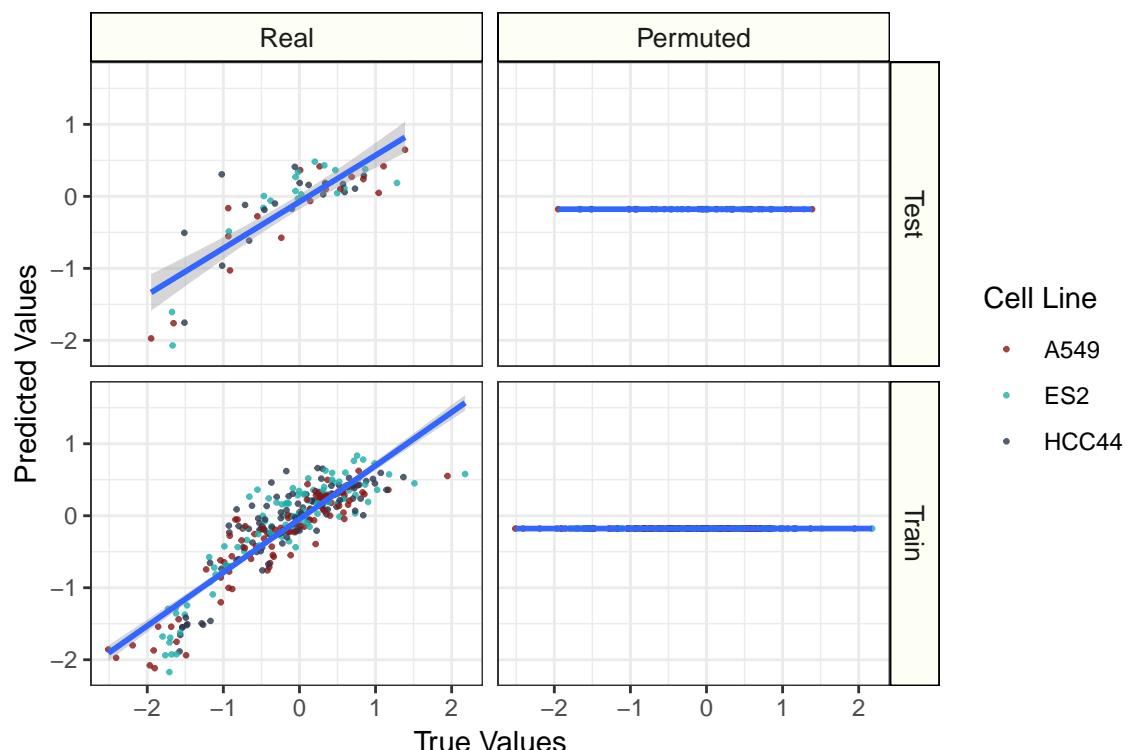


fit	shuffle	transform	r2	value
Train	Real	raw	0.44	0.32
Test	Real	raw	0.29	0.45
Train	Shuffle	raw	0.53	0.27
Test	Shuffle	raw	0.15	0.54



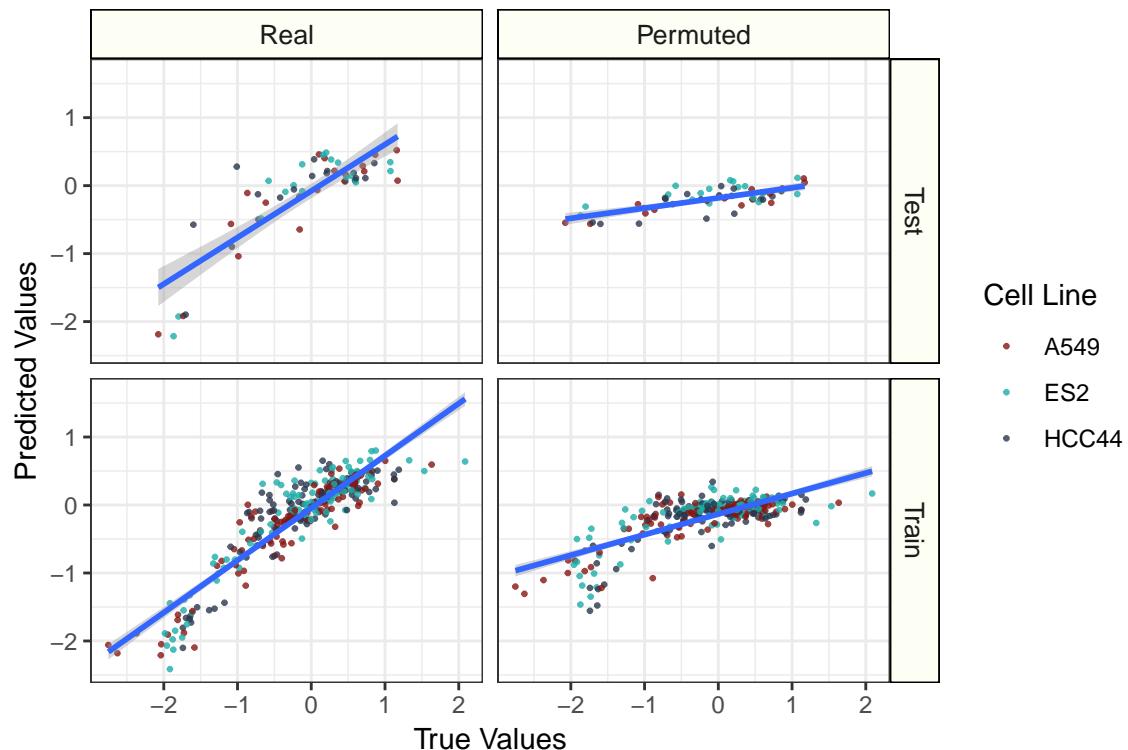
Performance: cc_s_intensity_nucleus_area_sum

Transform: raw

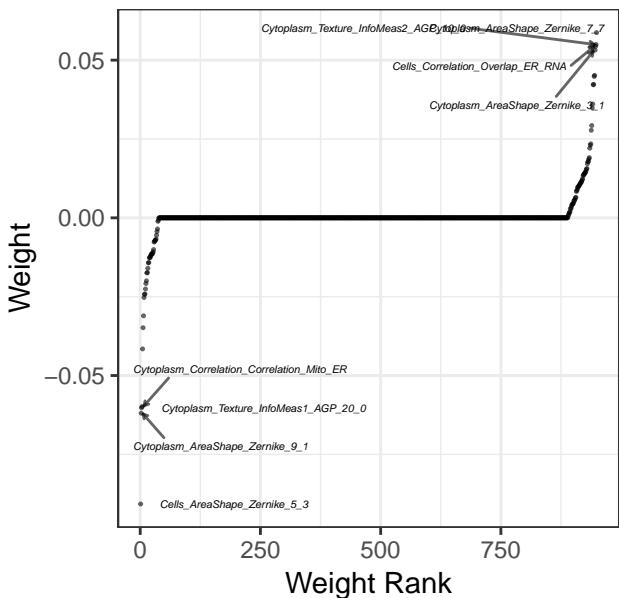


Performance: cc_s_n_objects

Transform: raw

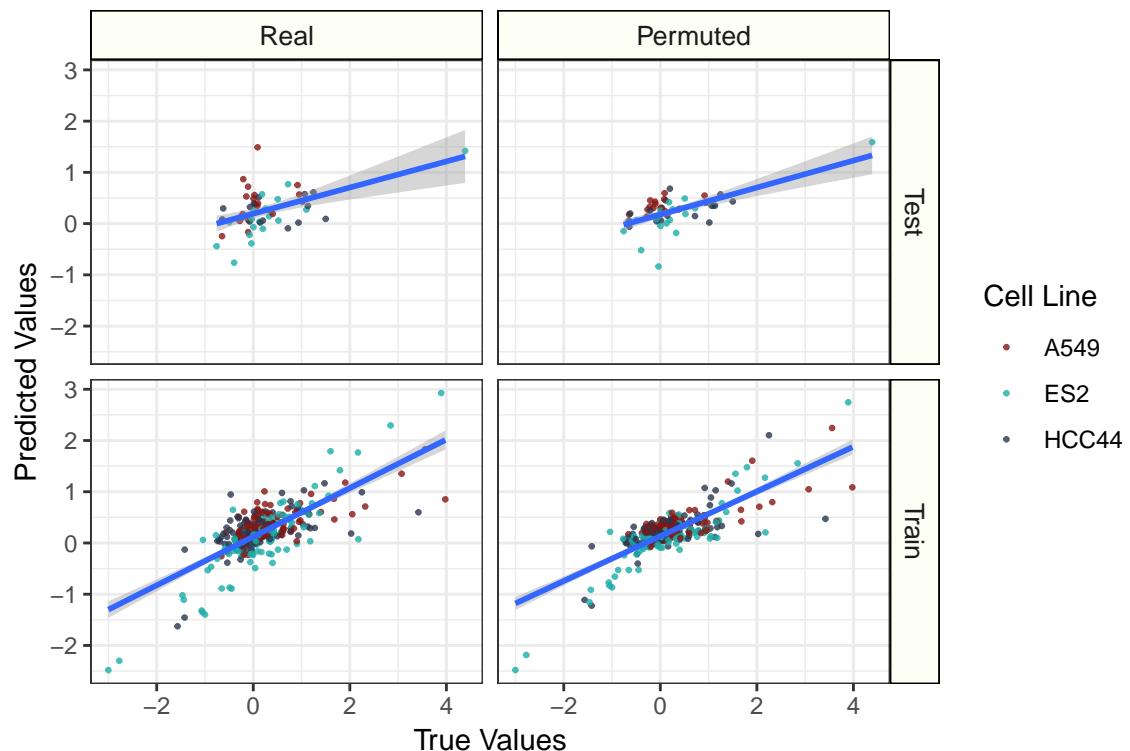


fit	shuffle	transform	r2	value
Train	Real	raw	0.81	0.12
Test	Real	raw	0.69	0.22
Train	Shuffle	raw	0.44	0.37
Test	Shuffle	raw	0.24	0.53

