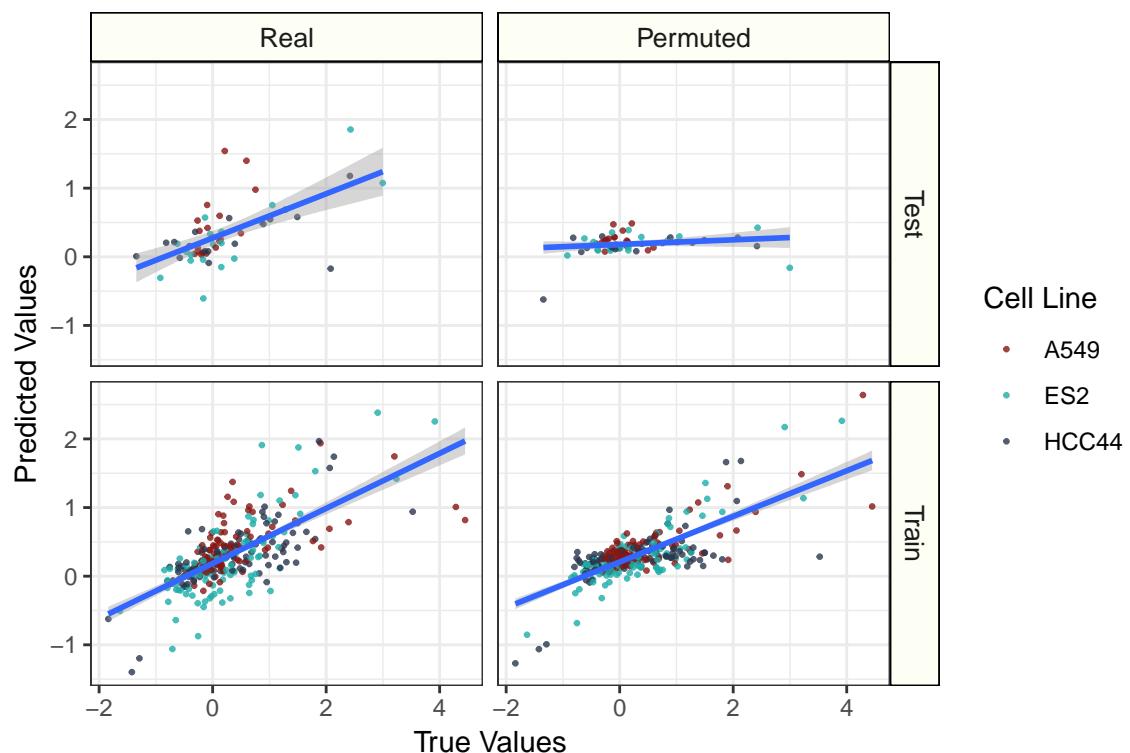
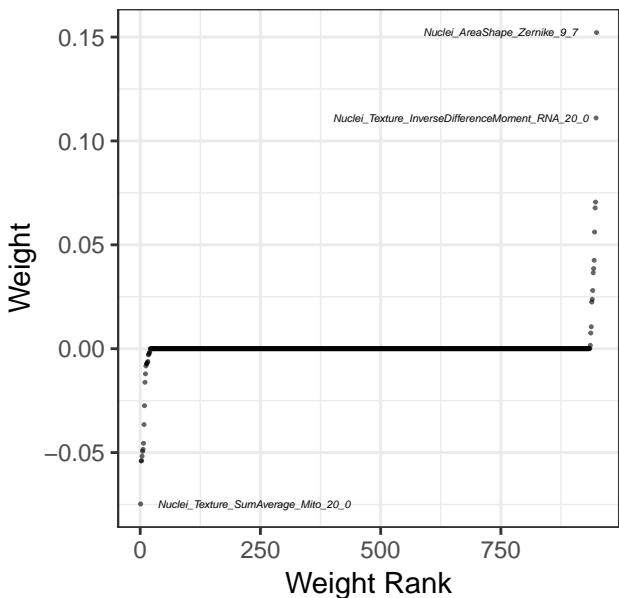


Performance: cc_all_high_h2ax

Transform: raw

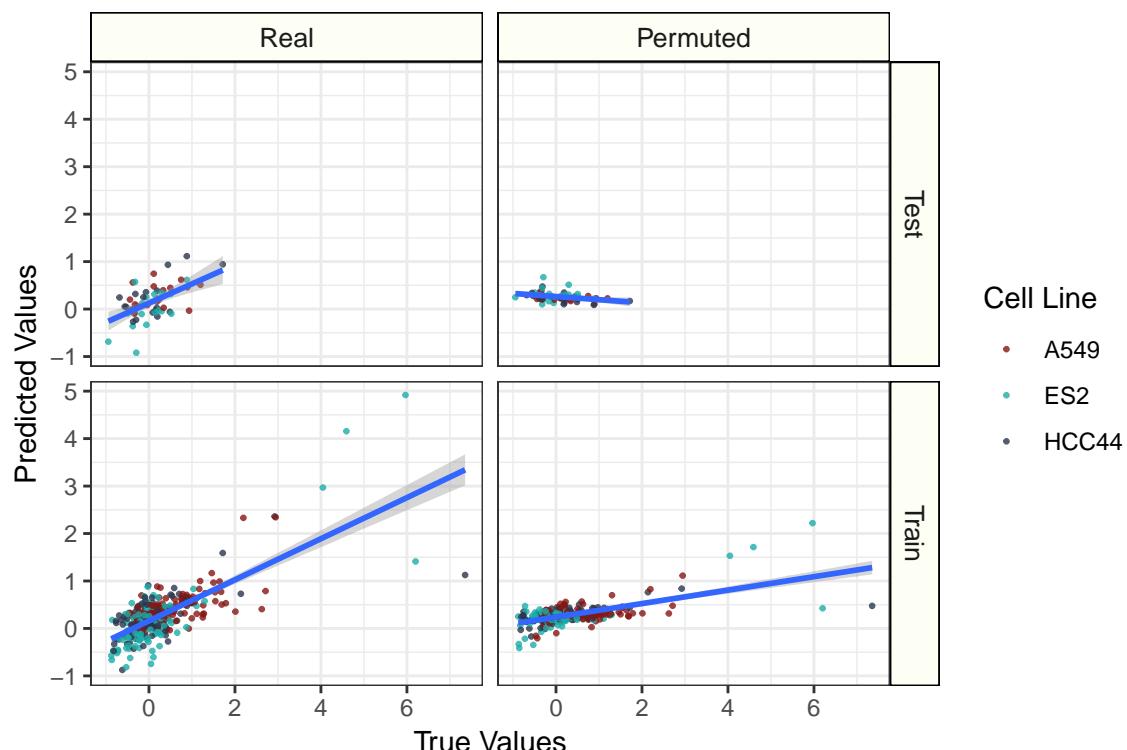


fit	shuffle	transform	r2	value
Train	Real	raw	0.47	0.37
Test	Real	raw	0.33	0.47
Train	Shuffle	raw	0.47	0.38
Test	Shuffle	raw	0.03	0.68

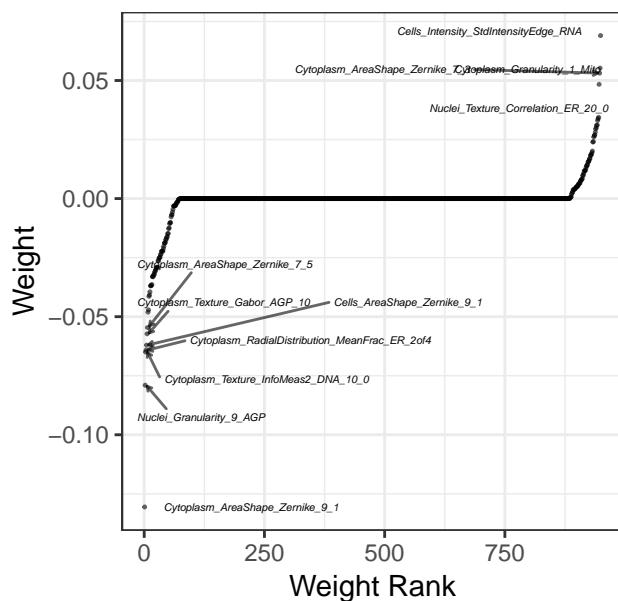


Performance: cc_all_large_notround_polinuclear_mean

Transform: raw

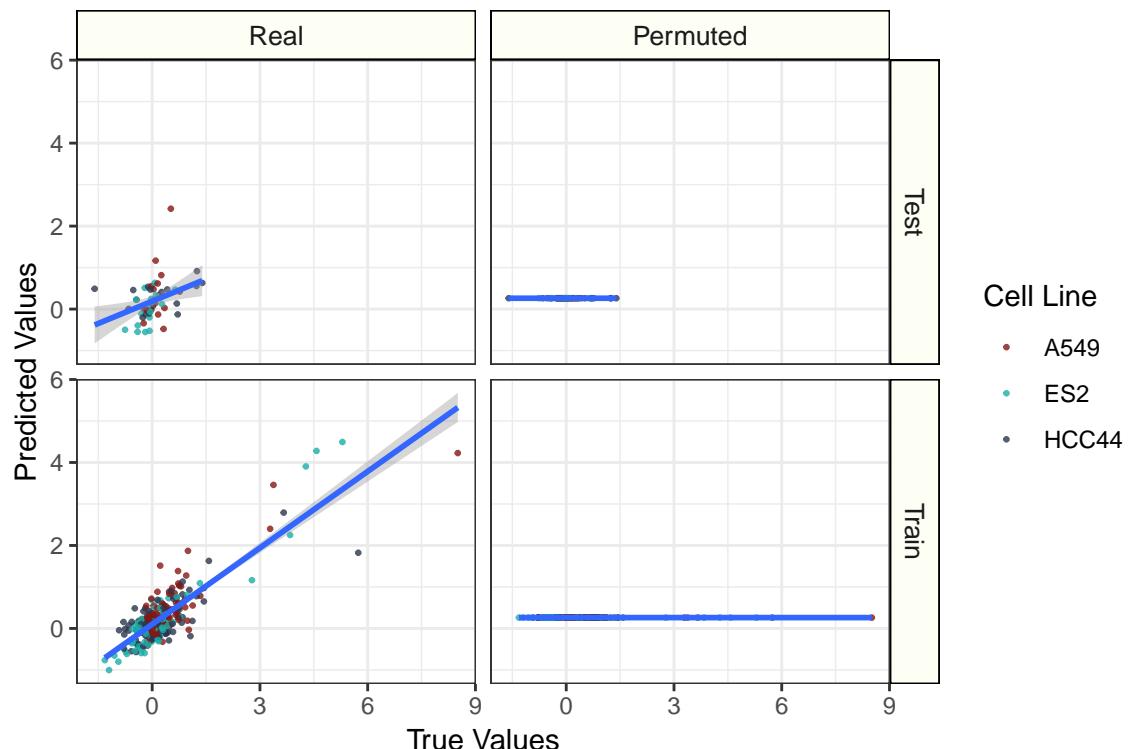


fit	shuffle	transform	r2	value
Train	Real	raw	0.52	0.45
Test	Real	raw	0.24	0.19
Train	Shuffle	raw	0.23	0.71
Test	Shuffle	raw	-0.29	0.32

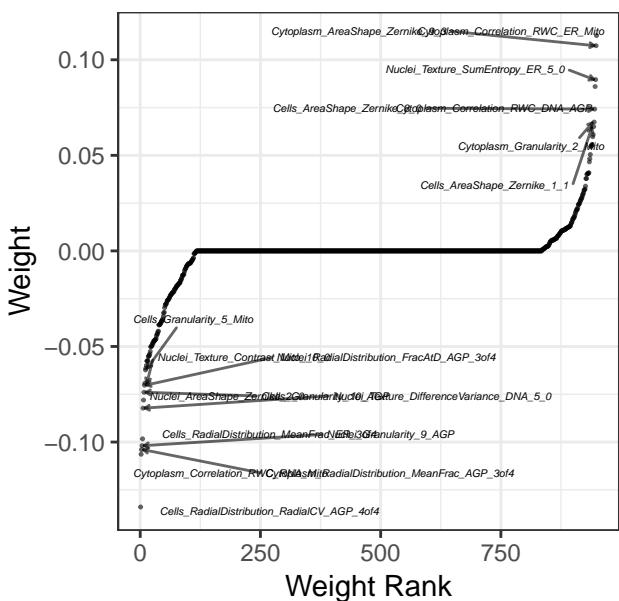


Performance: cc_all_large_round_polyplloid_mean

Transform: raw

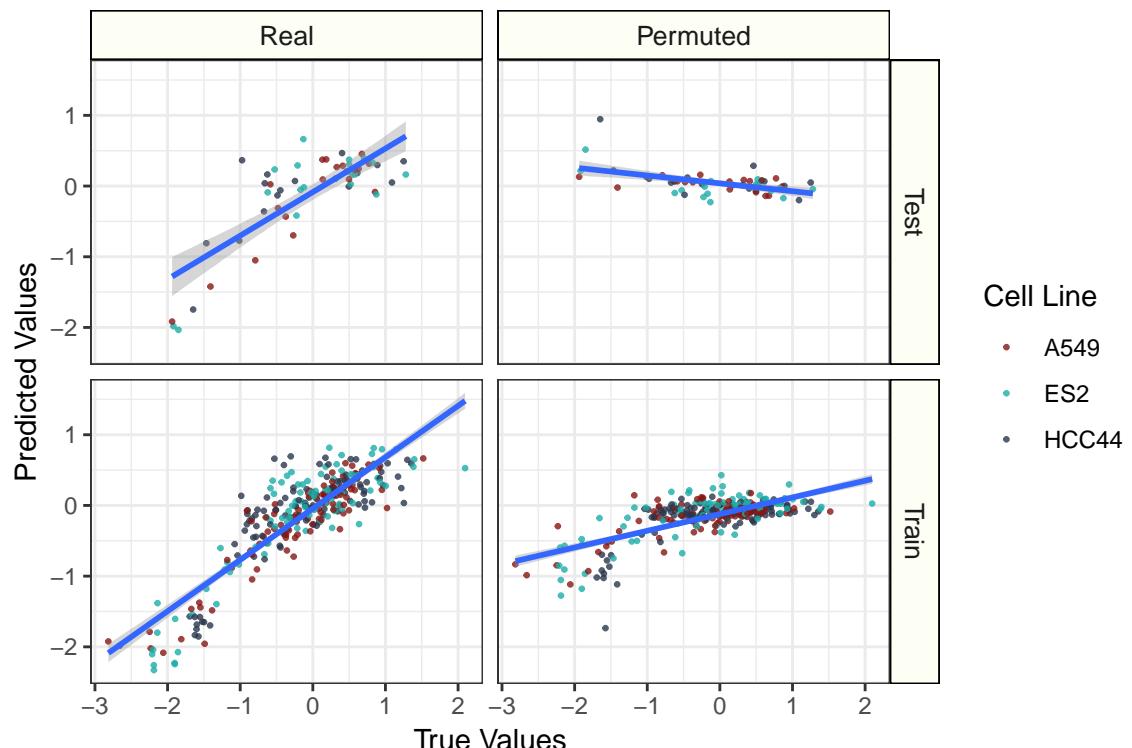


fit	shuffle	transform	r2	value
Train	Real	raw	0.71	0.26
Test	Real	raw	-0.39	0.33
Train	Shuffle	raw	0.00	0.91
Test	Shuffle	raw	-0.20	0.29

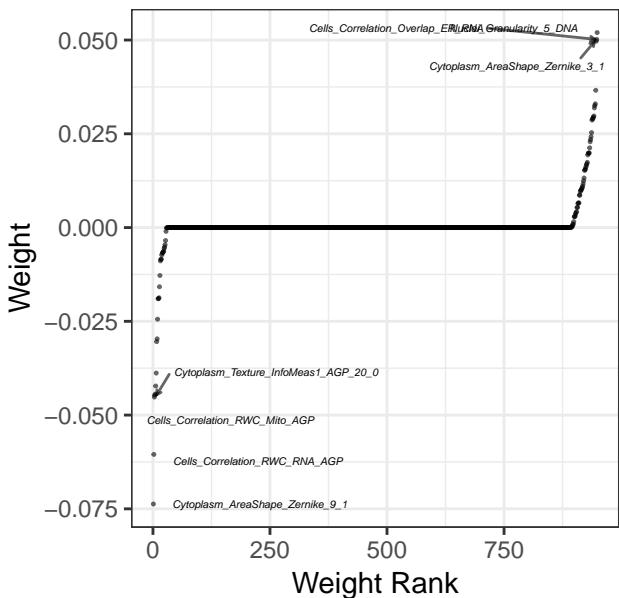


Performance: cc_all_n_objects

Transform: raw

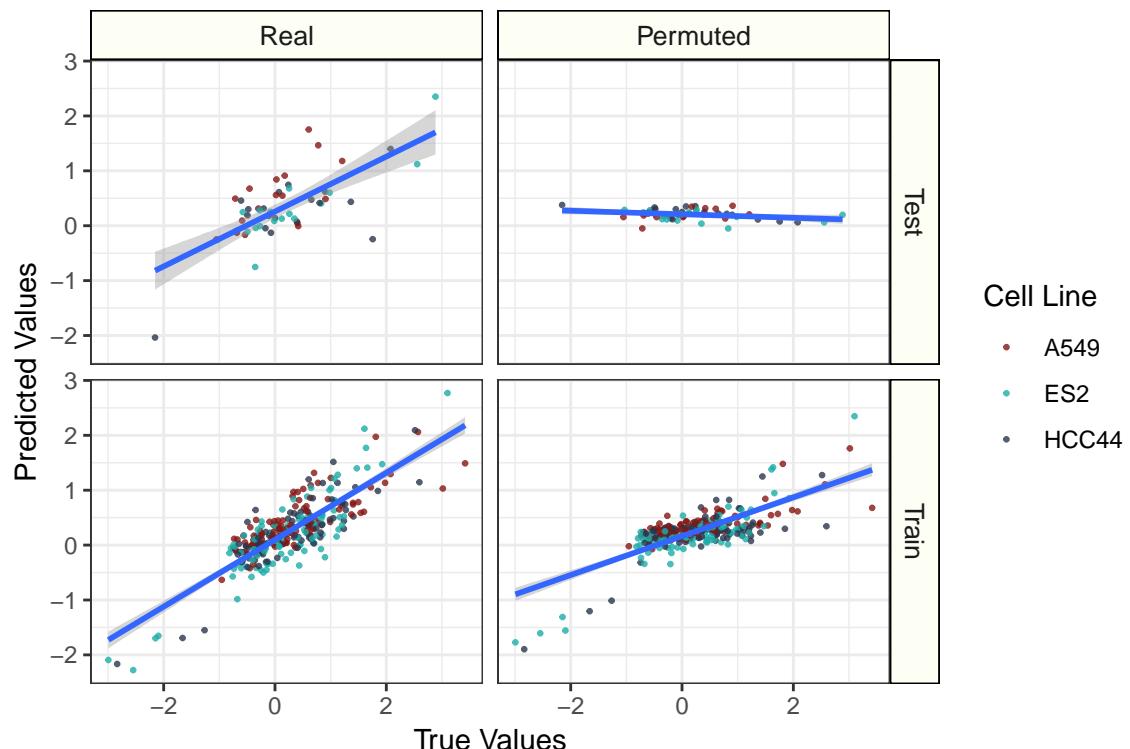


fit	shuffle	transform	r2	value
Train	Real	raw	0.76	0.16
Test	Real	raw	0.61	0.26
Train	Shuffle	raw	0.35	0.44
Test	Shuffle	raw	-0.28	0.86

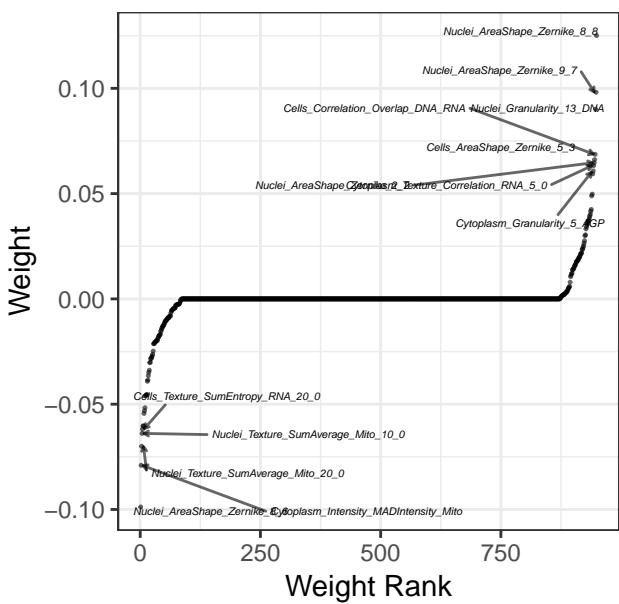


Performance: cc_all_n_spots_h2ax_mean

Transform: raw

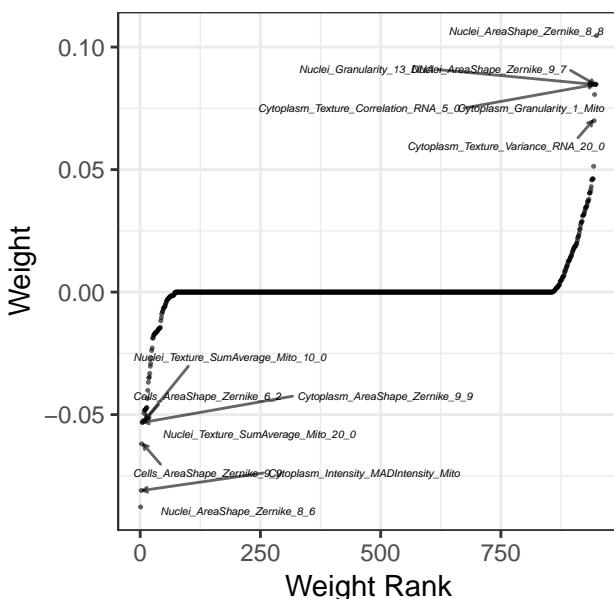
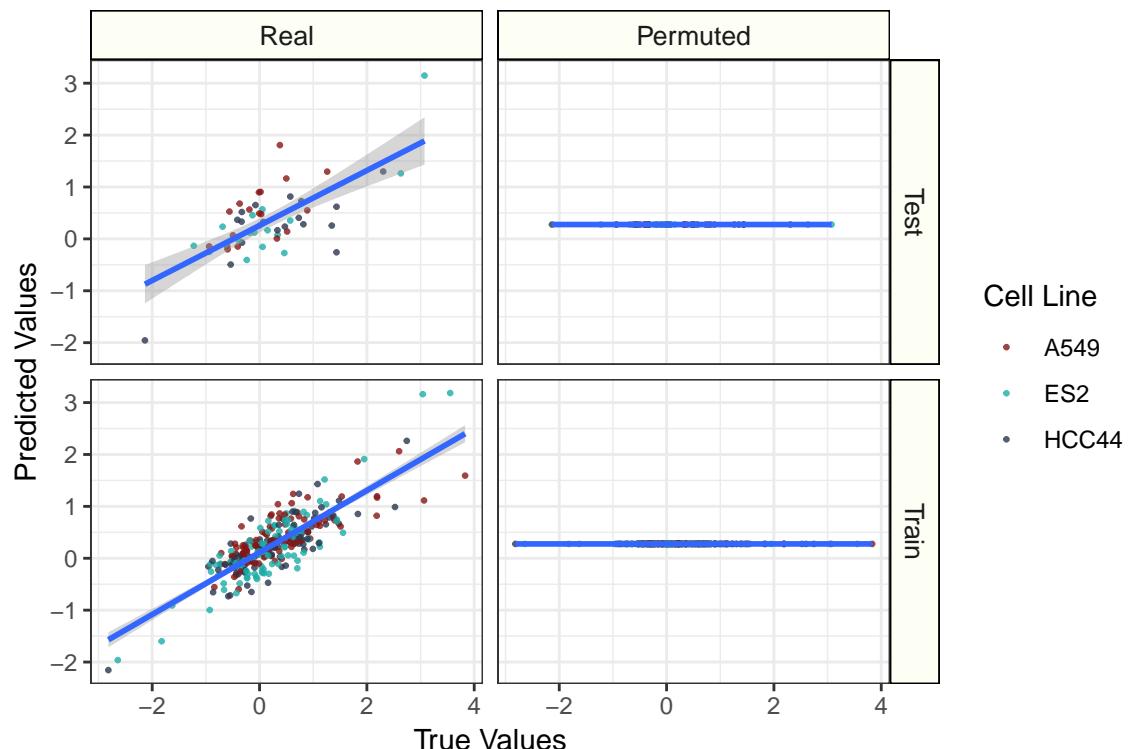


fit	shuffle	transform	r2	value
Train	Real	raw	0.69	0.21
Test	Real	raw	0.45	0.43
Train	Shuffle	raw	0.48	0.35
Test	Shuffle	raw	-0.08	0.85



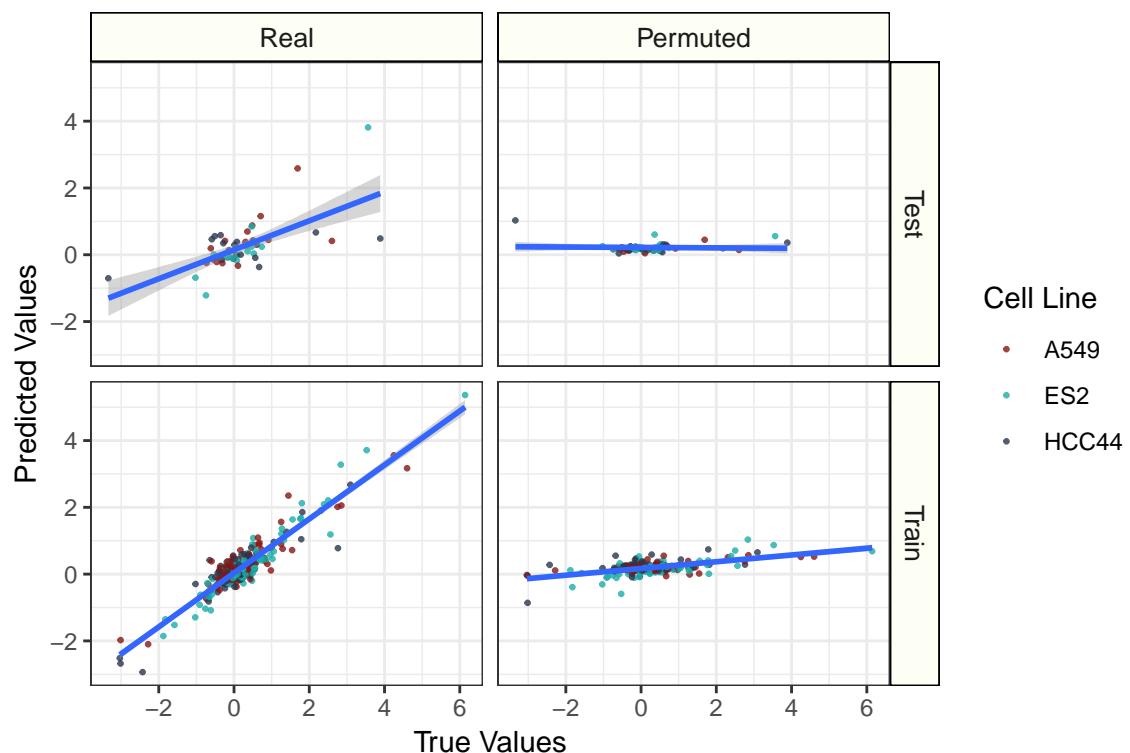
Performance: cc_all_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

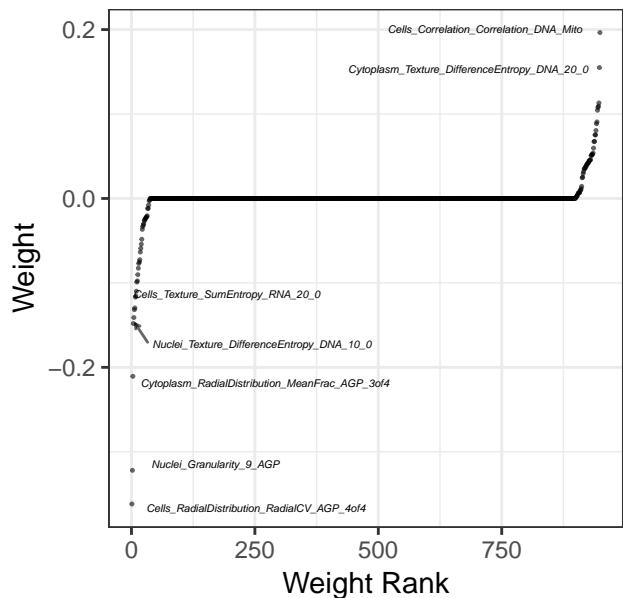


Performance: cc_all_nucleus_area_mean

Transform: raw

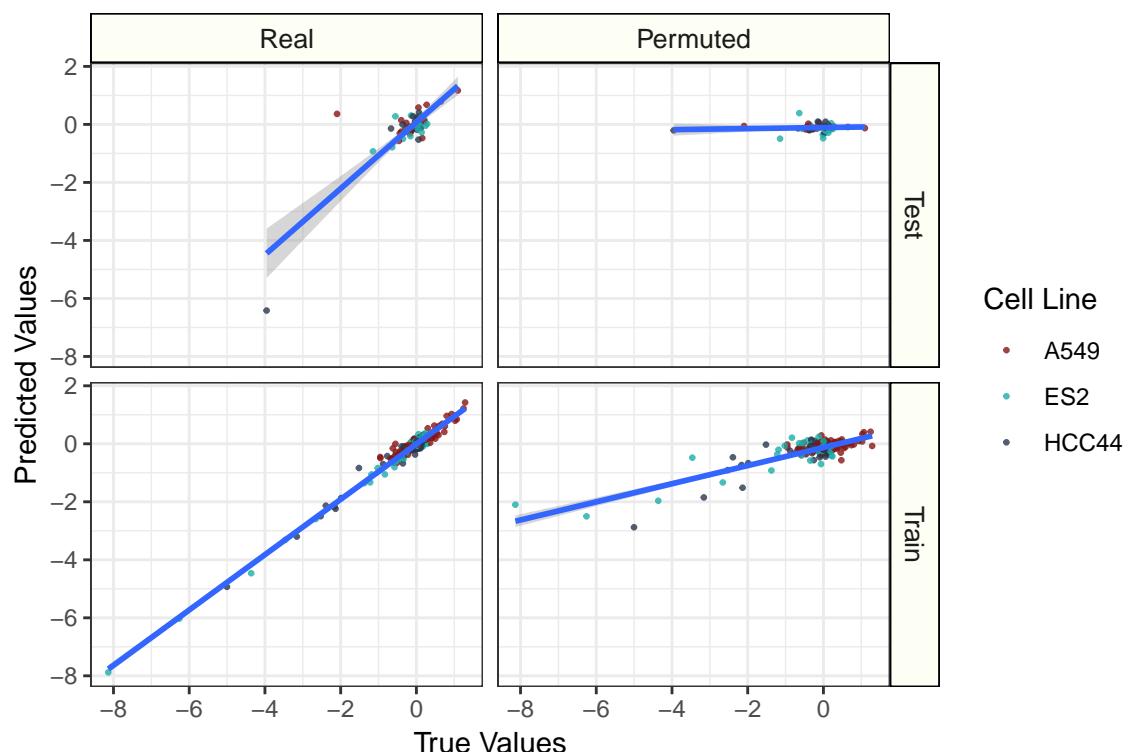


fit	shuffle	transform	r2	value
Train	Real	raw	0.86	0.12
Test	Real	raw	0.41	0.67
Train	Shuffle	raw	0.17	0.74
Test	Shuffle	raw	-0.03	1.17

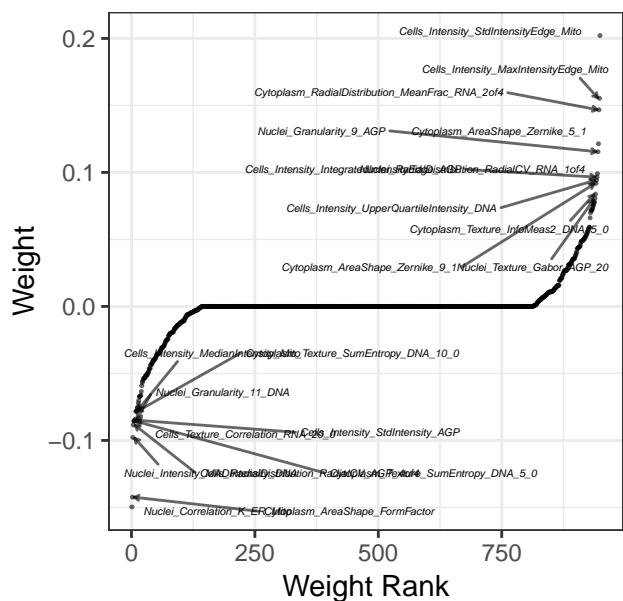


Performance: cc_all_nucleus_roundness_mean

Transform: raw

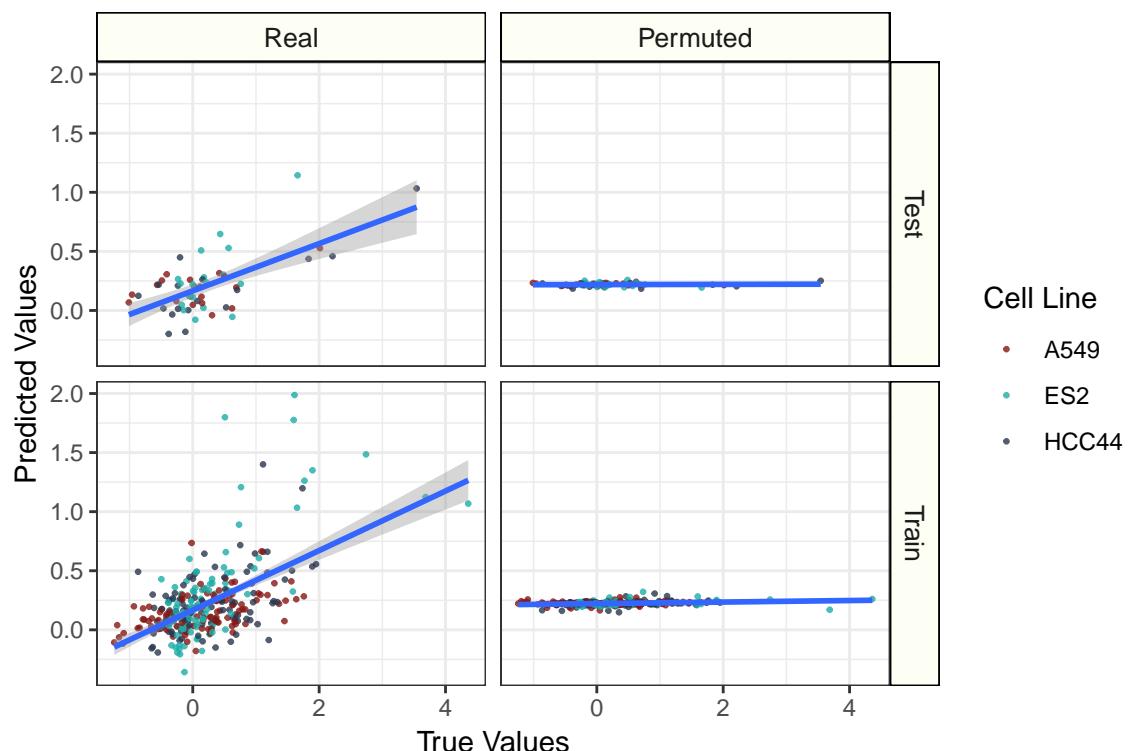


fit	shuffle	transform	r2	value
Train	Real	raw	0.97	0.02
Test	Real	raw	0.34	0.30
Train	Shuffle	raw	0.48	0.40
Test	Shuffle	raw	-0.01	0.46

