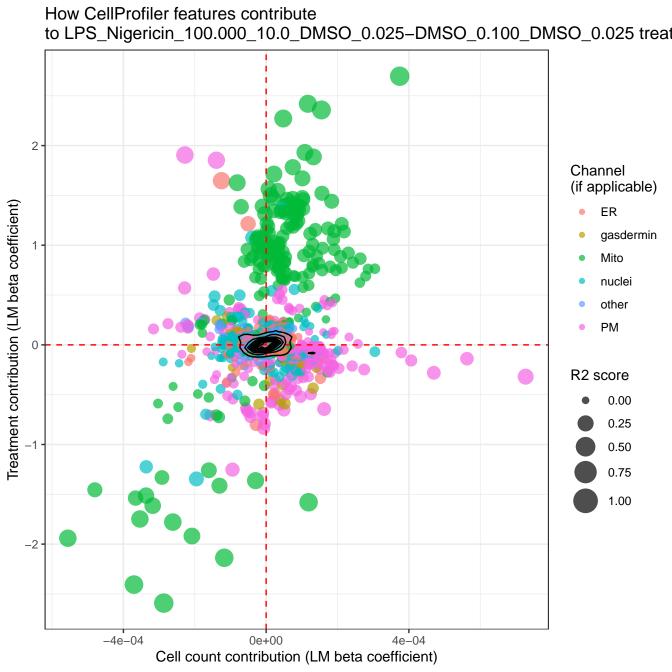
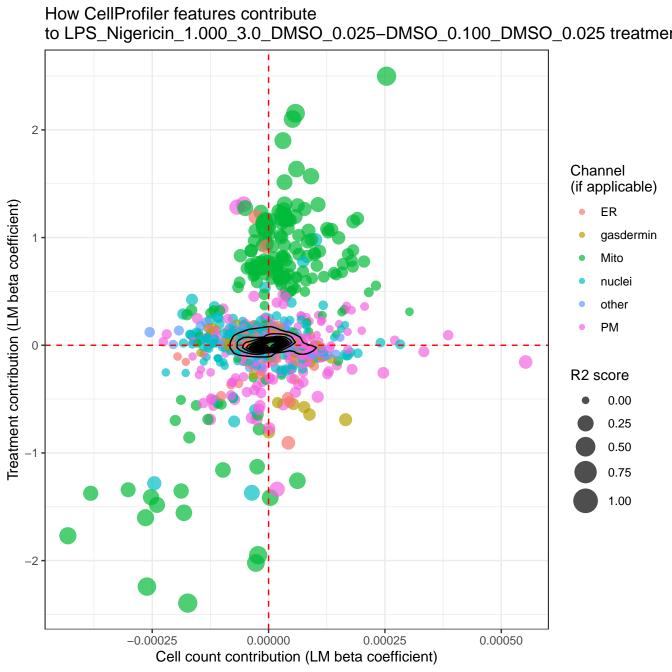
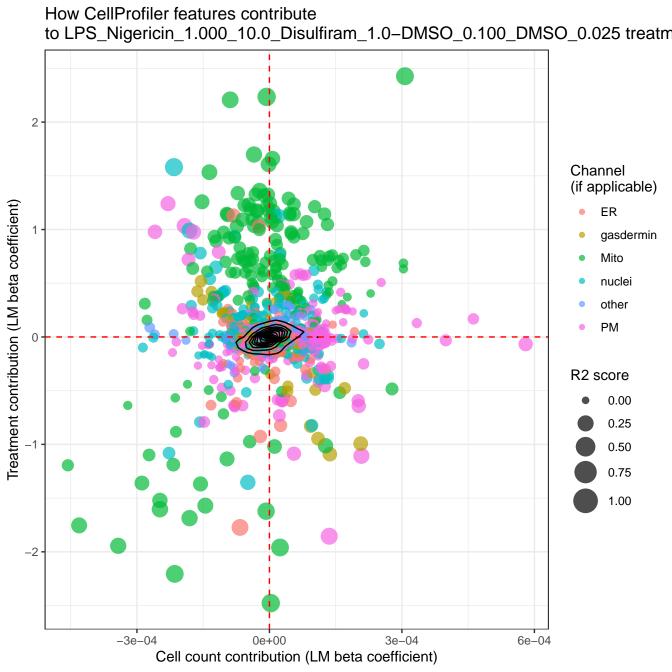
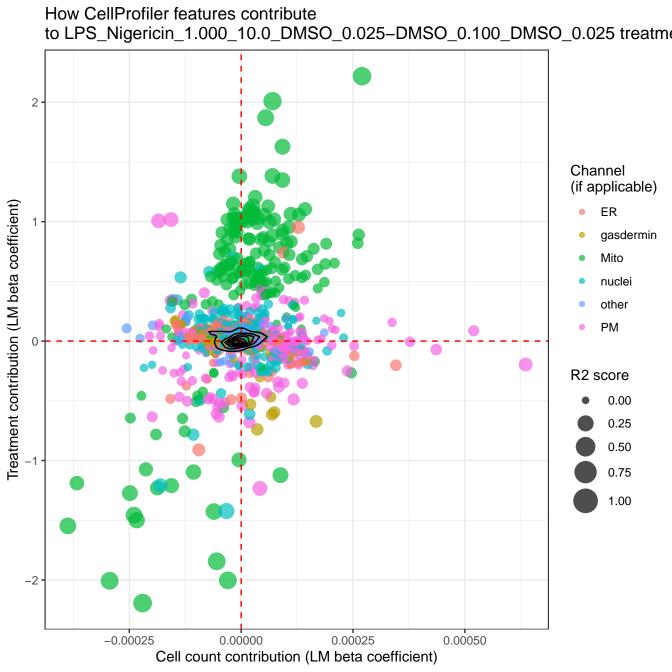
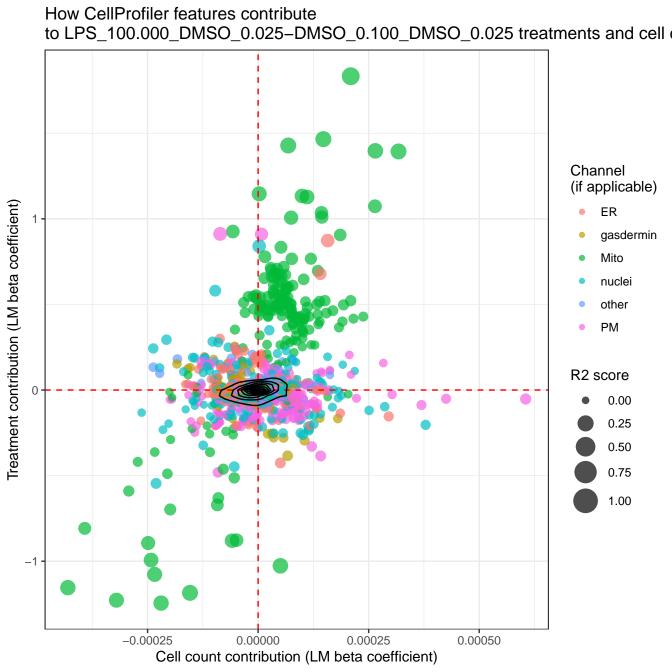
How CellProfiler features contribute to H2O2\_100.000\_Disulfiram\_1.0-DMSO\_0.100\_DMSO\_0.025 treatments and ce 2 -Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other PM R2 score 0.00 0.25 0.50 0.75 1.00 -2. -4e-04 0e+00 4e-04 Cell count contribution (LM beta coefficient)