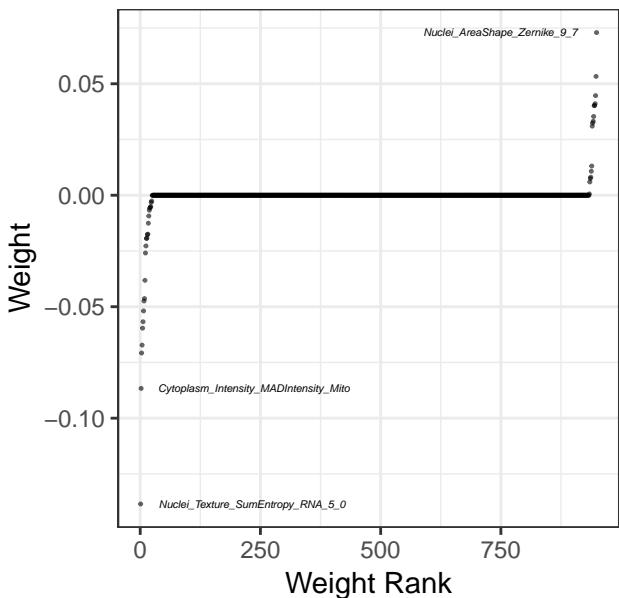


Performance: cc_s_n_spots_h2ax_mean

Transform: raw

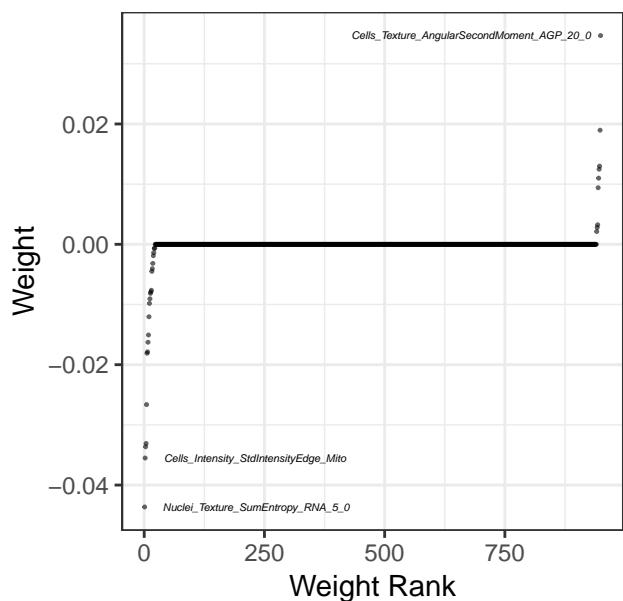
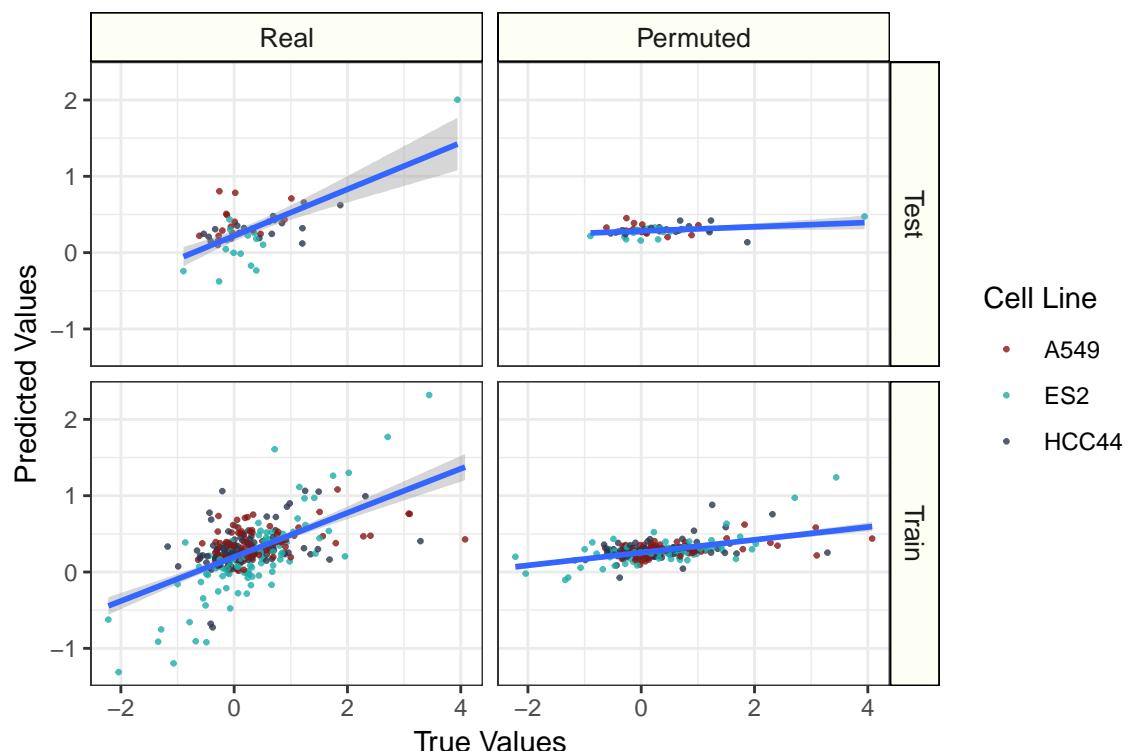


fit	shuffle	transform	r2	value
Train	Real	raw	0.54	0.31
Test	Real	raw	0.25	0.45
Train	Shuffle	raw	0.57	0.29
Test	Shuffle	raw	0.36	0.38



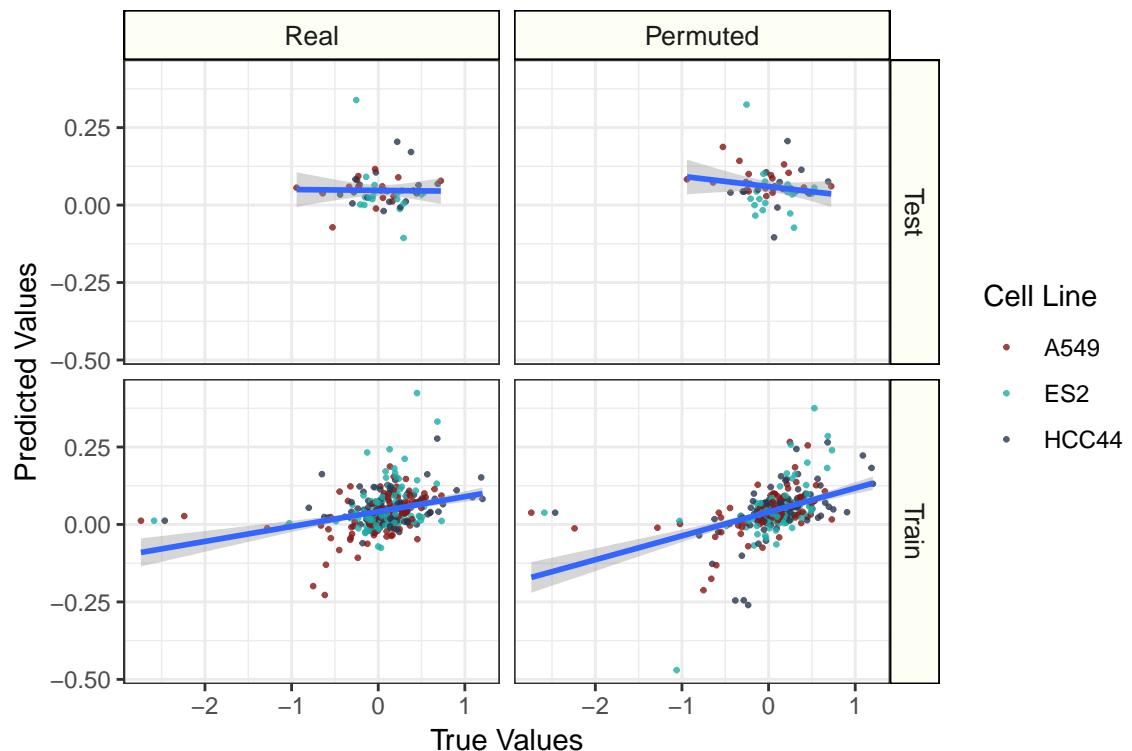
Performance: cc_s_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