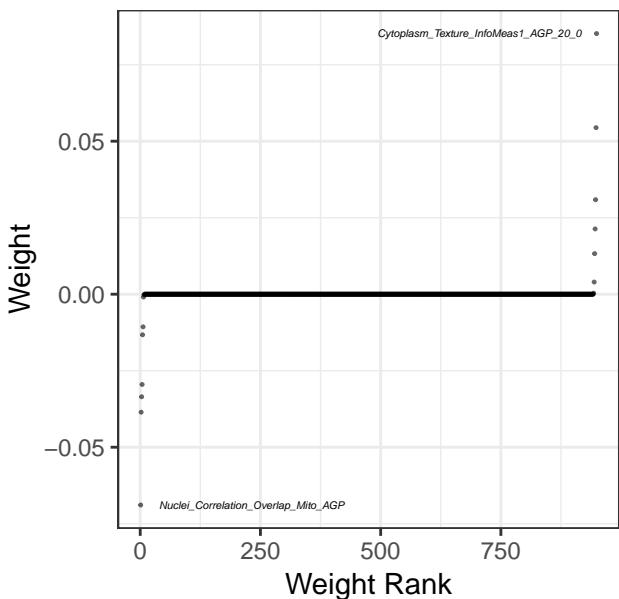


Performance: cc_cc_early_mitosis

Transform: raw

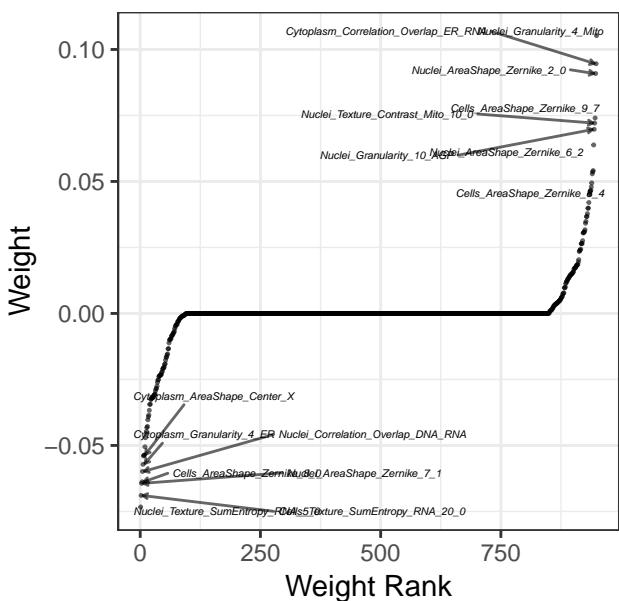
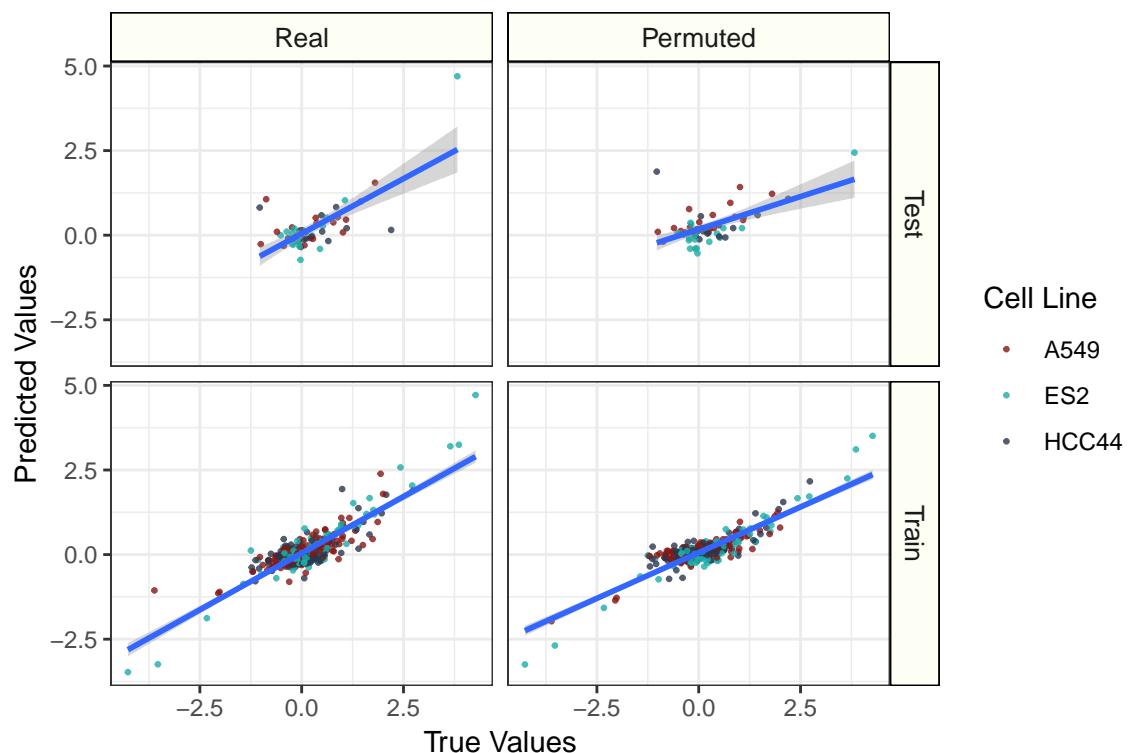


fit	shuffle	transform	r2	value
Train	Real	raw	0.31	0.34
Test	Real	raw	0.30	0.44
Train	Shuffle	raw	0.01	0.49
Test	Shuffle	raw	0.00	0.63



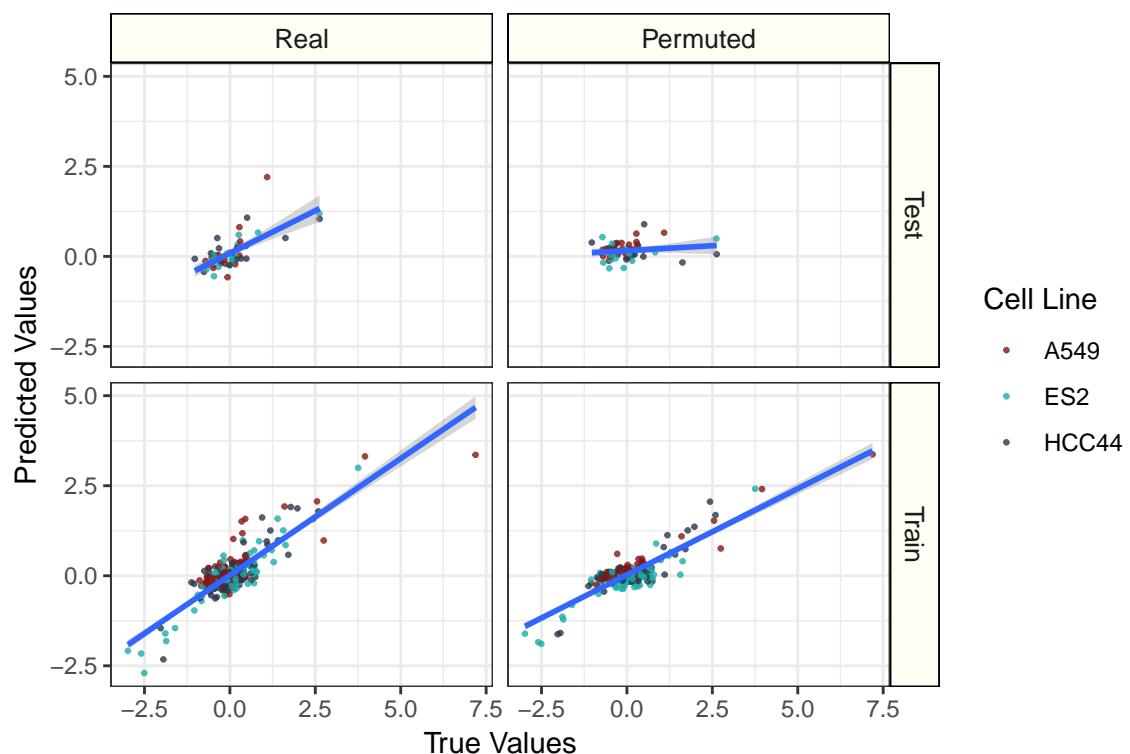
Performance: cc_cc_g1

Transform: raw

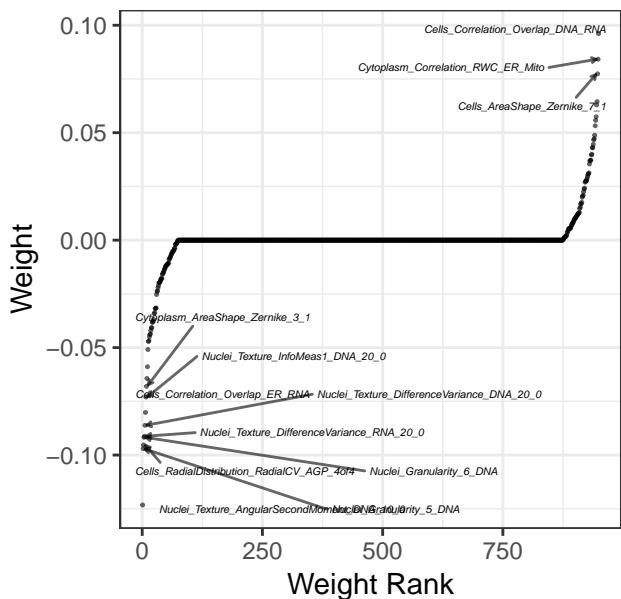


Performance: cc_cc_g2

Transform: raw

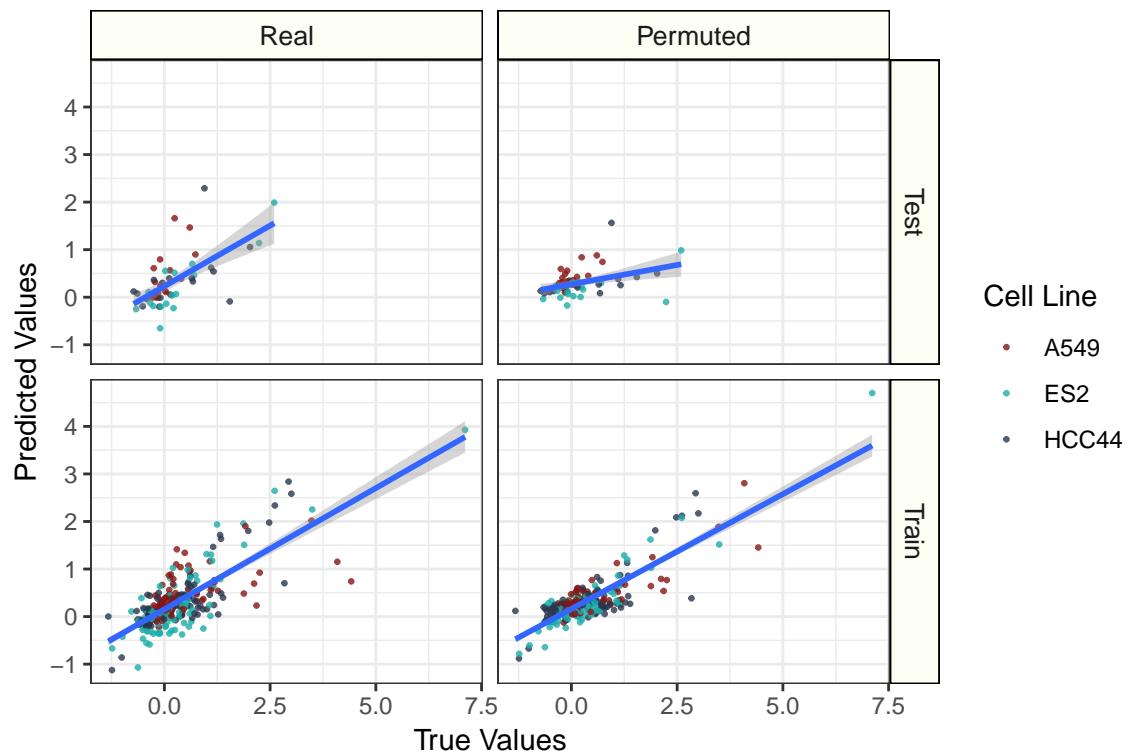


fit	shuffle	transform	r2	value
Train	Real	raw	0.73	0.20
Test	Real	raw	0.47	0.26
Train	Shuffle	raw	0.65	0.25
Test	Shuffle	raw	-0.05	0.52

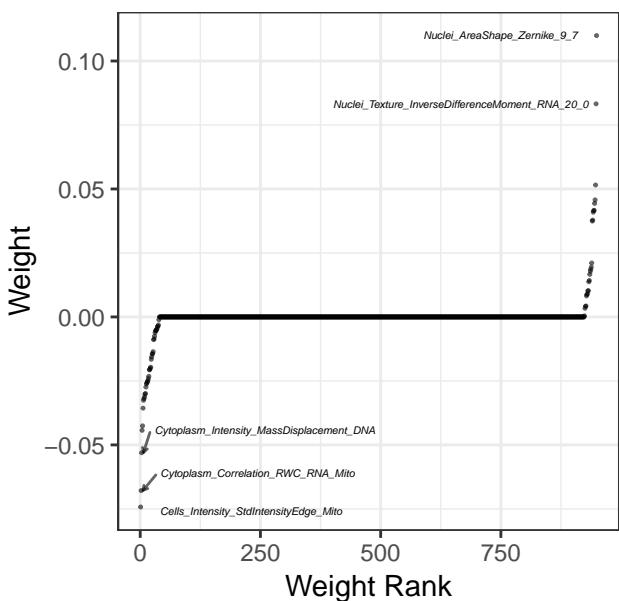


Performance: cc_cc_high_h2ax

Transform: raw

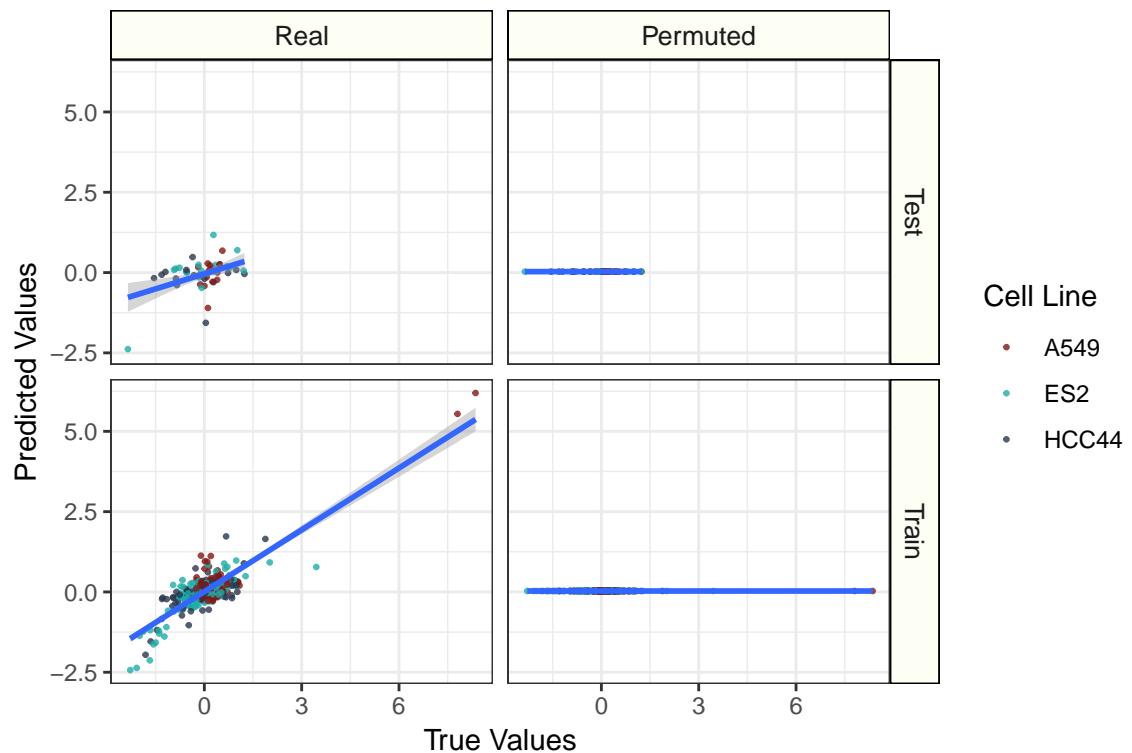


fit	shuffle	transform	r2	value
Train	Real	raw	0.58	0.33
Test	Real	raw	0.33	0.32
Train	Shuffle	raw	0.64	0.28
Test	Shuffle	raw	0.13	0.41

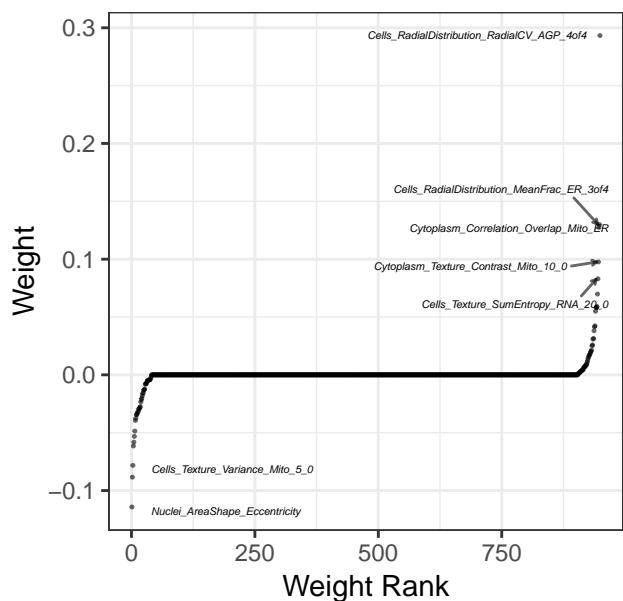


Performance: cc_cc_late_mitosis

Transform: raw

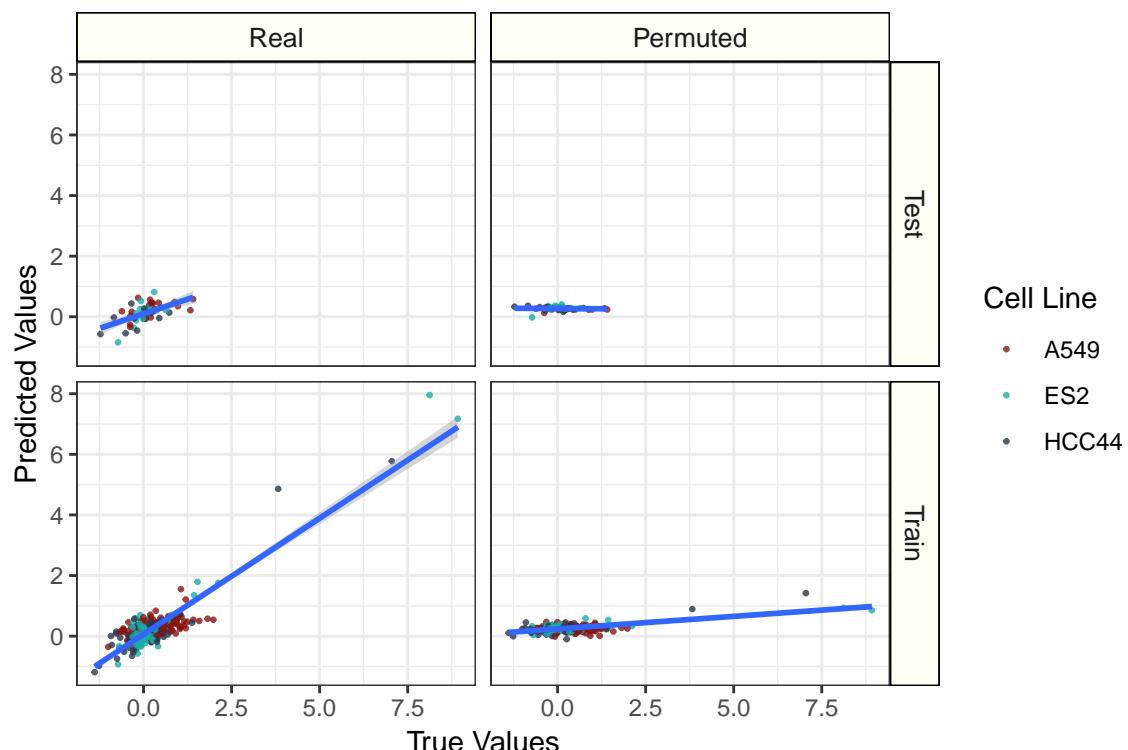


fit	shuffle	transform	r2	value
Train	Real	raw	0.72	0.24
Test	Real	raw	0.09	0.43
Train	Shuffle	raw	0.00	0.86
Test	Shuffle	raw	-0.01	0.47

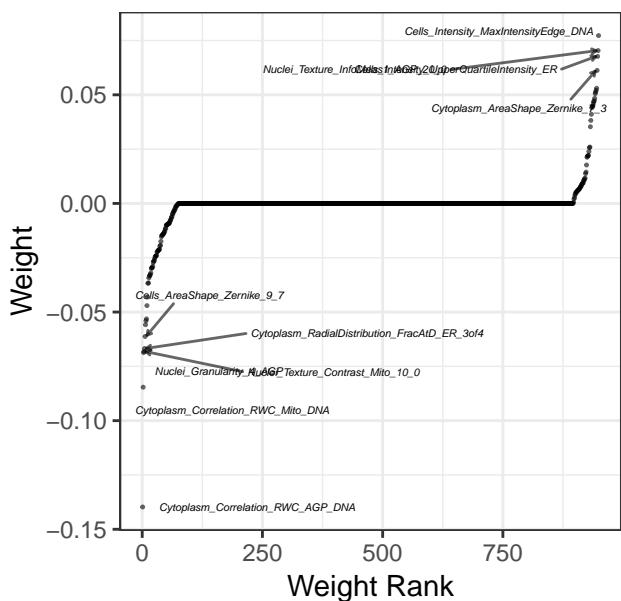


Performance: cc_cc_mitosis

Transform: raw

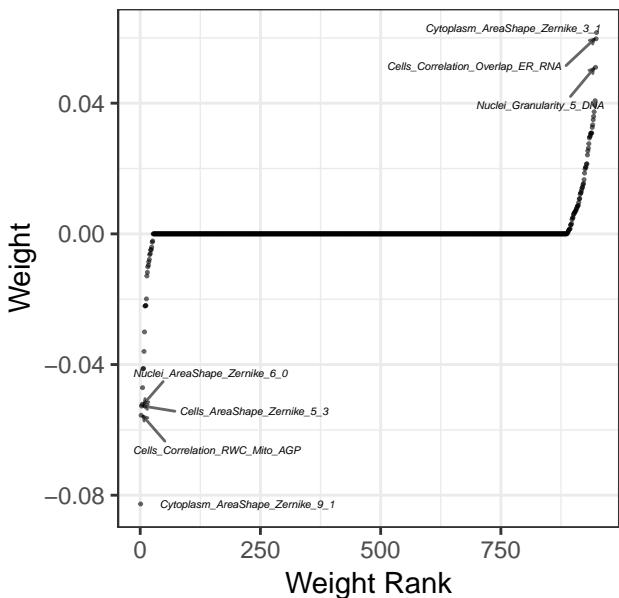
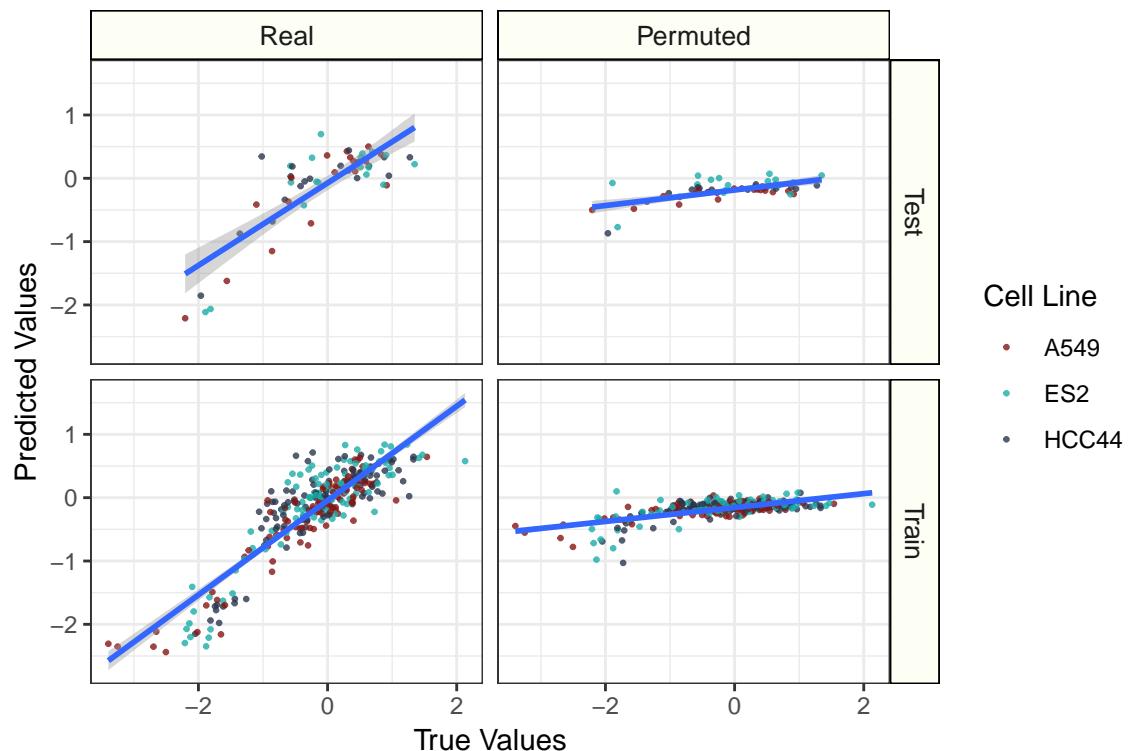


fit	shuffle	transform	r2	value
Train	Real	raw	0.82	0.17
Test	Real	raw	0.31	0.15
Train	Shuffle	raw	0.15	0.78
Test	Shuffle	raw	-0.21	0.27



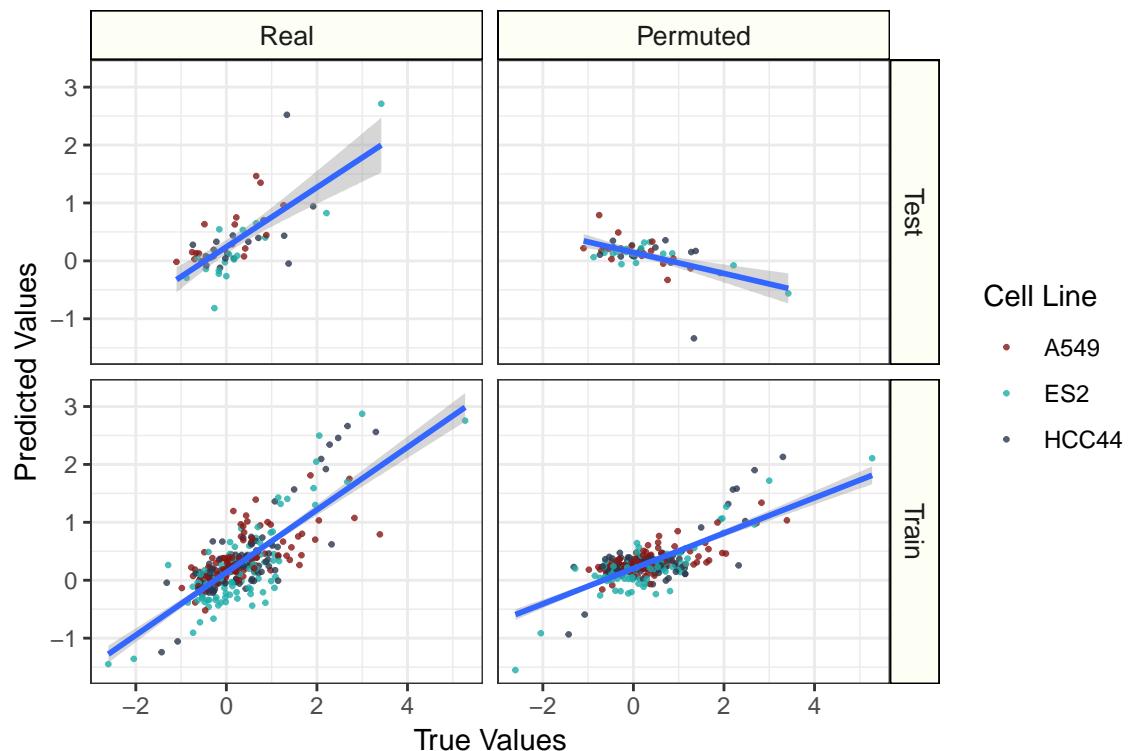
Performance: cc_cc_n_objects

Transform: raw

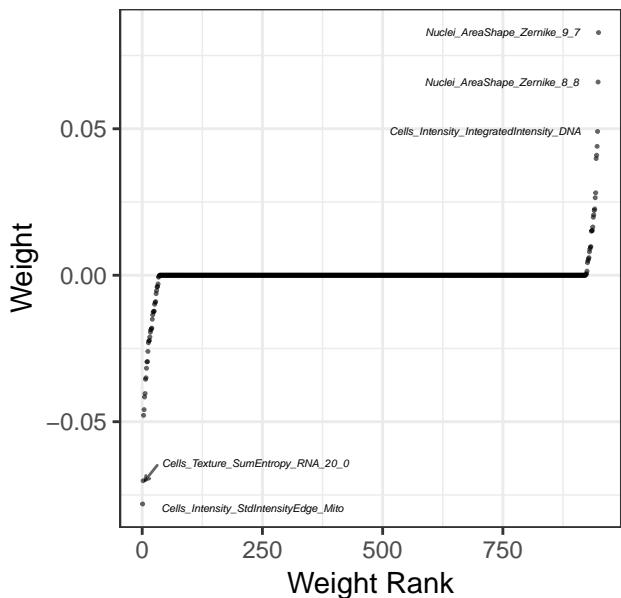


Performance: cc_cc_n_spots_h2ax_mean

Transform: raw

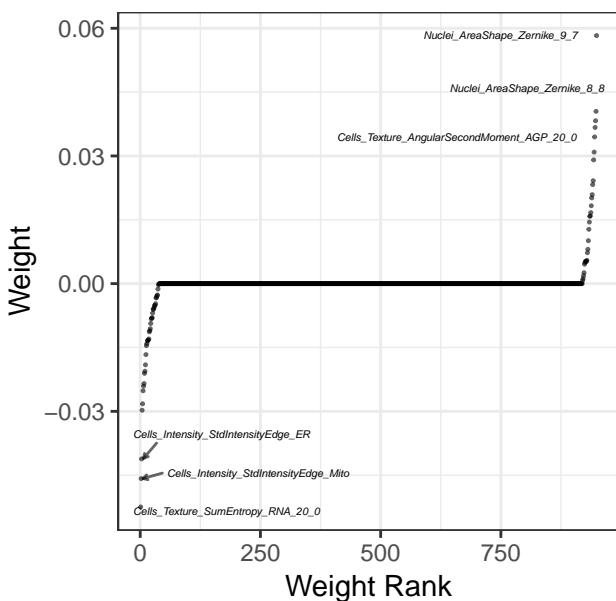
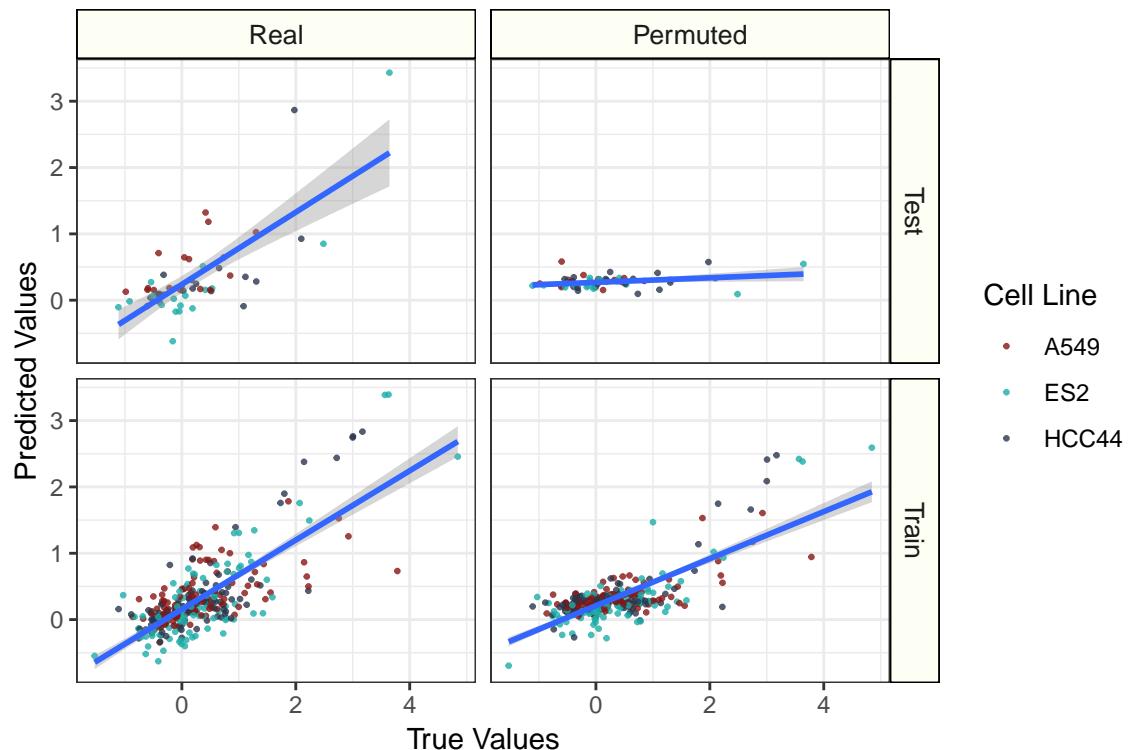


fit	shuffle	transform	r2	value
Train	Real	raw	0.60	0.30
Test	Real	raw	0.47	0.36
Train	Shuffle	raw	0.45	0.42
Test	Shuffle	raw	-0.48	1.01



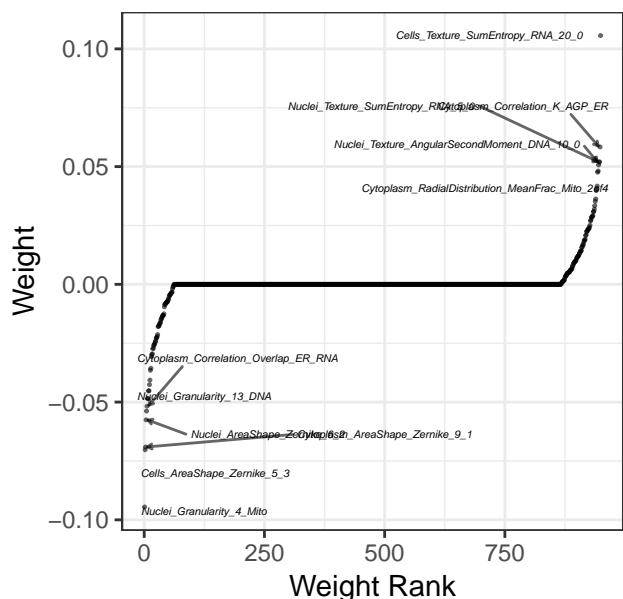
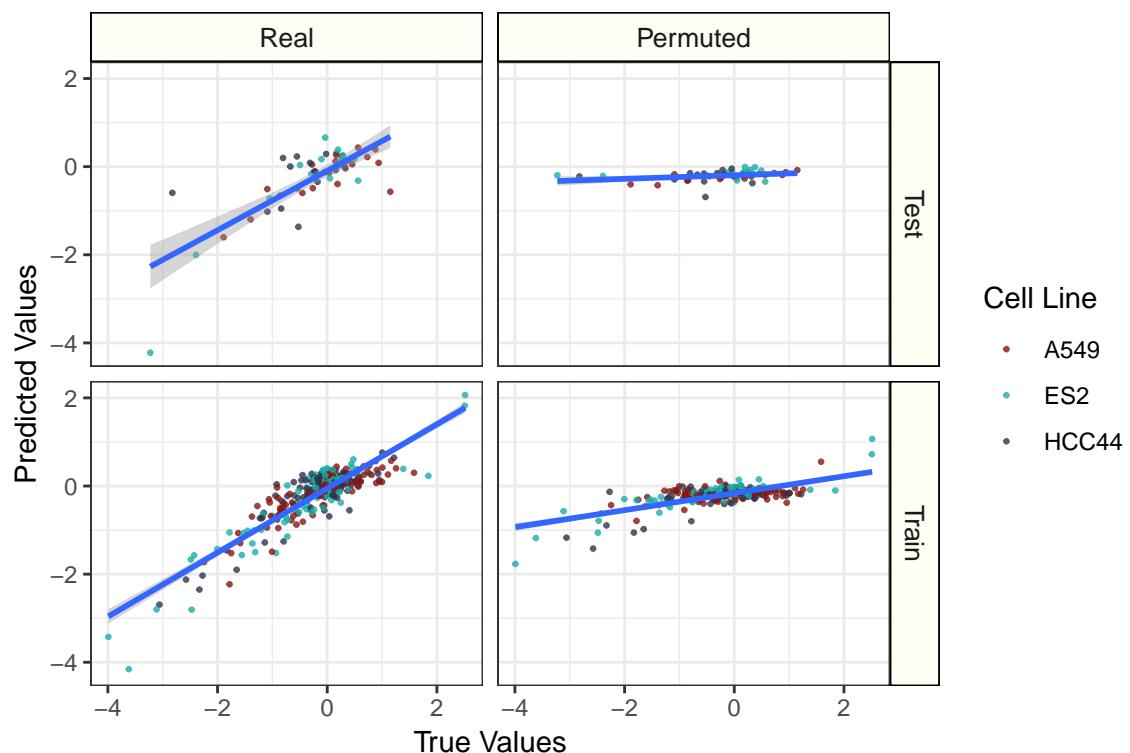
Performance: cc_cc_n_spots_h2ax_per_nucleus_area_mean