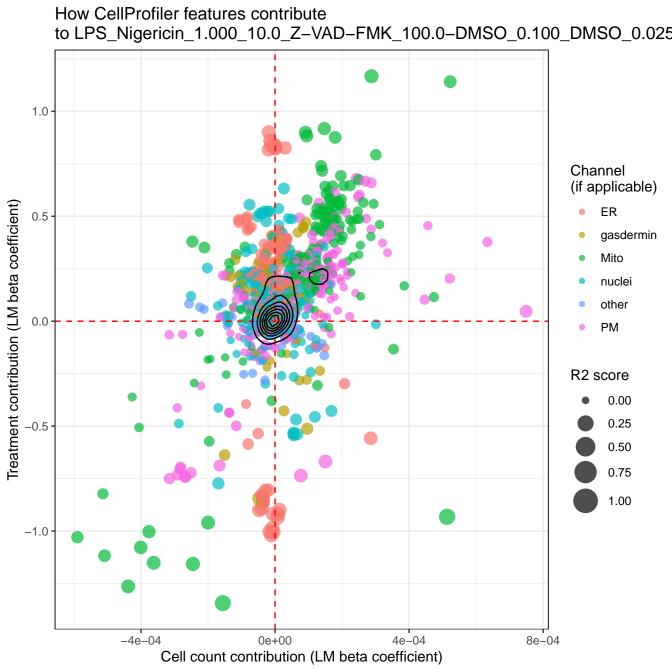


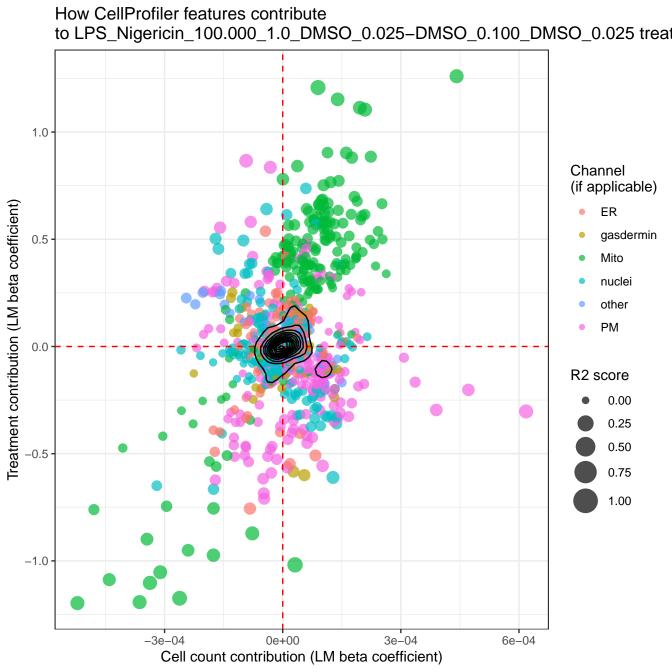


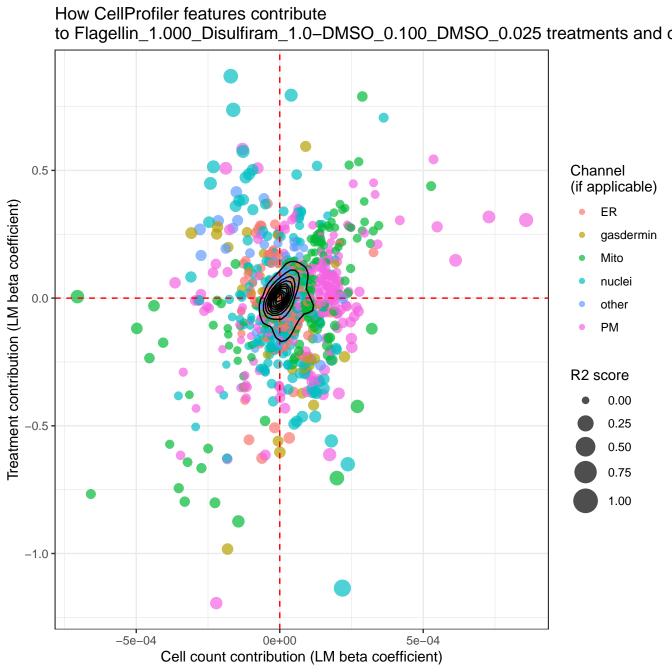


How CellProfiler features contribute to LPS\_Nigericin\_100.000\_3.0\_DMSO\_0.025-DMSO\_0.100\_DMSO\_0.025 treatments Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other PM R2 score 0.00 0.25 0.50 0.75 1.00 -1 · -2e-04 0e+00 2e-04 4e-04 Cell count contribution (LM beta coefficient)

How CellProfiler features contribute to Thapsigargin\_10.000\_DMSO\_0.025-DMSO\_0.100\_DMSO\_0.025 treatments 1.0 Channel 0.5 (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei 0.0 other PM R2 score 0.00 0.5 0.25 0.50 0.75 1.00 -1.0-4e-04 0e+00 4e-04 Cell count contribution (LM beta coefficient)