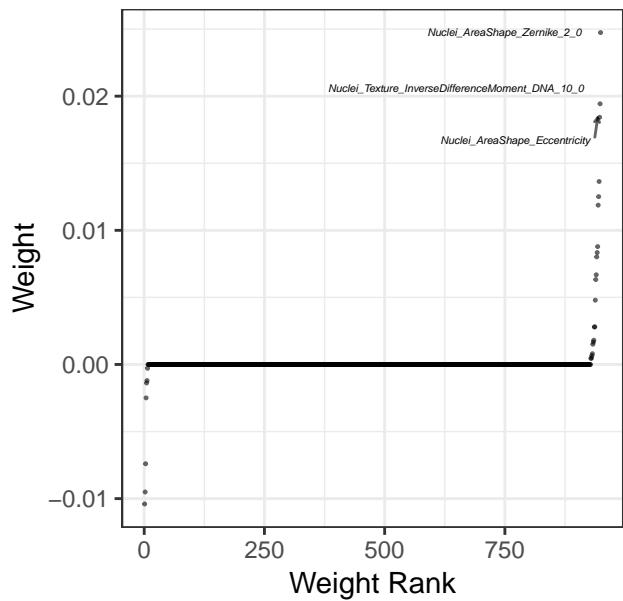


Performance: vb_infection_percentage

Transform: raw

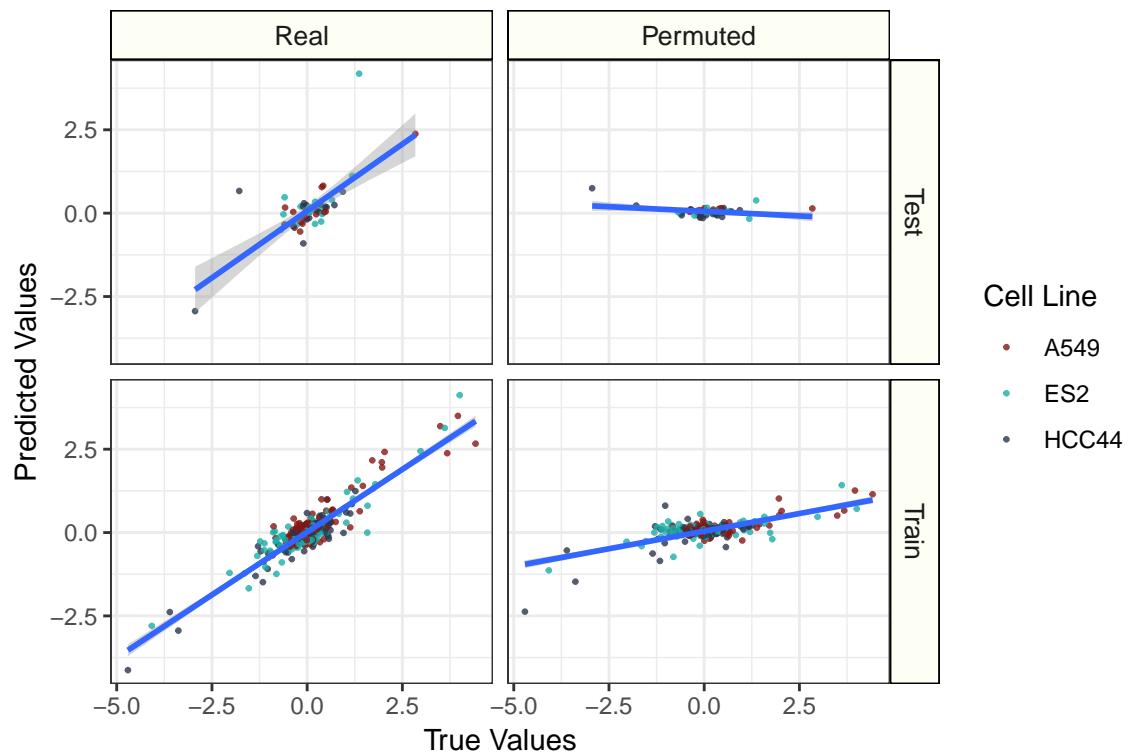


fit	shuffle	transform	r2	value
Train	Real	raw	0.07	0.18
Test	Real	raw	-0.05	0.11
Train	Shuffle	raw	0.12	0.17
Test	Shuffle	raw	-0.12	0.11

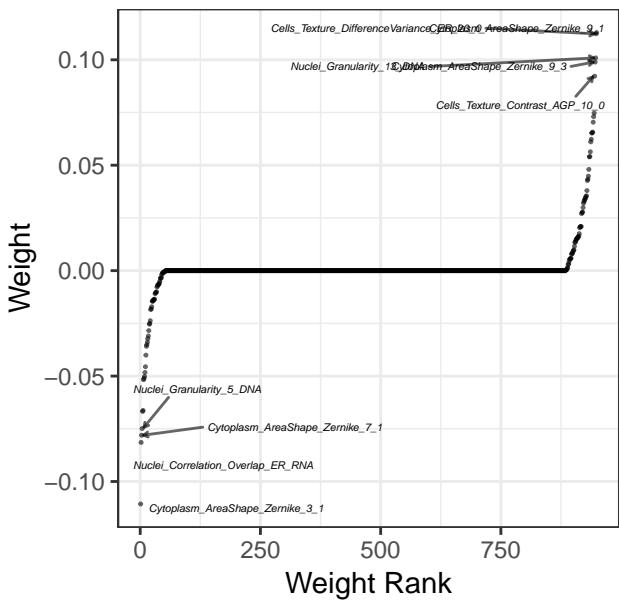


Performance: vb_live_cell_area

Transform: raw

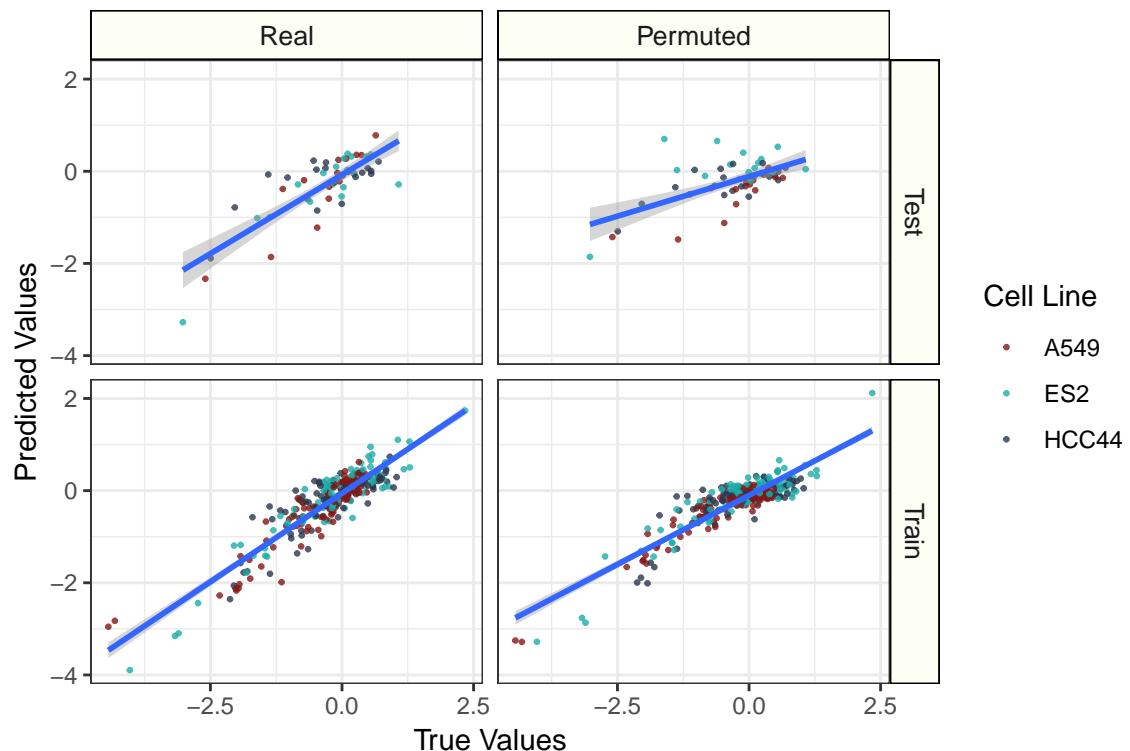


fit	shuffle	transform	r2	value
Train	Real	raw	0.83	0.15
Test	Real	raw	0.32	0.38
Train	Shuffle	raw	0.33	0.59
Test	Shuffle	raw	-0.14	0.63

