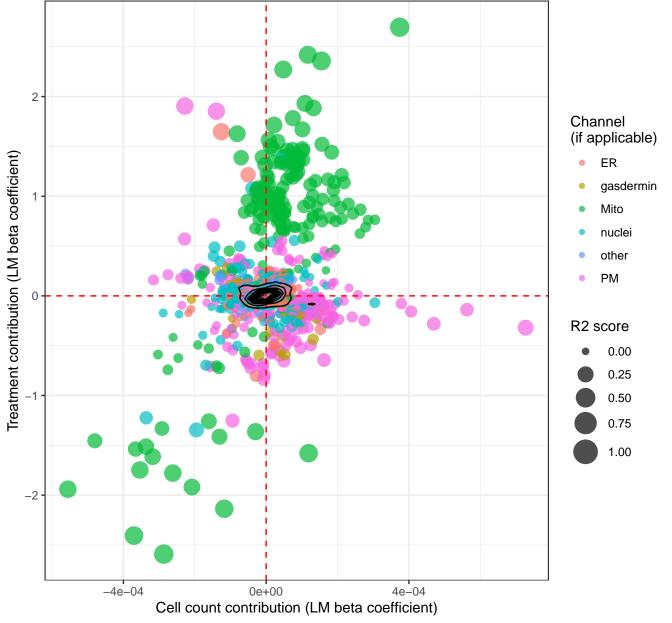
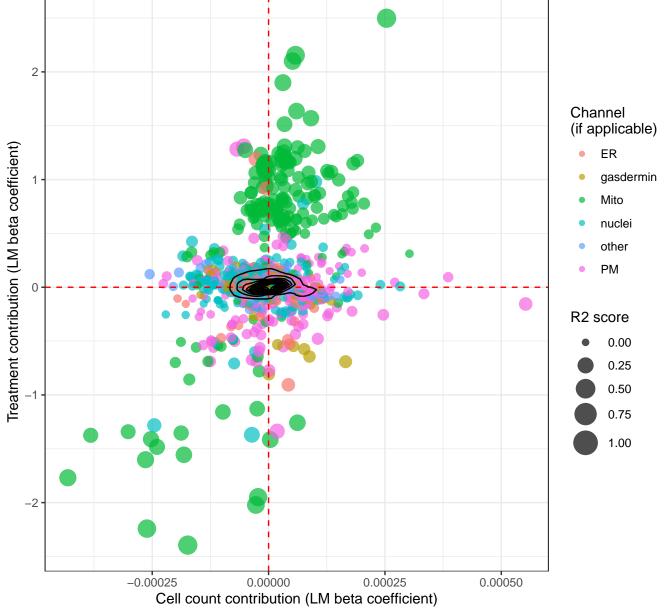
How CellProfiler features contribute to H2O2_100.000_Disulfiram_1.0 treatments and cell density 2 -Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other PM R2 score 0.00 0.25 0.50 0.75 1.00 -2 **-**-4e-04 0e+00 4e-04 Cell count contribution (LM beta coefficient)

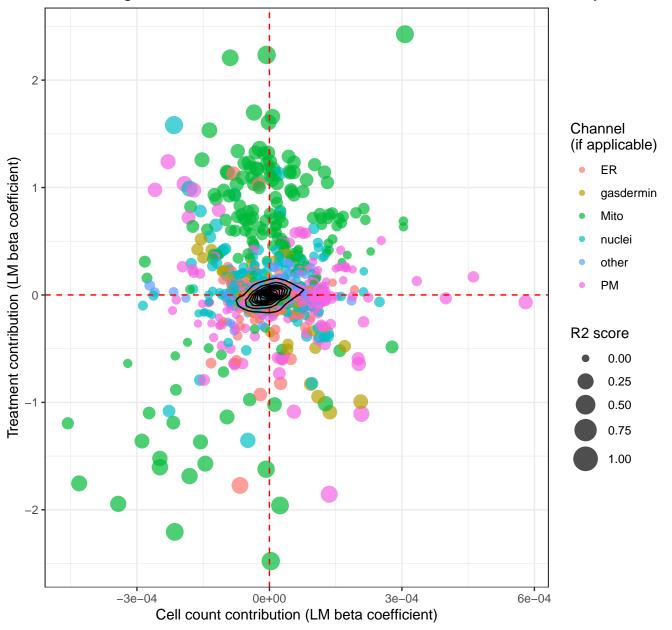
How CellProfiler features contribute to LPS_Nigericin_100.000_10.0_DMSO_0.025 treatments and cell density