Transform: raw



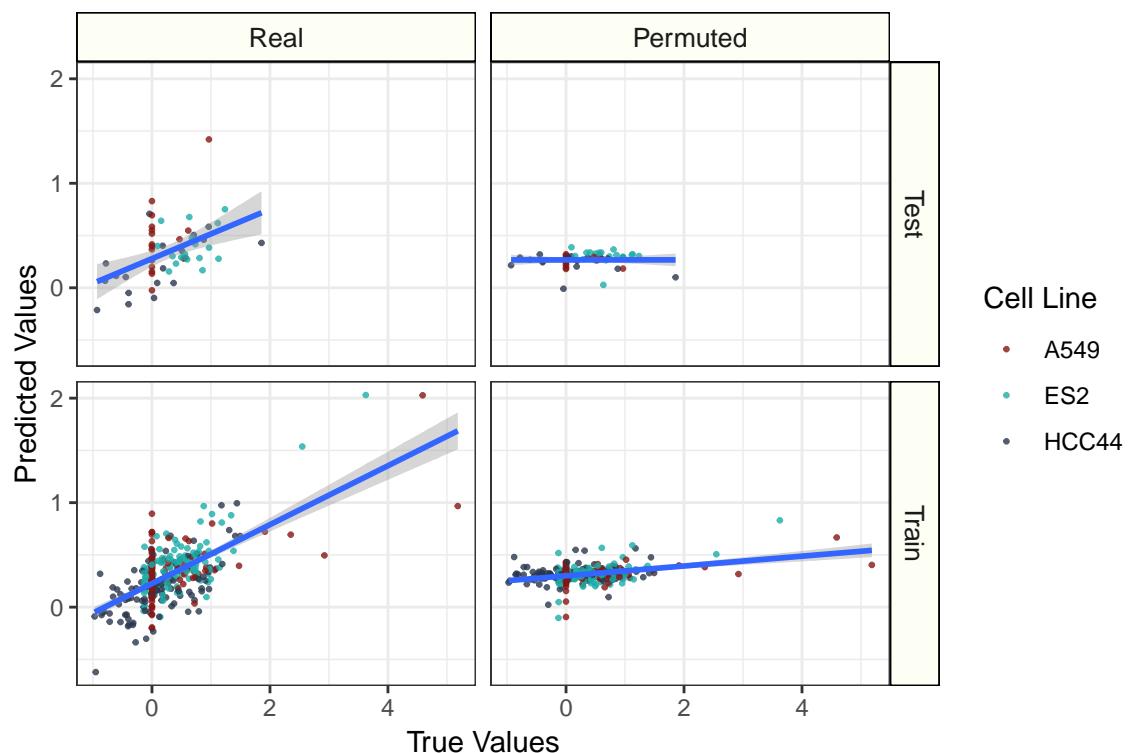
Performance: cc_cc_s

Transform: raw

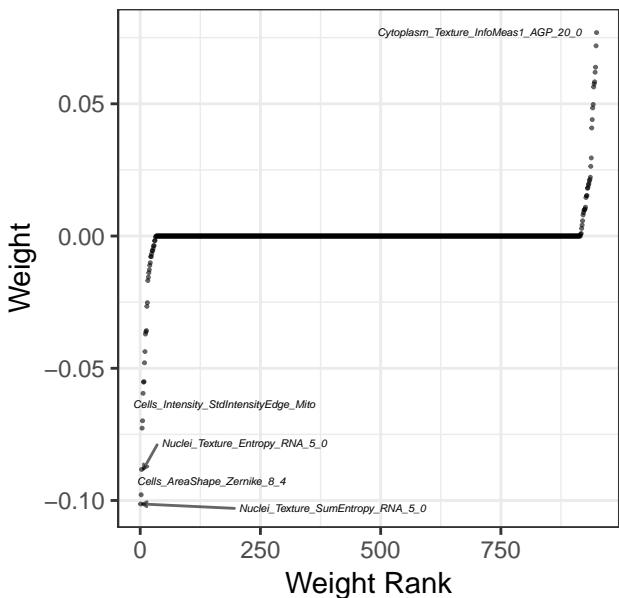


Performance: cc_early_mitosis_high_h2ax

Transform: raw

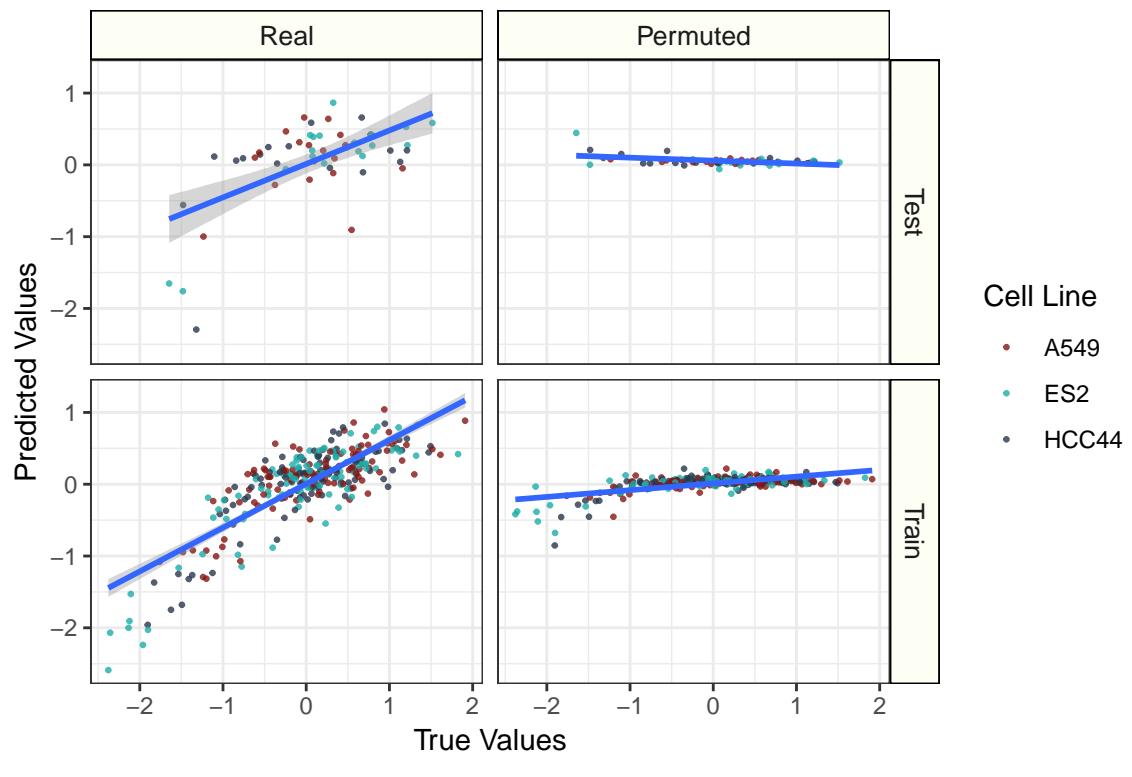


fit	shuffle	transform	r2	value
Train	Real	raw	0.38	0.28
Test	Real	raw	0.21	0.23
Train	Shuffle	raw	0.08	0.42
Test	Shuffle	raw	-0.02	0.30

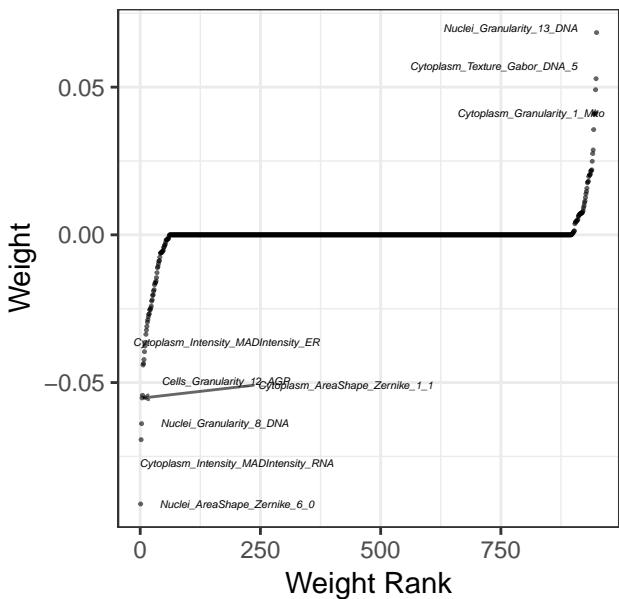


Performance: cc_early_mitosis_n_objects

Transform: raw

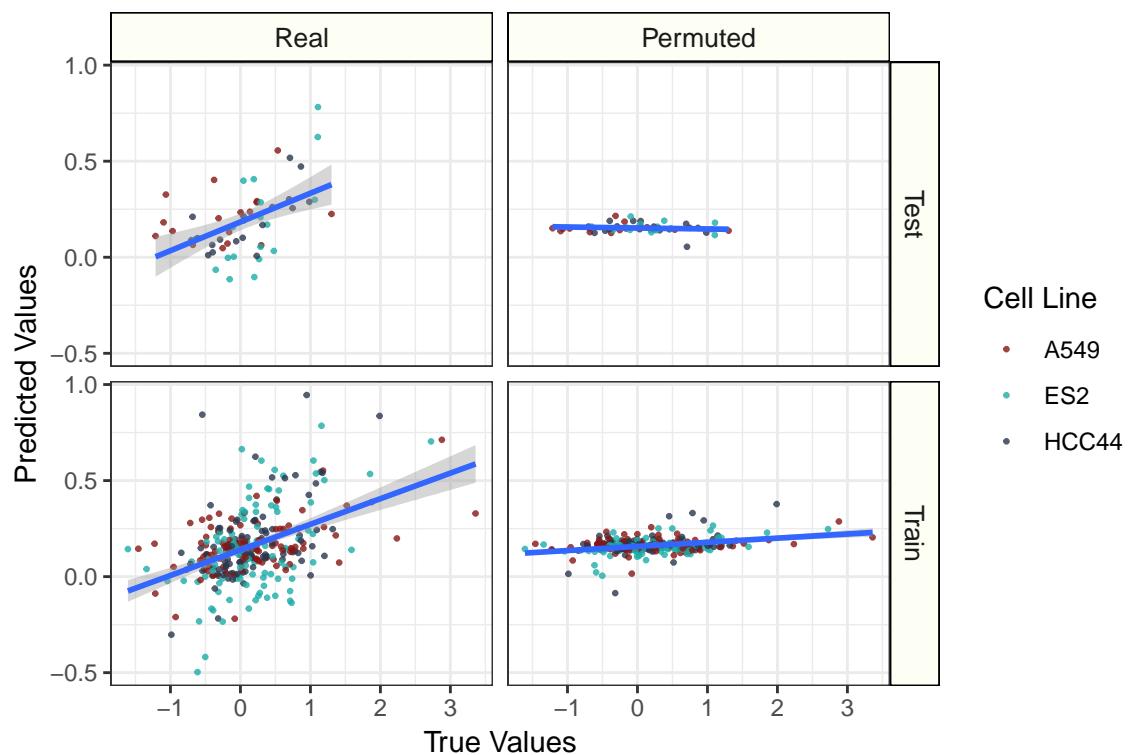


fit	shuffle	transform	r2	value
Train	Real	raw	0.66	0.20
Test	Real	raw	0.31	0.38
Train	Shuffle	raw	0.16	0.48
Test	Shuffle	raw	-0.09	0.61

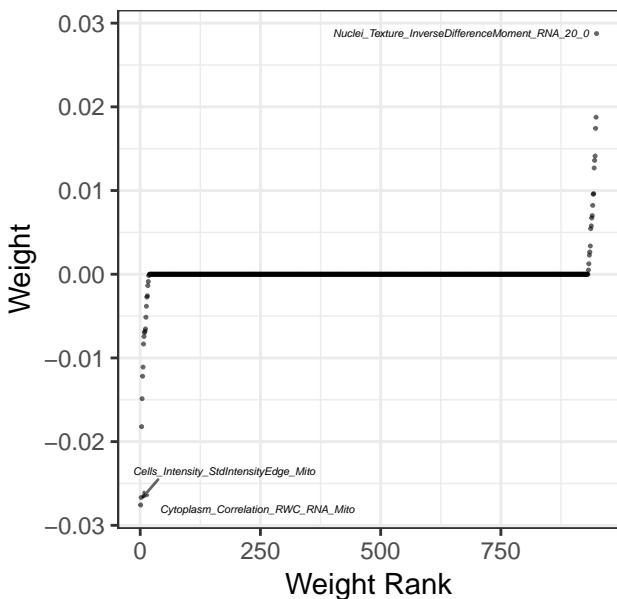


Performance: cc_early_mitosis_n_spots_h2ax_mean

Transform: raw

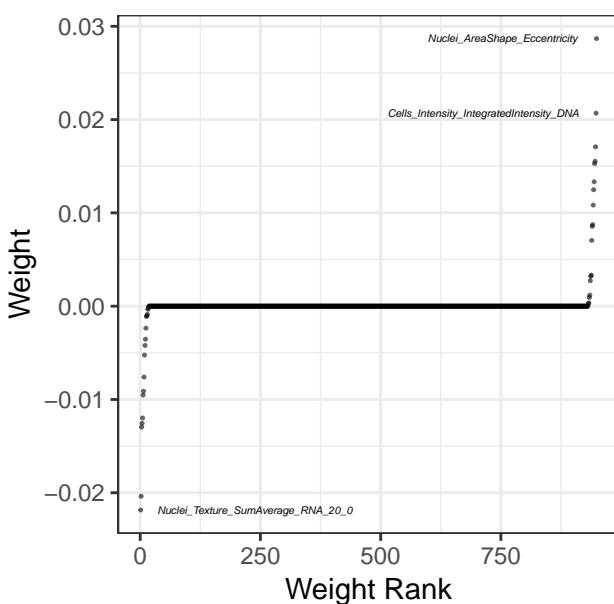
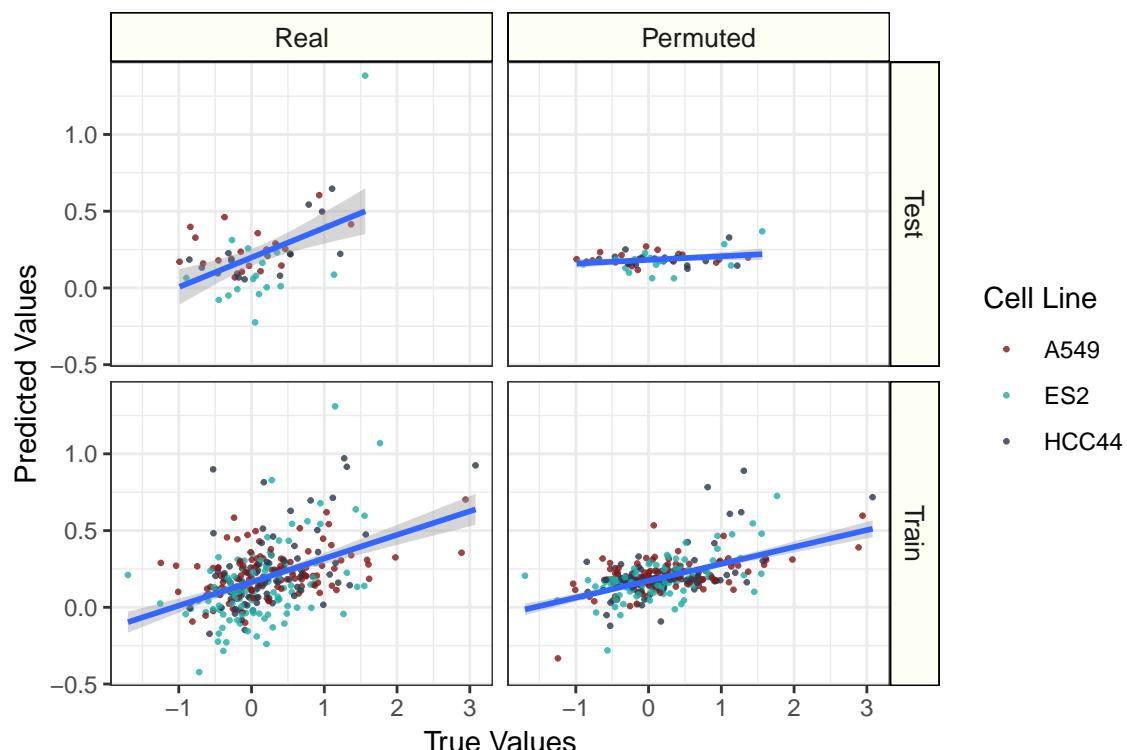


fit	shuffle	transform	r2	value
Train	Real	raw	0.18	0.34
Test	Real	raw	0.14	0.29
Train	Shuffle	raw	0.04	0.40
Test	Shuffle	raw	-0.05	0.35



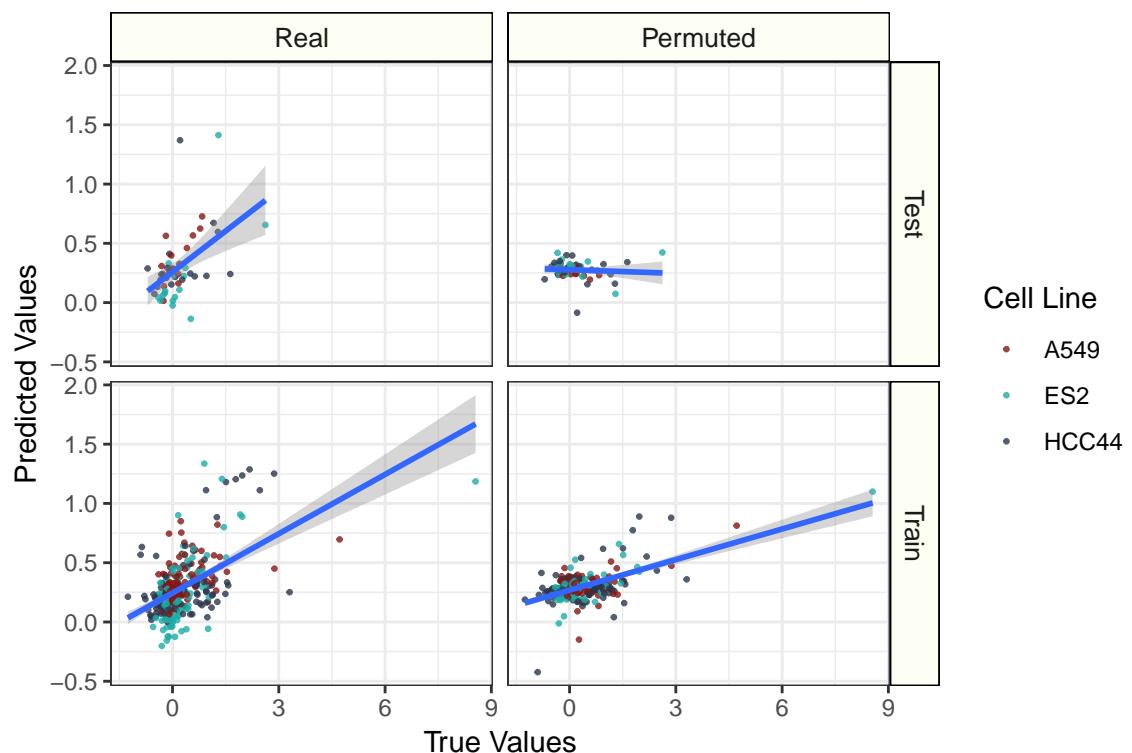
Performance: cc_early_mitosis_n_spots_h2ax_per_nucleus_are

Transform: raw

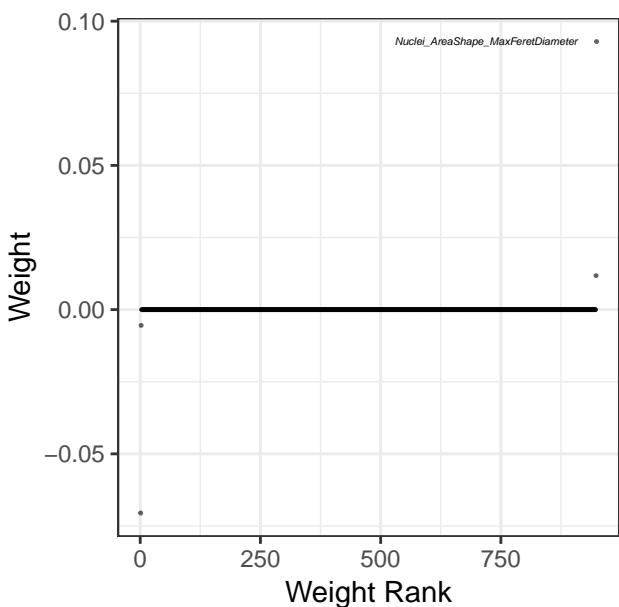


Performance: cc_g1_high_h2ax

Transform: raw

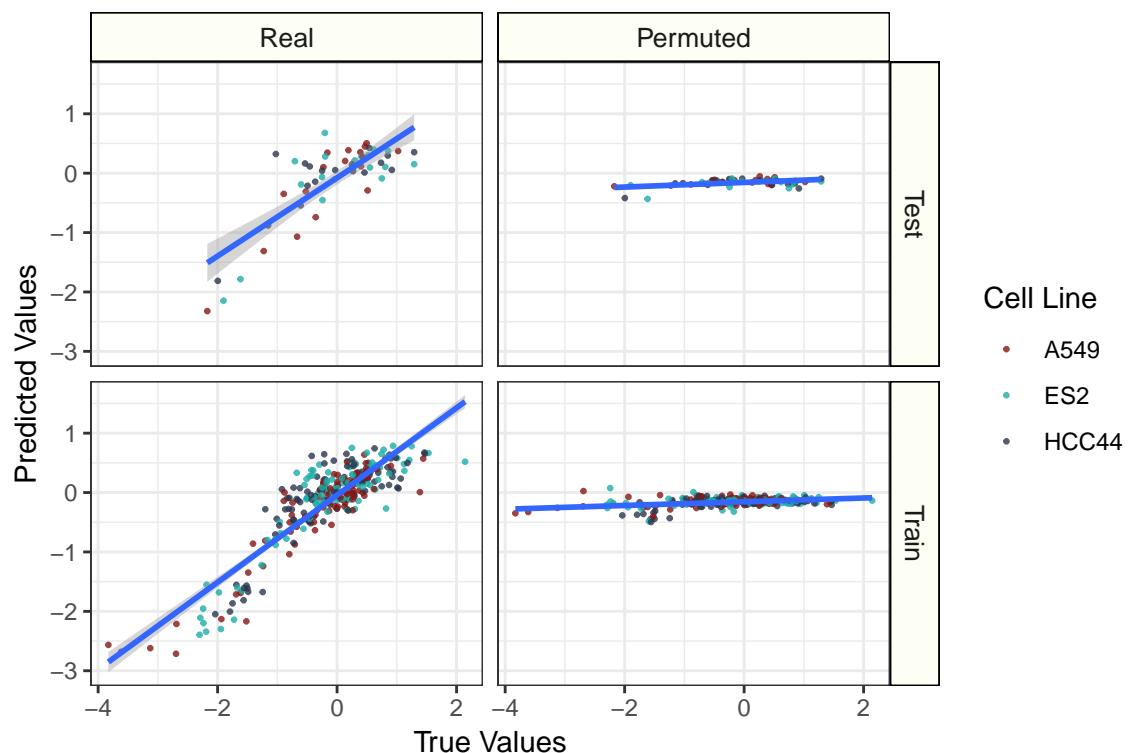


fit	shuffle	transform	r2	value
Train	Real	raw	0.24	0.52
Test	Real	raw	0.19	0.29
Train	Shuffle	raw	0.15	0.58
Test	Shuffle	raw	-0.08	0.38

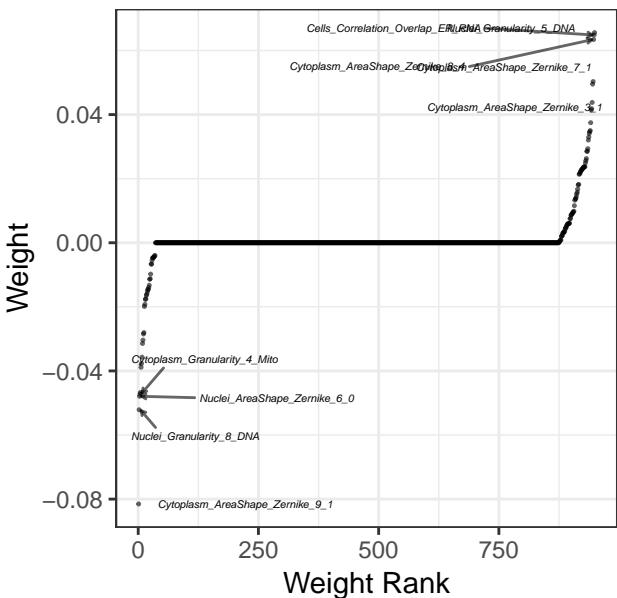


Performance: cc_g1_n_objects

Transform: raw

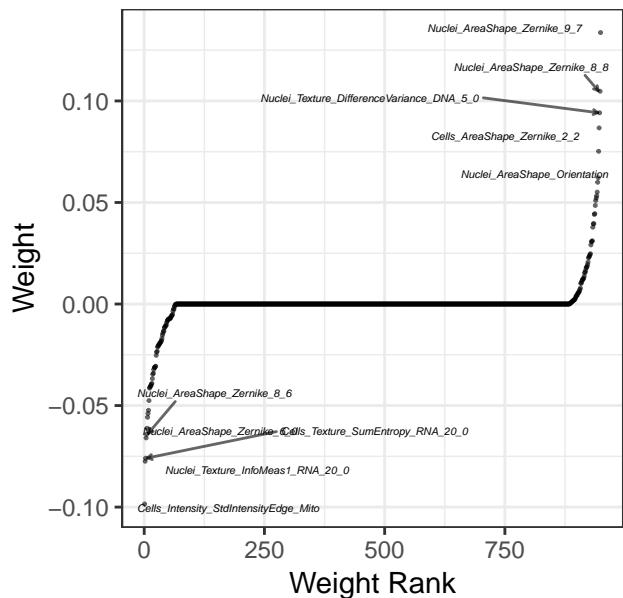
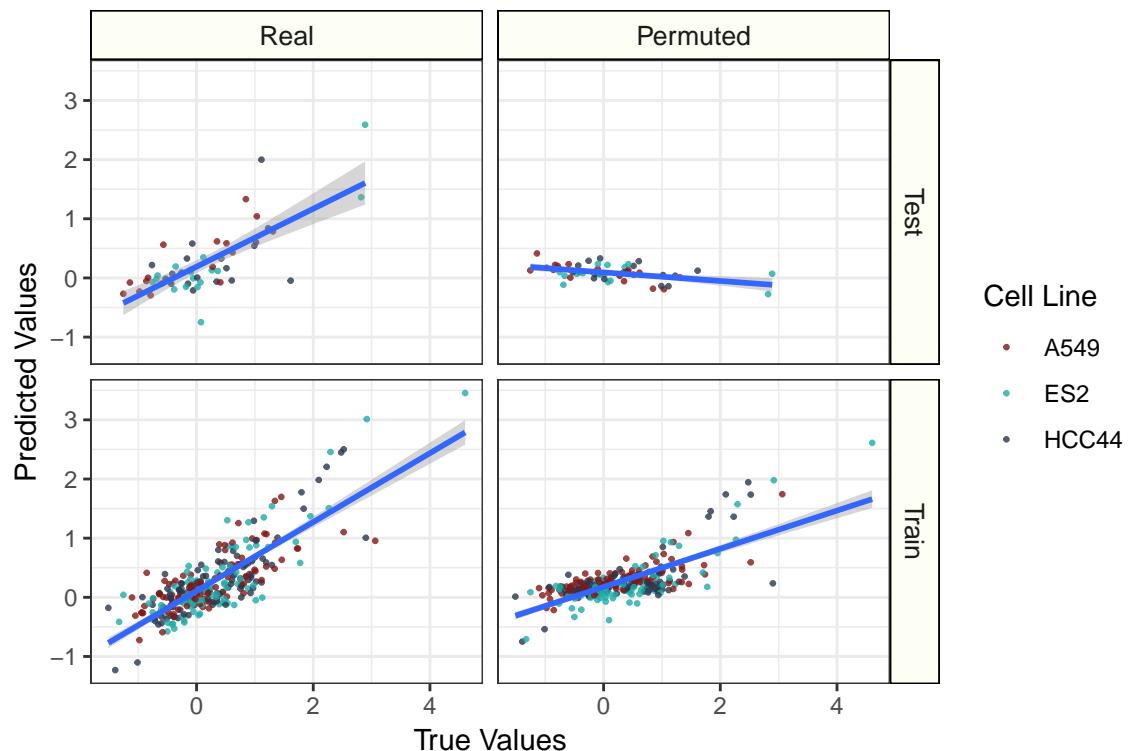


fit	shuffle	transform	r2	value
Train	Real	raw	0.77	0.17
Test	Real	raw	0.62	0.24
Train	Shuffle	raw	0.06	0.70
Test	Shuffle	raw	0.06	0.60



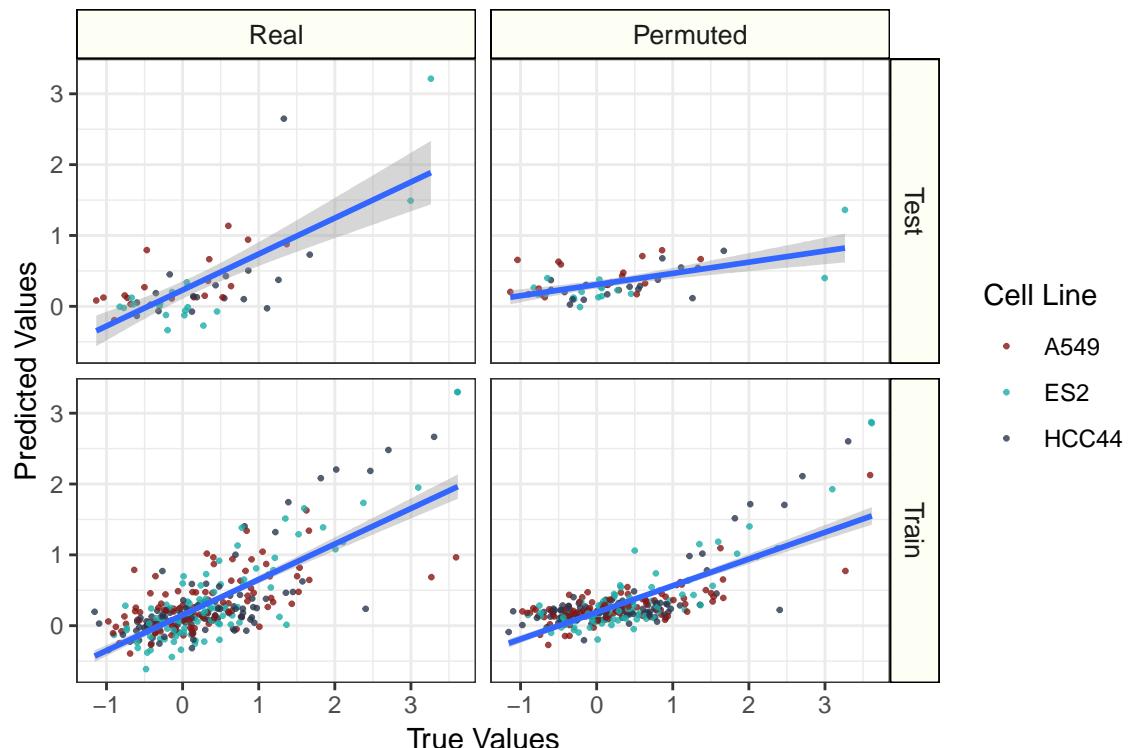
Performance: cc_g1_n_spots_h2ax_mean

Transform: raw

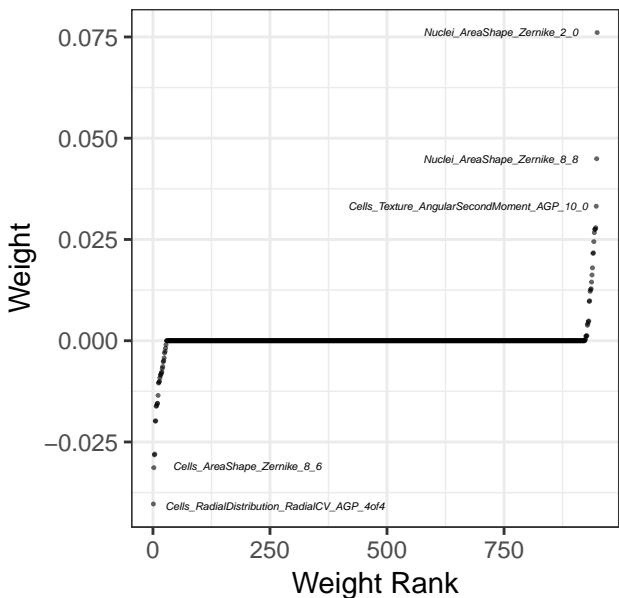


Performance: cc_g1_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

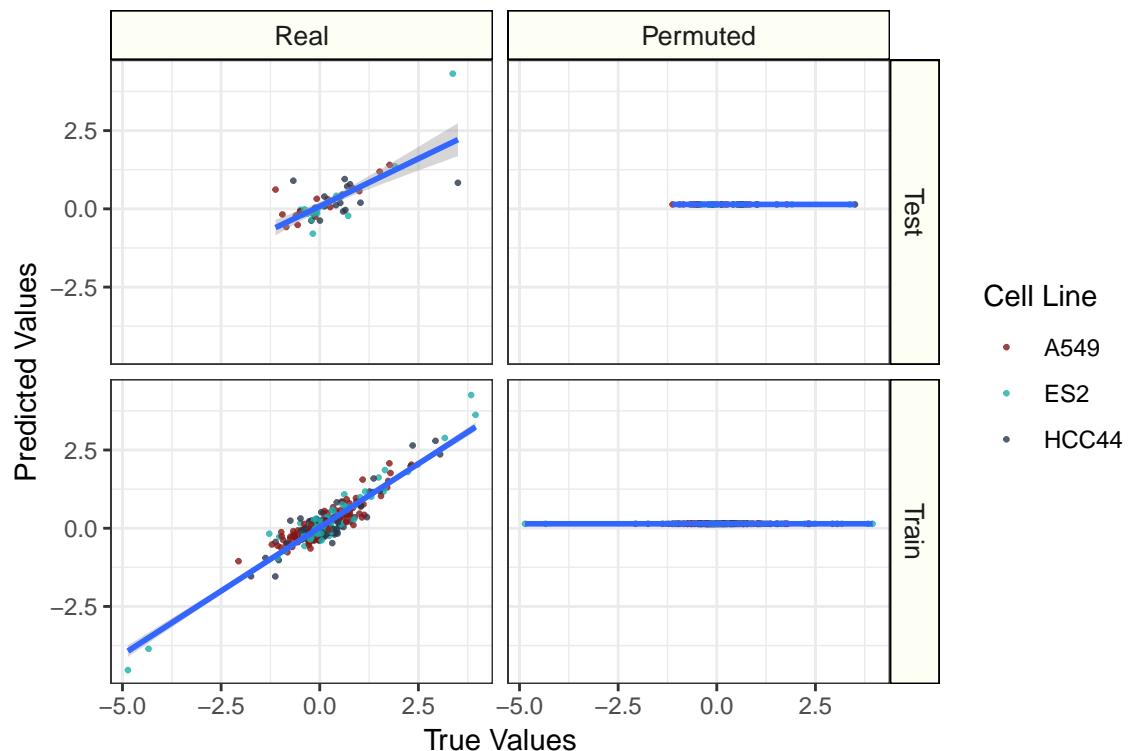


fit	shuffle	transform	r2	value
Train	Real	raw	0.55	0.29
Test	Real	raw	0.48	0.40
Train	Shuffle	raw	0.51	0.32
Test	Shuffle	raw	0.20	0.61