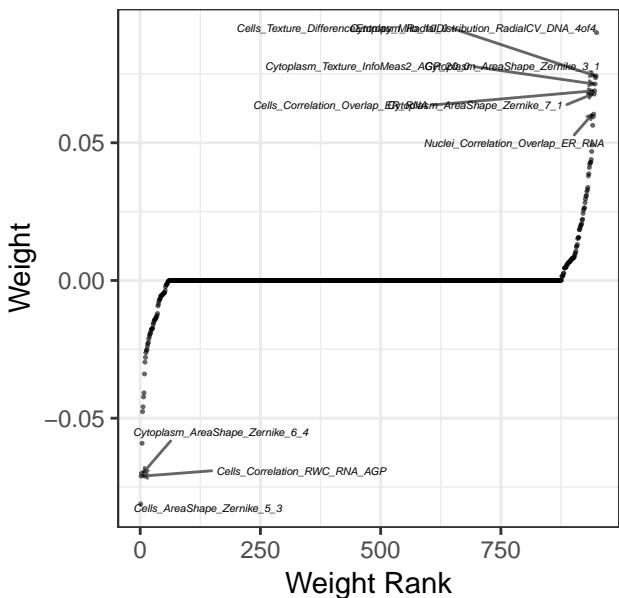


Performance: vb_live_cell_roundness

Transform: raw

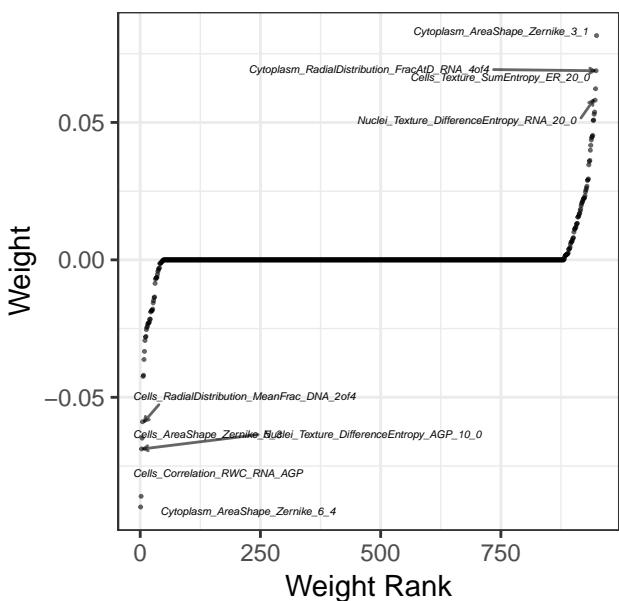
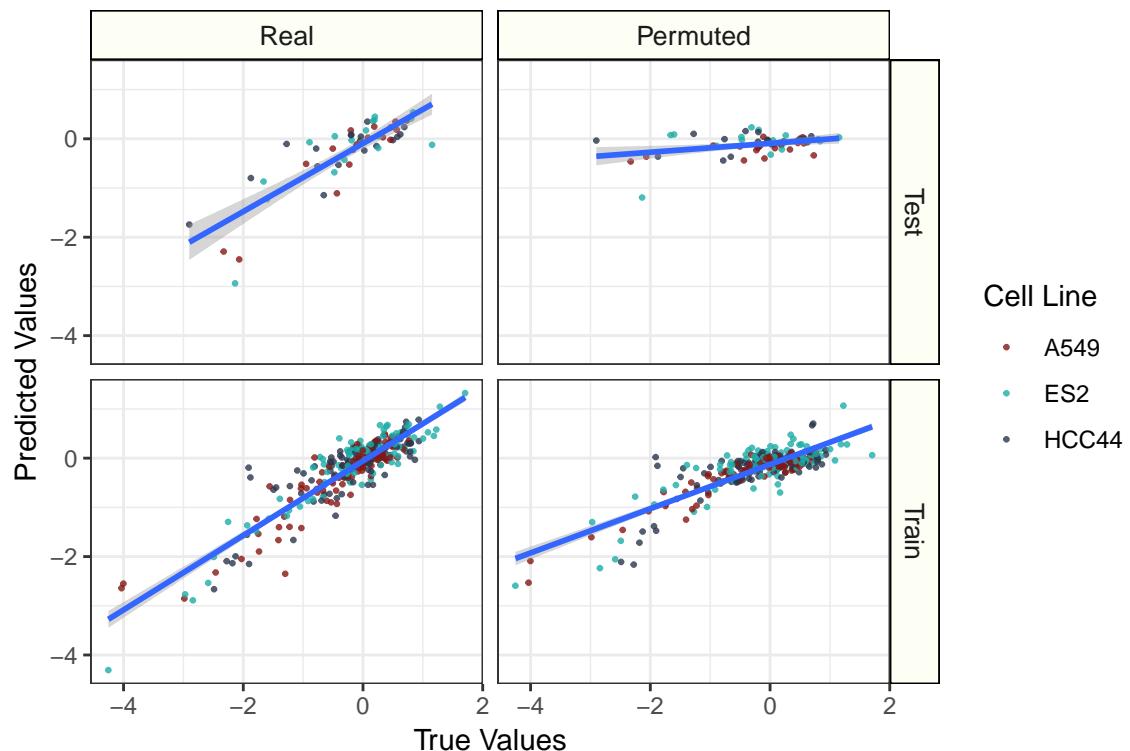


fit	shuffle	transform	r2	value
Train	Real	raw	0.82	0.13
Test	Real	raw	0.65	0.25
Train	Shuffle	raw	0.75	0.19
Test	Shuffle	raw	0.35	0.47