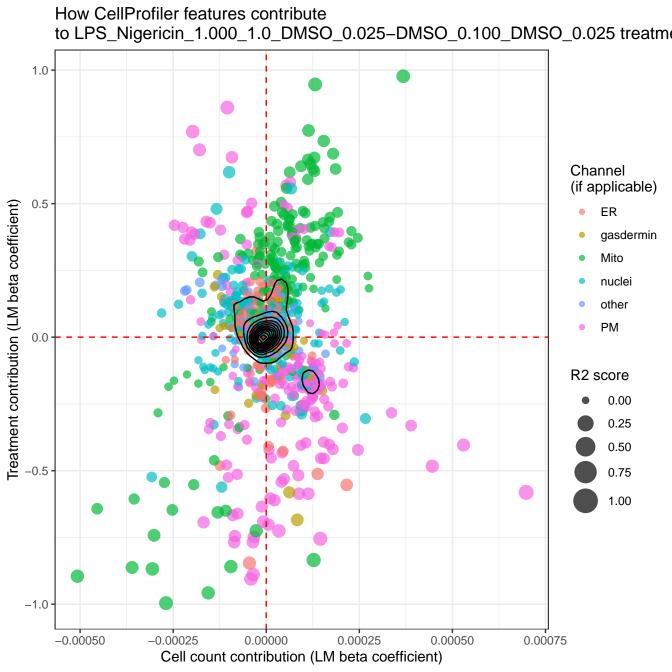


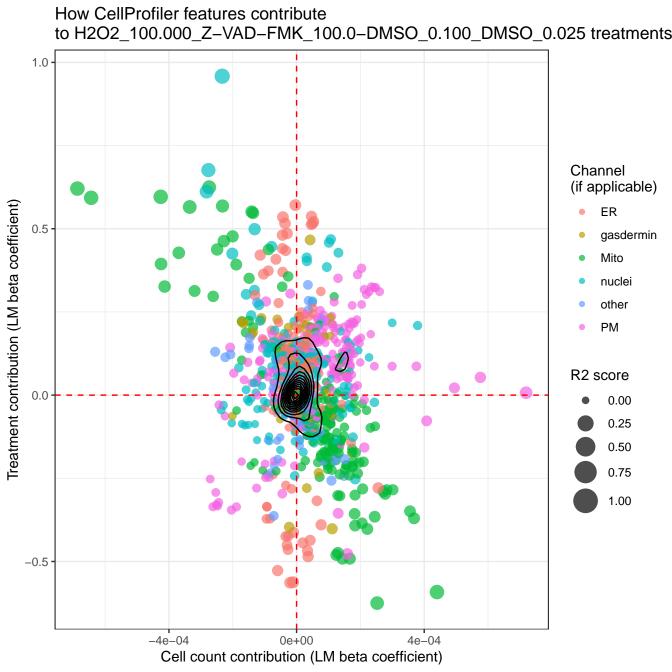
How CellProfiler features contribute to LPS\_10.000\_Disulfiram\_0.1-DMSO\_0.100\_DMSO\_0.025 treatments and cell 1.0 Channel (if applicable) 0.5 Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other 0.0 PM R2 score 0.00 0.25 -0.5 0.50 0.75 1.00 -1.0-0.00050-0.00025 0.00000 0.00025 0.00050 0.00075 Cell count contribution (LM beta coefficient)

How CellProfiler features contribute to LPS\_10.000\_Disulfiram\_2.5-DMSO\_0.100\_DMSO\_0.025 treatments and cell 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)