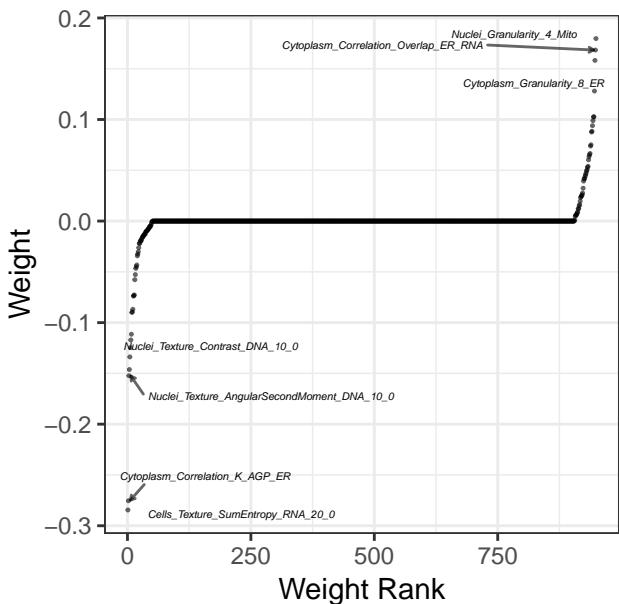


Performance: cc_g1_plus_g2_count

Transform: raw

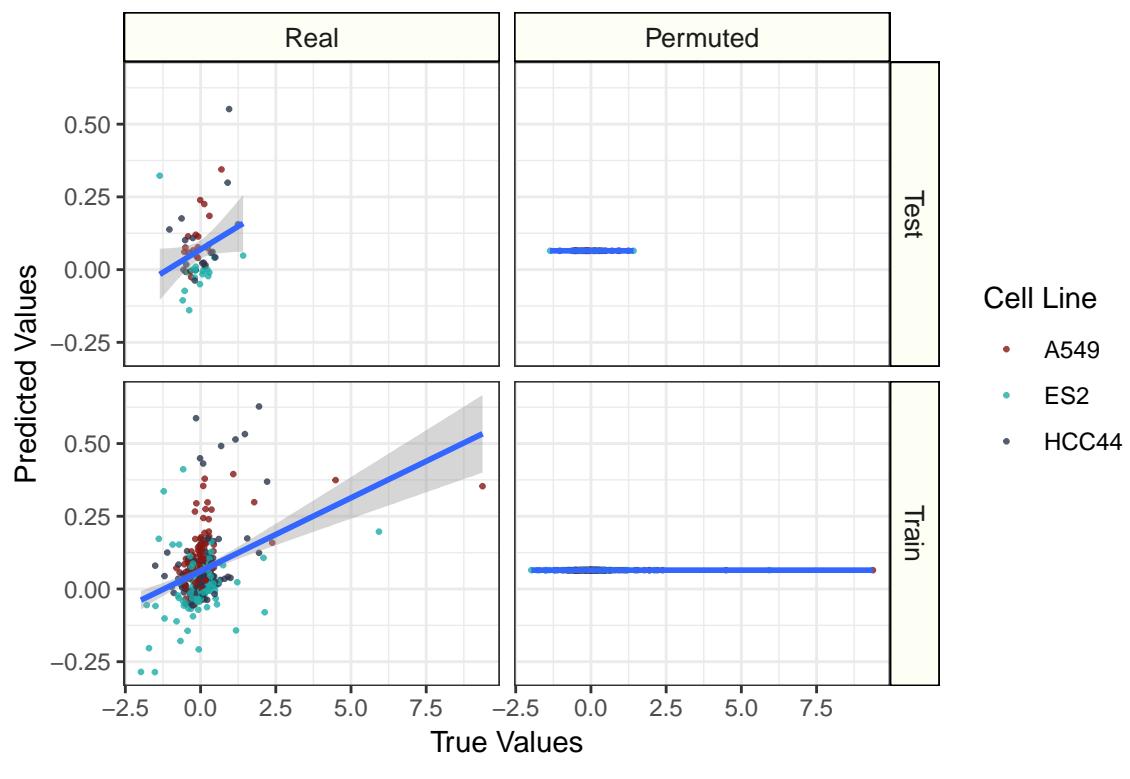


fit	shuffle	transform	r2	value
Train	Real	raw	0.86	0.11
Test	Real	raw	0.53	0.37
Train	Shuffle	raw	0.00	0.78
Test	Shuffle	raw	-0.01	0.79

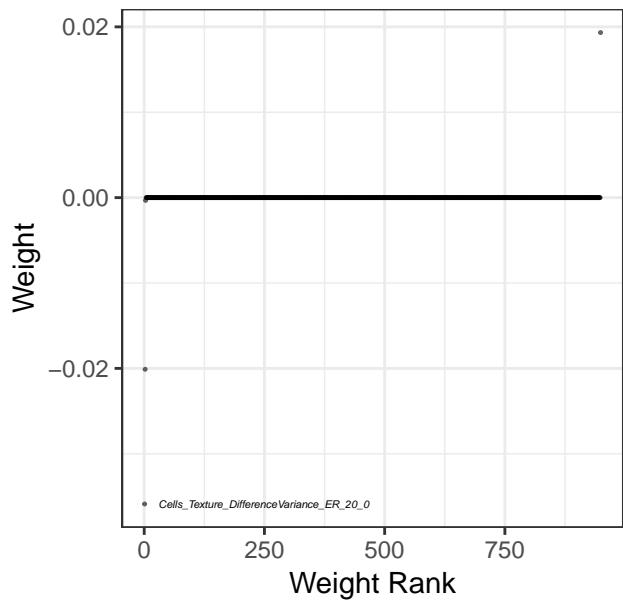


Performance: cc_g2_g1_count

Transform: raw

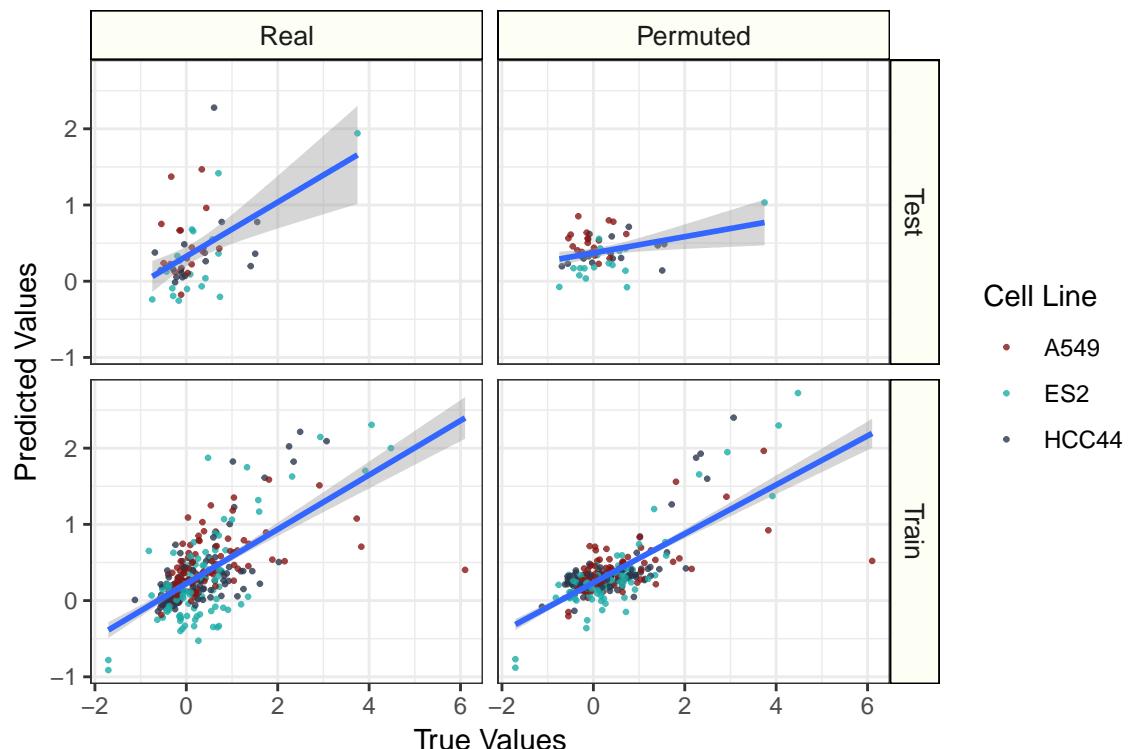


fit	shuffle	transform	r2	value
Train	Real	raw	0.08	0.72
Test	Real	raw	0.02	0.25
Train	Shuffle	raw	0.00	0.79
Test	Shuffle	raw	-0.05	0.27

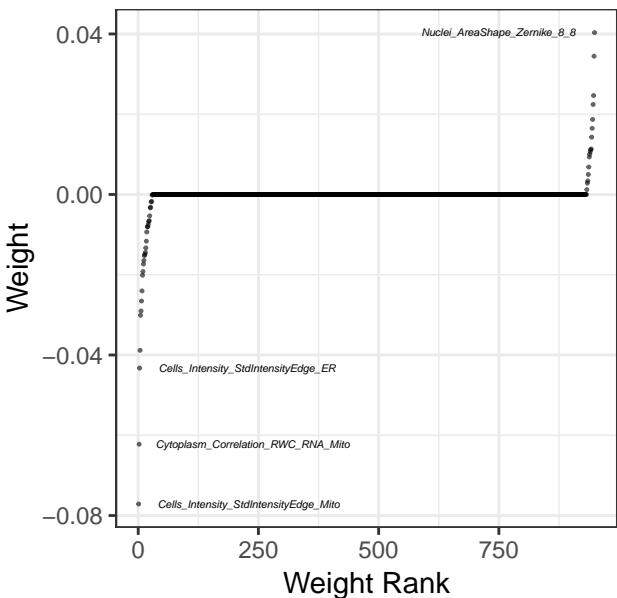


Performance: cc_g2_high_h2ax

Transform: raw

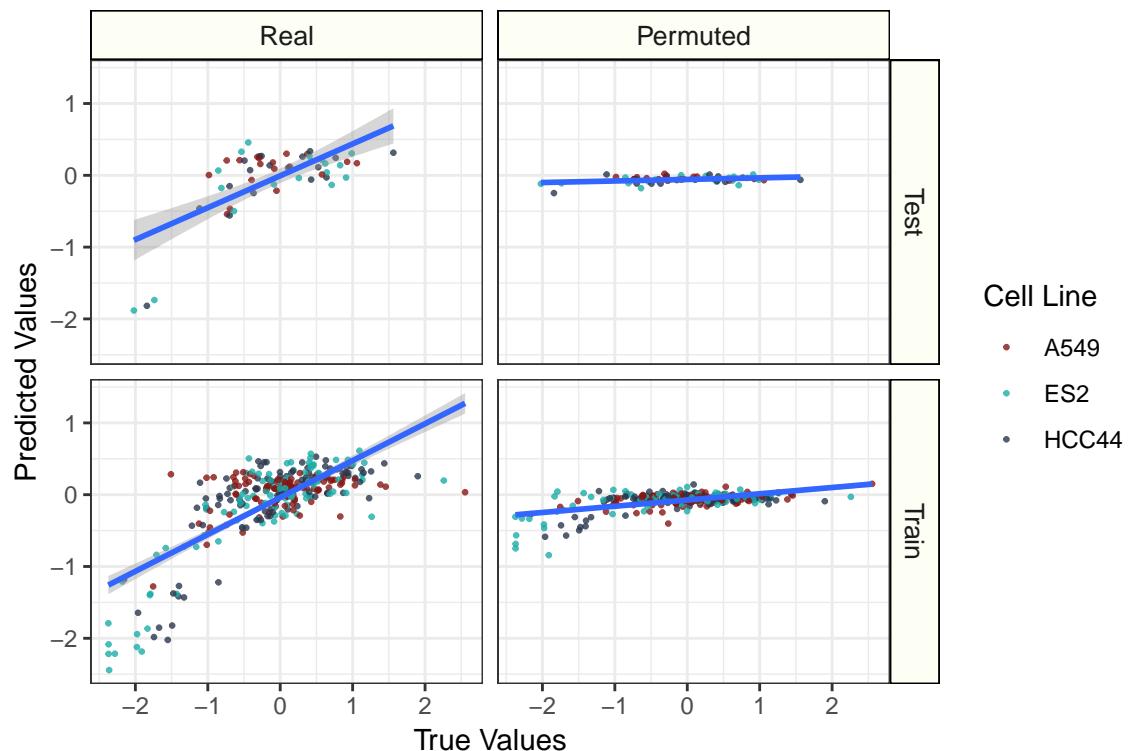


fit	shuffle	transform	r2	value
Train	Real	raw	0.41	0.46
Test	Real	raw	0.09	0.45
Train	Shuffle	raw	0.46	0.43
Test	Shuffle	raw	0.02	0.49

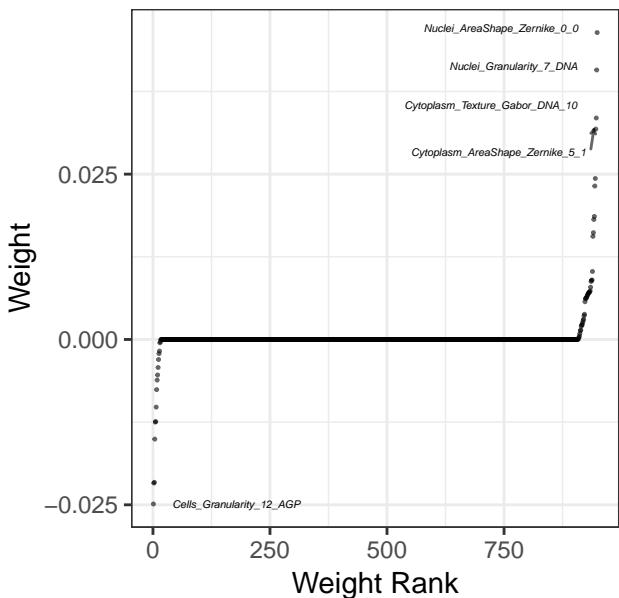


Performance: cc_g2_n_objects

Transform: raw

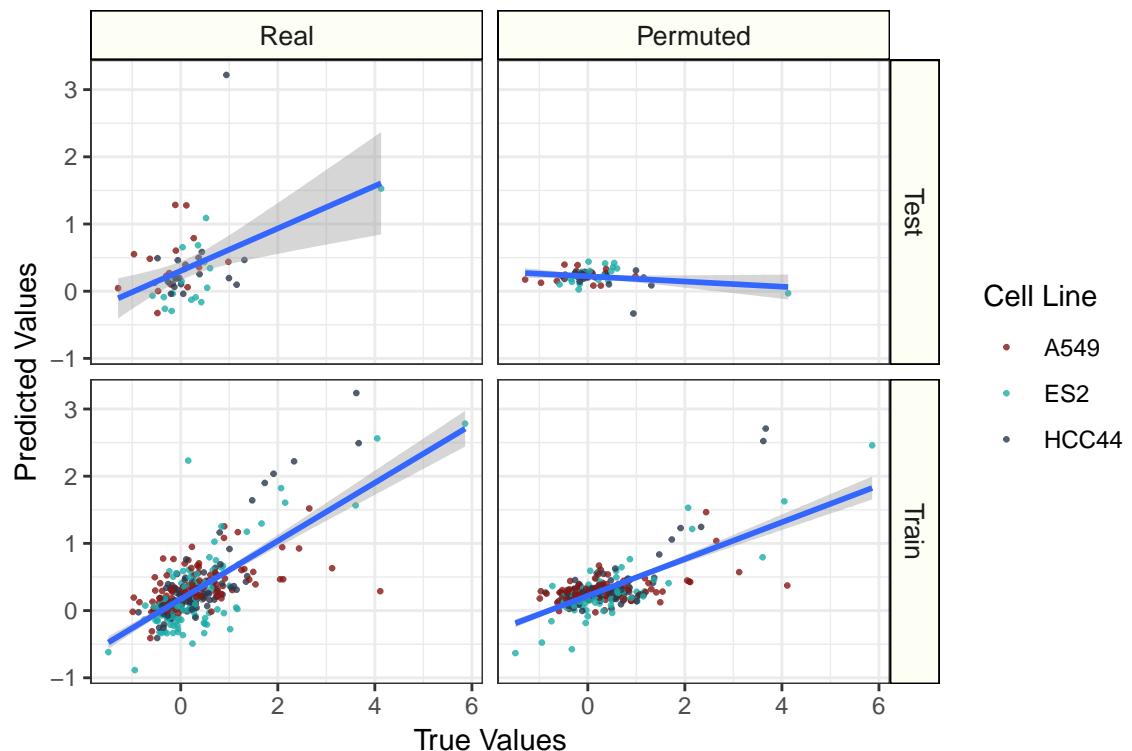


fit	shuffle	transform	r2	value
Train	Real	raw	0.55	0.29
Test	Real	raw	0.45	0.30
Train	Shuffle	raw	0.15	0.56
Test	Shuffle	raw	0.04	0.53

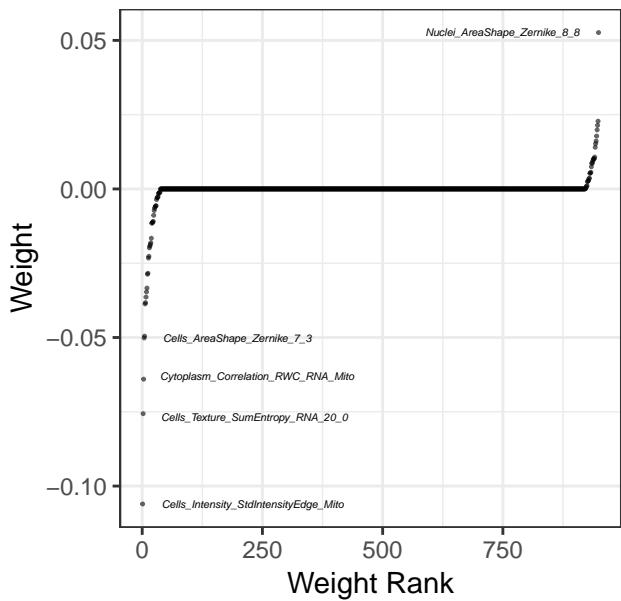


Performance: cc_g2_n_spots_h2ax_mean

Transform: raw

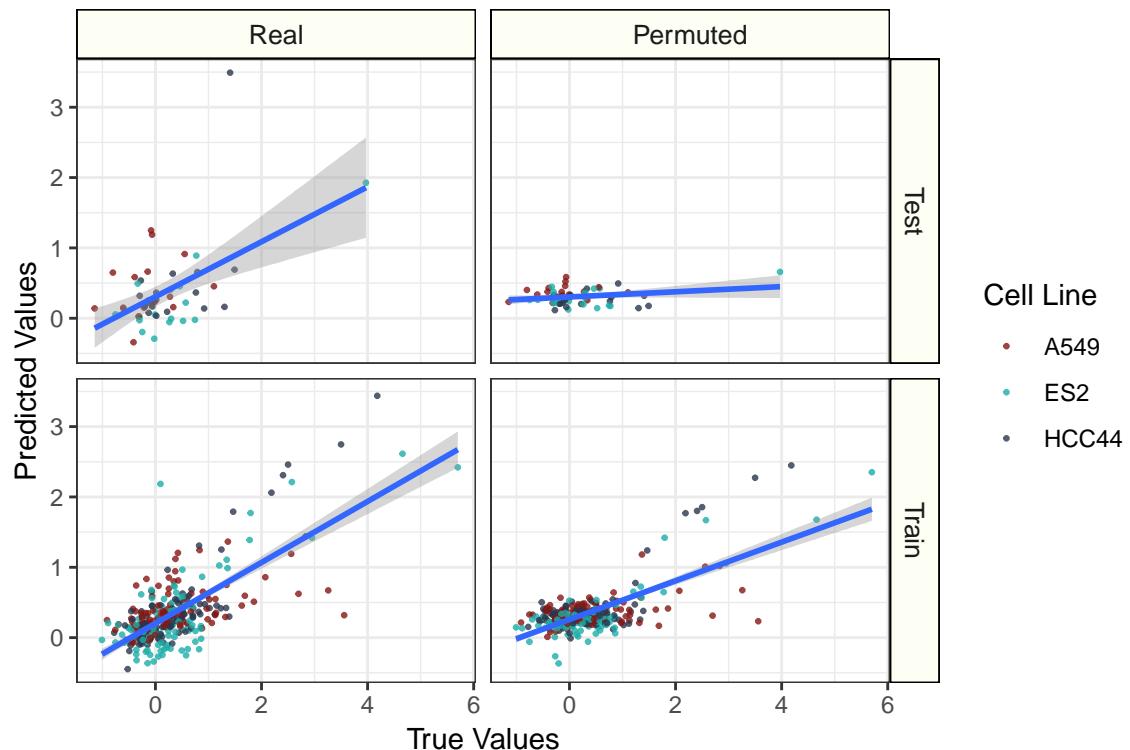


fit	shuffle	transform	r2	value
Train	Real	raw	0.50	0.36
Test	Real	raw	-0.01	0.55
Train	Shuffle	raw	0.40	0.43
Test	Shuffle	raw	-0.12	0.62

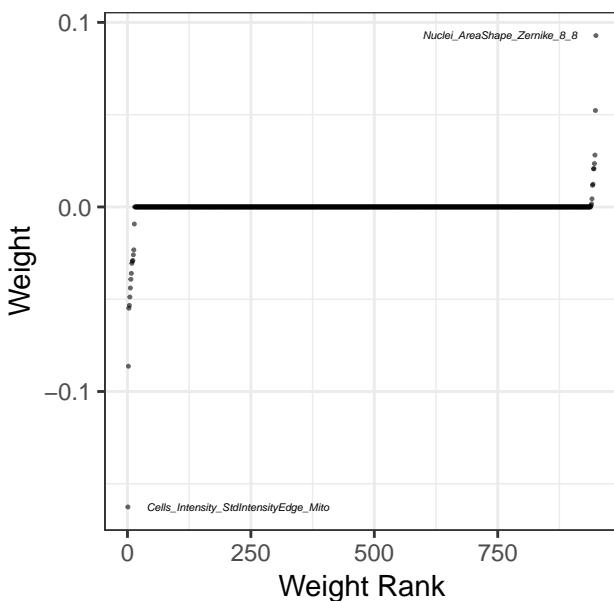


Performance: cc_g2_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

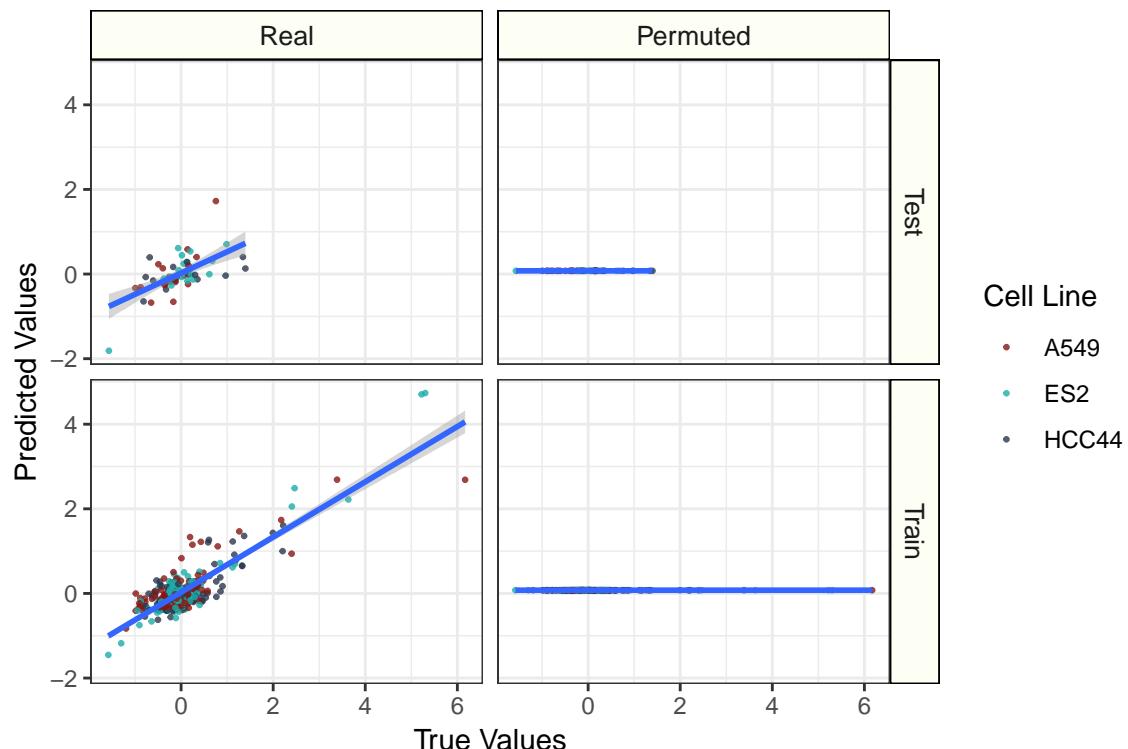


fit	shuffle	transform	r2	value
Train	Real	raw	0.50	0.36
Test	Real	raw	0.12	0.50
Train	Shuffle	raw	0.40	0.43
Test	Shuffle	raw	0.01	0.56

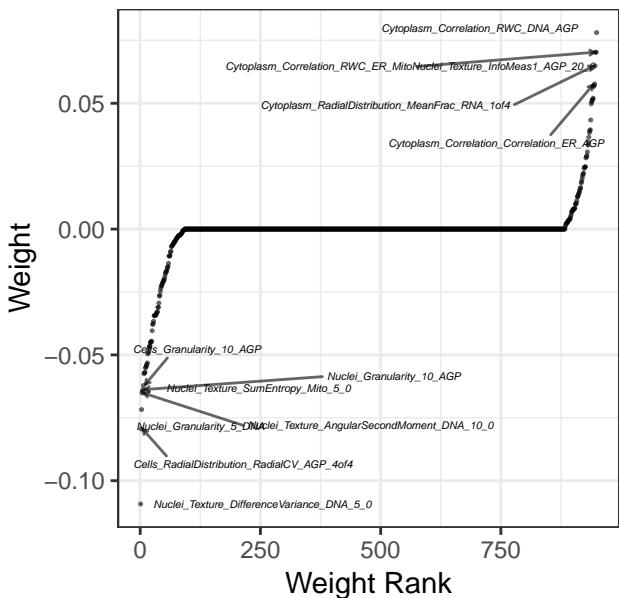


Performance: cc_g2_plus_all_m_count

Transform: raw

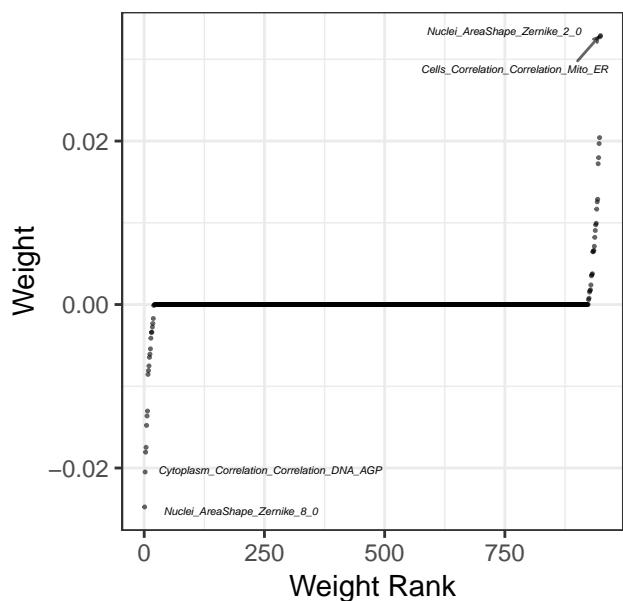
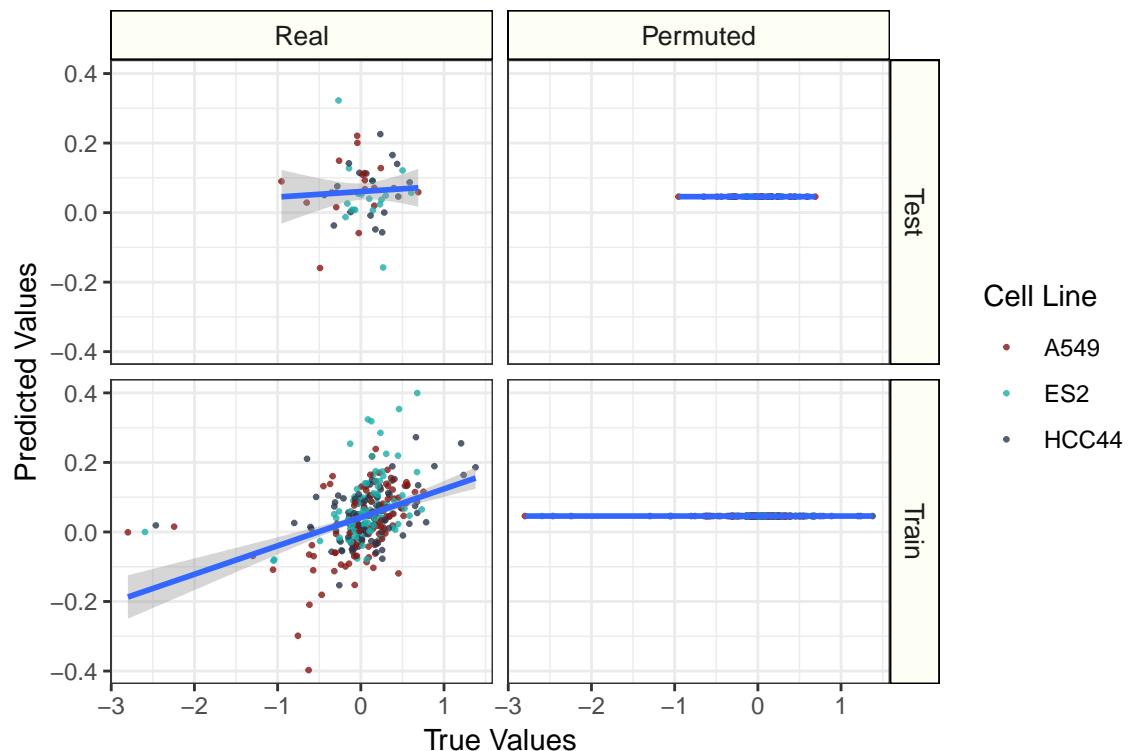


fit	shuffle	transform	r2	value
Train	Real	raw	0.73	0.19
Test	Real	raw	0.32	0.21
Train	Shuffle	raw	0.00	0.70
Test	Shuffle	raw	-0.04	0.32



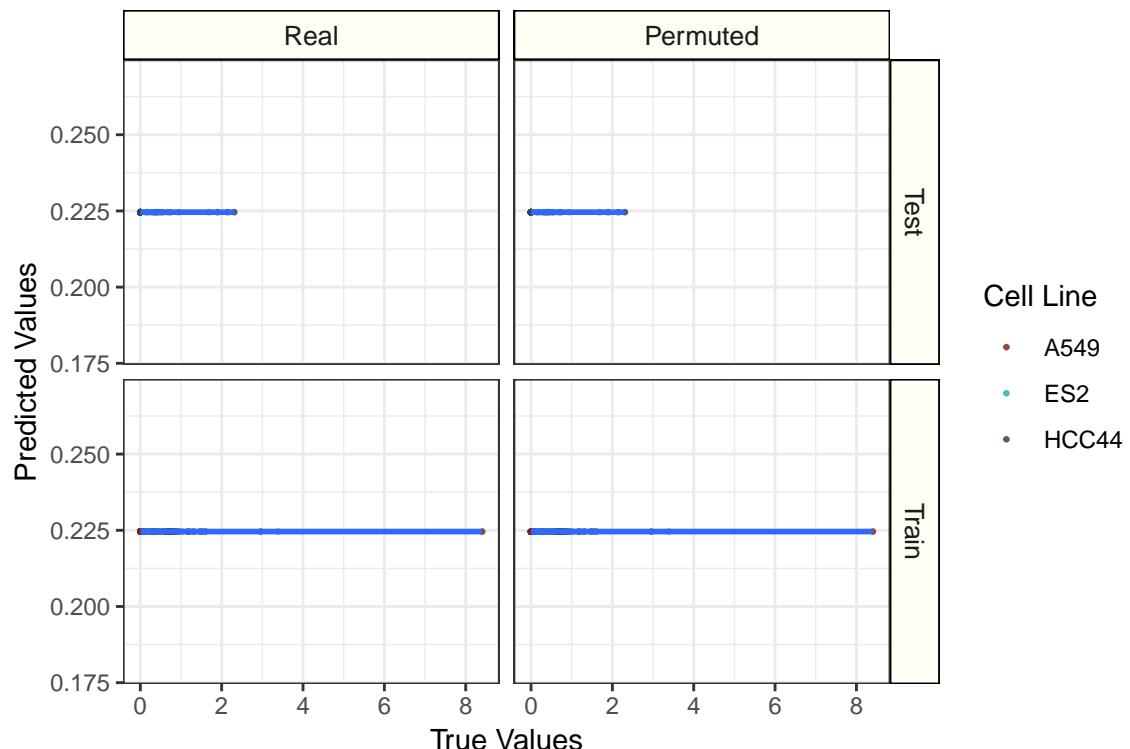
Performance: cc_infection_percentage

Transform: raw

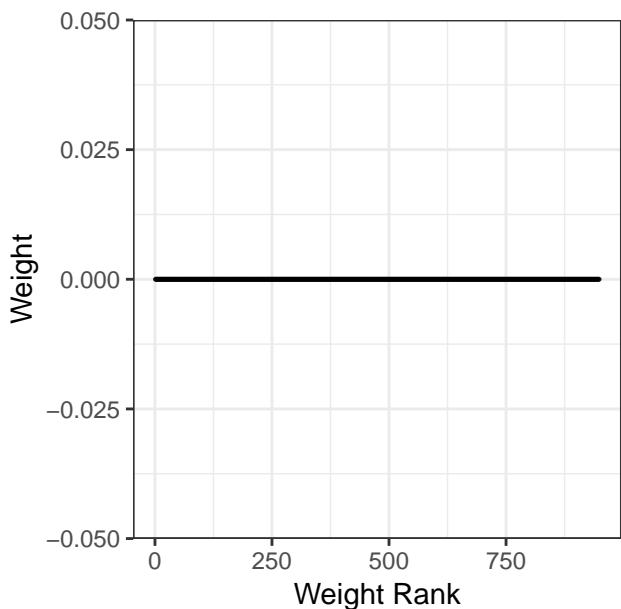


Performance: cc_late_mitosis_high_h2ax

Transform: raw

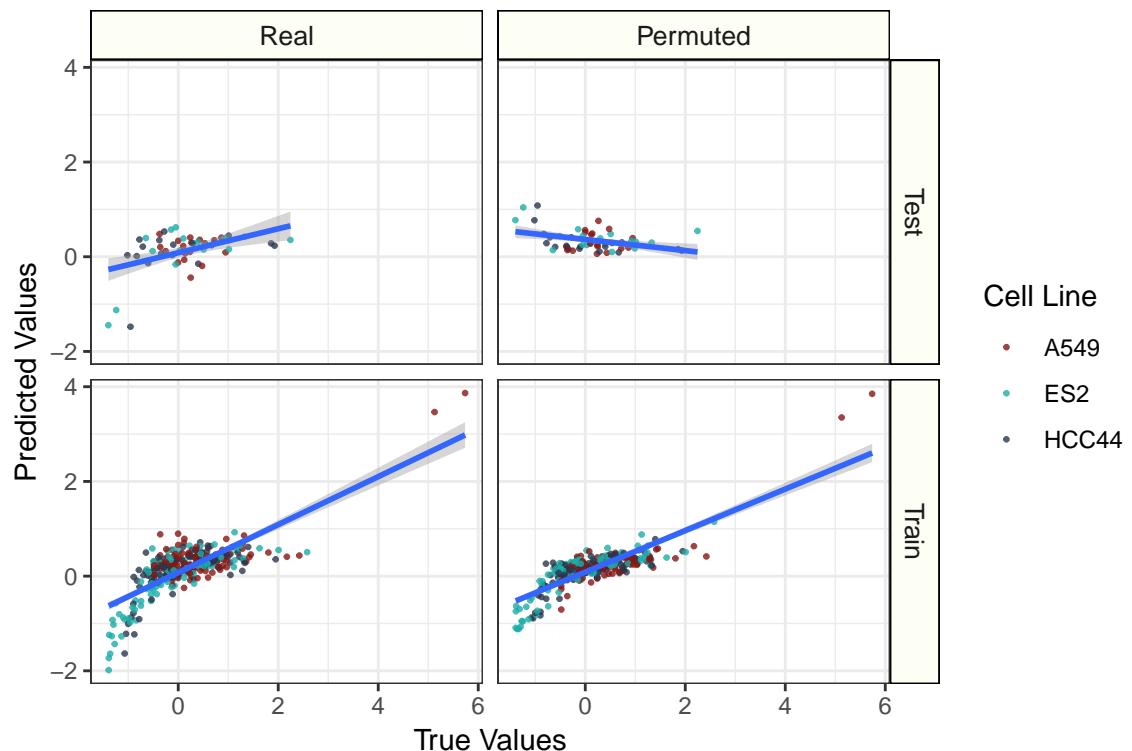


fit	shuffle	transform	r2	value
Train	Real	raw	0.00	0.39
Test	Real	raw	-0.01	0.30
Train	Shuffle	raw	0.00	0.39
Test	Shuffle	raw	-0.01	0.30