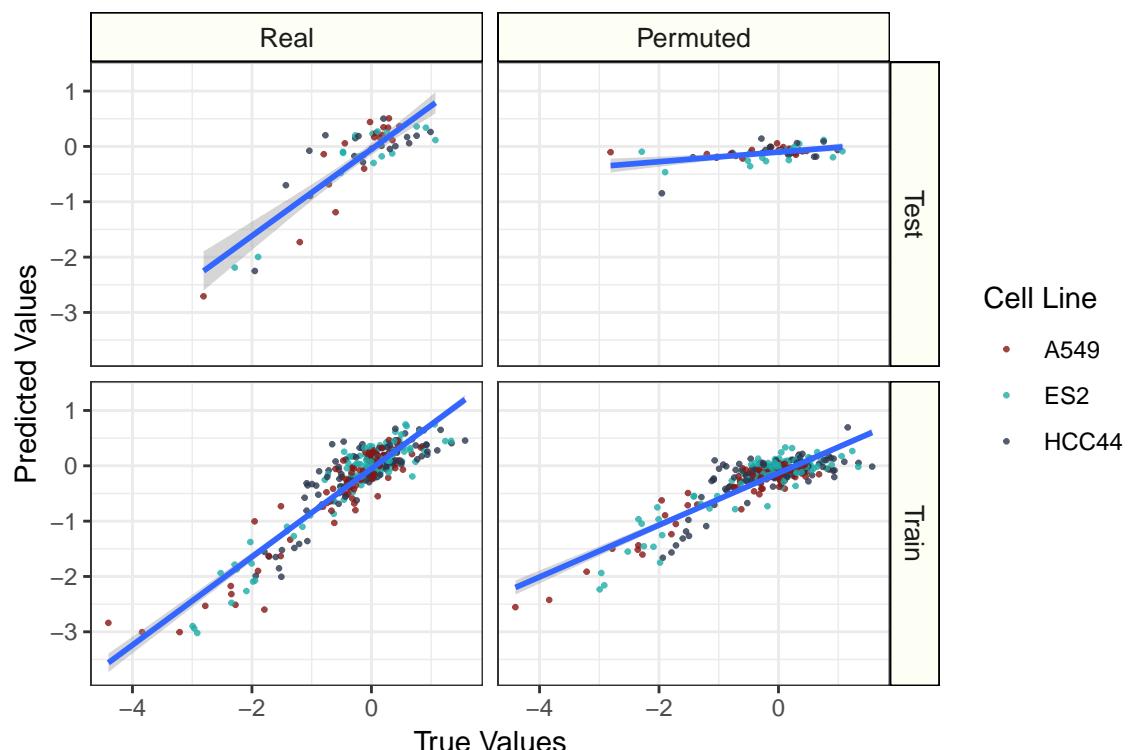
Performance: vb_live_cell_width_length

Transform: raw

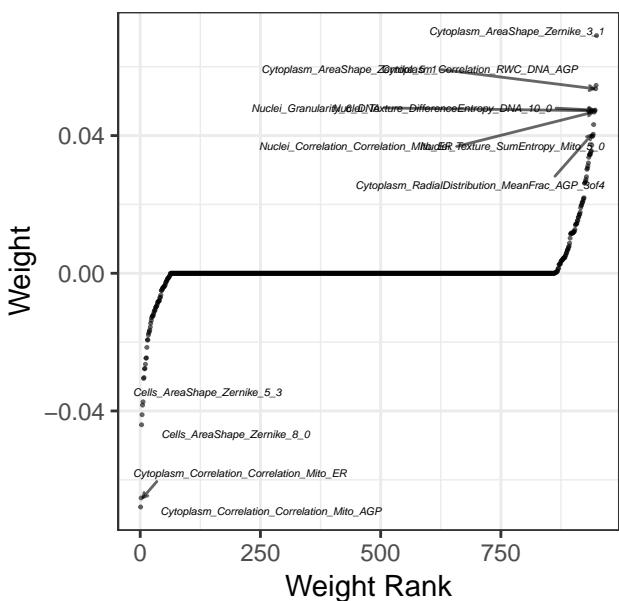


Performance: vb_num_live_cells

Transform: raw

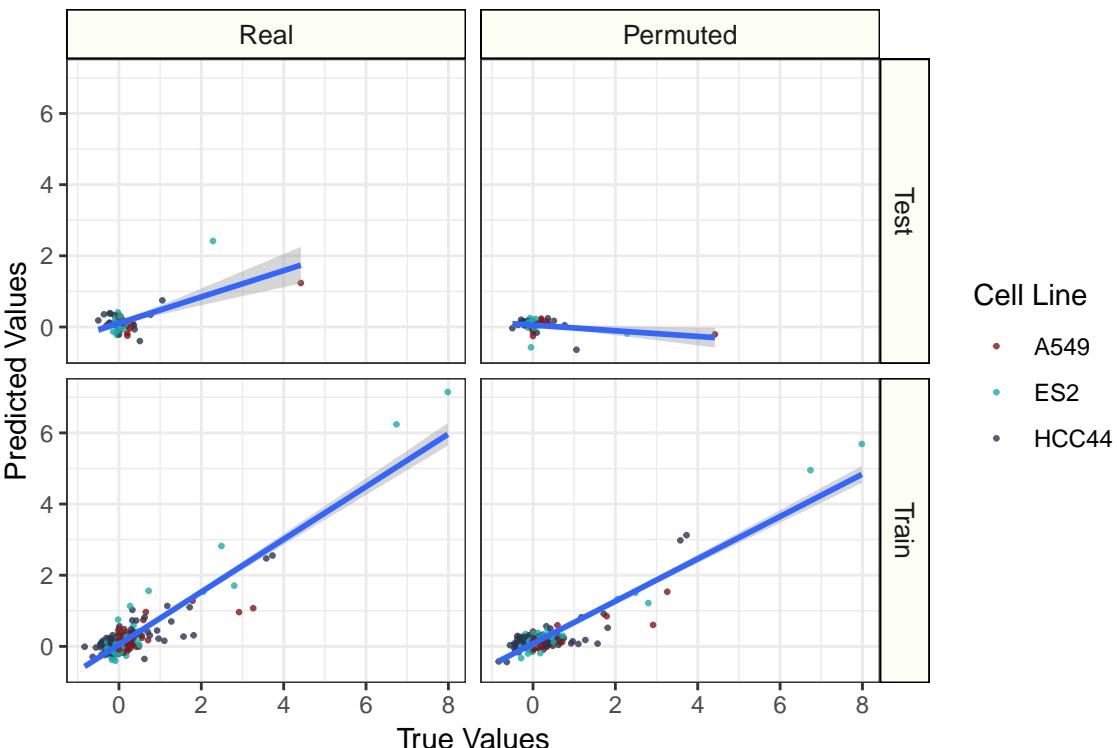


fit	shuffle	transform	r2	value
Train	Real	raw	0.84	0.12
Test	Real	raw	0.74	0.16
Train	Shuffle	raw	0.65	0.27
Test	Shuffle	raw	0.13	0.54

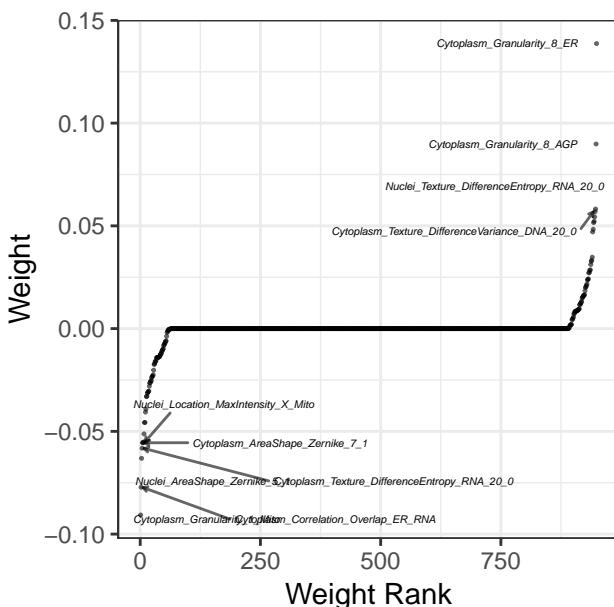


Performance: vb_percent_all_apoptosis

Transform: raw

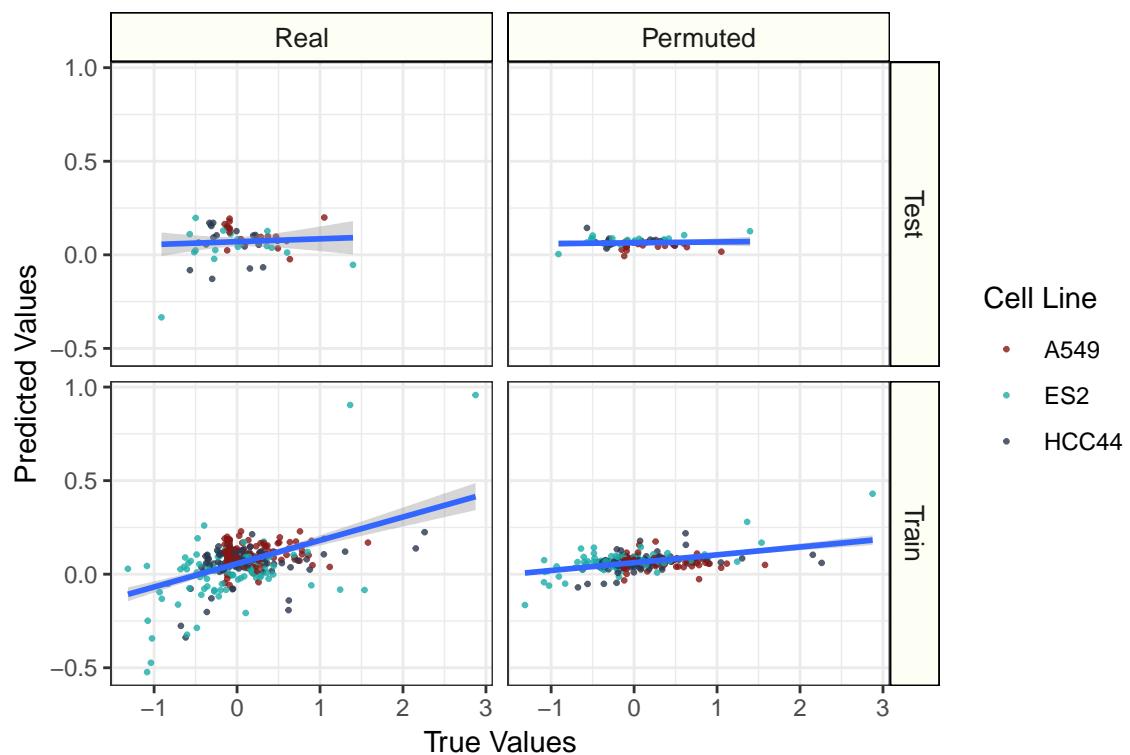


fit	shuffle	transform	r2	value
Train	Real	raw	0.81	0.12
Test	Real	raw	0.42	0.29
Train	Shuffle	raw	0.76	0.15
Test	Shuffle	raw	-0.24	0.62