How CellProfiler features contribute to LPS_Nigericin_1.000_3.0_DMSO_0.025 treatments and cell density



How CellProfiler features contribute to LPS_Nigericin_1.000_10.0_Disulfiram_1.0 treatments and cell density



How CellProfiler features contribute to LPS_Nigericin_1.000_10.0_DMSO_0.025 treatments and cell density 2 -Channel (if applicable) ER gasdermin Mito nuclei other PM R2 score 0.00 0.25 0.50 0.75 1.00 -2 -0.000250.00000 0.00025 0.00050

Cell count contribution (LM beta coefficient)

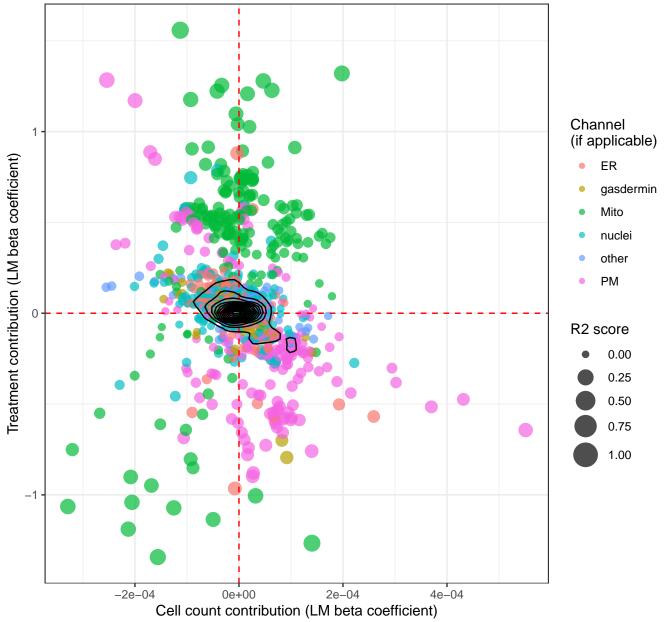
Treatment contribution (LM beta coefficient)

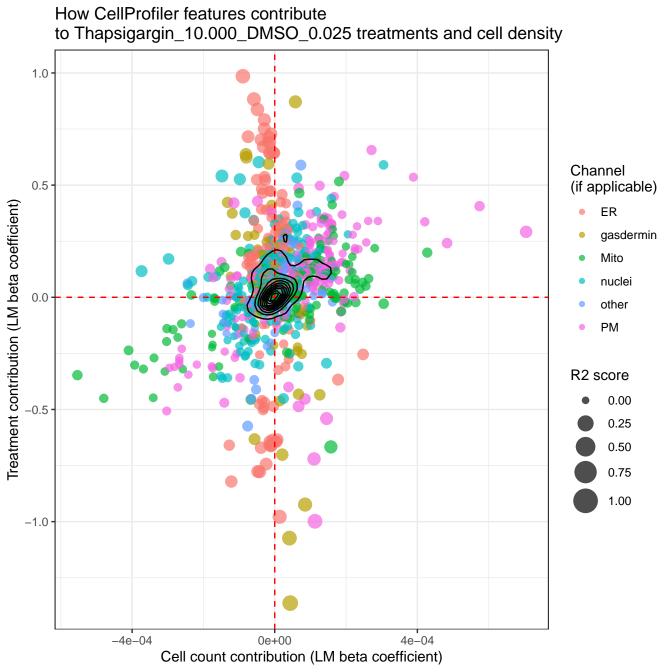
How CellProfiler features contribute to LPS_100.000_DMSO_0.025 treatments and cell density Channel (if applicable) ER gasdermin Mito nuclei other PM R2 score 0.00 0.25 0.50 0.75 1.00 -0.00025 0.00000 0.00025 0.00050

Cell count contribution (LM beta coefficient)

Treatment contribution (LM beta coefficient)

How CellProfiler features contribute to LPS_Nigericin_100.000_3.0_DMSO_0.025 treatments and cell density