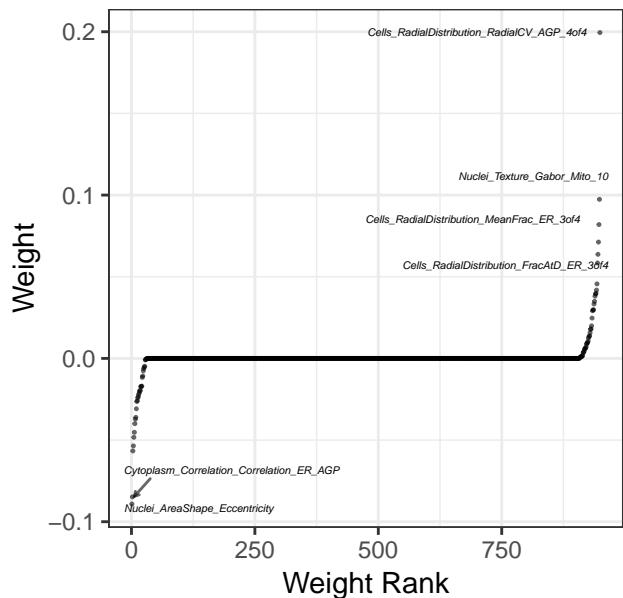


Performance: cc_late_mitosis_n_objects

Transform: raw

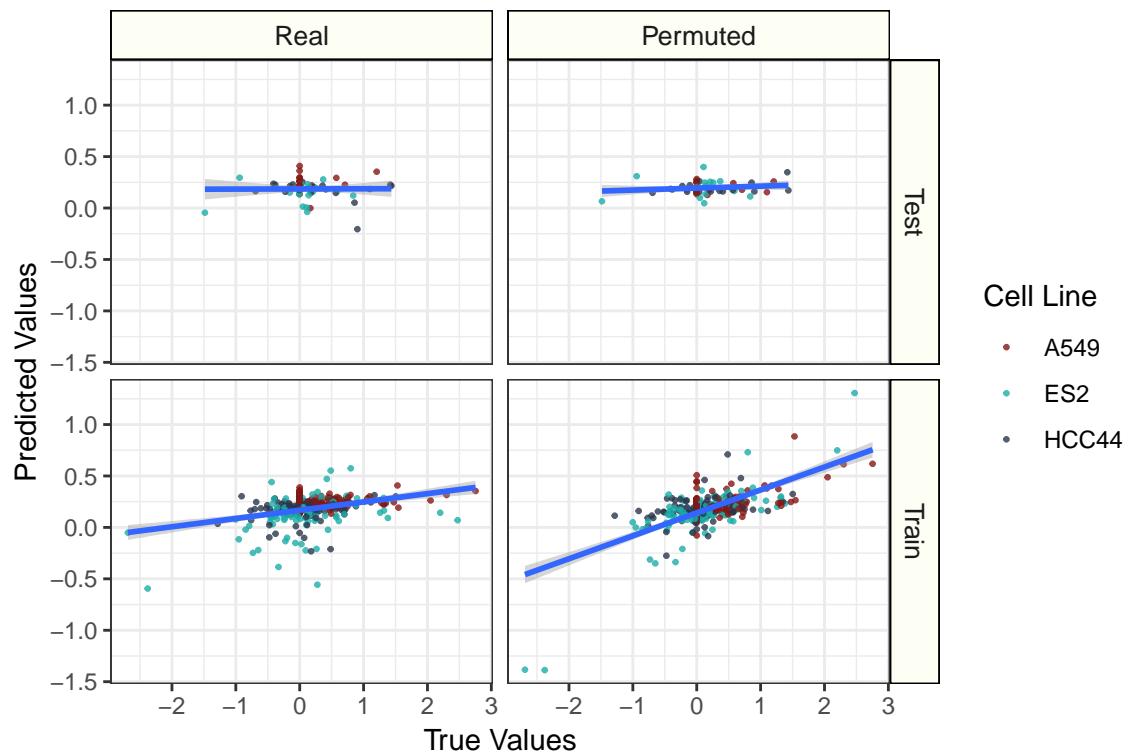


fit	shuffle	transform	r2	value
Train	Real	raw	0.58	0.29
Test	Real	raw	0.20	0.45
Train	Shuffle	raw	0.59	0.28
Test	Shuffle	raw	-0.38	0.77

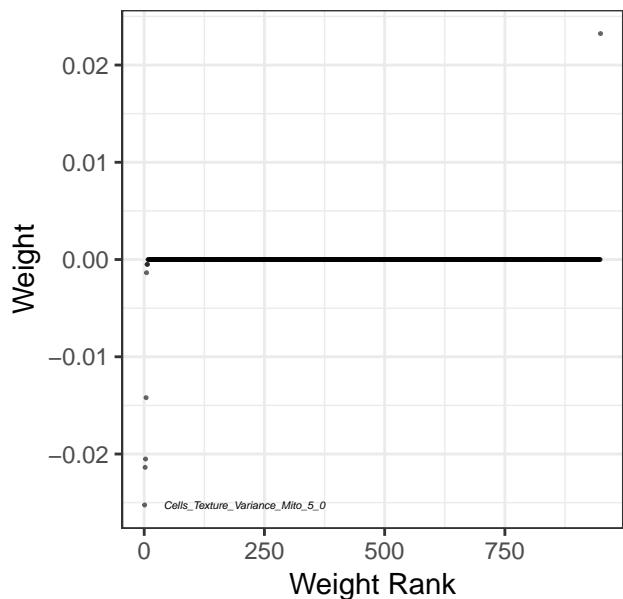


Performance: cc_late_mitosis_n_spots_h2ax_mean

Transform: raw

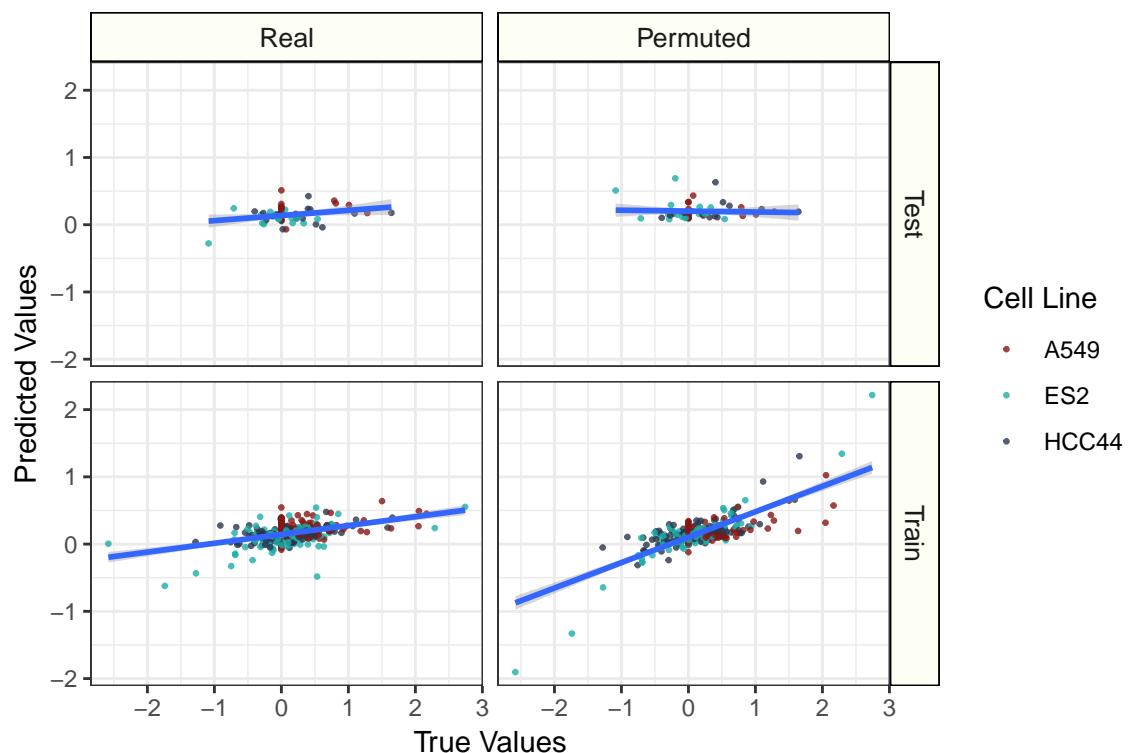


fit	shuffle	transform	r2	value
Train	Real	raw	0.11	0.31
Test	Real	raw	-0.04	0.27
Train	Shuffle	raw	0.33	0.23
Test	Shuffle	raw	0.02	0.25

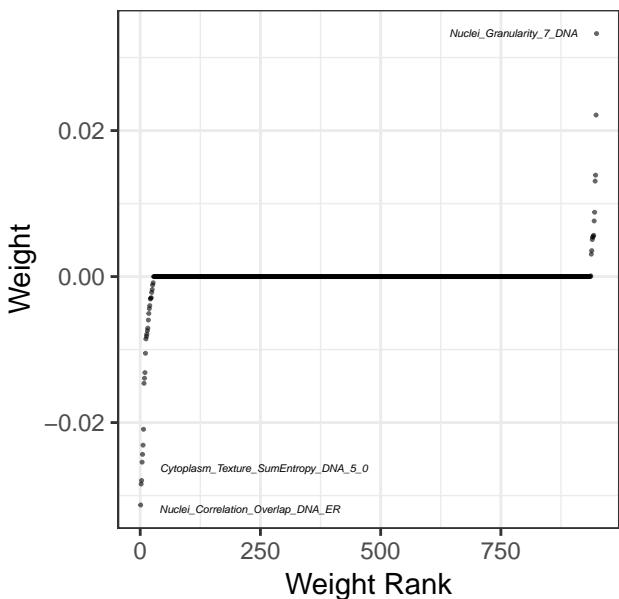


Performance: cc_late_mitosis_n_spots_h2ax_per_nucleus_area

Transform: raw

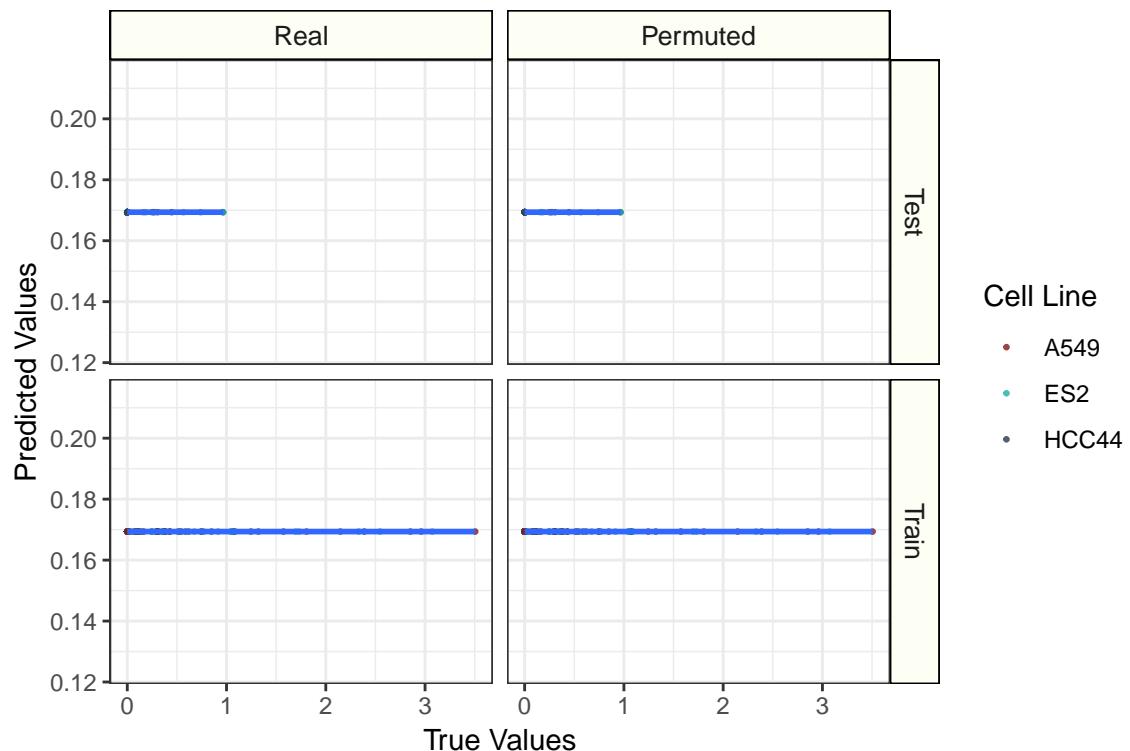


fit	shuffle	transform	r2	value
Train	Real	raw	0.18	0.23
Test	Real	raw	0.07	0.19
Train	Shuffle	raw	0.51	0.14
Test	Shuffle	raw	-0.12	0.23

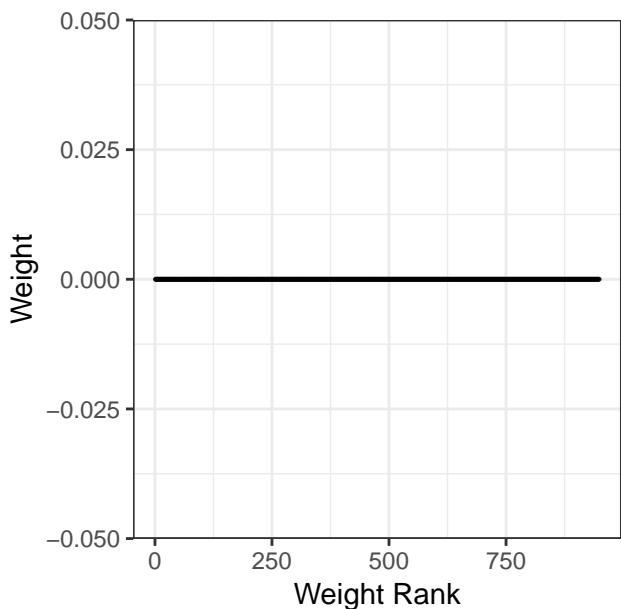


Performance: cc_mitosis_high_h2ax

Transform: raw

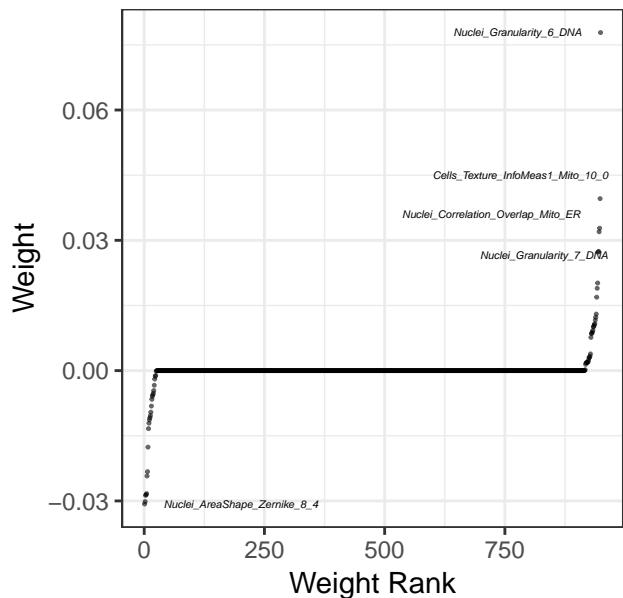
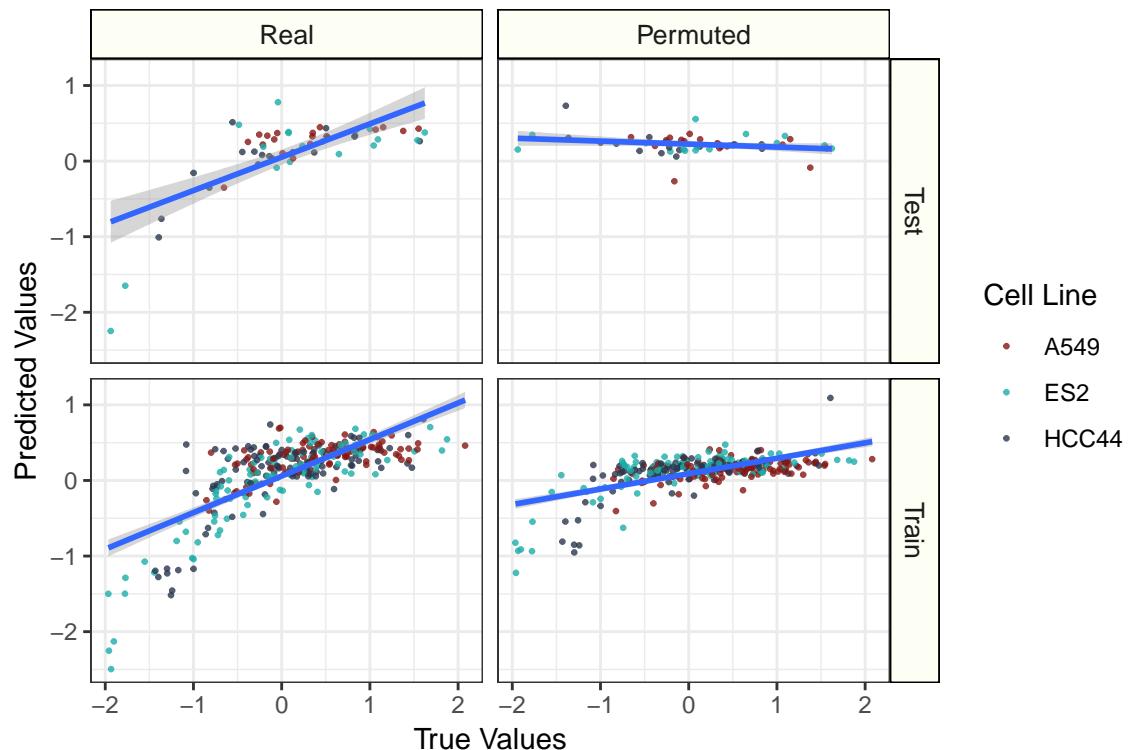


fit	shuffle	transform	r2	value
Train	Real	raw	0.00	0.27
Test	Real	raw	-0.12	0.06
Train	Shuffle	raw	0.00	0.27
Test	Shuffle	raw	-0.12	0.06



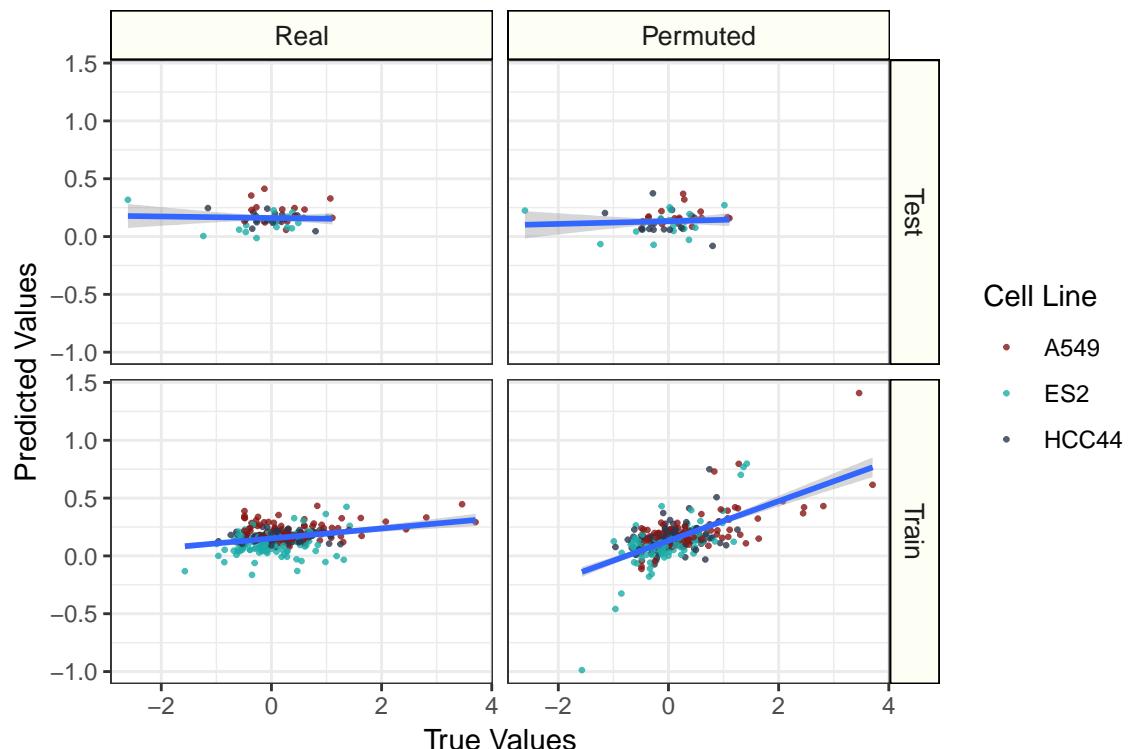
Performance: cc_mitosis_n_objects

Transform: raw

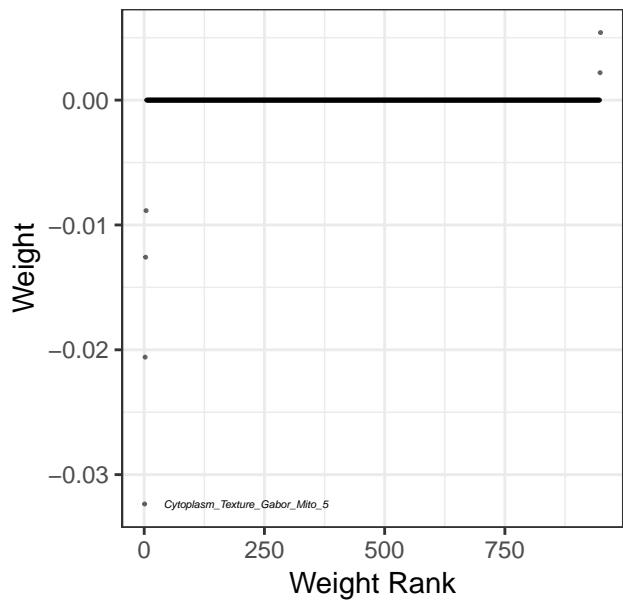


Performance: cc_mitosis_n_spots_h2ax_mean

Transform: raw

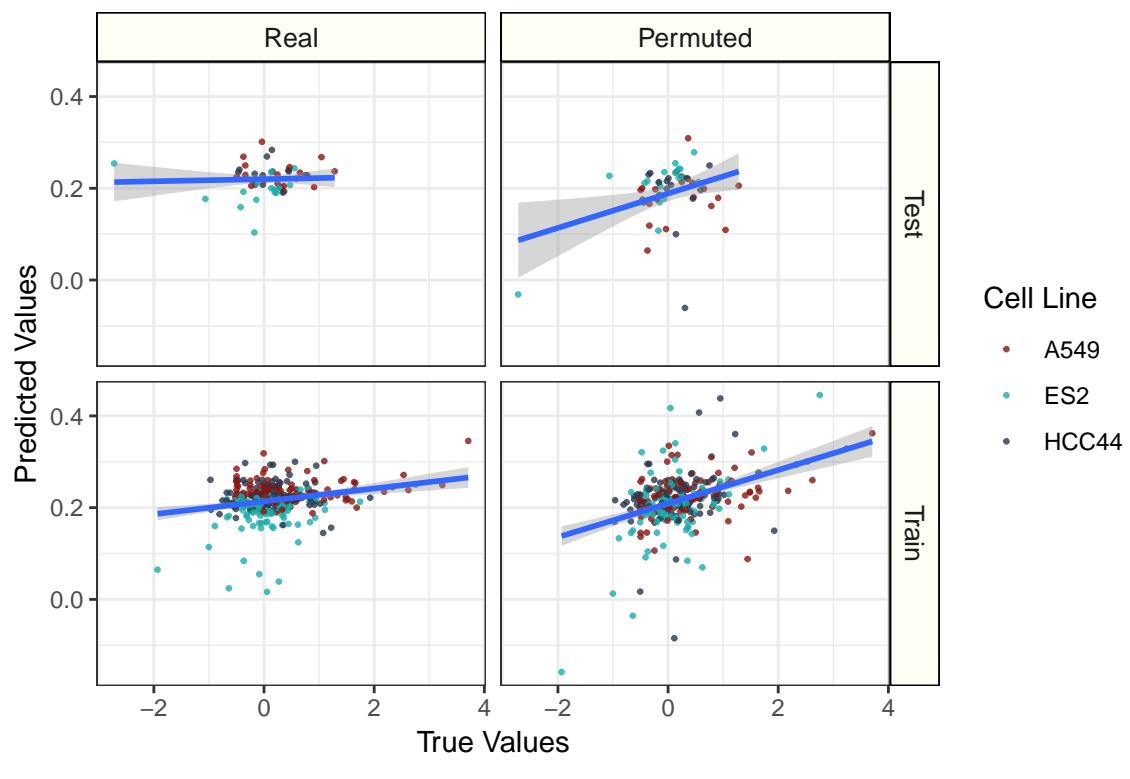


fit	shuffle	transform	r2	value
Train	Real	raw	0.07	0.39
Test	Real	raw	-0.12	0.38
Train	Shuffle	raw	0.27	0.30
Test	Shuffle	raw	-0.07	0.37

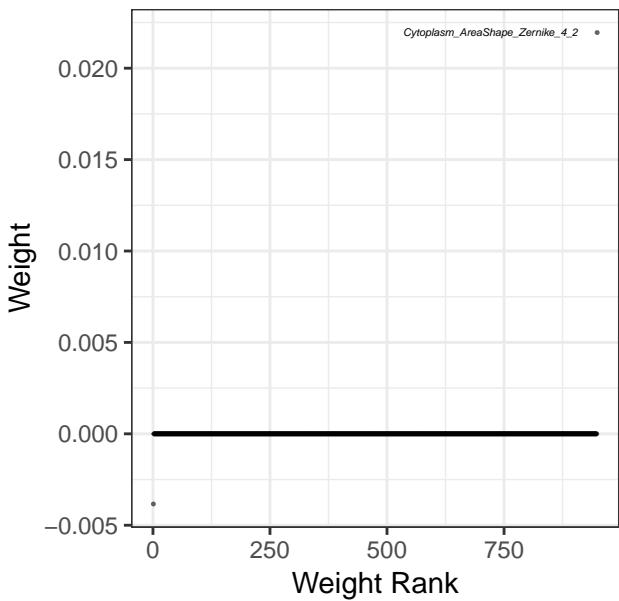


Performance: cc_mitosis_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

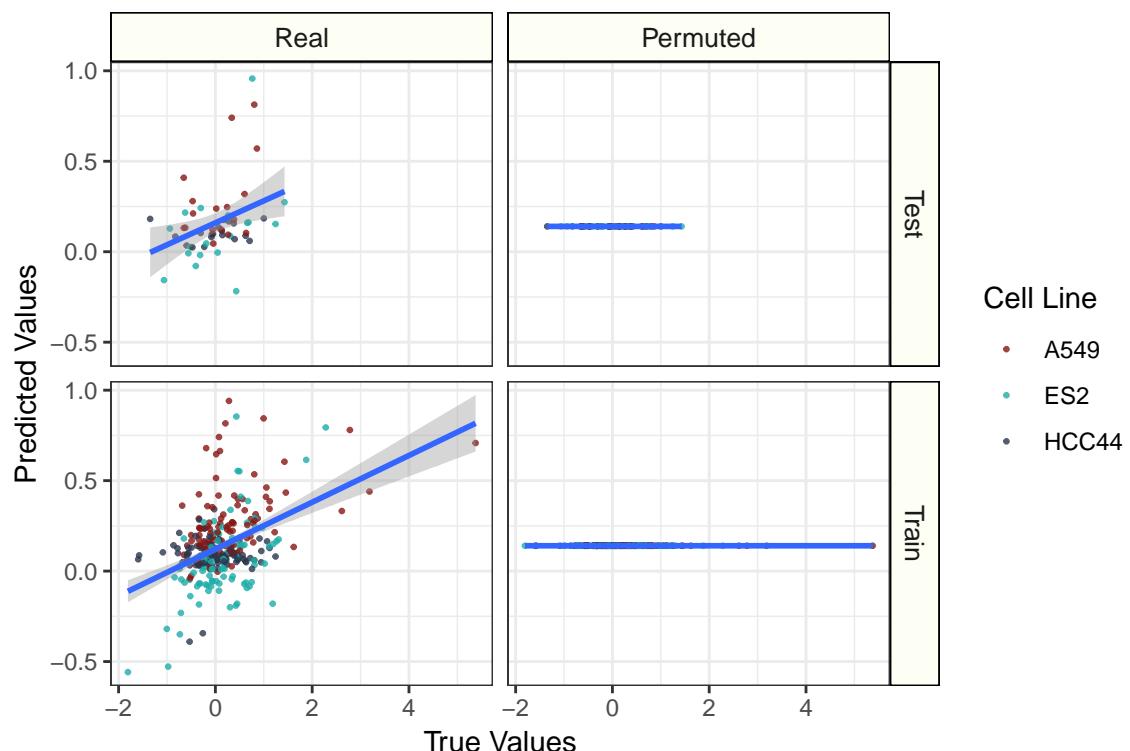


fit	shuffle	transform	r2	value
Train	Real	raw	0.02	0.44
Test	Real	raw	-0.10	0.37
Train	Shuffle	raw	0.06	0.42
Test	Shuffle	raw	-0.01	0.34

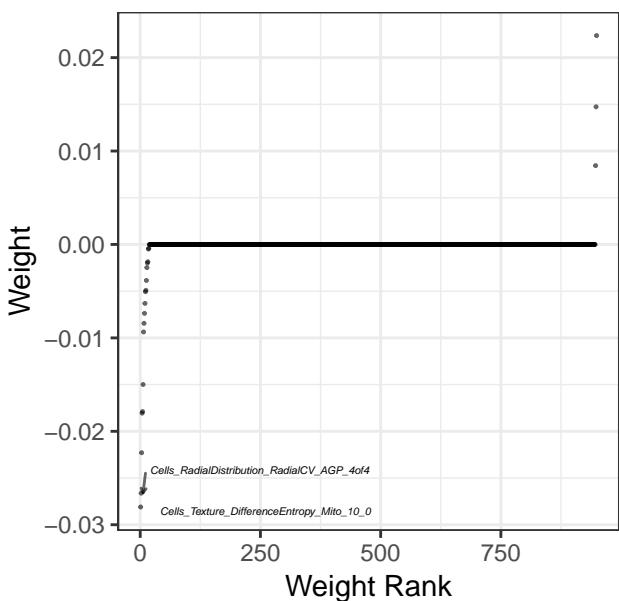


Performance: cc_polynuclear_high_h2ax

Transform: raw

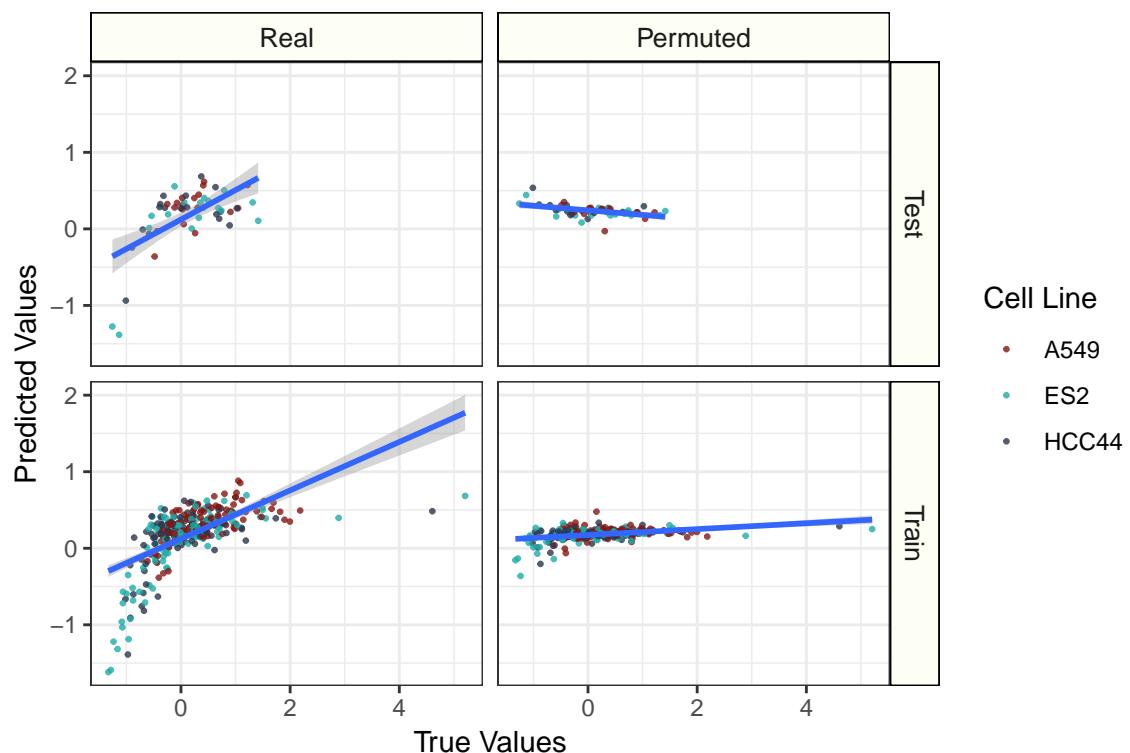


fit	shuffle	transform	r2	value
Train	Real	raw	0.17	0.39
Test	Real	raw	0.08	0.32
Train	Shuffle	raw	0.00	0.47
Test	Shuffle	raw	-0.03	0.35