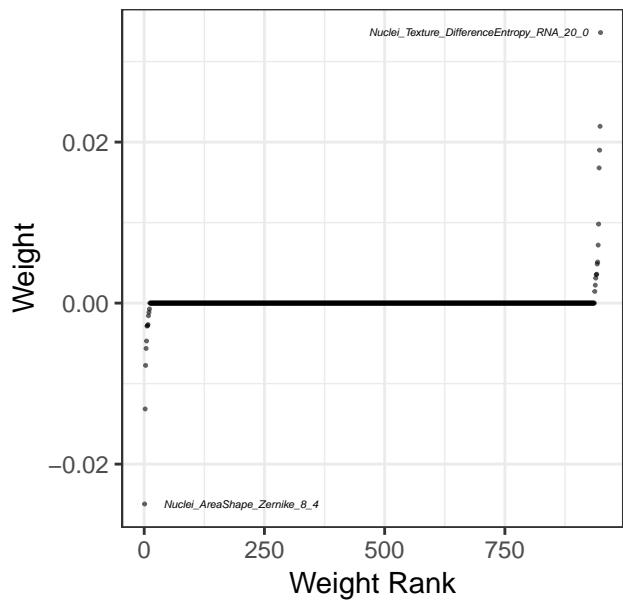


Performance: vb_percent_caspase_dead_only

Transform: raw

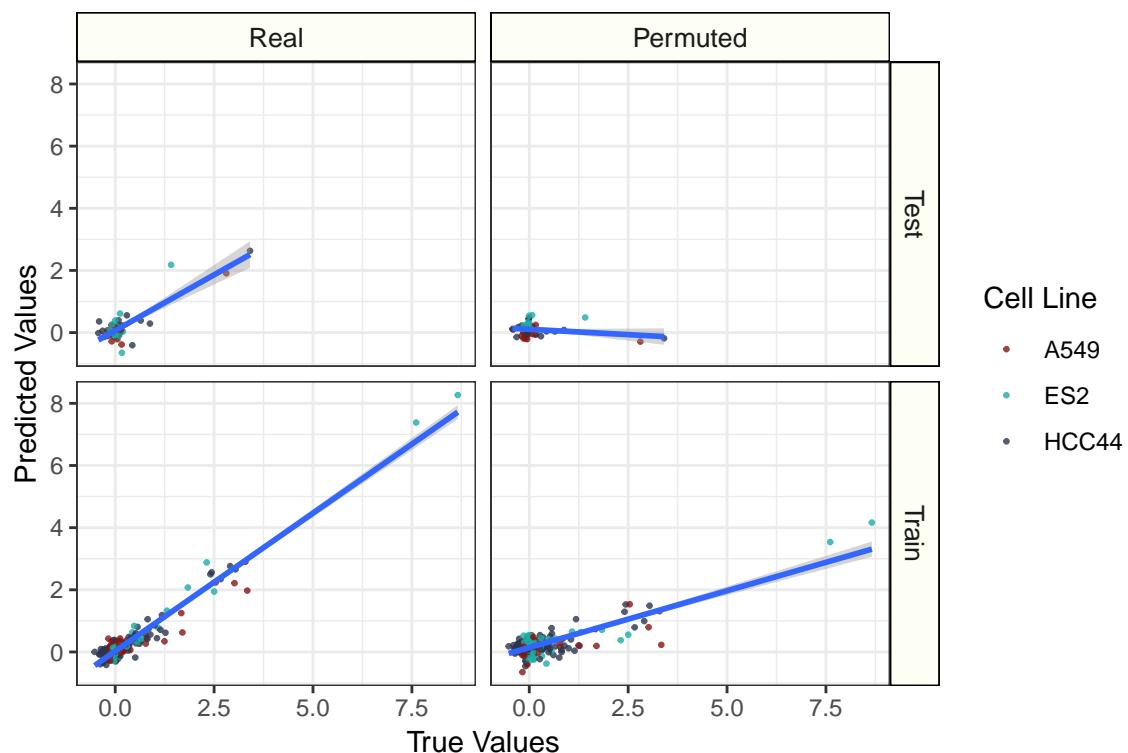


fit	shuffle	transform	r2	value
Train	Real	raw	0.18	0.19
Test	Real	raw	-0.03	0.18
Train	Shuffle	raw	0.08	0.22
Test	Shuffle	raw	-0.01	0.18

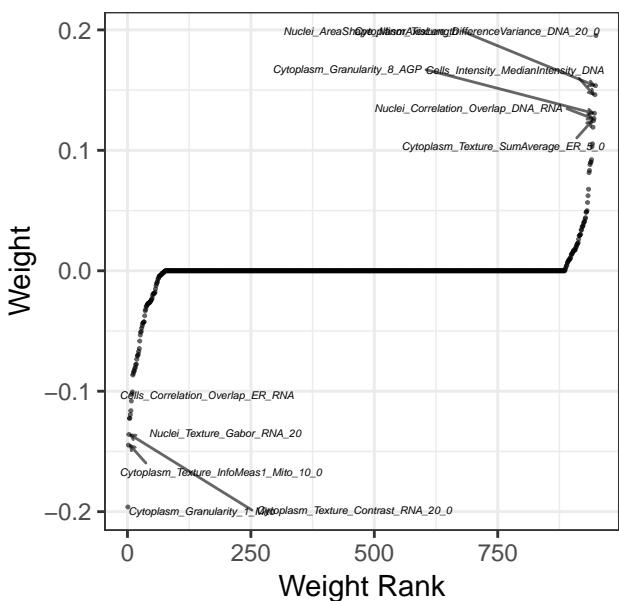


Performance: vb_percent_dead

Transform: raw

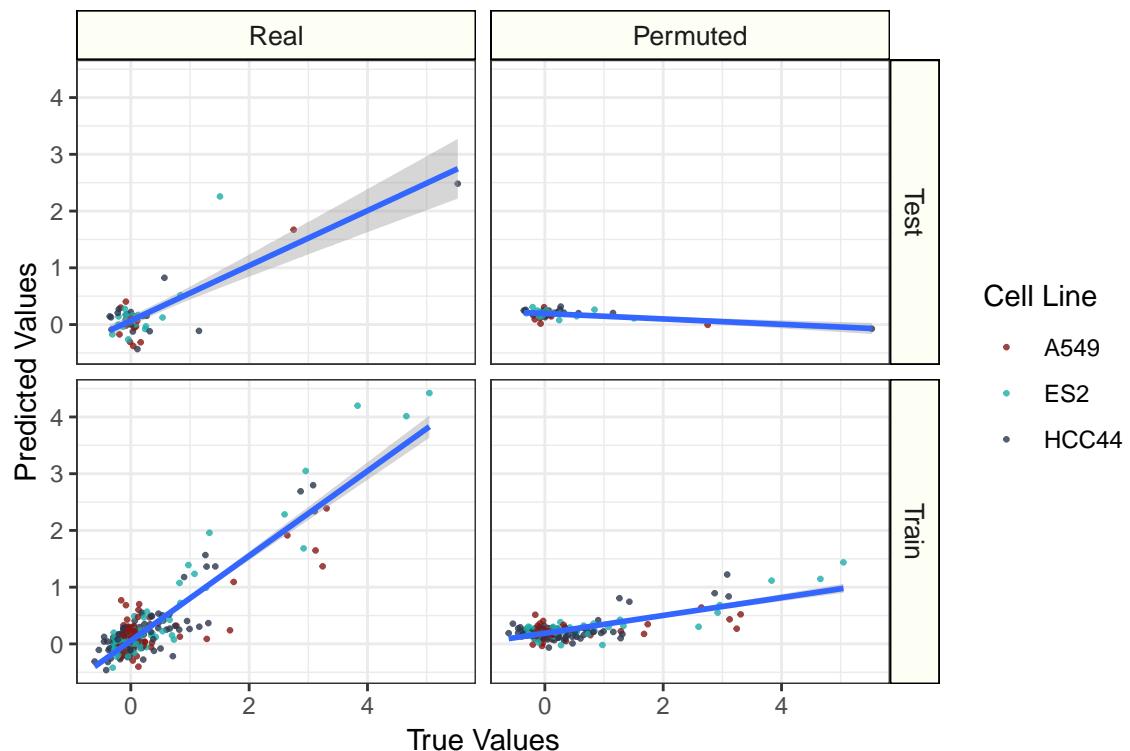


fit	shuffle	transform	r2	value
Train	Real	raw	0.93	0.05
Test	Real	raw	0.70	0.12
Train	Shuffle	raw	0.53	0.37
Test	Shuffle	raw	-0.23	0.52