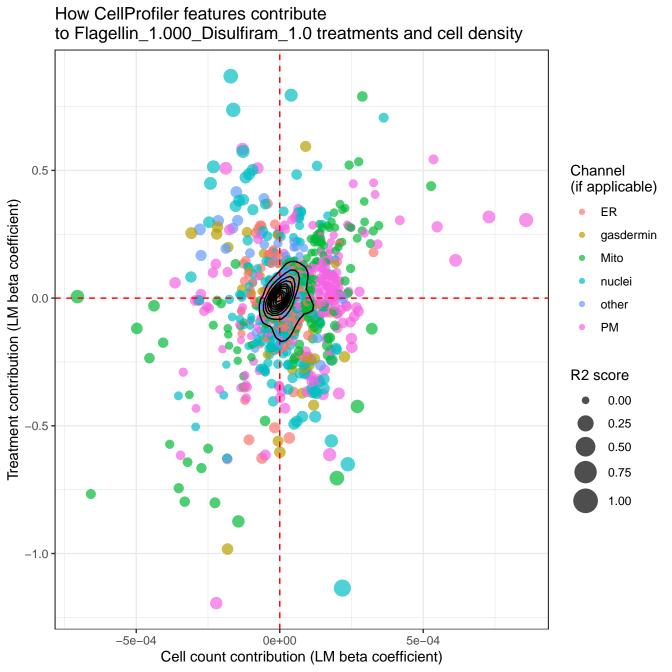


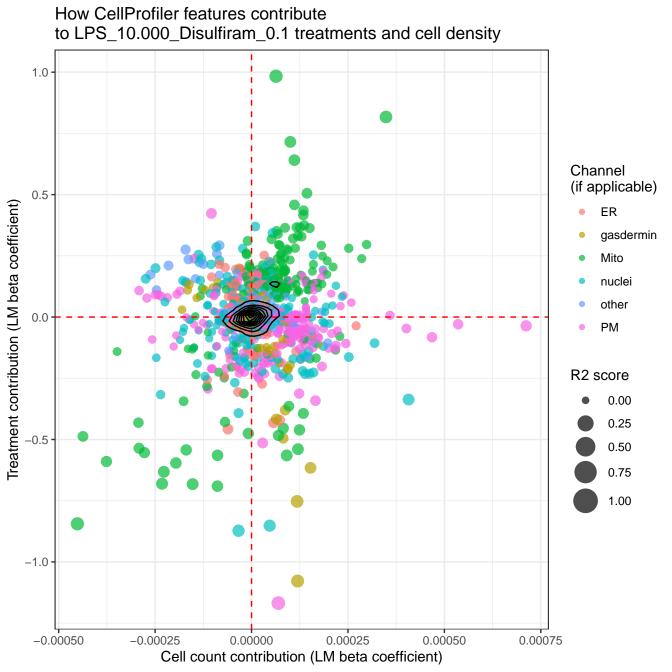


How CellProfiler features contribute to LPS_Nigericin_1.000_10.0_Z-VAD-FMK_100.0 treatments and cell density 1.0 Channel (if applicable) ER 0.5 gasdermin Mito nuclei other 0.0 PM R2 score 0.00 0.25 -0.5 0.50 0.75 1.00 -1.0 --4e-04 0e+00 4e-04 8e-04 Cell count contribution (LM beta coefficient)

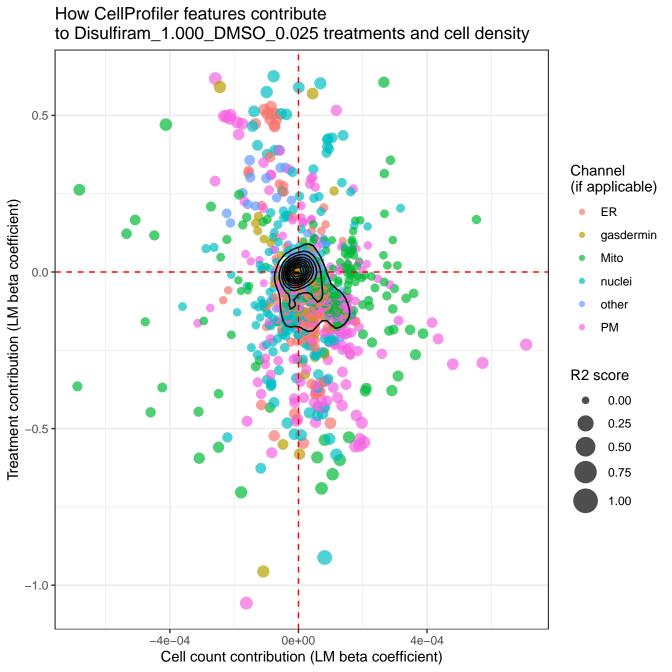
Treatment contribution (LM beta coefficient)