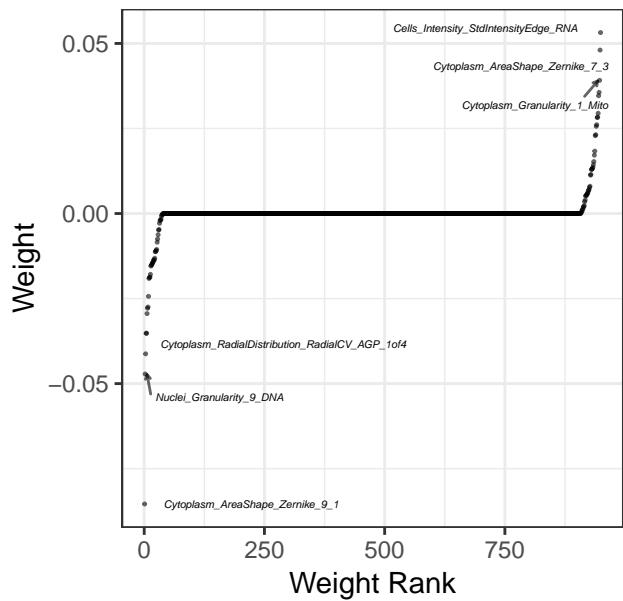


Performance: cc_polynuclear_n_objects

Transform: raw

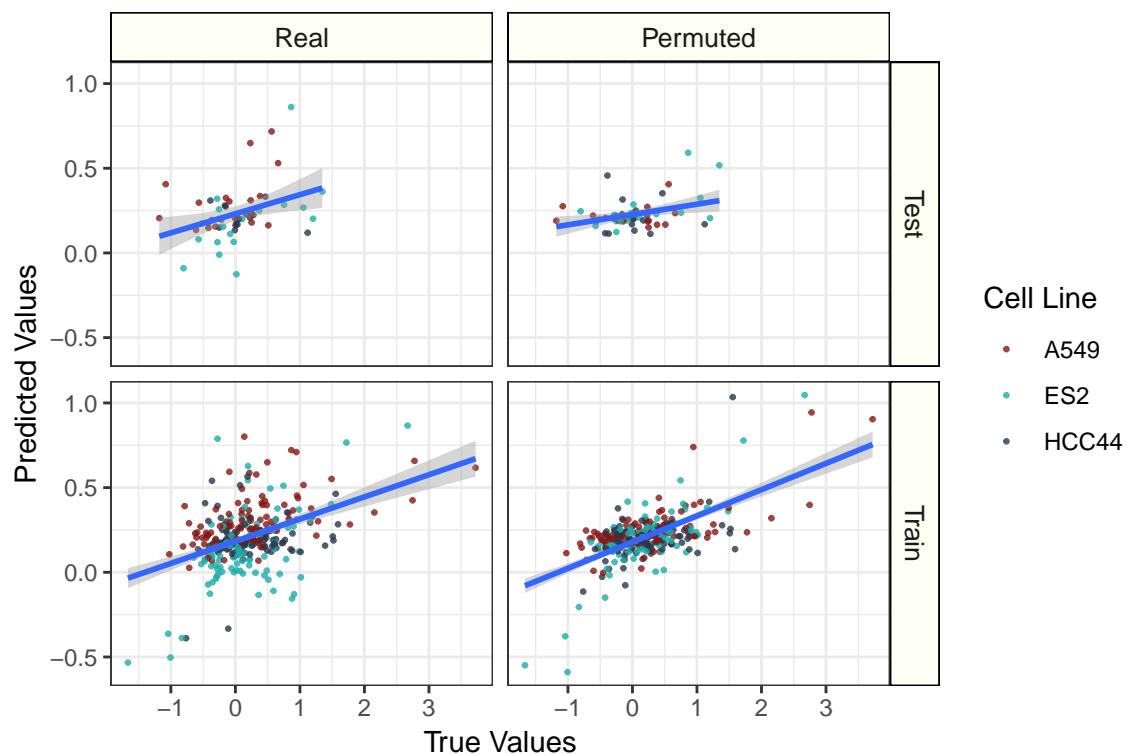


fit	shuffle	transform	r2	value
Train	Real	raw	0.37	0.36
Test	Real	raw	0.35	0.24
Train	Shuffle	raw	0.07	0.54
Test	Shuffle	raw	-0.16	0.43

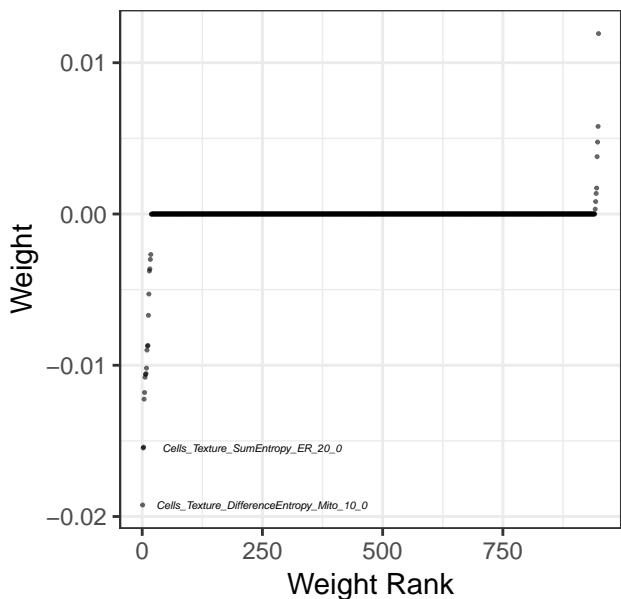


Performance: cc_polynuclear_n_spots_h2ax_mean

Transform: raw

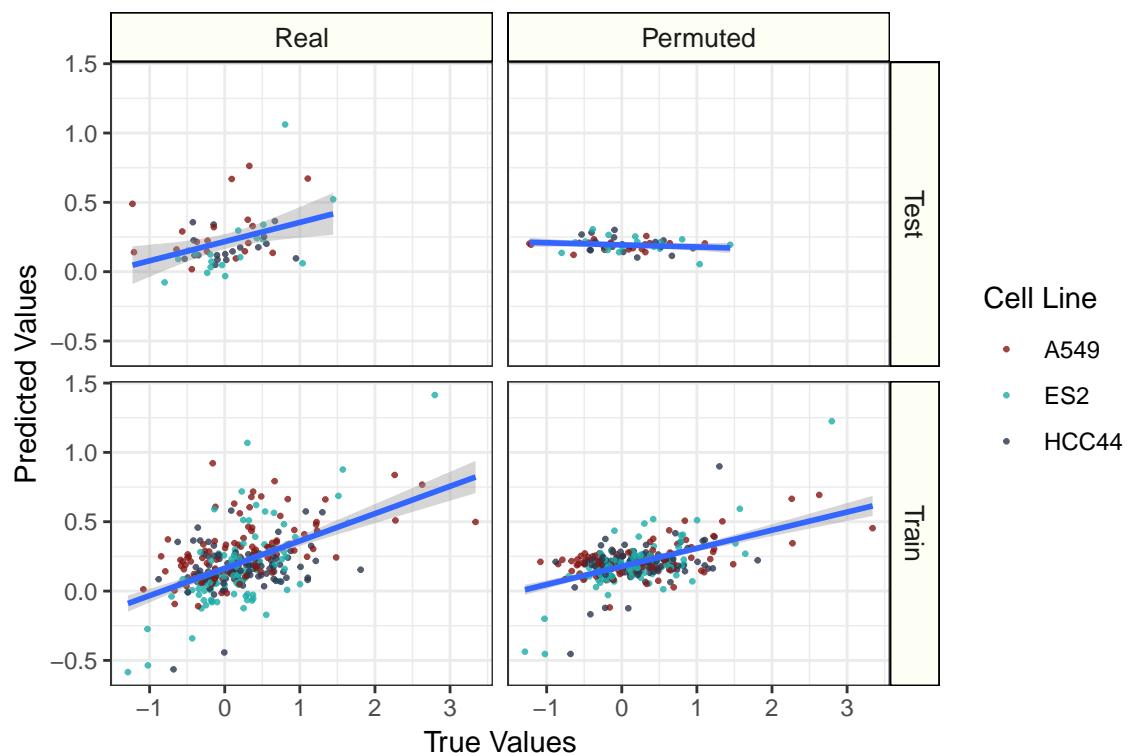


fit	shuffle	transform	r2	value
Train	Real	raw	0.18	0.32
Test	Real	raw	-0.03	0.28
Train	Shuffle	raw	0.25	0.29
Test	Shuffle	raw	-0.06	0.28

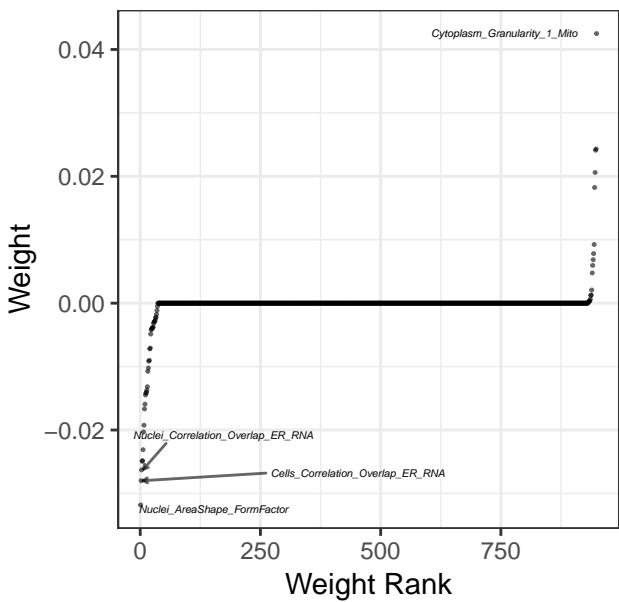


Performance: cc_polynuclear_n_spots_h2ax_per_nucleus_area

Transform: raw

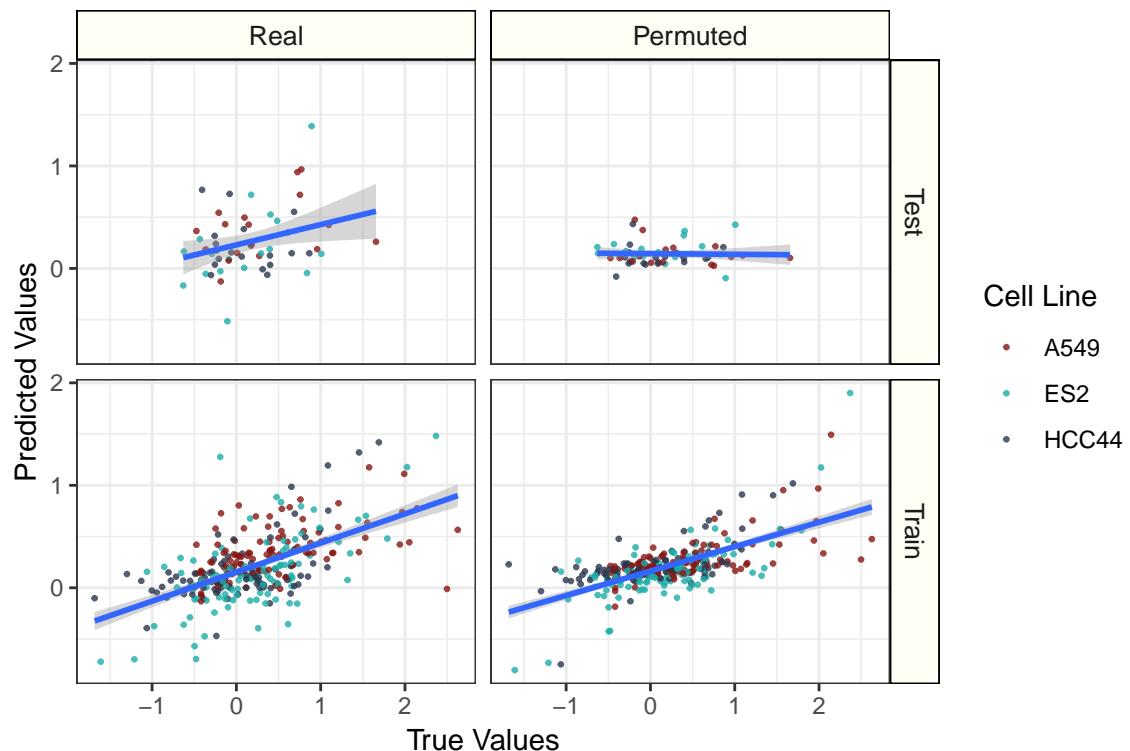


fit	shuffle	transform	r2	value
Train	Real	raw	0.25	0.27
Test	Real	raw	0.00	0.29
Train	Shuffle	raw	0.20	0.29
Test	Shuffle	raw	-0.13	0.33

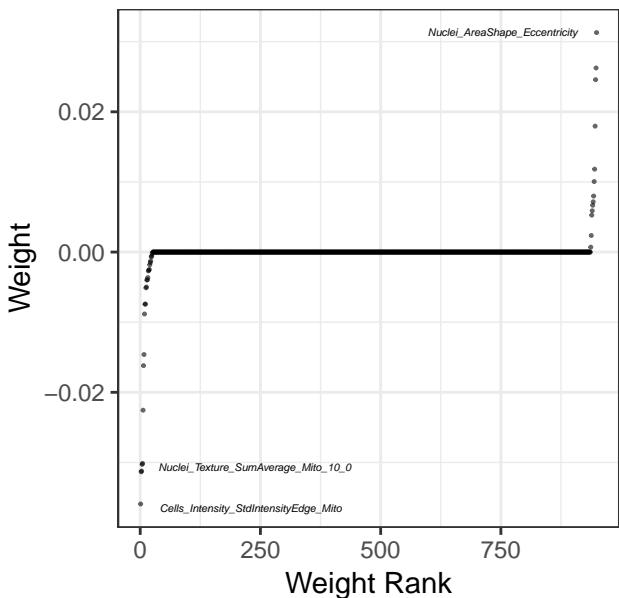


Performance: cc_polyplloid_high_h2ax

Transform: raw

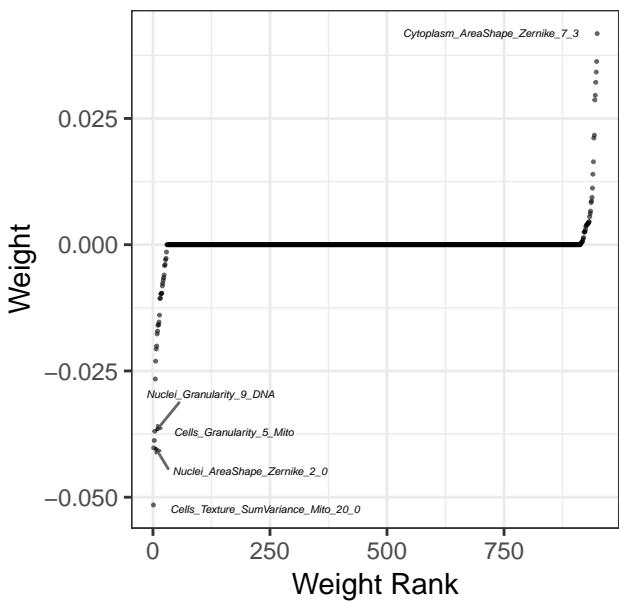
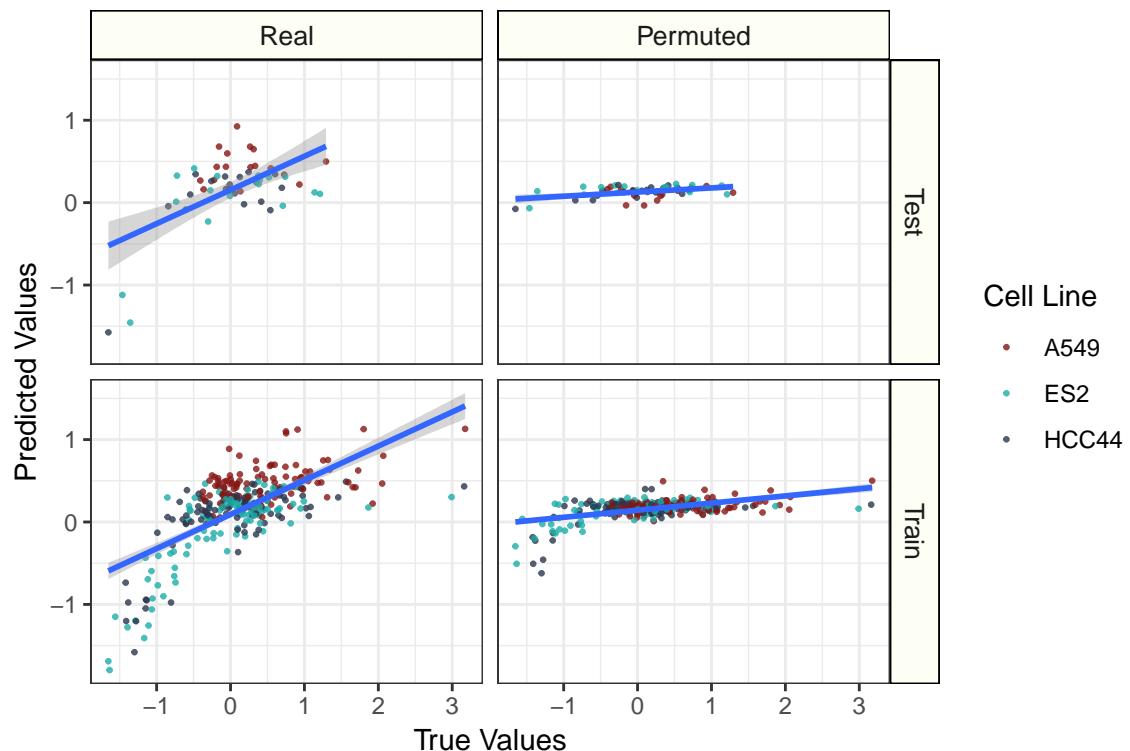


fit	shuffle	transform	r2	value
Train	Real	raw	0.34	0.30
Test	Real	raw	-0.06	0.26
Train	Shuffle	raw	0.34	0.30
Test	Shuffle	raw	-0.07	0.26



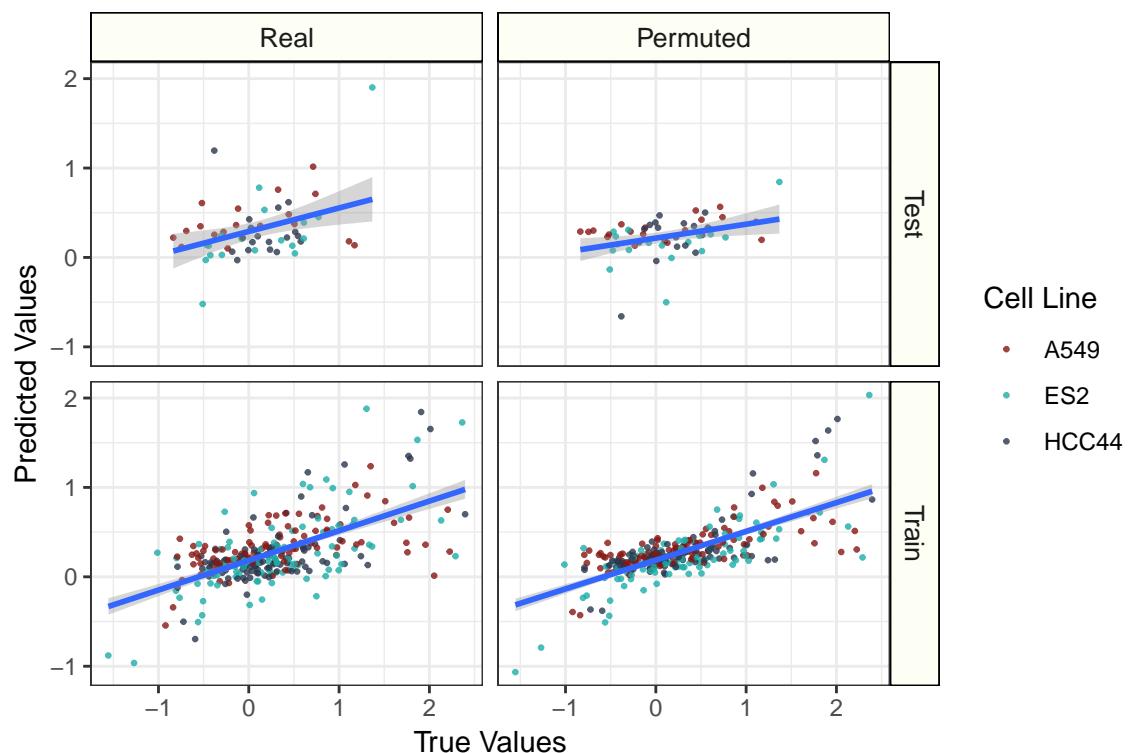
Performance: cc_polyplloid_n_objects

Transform: raw

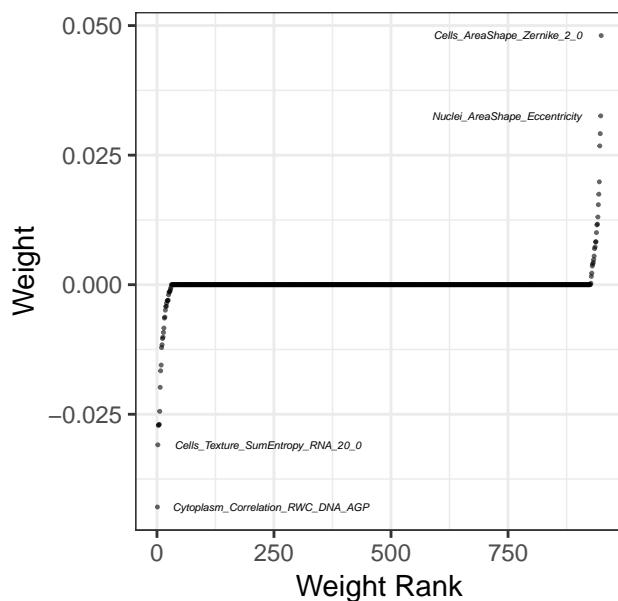


Performance: cc_polyplloid_n_spots_h2ax_mean

Transform: raw

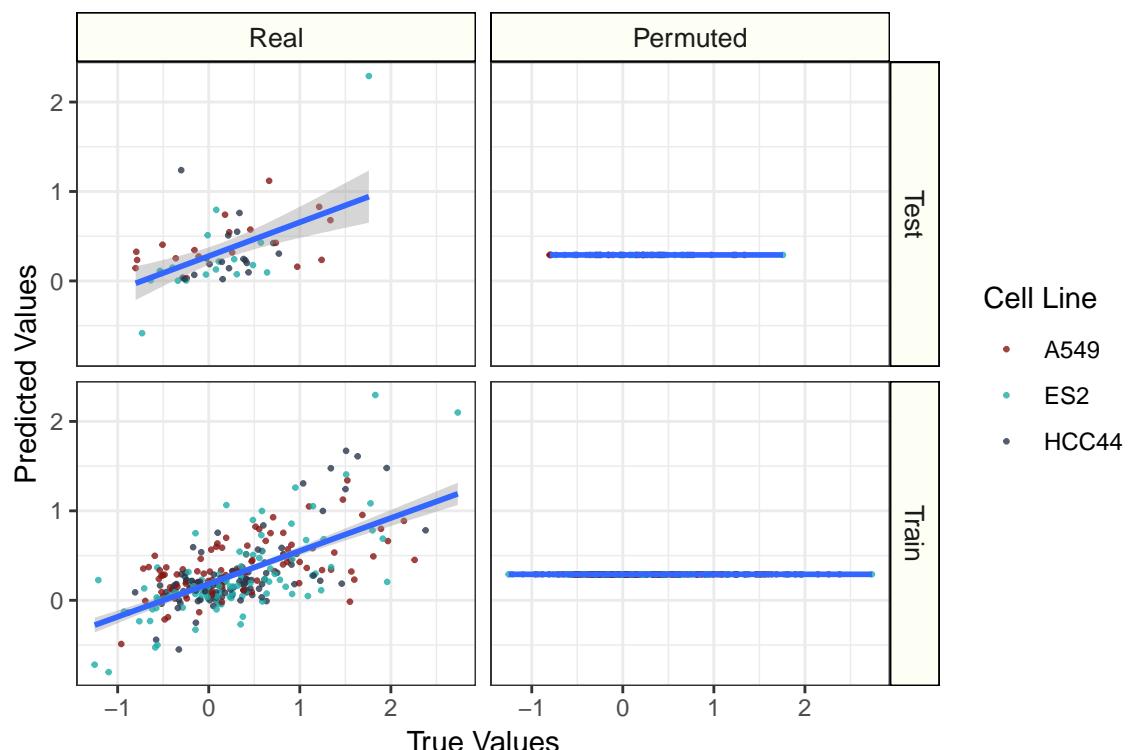


fit	shuffle	transform	r2	value
Train	Real	raw	0.38	0.29
Test	Real	raw	-0.16	0.28
Train	Shuffle	raw	0.44	0.26
Test	Shuffle	raw	0.03	0.24

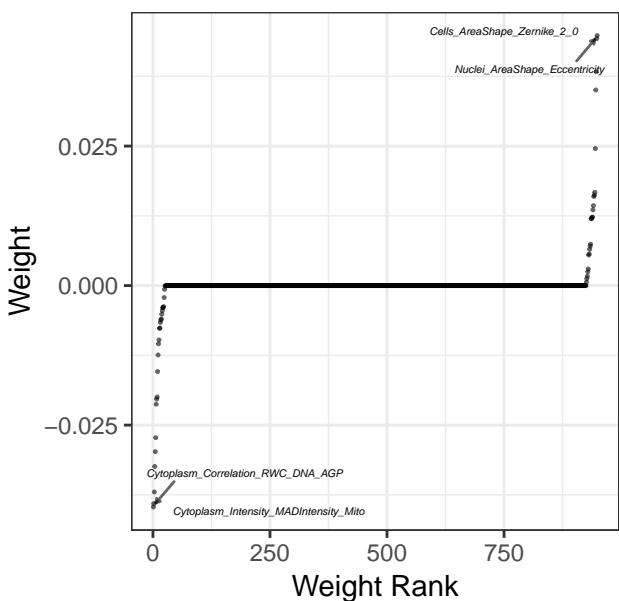


Performance: cc_polyplloid_n_spots_h2ax_per_nucleus_area_m

Transform: raw

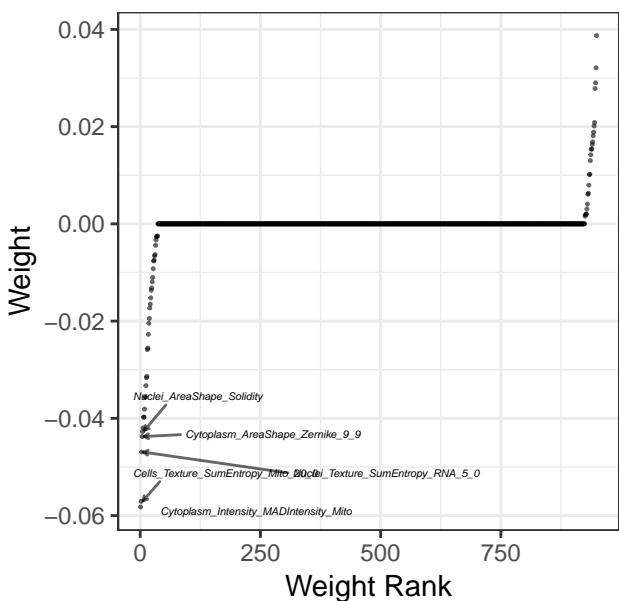
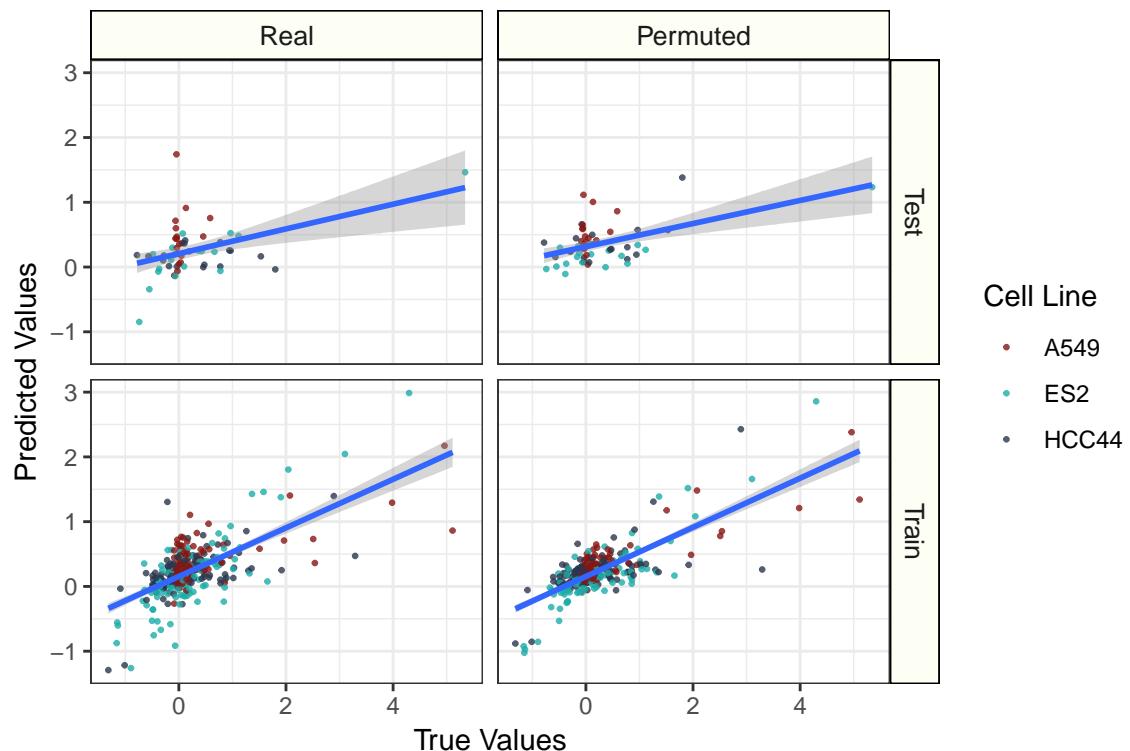


fit	shuffle	transform	r2	value
Train	Real	raw	0.41	0.27
Test	Real	raw	0.12	0.27
Train	Shuffle	raw	0.00	0.46
Test	Shuffle	raw	-0.07	0.33



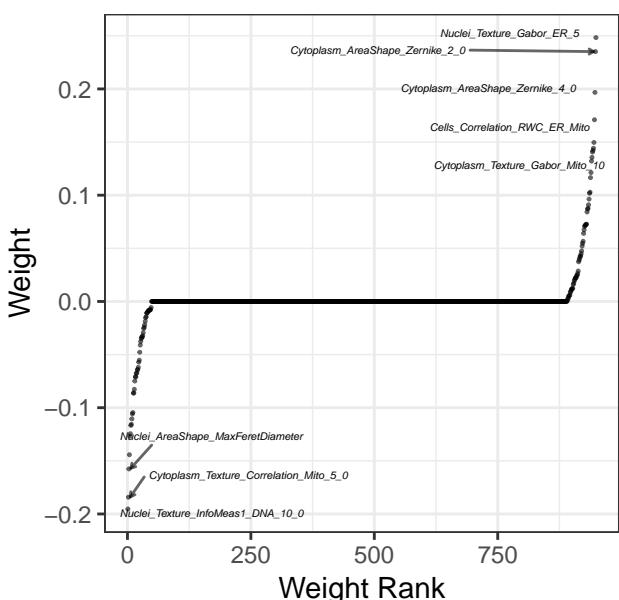
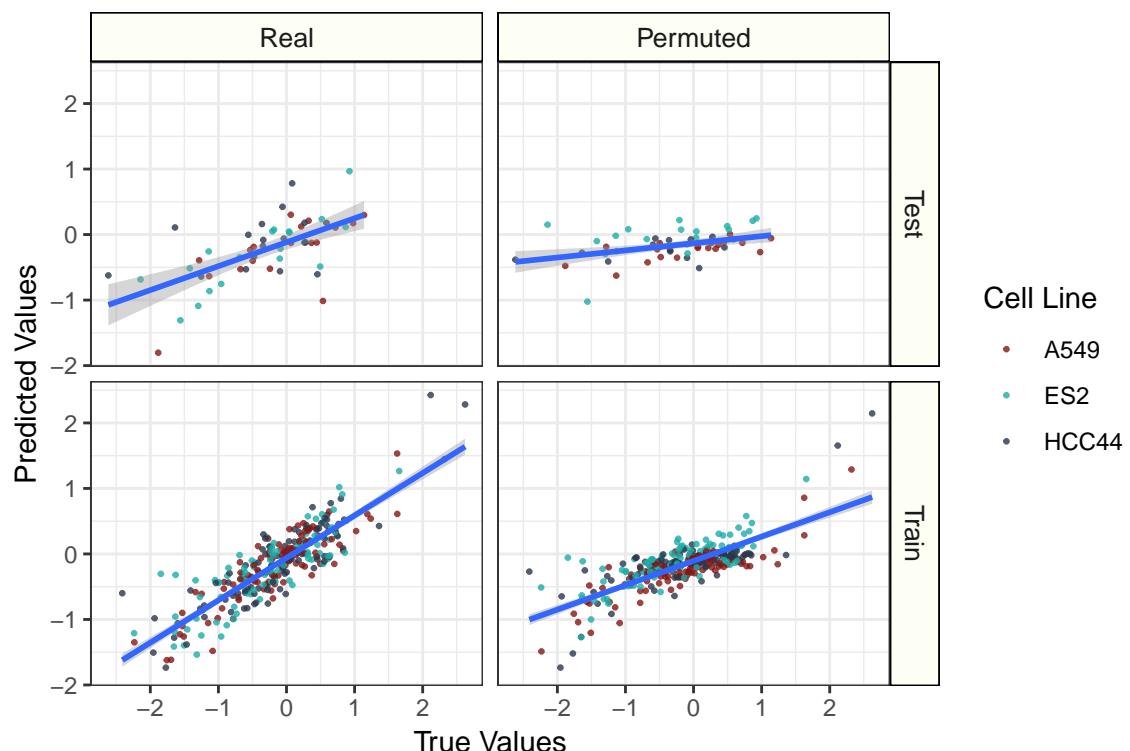
Performance: cc_s_high_h2ax