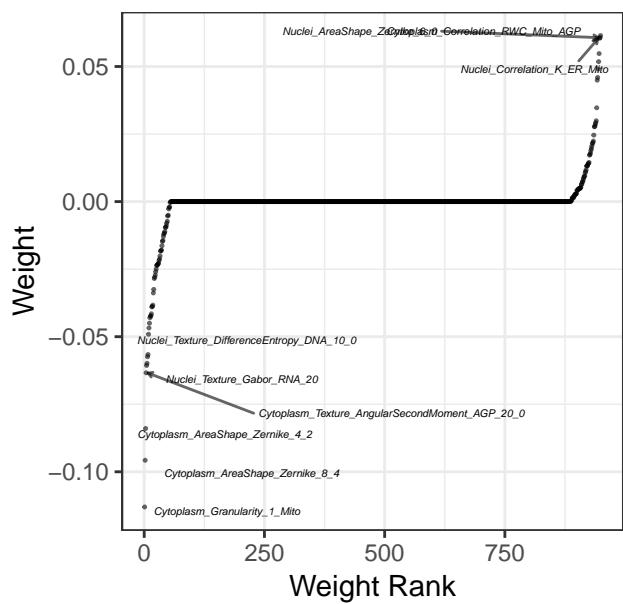


Performance: vb_percent_dead_only

Transform: raw

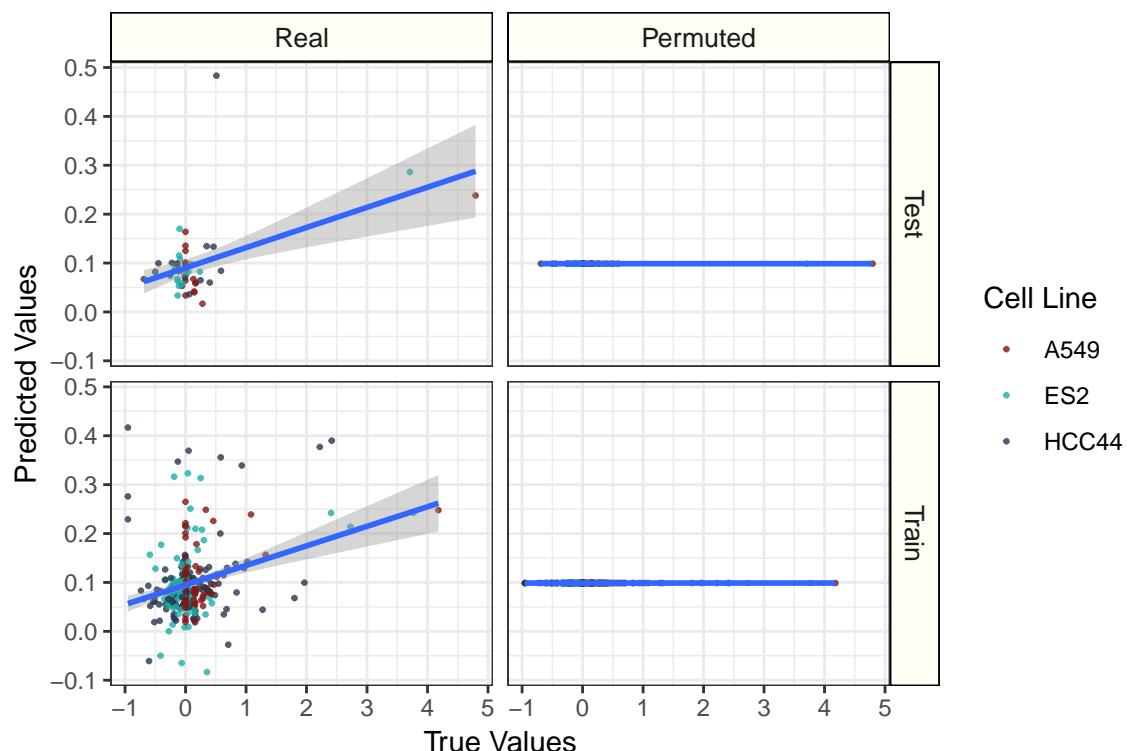


fit	shuffle	transform	r2	value
Train	Real	raw	0.81	0.11
Test	Real	raw	0.61	0.30
Train	Shuffle	raw	0.27	0.43
Test	Shuffle	raw	-0.10	0.85

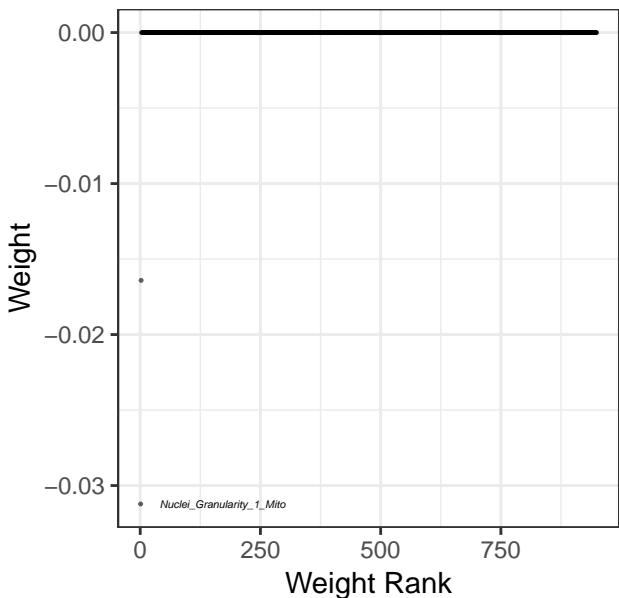


Performance: vb_percent_early_apoptosis

Transform: raw

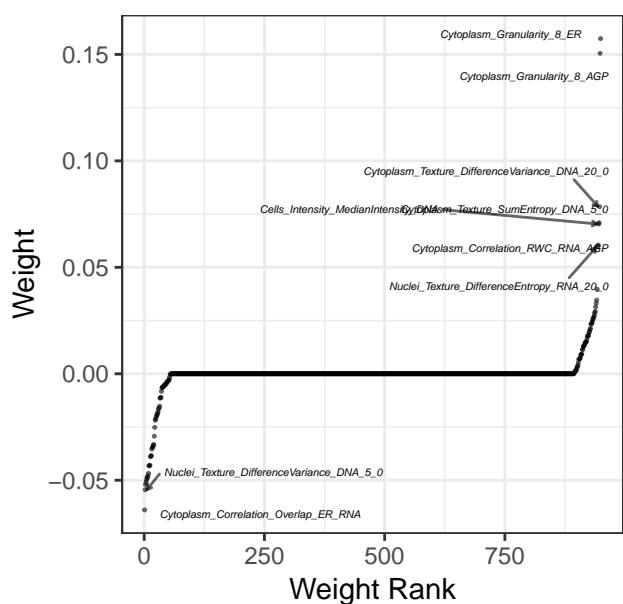
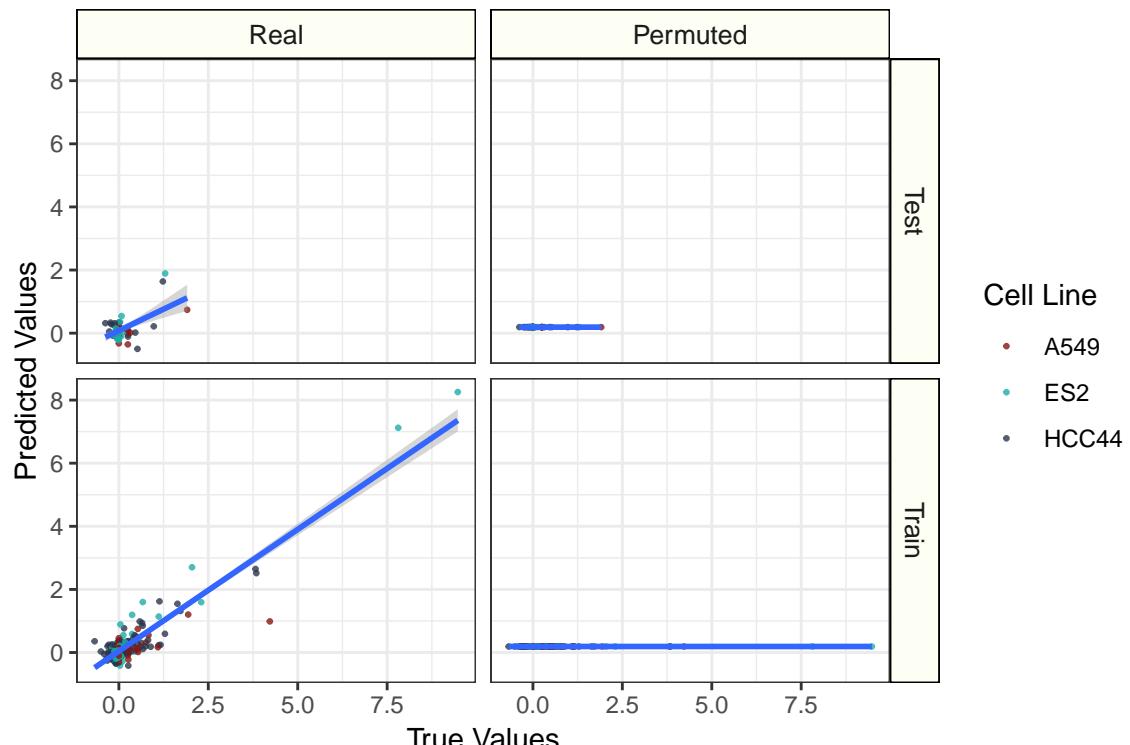


fit	shuffle	transform	r2	value
Train	Real	raw	0.06	0.27
Test	Real	raw	0.07	0.66
Train	Shuffle	raw	0.00	0.29
Test	Shuffle	raw	0.00	0.71