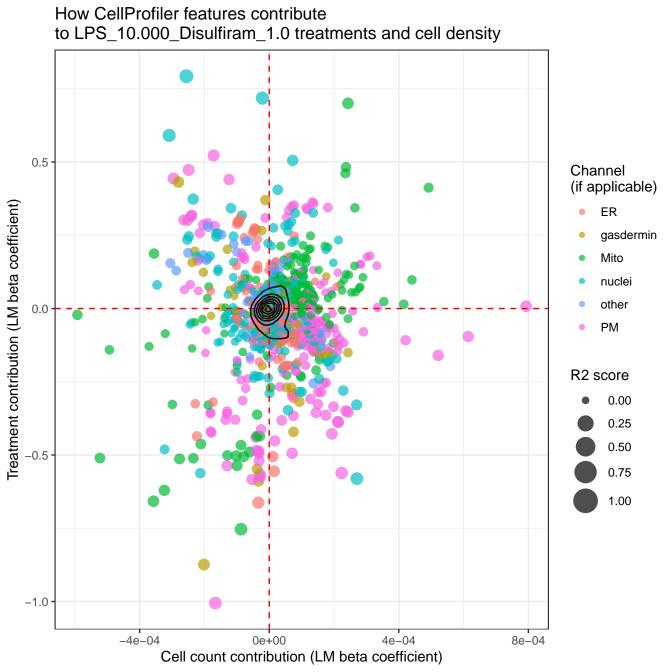
How CellProfiler features contribute to LPS_Nigericin_100.000_1.0_DMSO_0.025 treatments and cell density 1.0 Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin 0.5 Mito nuclei other PM 0.0 R2 score 0.00 0.25 0.50 -0.5 0.75 1.00 -1.0-3e-04 0e+00 3e-04 6e-04 Cell count contribution (LM beta coefficient)