Transform: raw



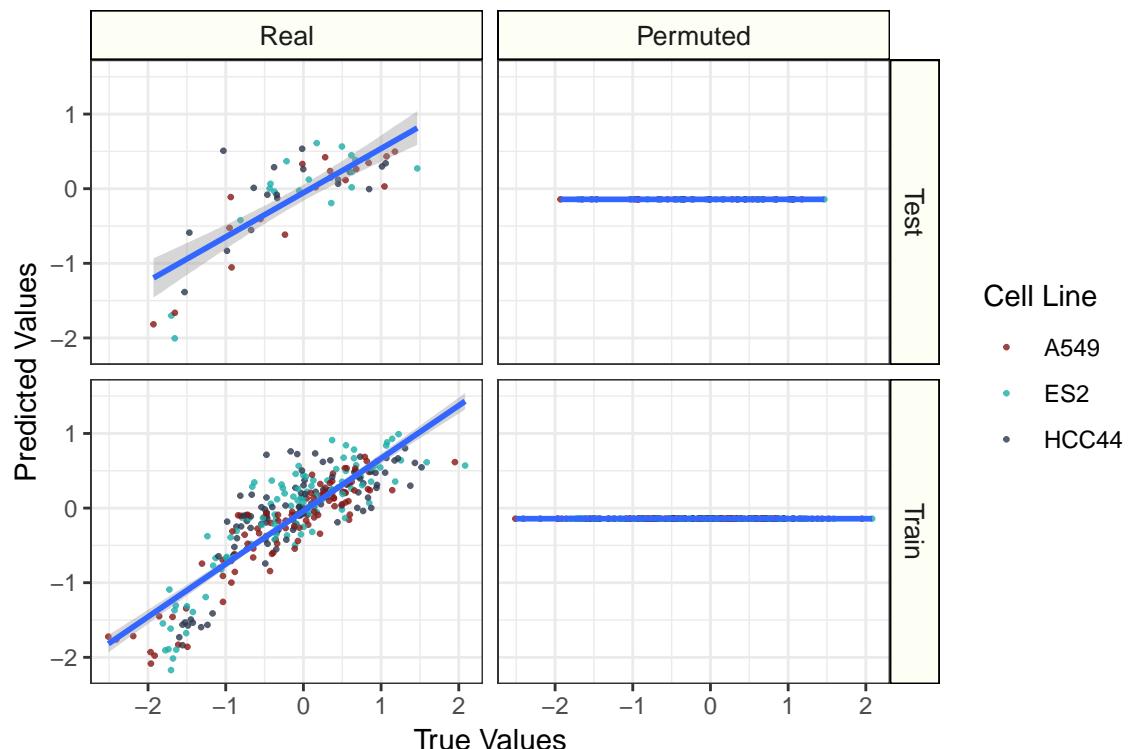
Performance: cc_s_intensity_nucleus_area_mean

Transform: raw

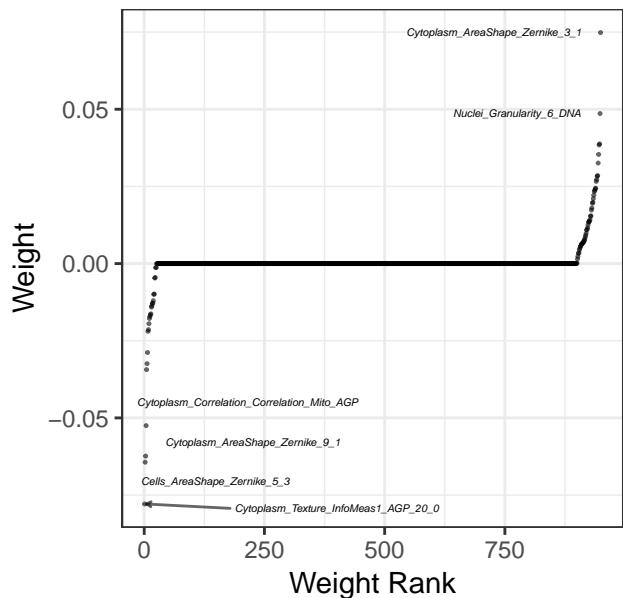


Performance: cc_s_intensity_nucleus_area_sum

Transform: raw

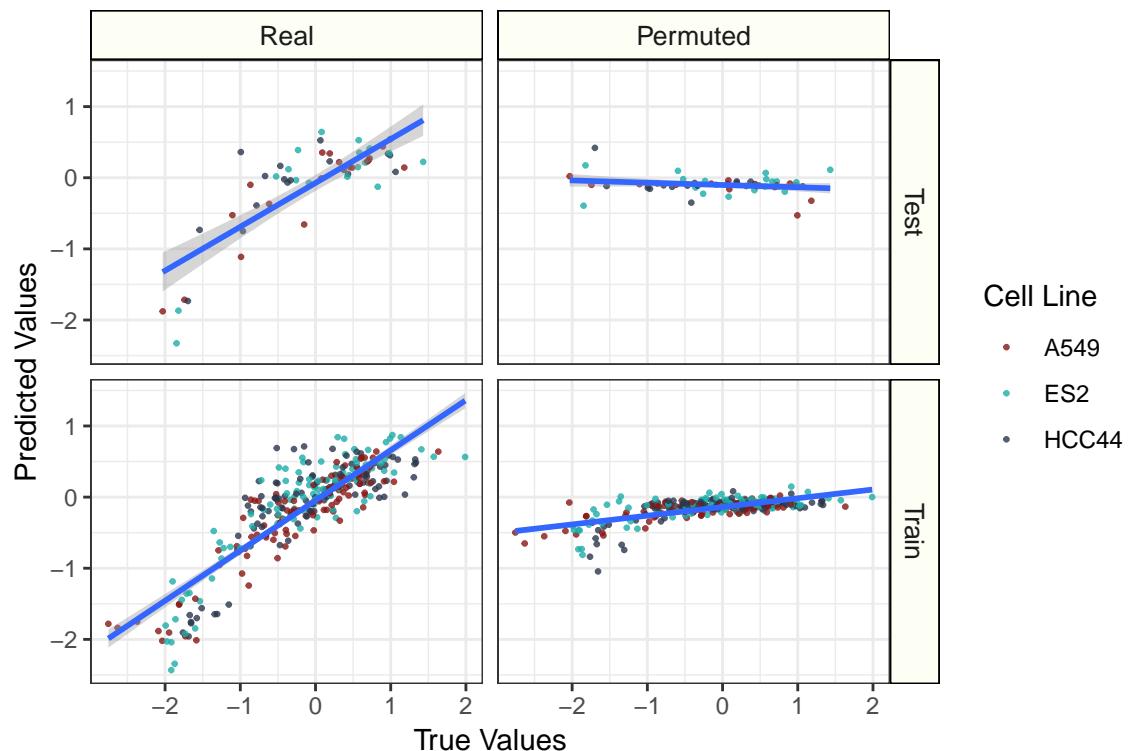


fit	shuffle	transform	r2	value
Train	Real	raw	0.75	0.17
Test	Real	raw	0.62	0.27
Train	Shuffle	raw	0.00	0.68
Test	Shuffle	raw	-0.01	0.71

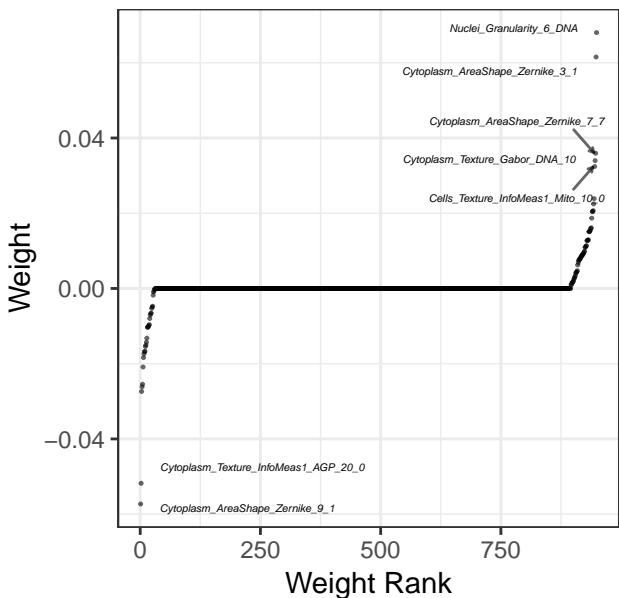


Performance: cc_s_n_objects

Transform: raw

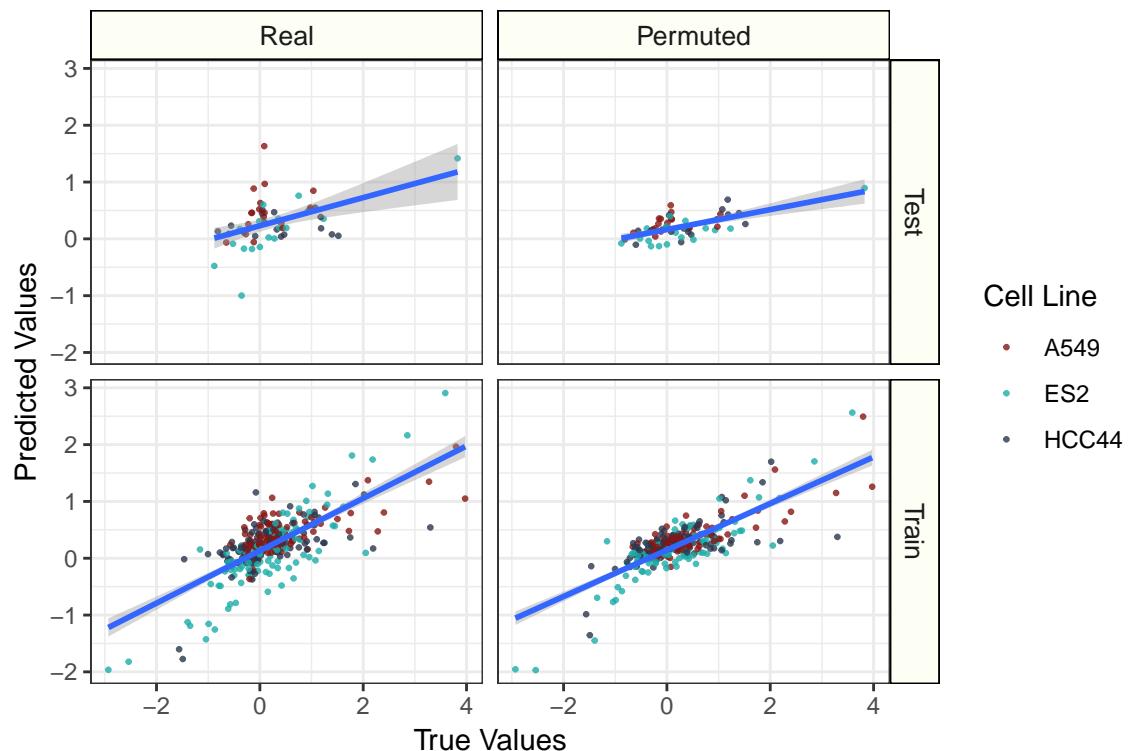


fit	shuffle	transform	r2	value
Train	Real	raw	0.74	0.18
Test	Real	raw	0.64	0.28
Train	Shuffle	raw	0.21	0.56
Test	Shuffle	raw	-0.09	0.83

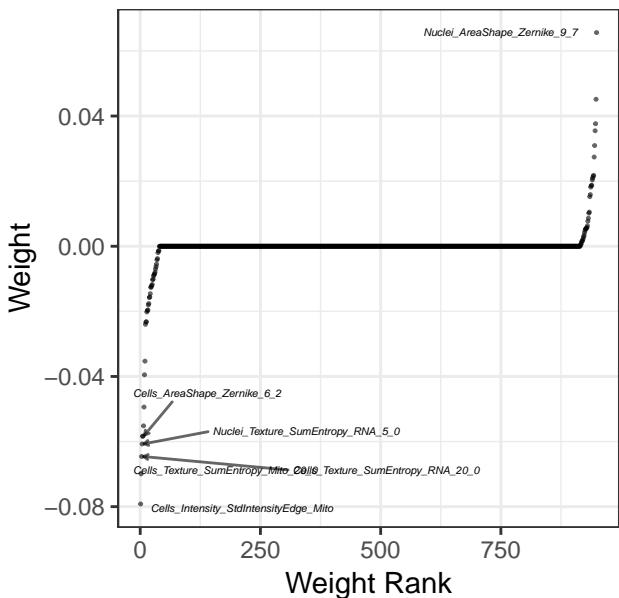


Performance: cc_s_n_spots_h2ax_mean

Transform: raw

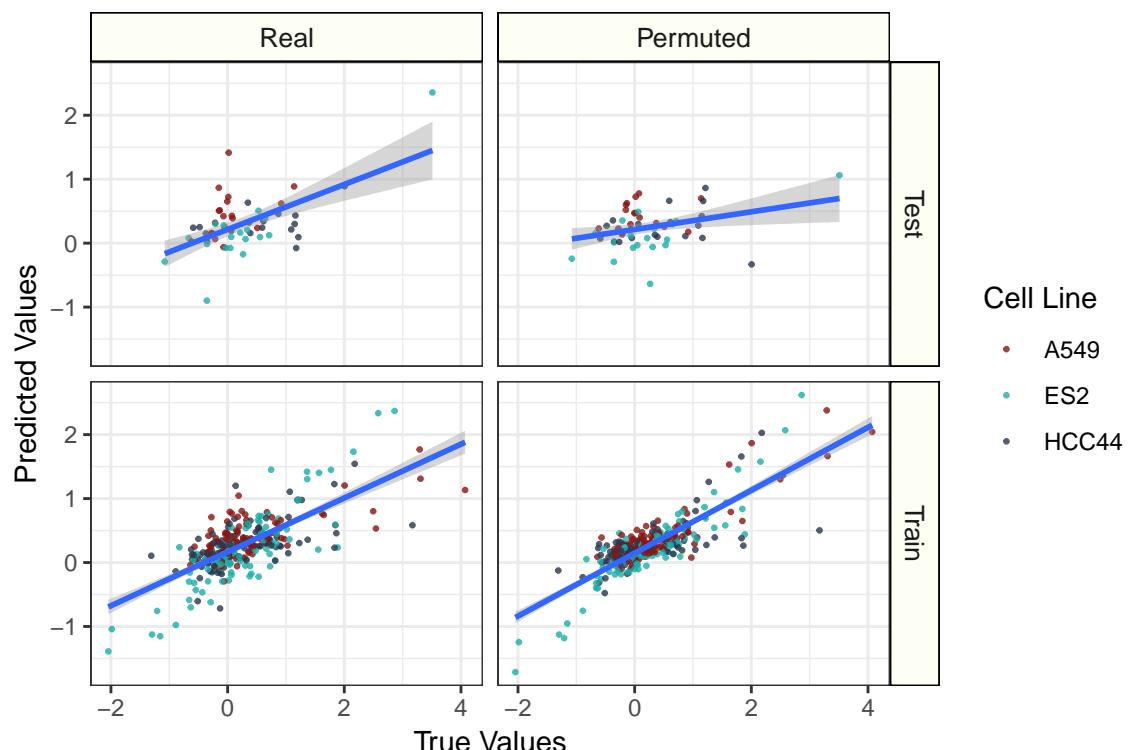


fit	shuffle	transform	r2	value
Train	Real	raw	0.52	0.33
Test	Real	raw	0.20	0.45
Train	Shuffle	raw	0.55	0.32
Test	Shuffle	raw	0.27	0.41

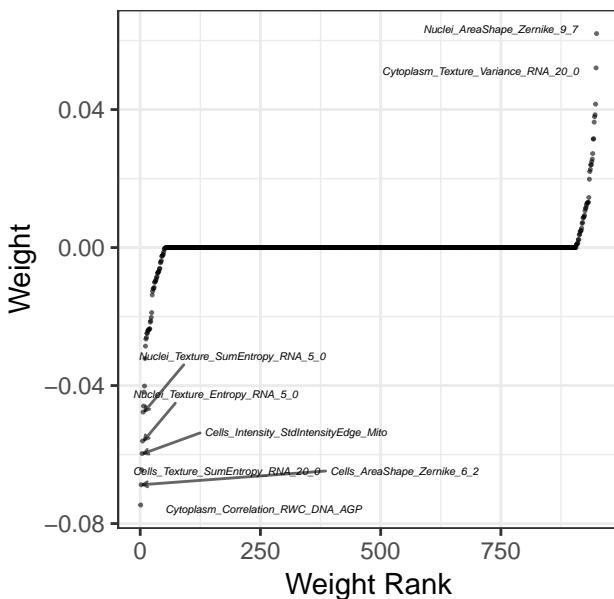


Performance: cc_s_n_spots_h2ax_per_nucleus_area_mean

Transform: raw

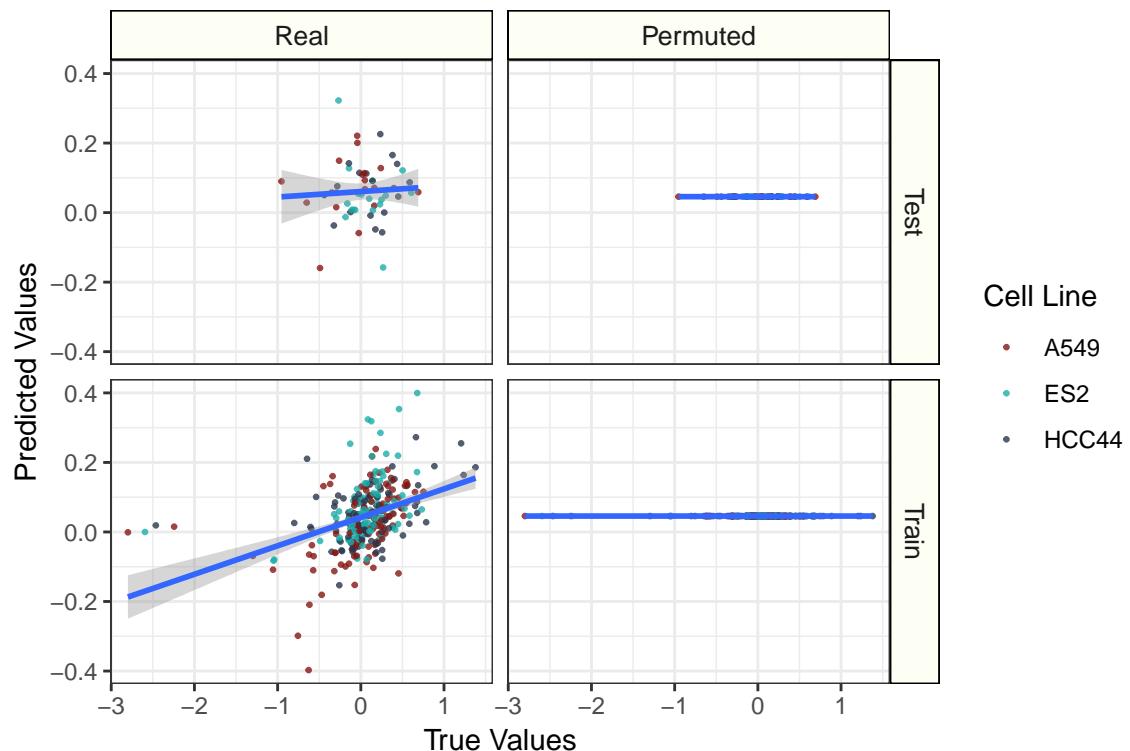


fit	shuffle	transform	r2	value
Train	Real	raw	0.50	0.30
Test	Real	raw	0.34	0.35
Train	Shuffle	raw	0.64	0.22
Test	Shuffle	raw	0.10	0.48

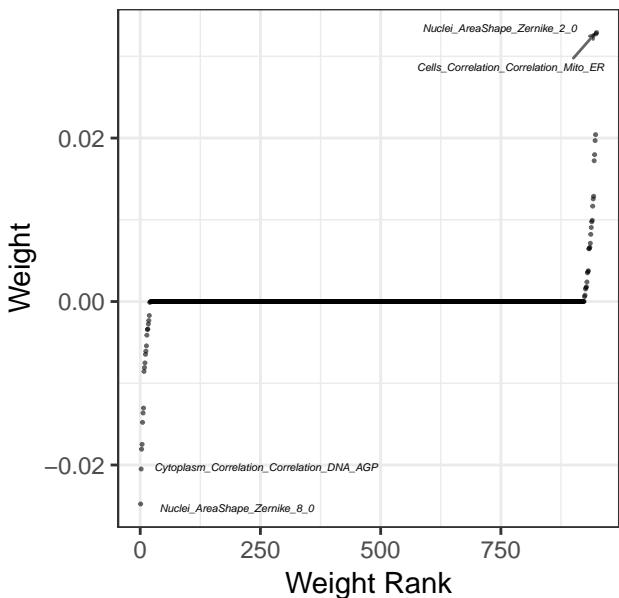


Performance: vb_infection_percentage

Transform: raw

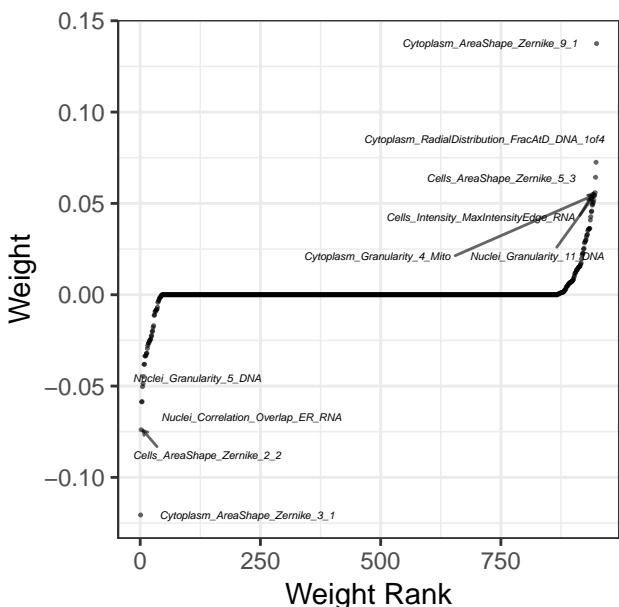
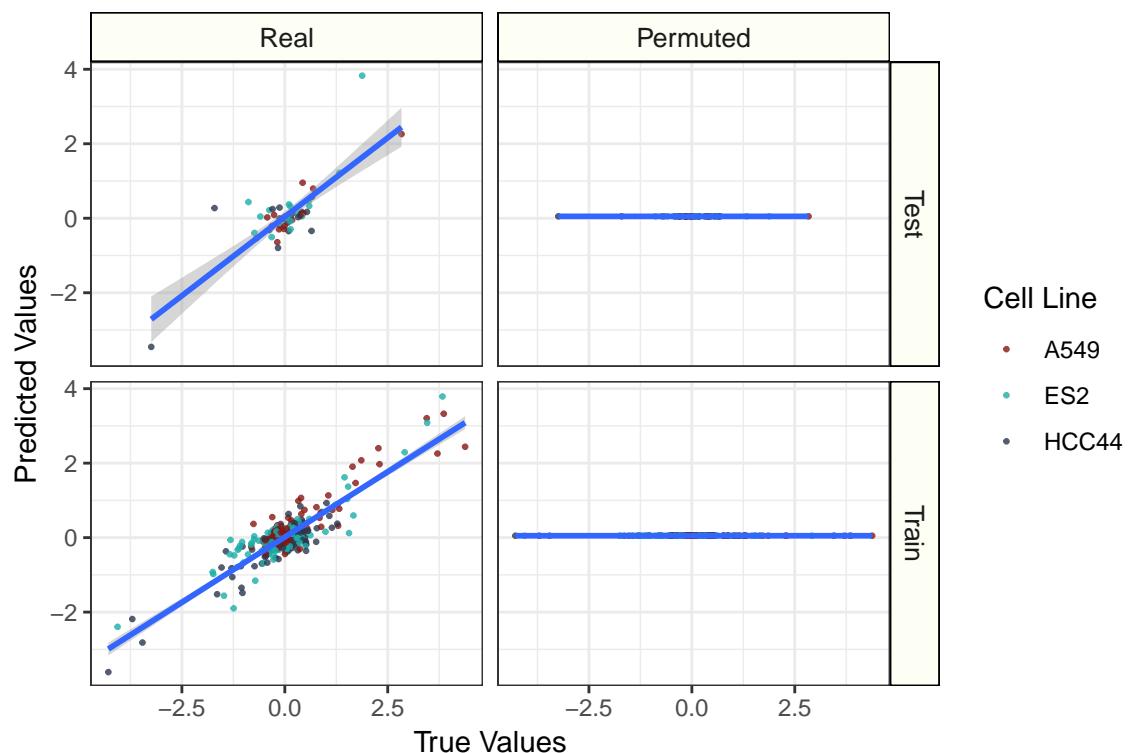


fit	shuffle	transform	r2	value
Train	Real	raw	0.12	0.18
Test	Real	raw	-0.05	0.10
Train	Shuffle	raw	0.00	0.20
Test	Shuffle	raw	0.00	0.10



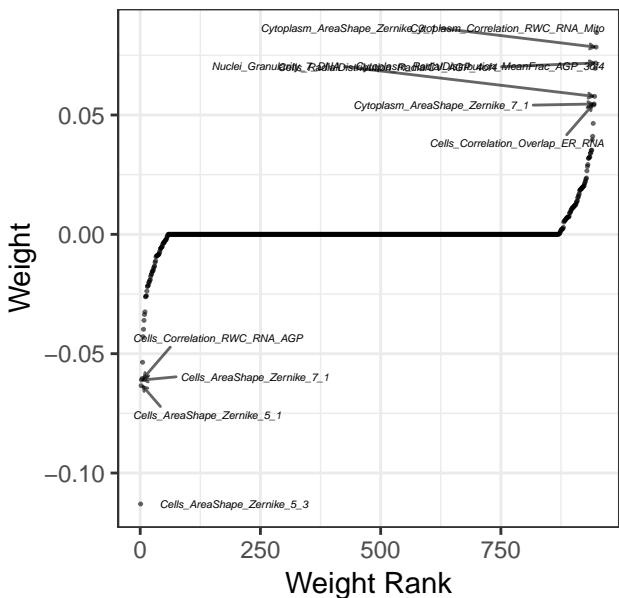
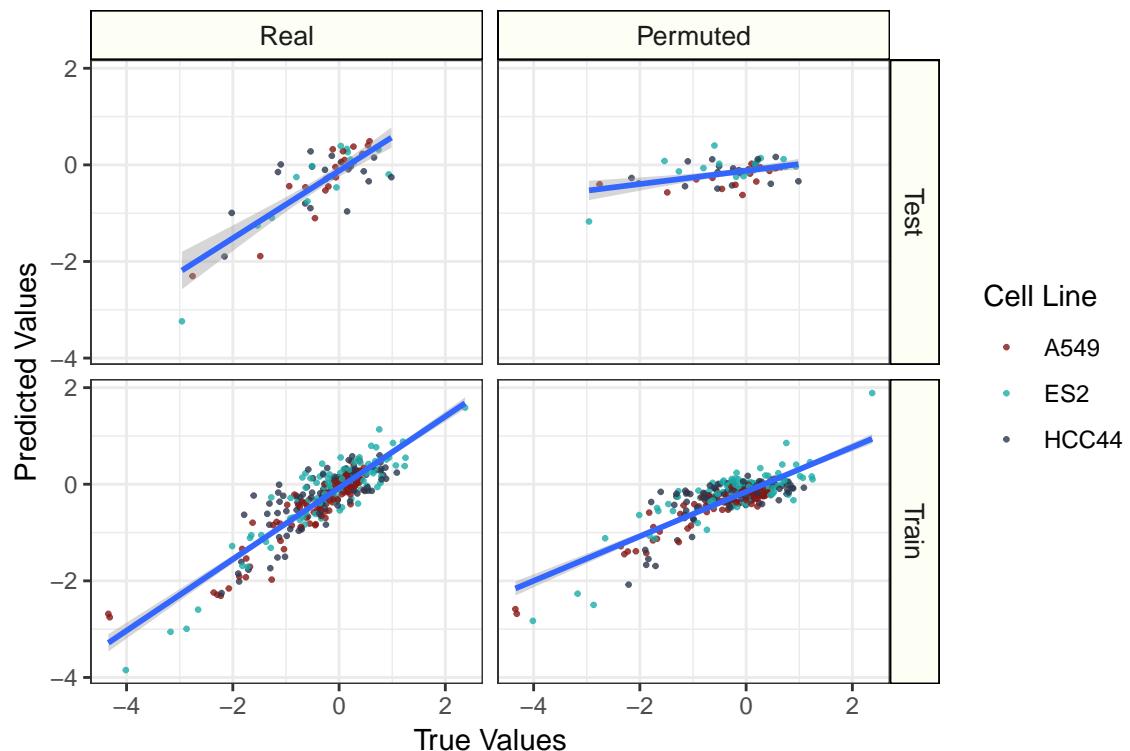
Performance: vb_live_cell_area

Transform: raw



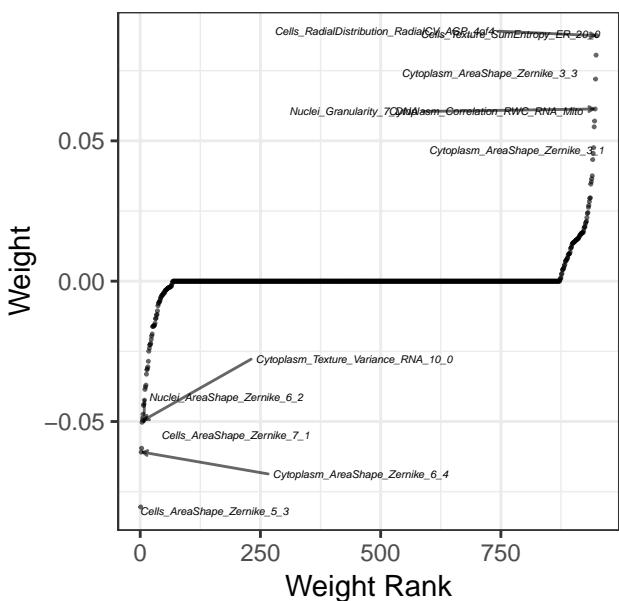
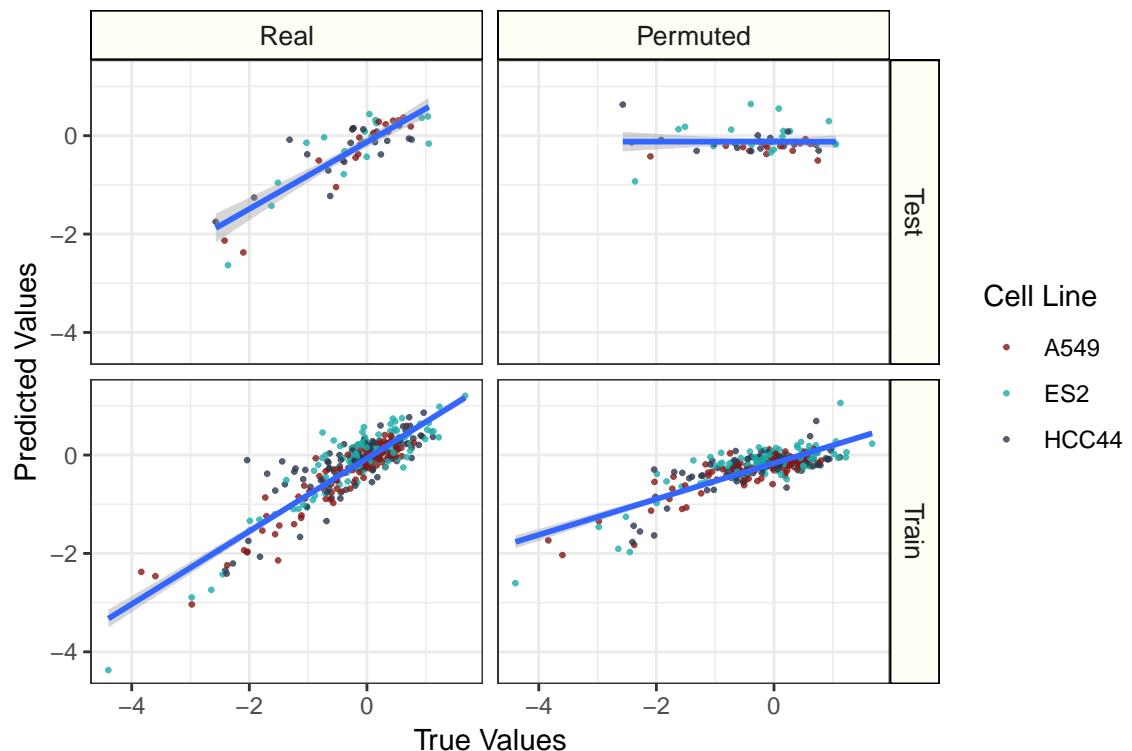
Performance: vb_live_cell_roundness

Transform: raw



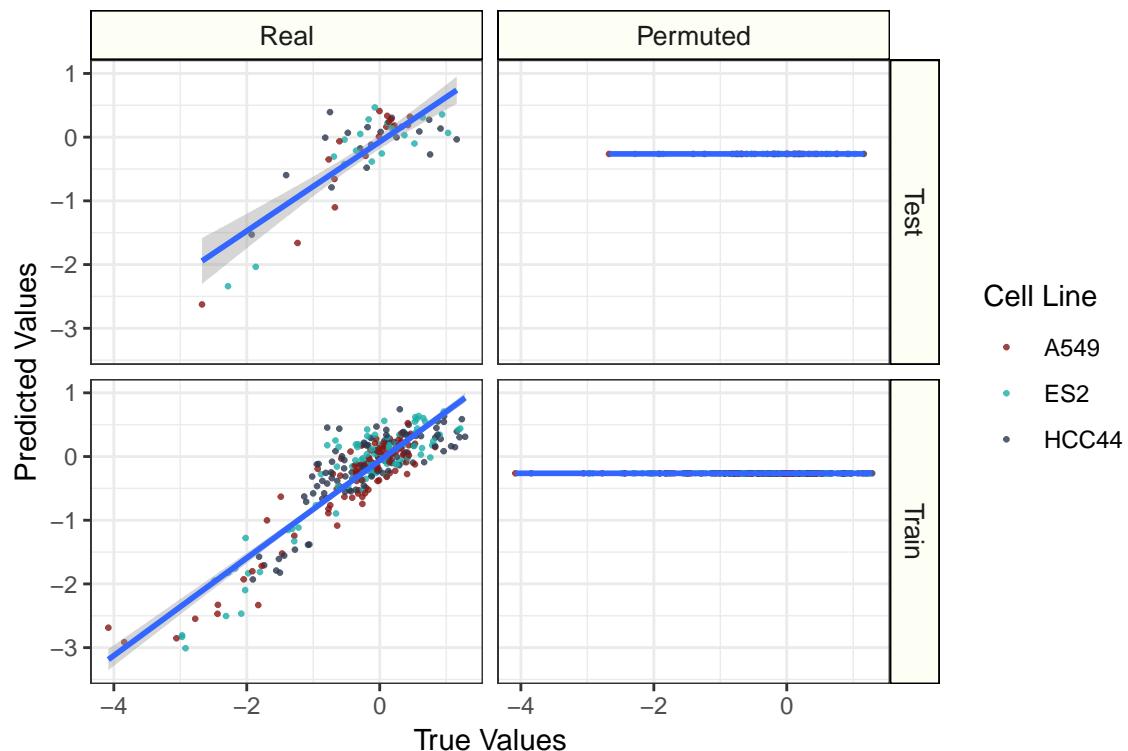
Performance: vb_live_cell_width_length

Transform: raw

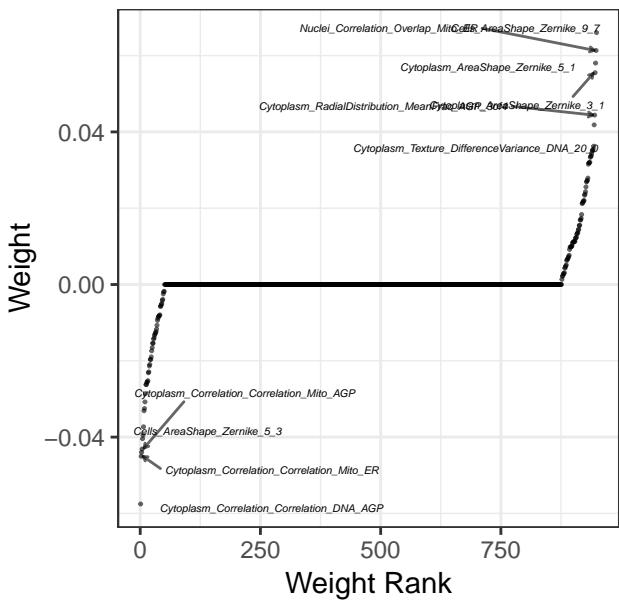


Performance: vb_num_live_cells

Transform: raw

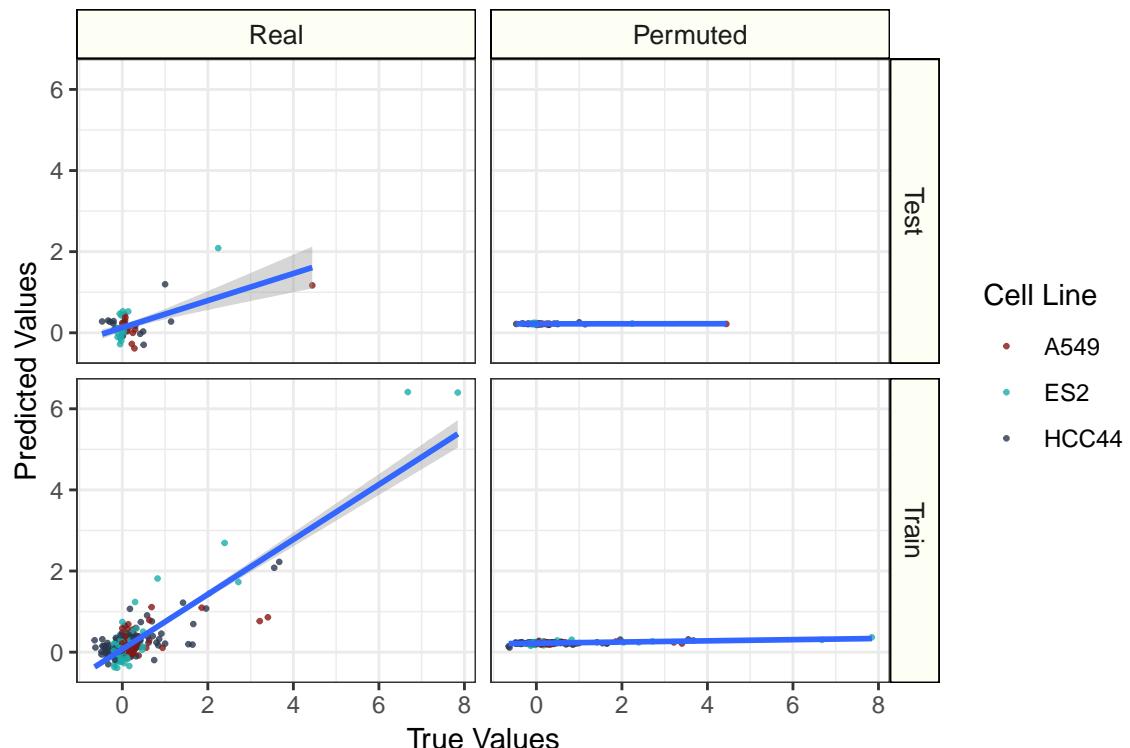


fit	shuffle	transform	r2	value
Train	Real	raw	0.81	0.14
Test	Real	raw	0.67	0.21
Train	Shuffle	raw	0.00	0.75
Test	Shuffle	raw	-0.01	0.63