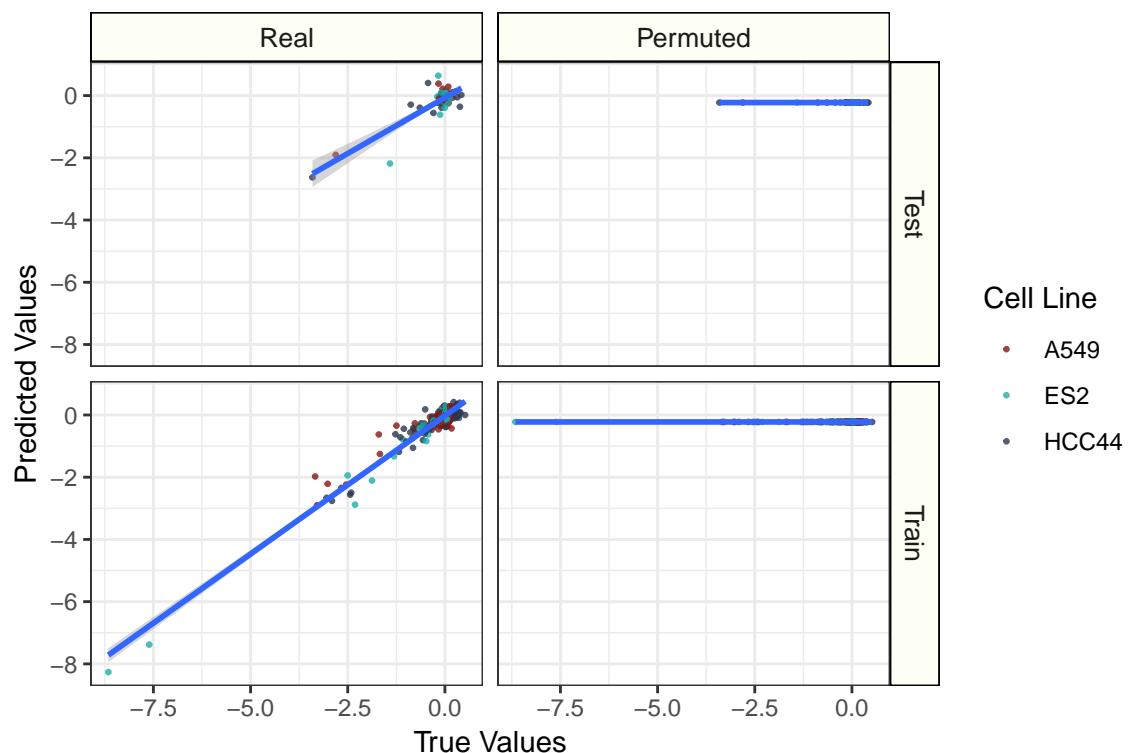
Performance: vb_percent_late_apoptosis

Transform: raw

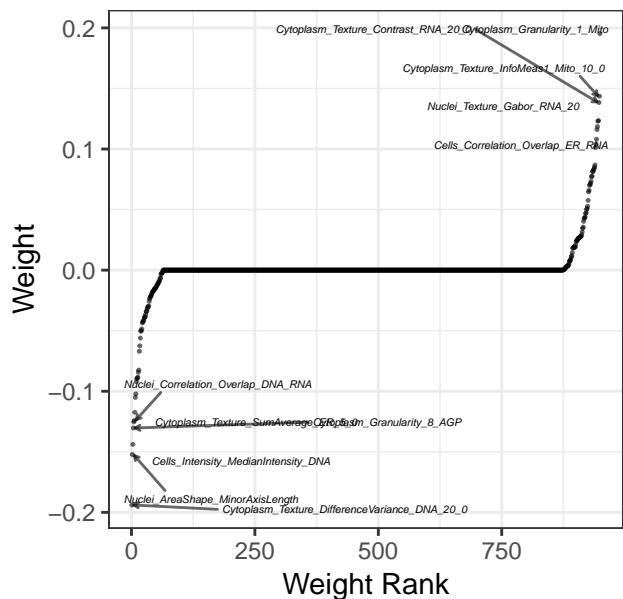


Performance: vb_percent_live

Transform: raw

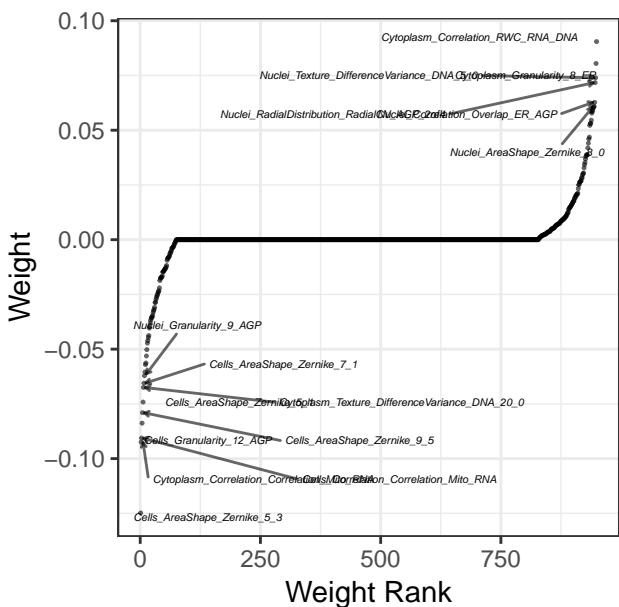
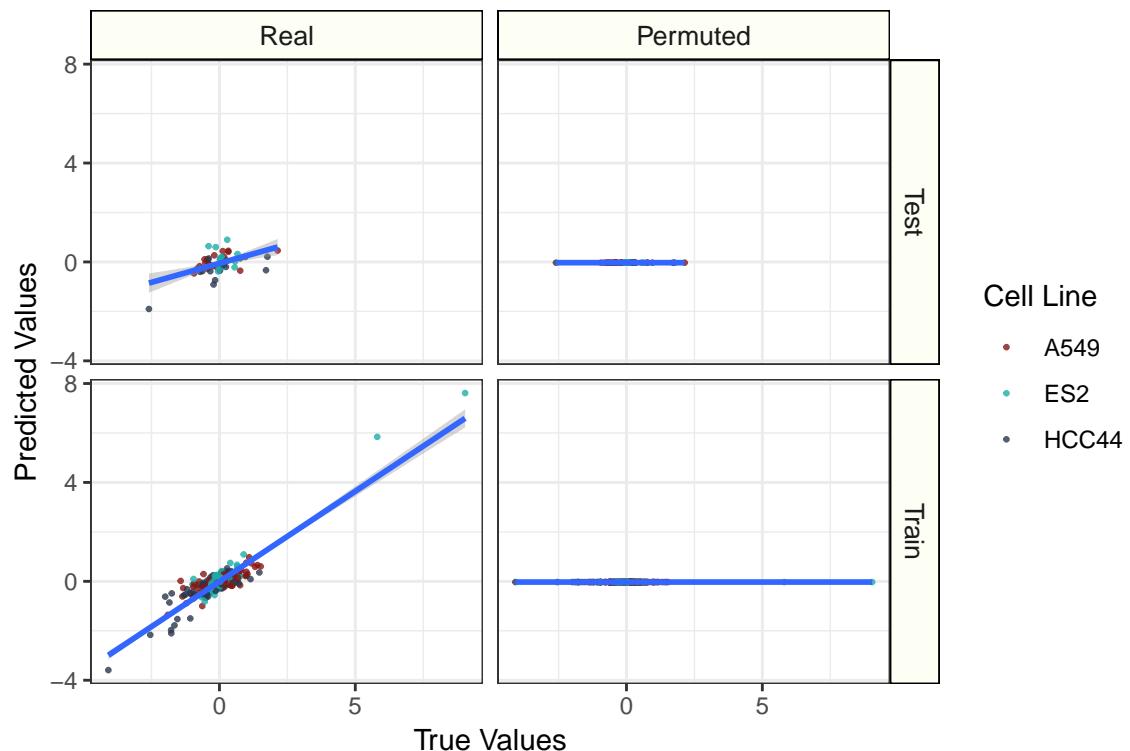


fit	shuffle	transform	r2	value
Train	Real	raw	0.93	0.05
Test	Real	raw	0.71	0.12
Train	Shuffle	raw	0.00	0.78
Test	Shuffle	raw	-0.01	0.42



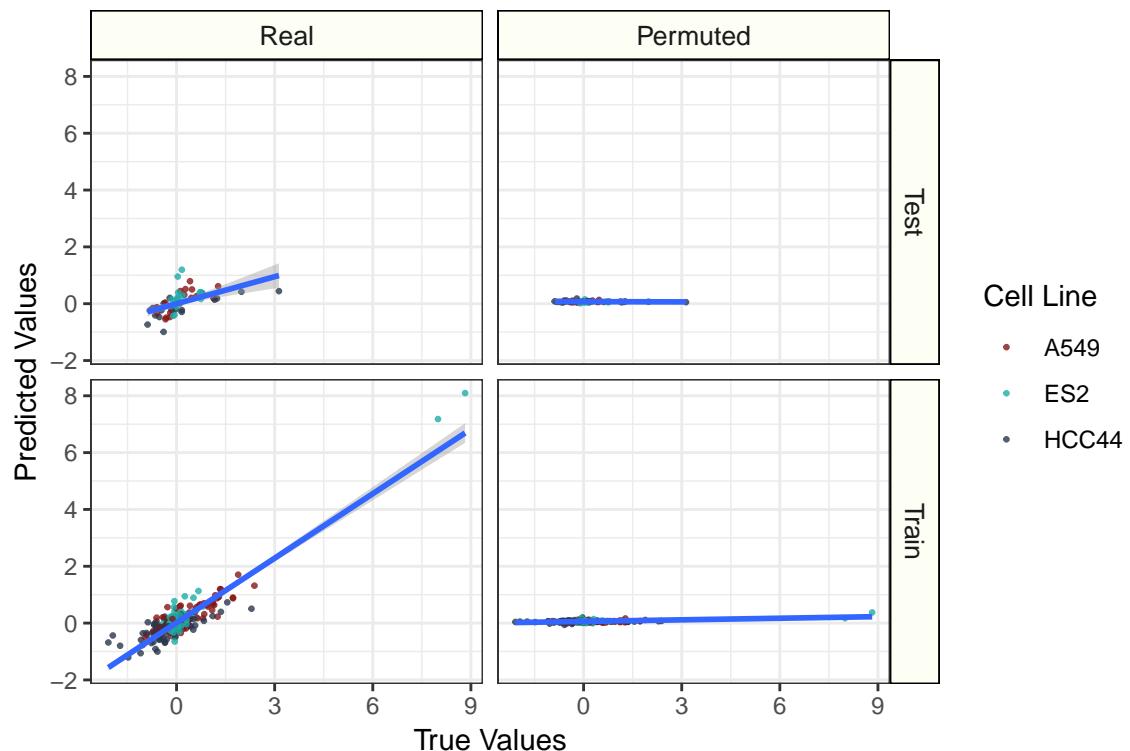
Performance: vb_ros_back_mean

Transform: raw



Performance: vb_ros_mean

Transform: raw



fit	shuffle	transform	r2	value
Train	Real	raw	0.83	0.14
Test	Real	raw	0.27	0.37
Train	Shuffle	raw	0.04	0.79
Test	Shuffle	raw	-0.02	0.52