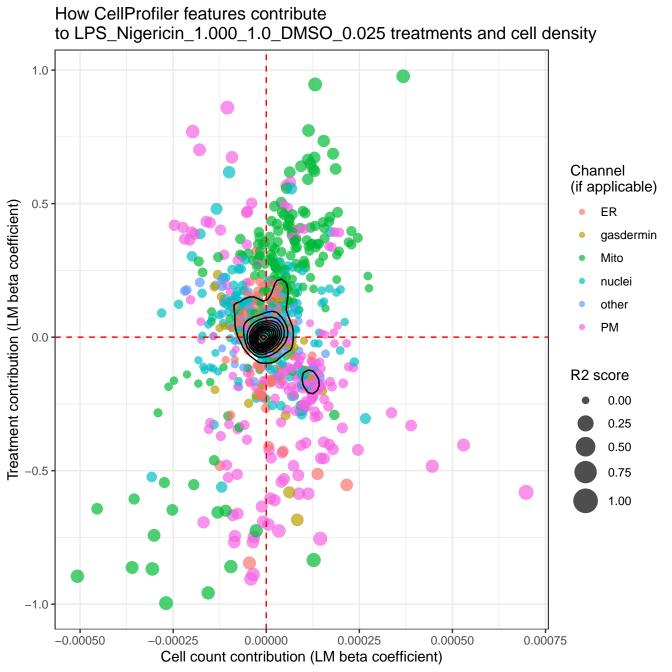


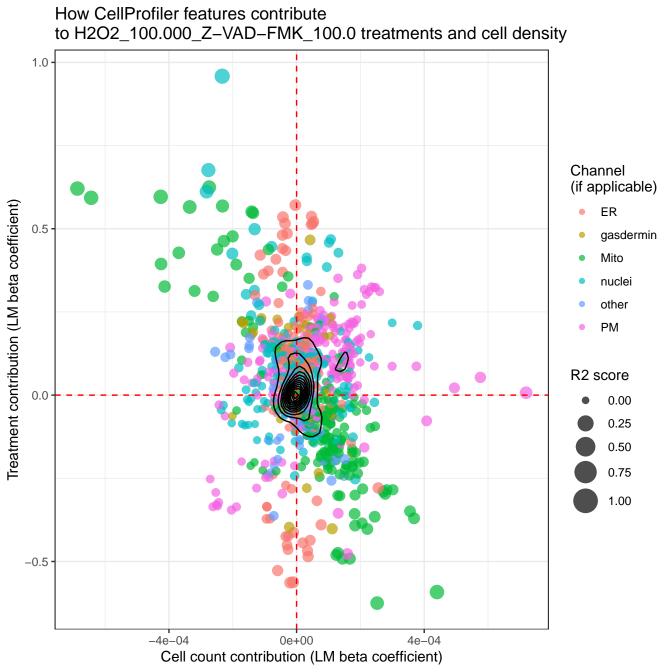


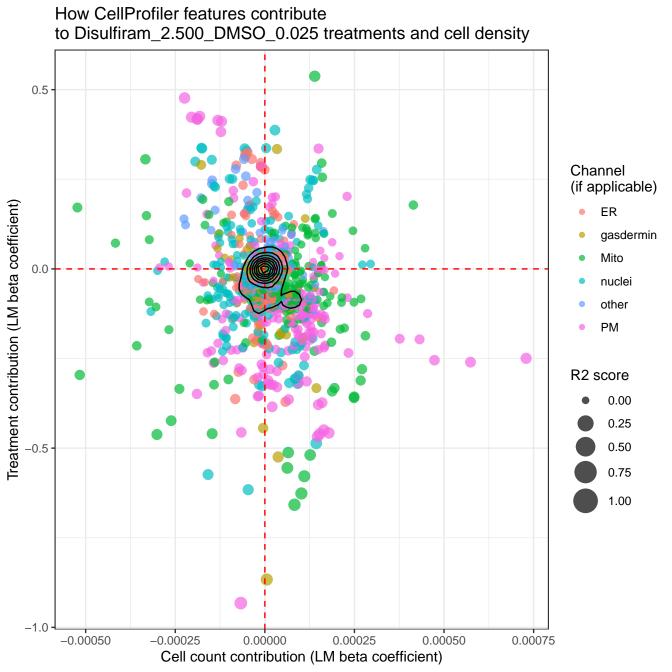
How CellProfiler features contribute to LPS_10.000_Disulfiram_2.5 treatments and cell density Channel 0.5 (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other 0.0 PM R2 score 0.00 0.25 0.5 0.50 0.75 1.00 -1.0-6e-04 -3e-04 0e+00 3e-04 6e-04 Cell count contribution (LM beta coefficient)