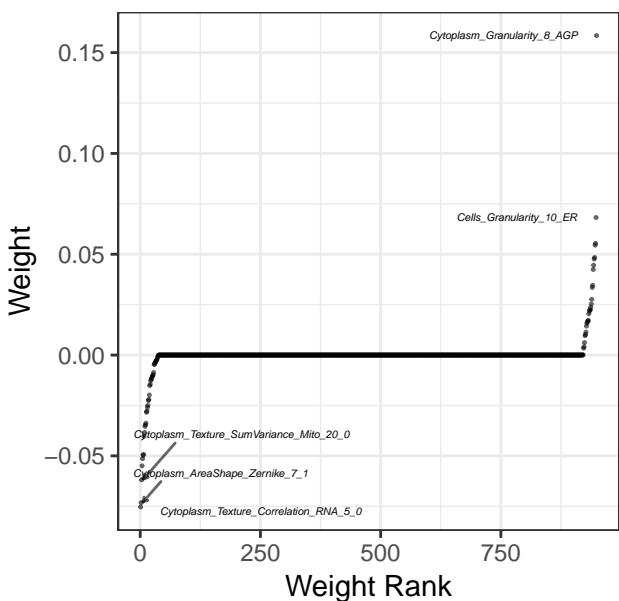


Performance: vb_percent_all_apoptosis

Transform: raw

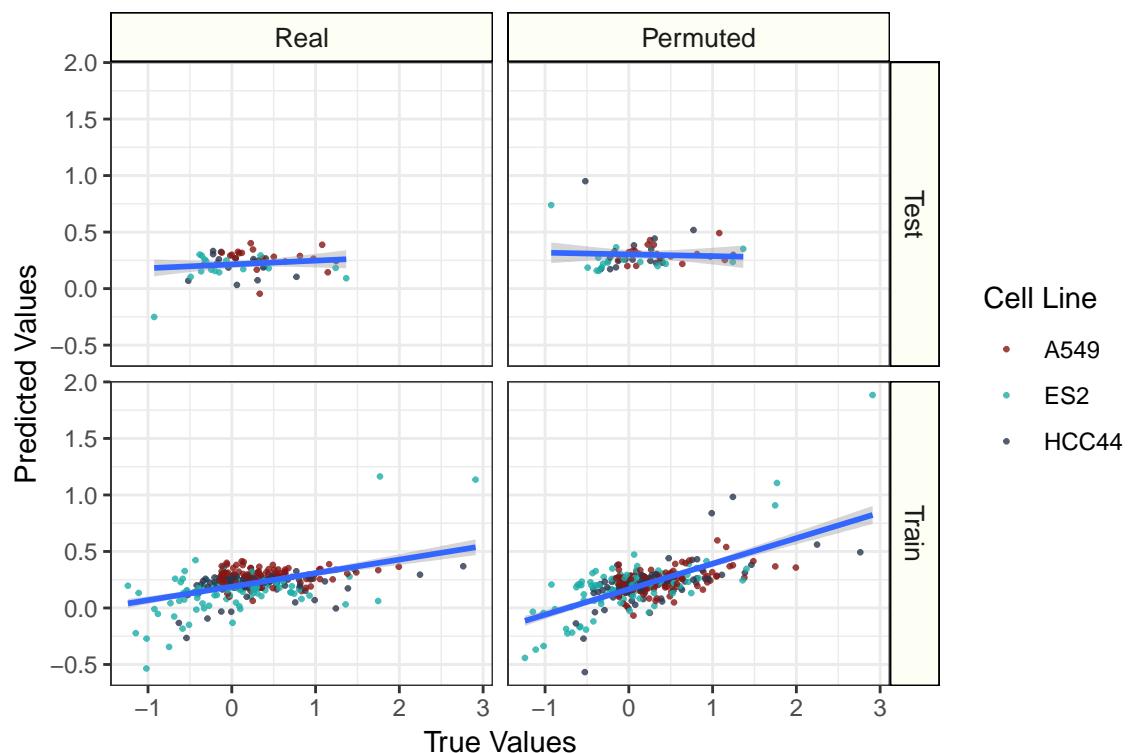


fit	shuffle	transform	r2	value
Train	Real	raw	0.74	0.16
Test	Real	raw	0.37	0.31
Train	Shuffle	raw	0.03	0.62
Test	Shuffle	raw	0.00	0.50

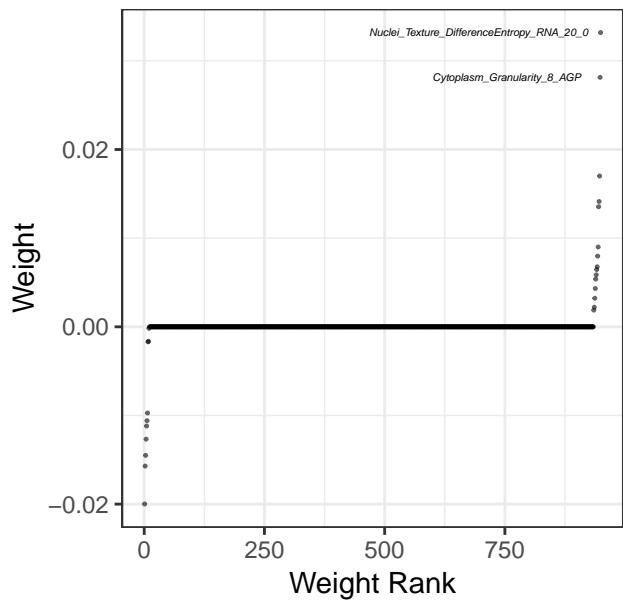


Performance: vb_percent_caspase_dead_only

Transform: raw

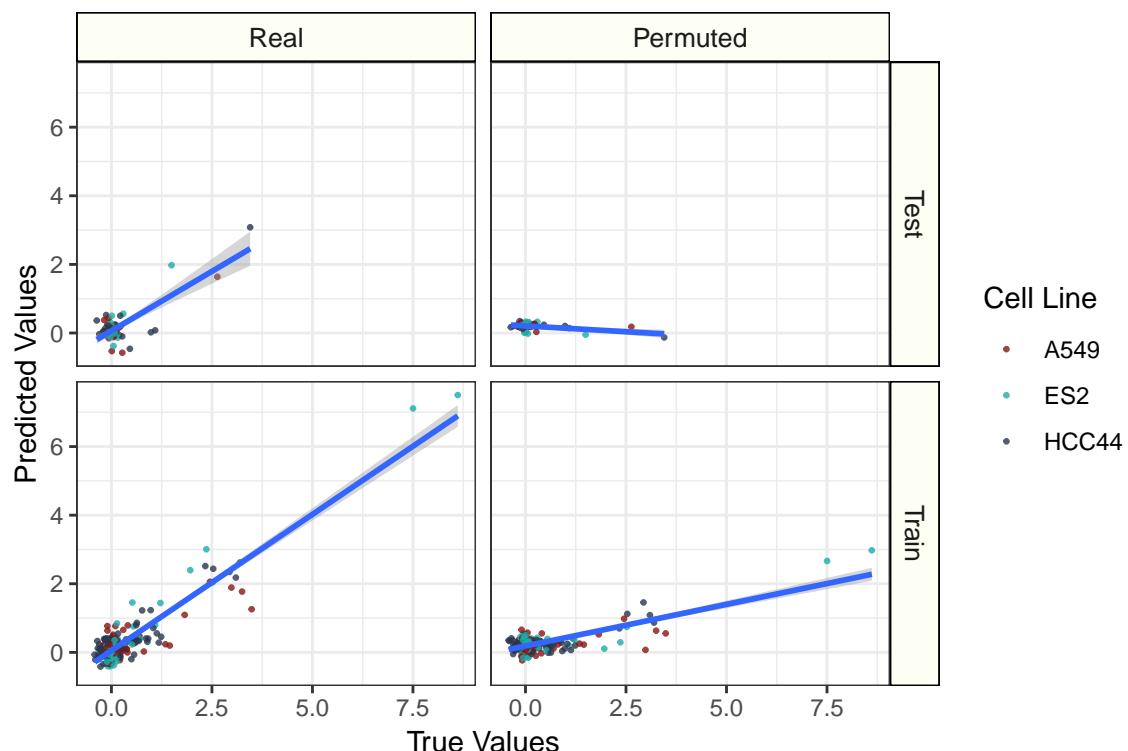


fit	shuffle	transform	r2	value
Train	Real	raw	0.17	0.26
Test	Real	raw	0.00	0.24
Train	Shuffle	raw	0.34	0.21
Test	Shuffle	raw	-0.18	0.28

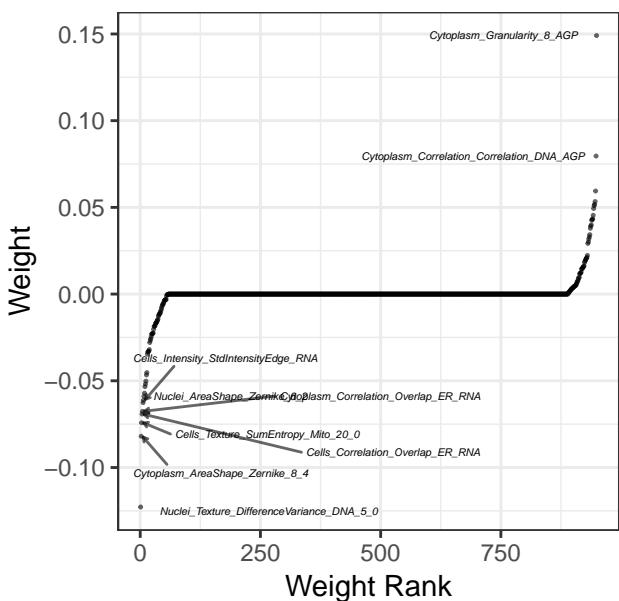


Performance: vb_percent_dead

Transform: raw

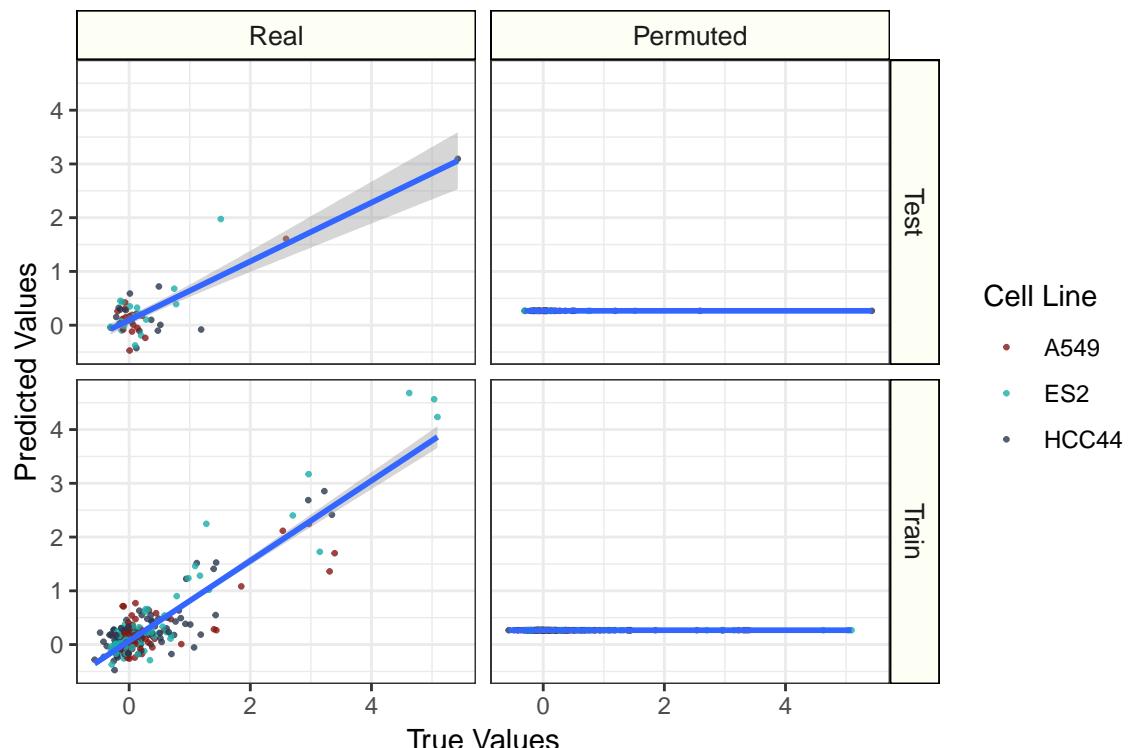


fit	shuffle	transform	r2	value
Train	Real	raw	0.85	0.12
Test	Real	raw	0.62	0.16
Train	Shuffle	raw	0.39	0.48
Test	Shuffle	raw	-0.16	0.48

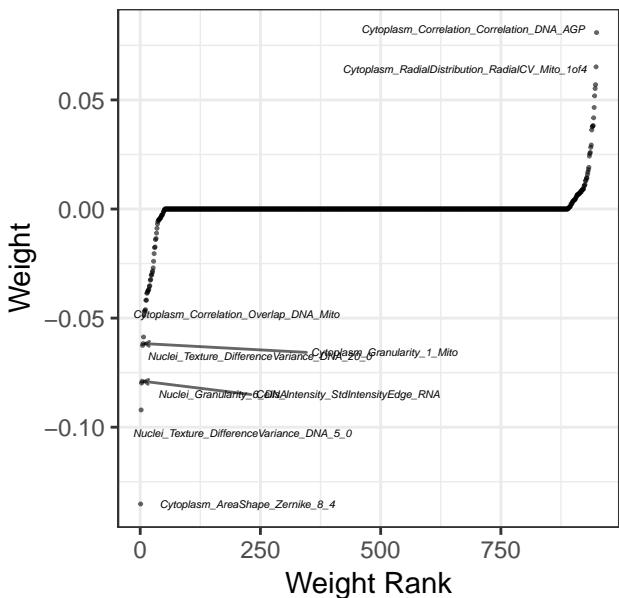


Performance: vb_percent_dead_only

Transform: raw

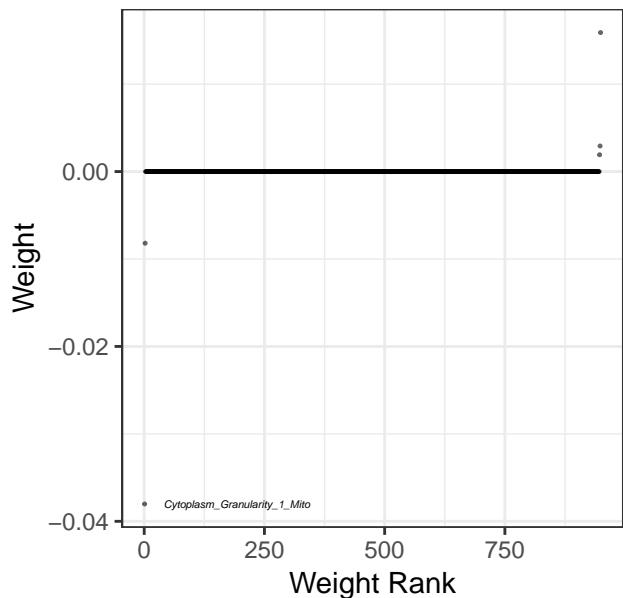
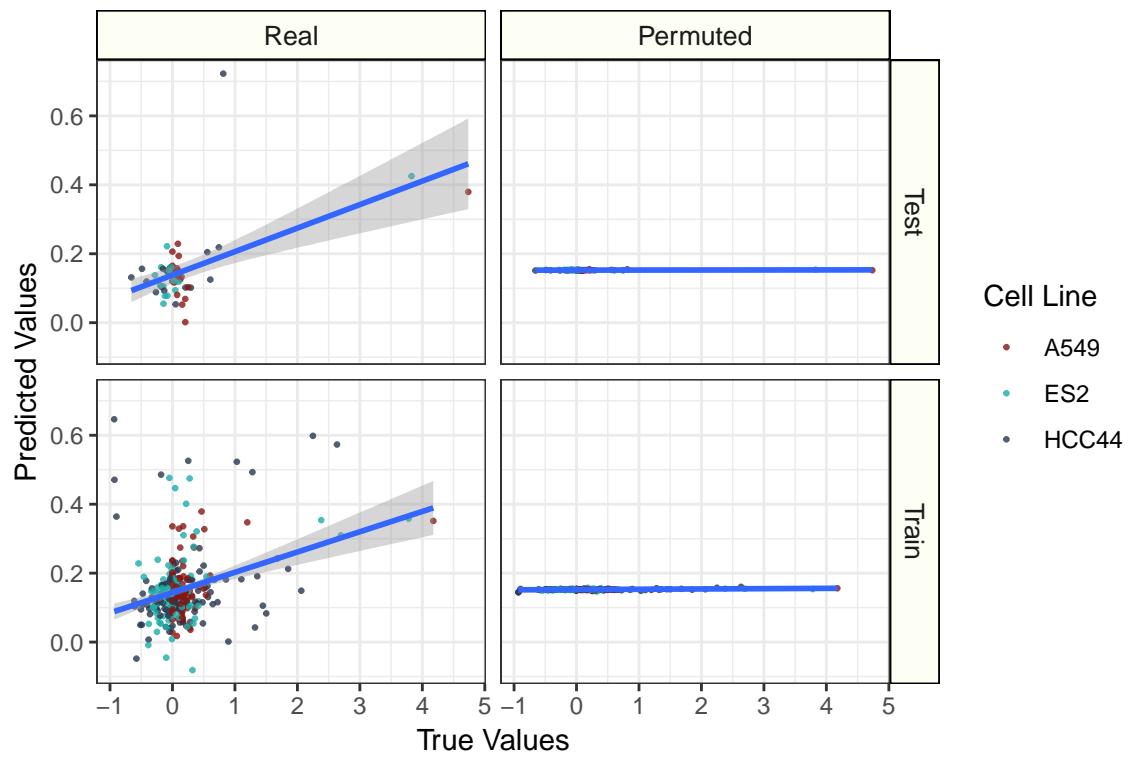


fit	shuffle	transform	r2	value
Train	Real	raw	0.80	0.12
Test	Real	raw	0.66	0.25
Train	Shuffle	raw	0.00	0.63
Test	Shuffle	raw	0.00	0.73



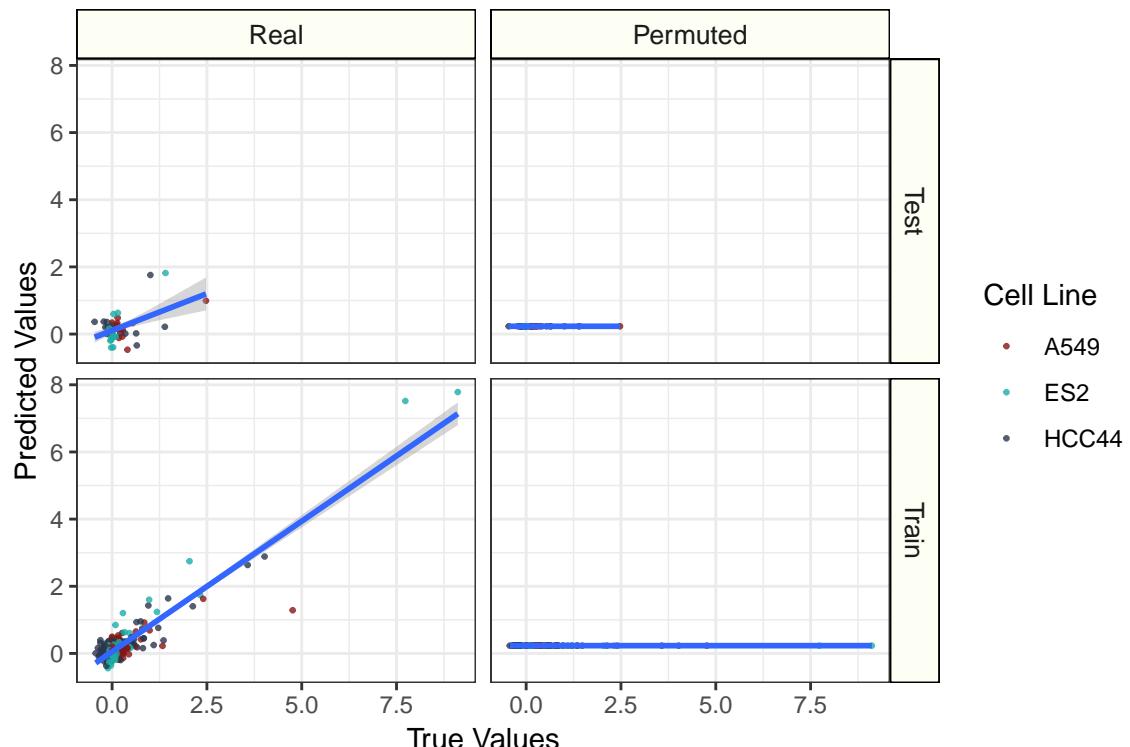
Performance: vb_percent_early_apoptosis

Transform: raw

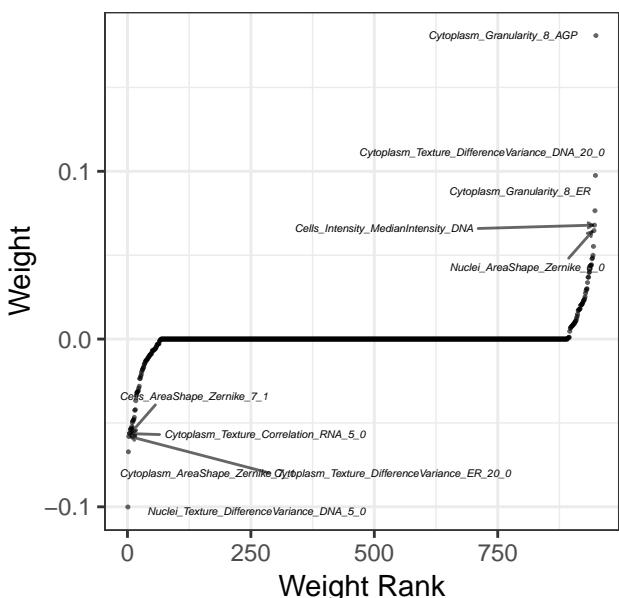


Performance: vb_percent_late_apoptosis

Transform: raw

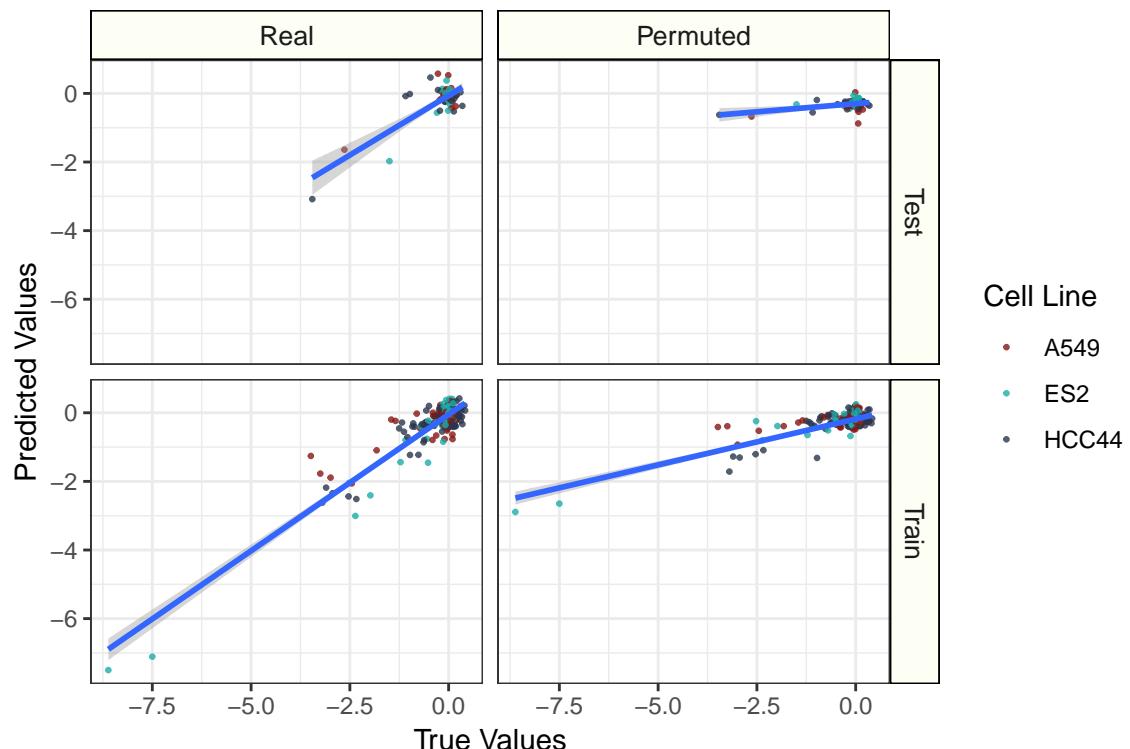


fit	shuffle	transform	r2	value
Train	Real	raw	0.84	0.12
Test	Real	raw	0.12	0.19
Train	Shuffle	raw	0.00	0.75
Test	Shuffle	raw	-0.01	0.22

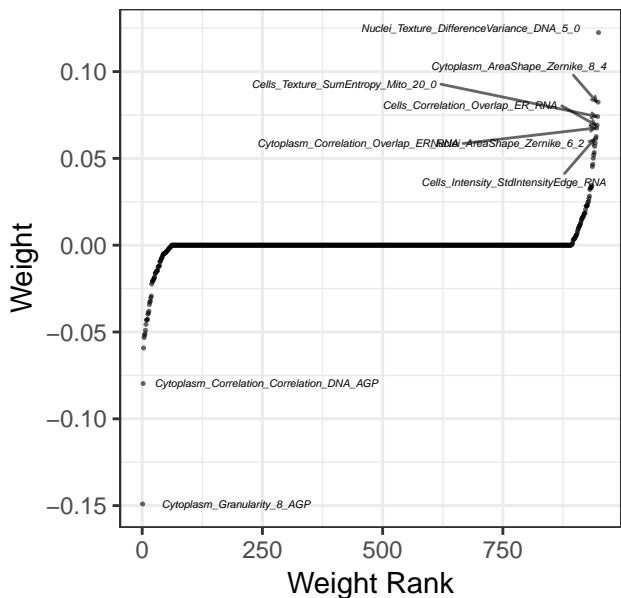


Performance: vb_percent_live

Transform: raw

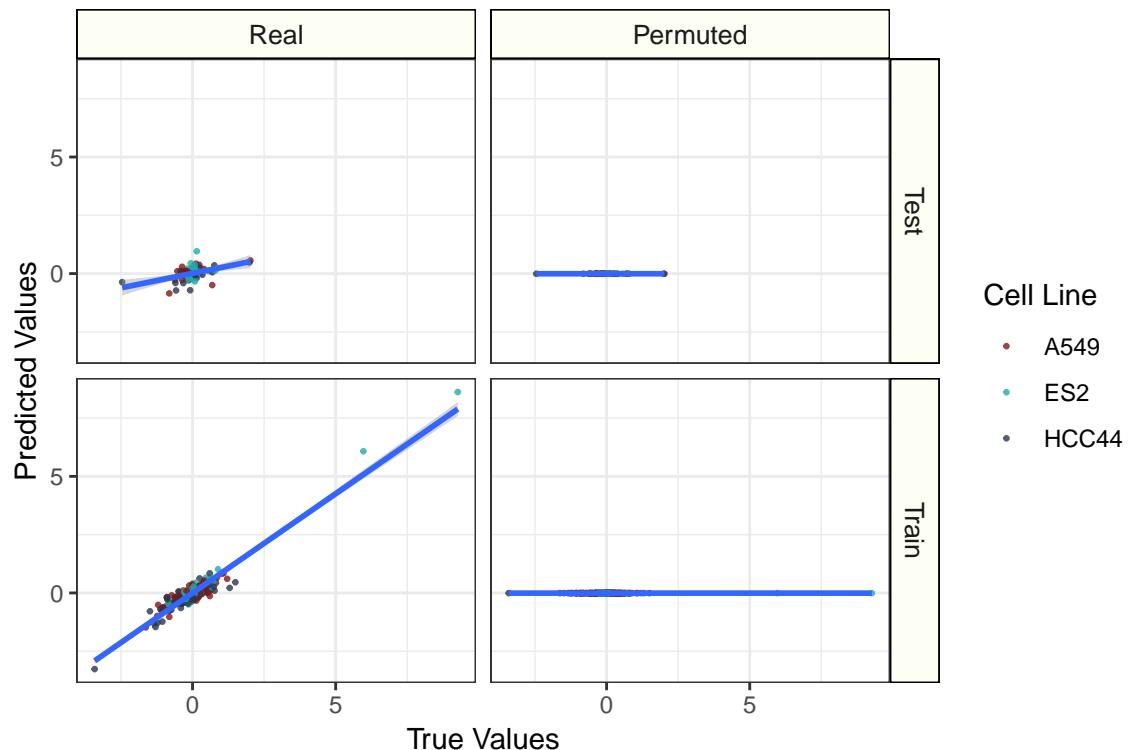


fit	shuffle	transform	r2	value
Train	Real	raw	0.85	0.12
Test	Real	raw	0.62	0.16
Train	Shuffle	raw	0.42	0.45
Test	Shuffle	raw	0.10	0.38

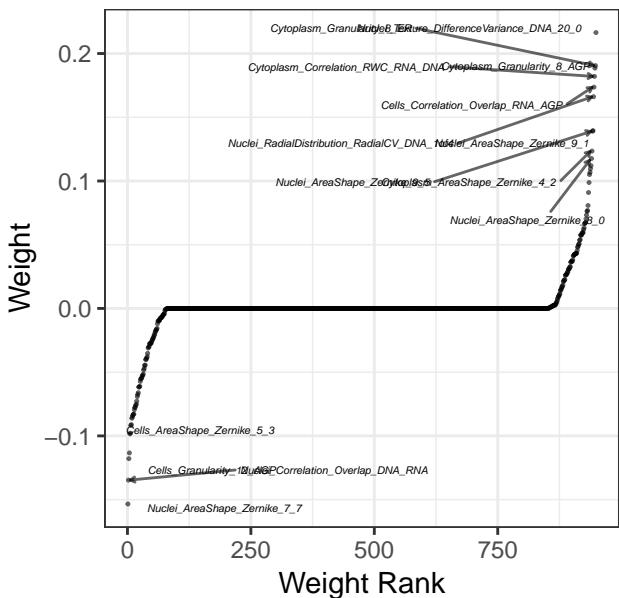


Performance: vb_ros_back_mean

Transform: raw

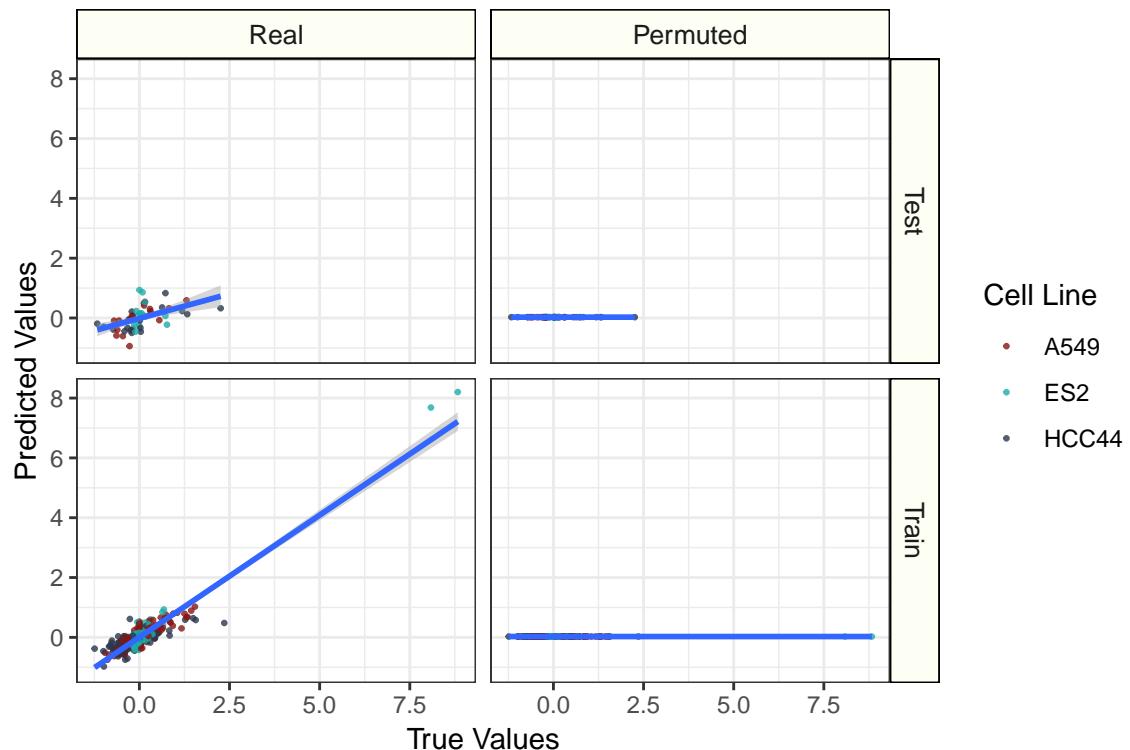


fit	shuffle	transform	r2	value
Train	Real	raw	0.90	0.07
Test	Real	raw	0.21	0.30
Train	Shuffle	raw	0.00	0.65
Test	Shuffle	raw	0.00	0.38



Performance: vb_ros_mean

Transform: raw



fit	shuffle	transform	r2	value
Train	Real	raw	0.87	0.09
Test	Real	raw	0.21	0.27
Train	Shuffle	raw	0.00	0.68
Test	Shuffle	raw	0.00	0.35