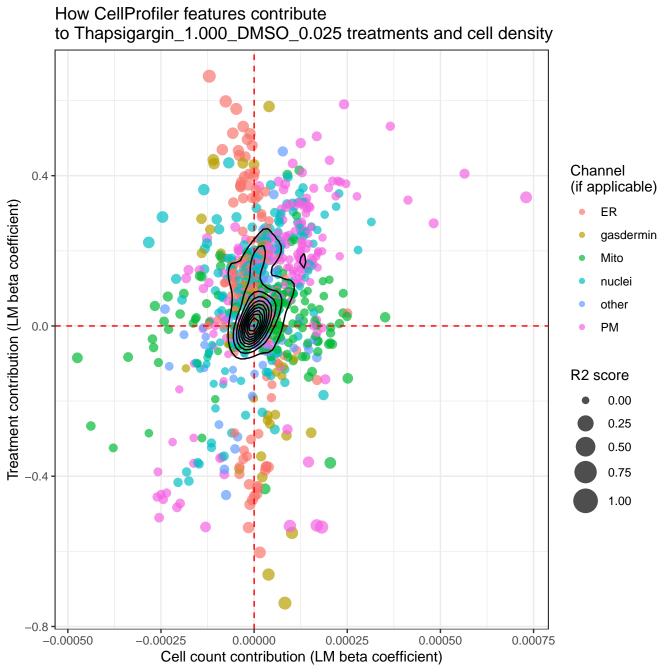


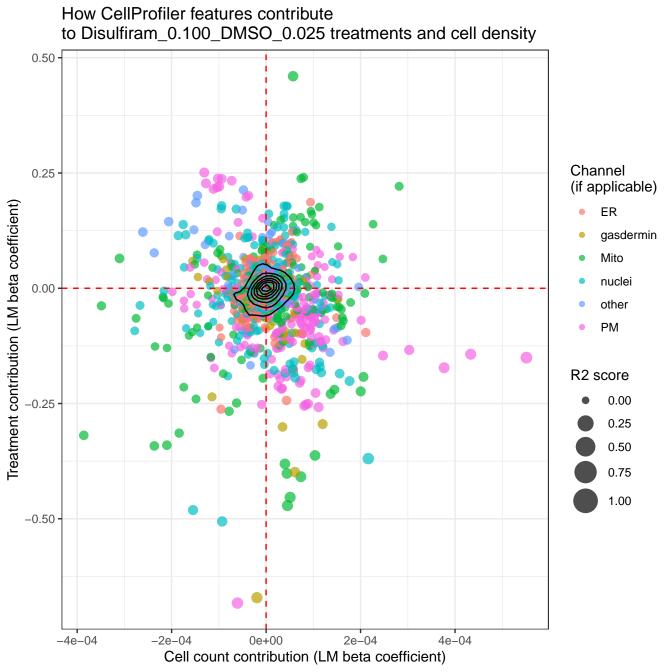




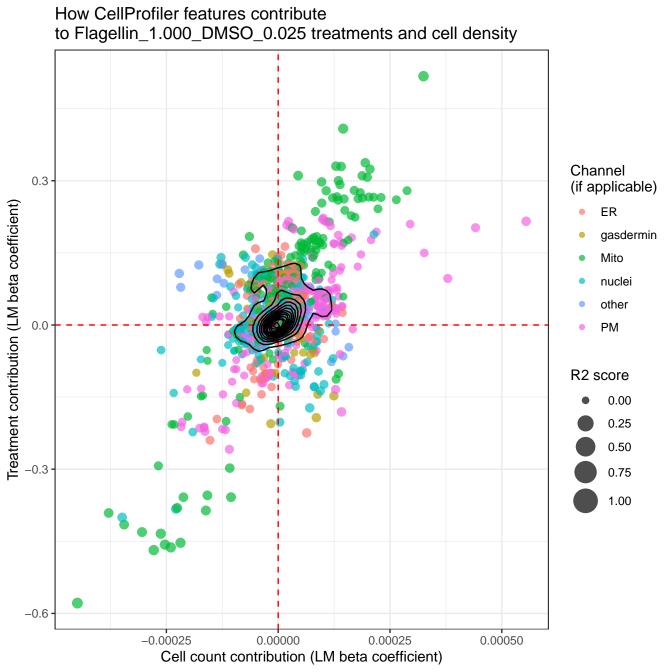


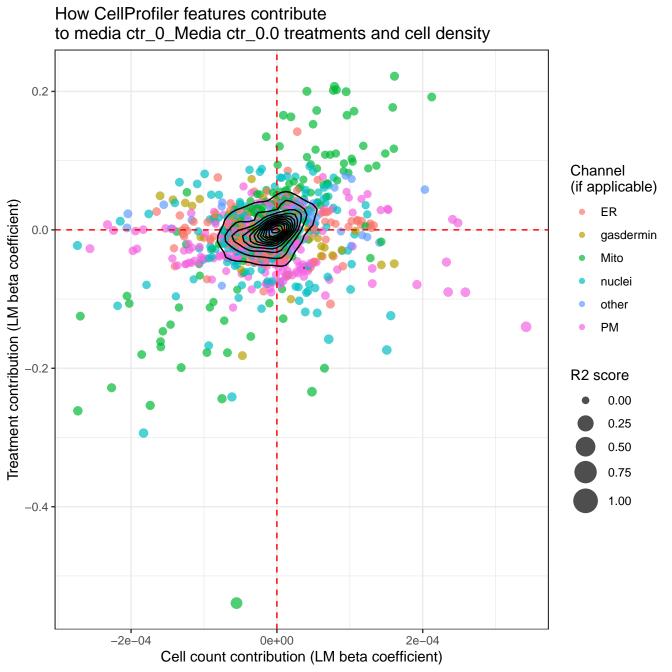


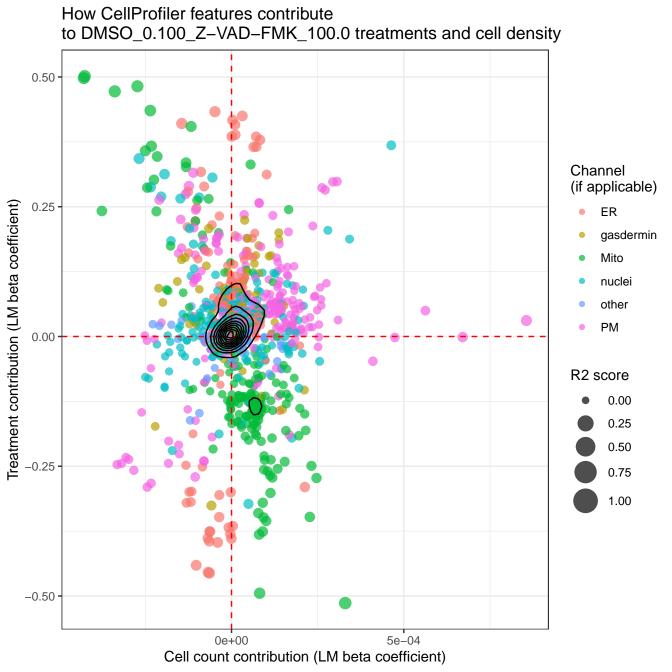


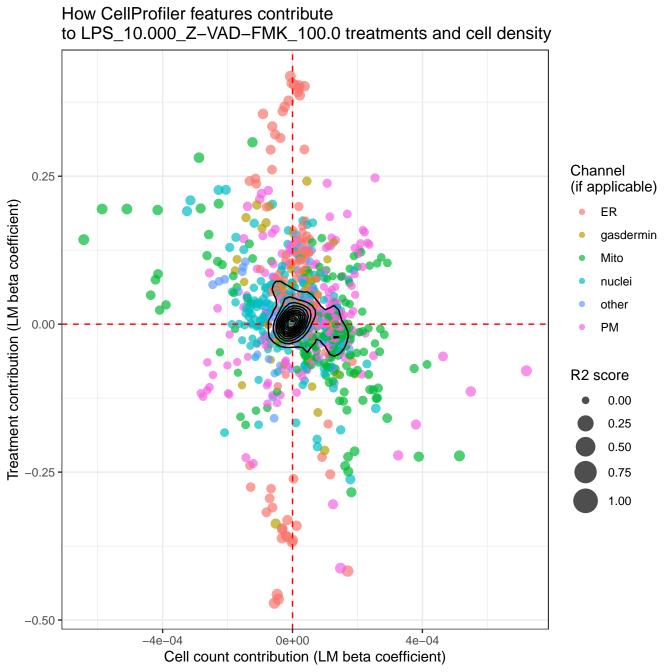


How CellProfiler features contribute to Flagellin_0.100_DMSO_0.025 treatments and cell density 0.4 Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other PM 0.0 R2 score 0.00 0.25 0.50 0.75 -0.4 -1.00 0.00000 -0.00050-0.00025 0.00025 0.00050 Cell count contribution (LM beta coefficient)









How CellProfiler features contribute to DMSO_0.100_DMSO_1.0 treatments and cell density 0.4 Channel (if applicable) Treatment contribution (LM beta coefficient) ER 0.2 gasdermin Mito nuclei other PM 0.0 R2 score 0.00 0.25 0.50 0.75 -0.21.00 -4e-04 -2e-04 0e+00 2e-04 4e-04 6e-04 Cell count contribution (LM beta coefficient)

How CellProfiler features contribute to LPS_10.000_DMSO_0.025 treatments and cell density 0.4 Channel (if applicable) 0.2 -Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other 0.0 PM R2 score 0.00 0.25 0.50 -0.2 0.75 1.00 -0.4-0.00050-0.000250.00000 0.00025 0.00050 0.00075 Cell count contribution (LM beta coefficient)