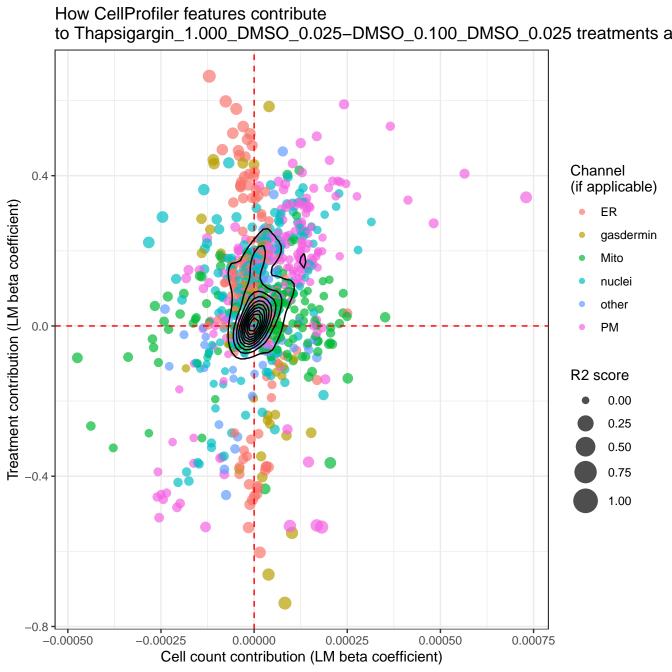
How CellProfiler features contribute to Disulfiram\_1.000\_DMSO\_0.025-DMSO\_0.100\_DMSO\_0.025 treatments and 0.5 Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito 0.0 nuclei other PM R2 score 0.00 0.25 -0.5 0.50 0.75 1.00 -1.0 --4e-04 0e+00 4e-04 Cell count contribution (LM beta coefficient)

How CellProfiler features contribute to LPS\_10.000\_Disulfiram\_1.0-DMSO\_0.100\_DMSO\_0.025 treatments and cell 0.5 Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other 0.0 PM R2 score 0.00 0.25 0.50 -0.5 0.75 1.00 -1.0 **-**-4e-04 0e+00 4e-04 8e-04 Cell count contribution (LM beta coefficient)

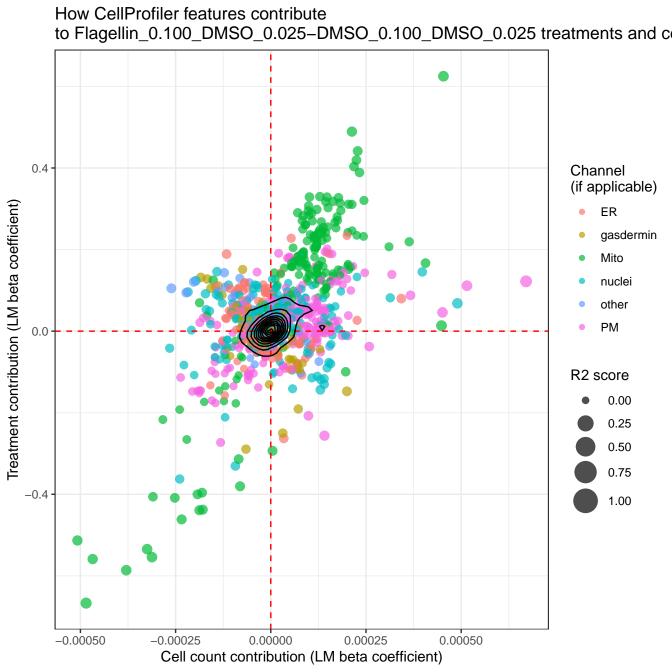


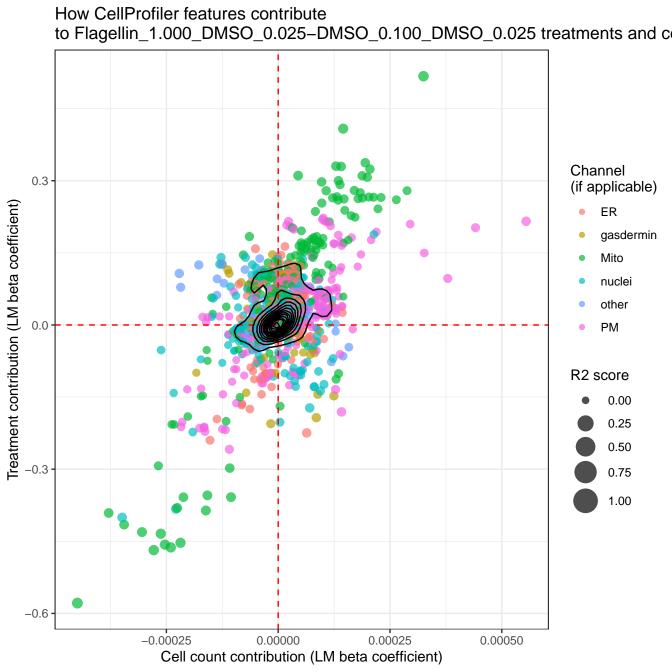


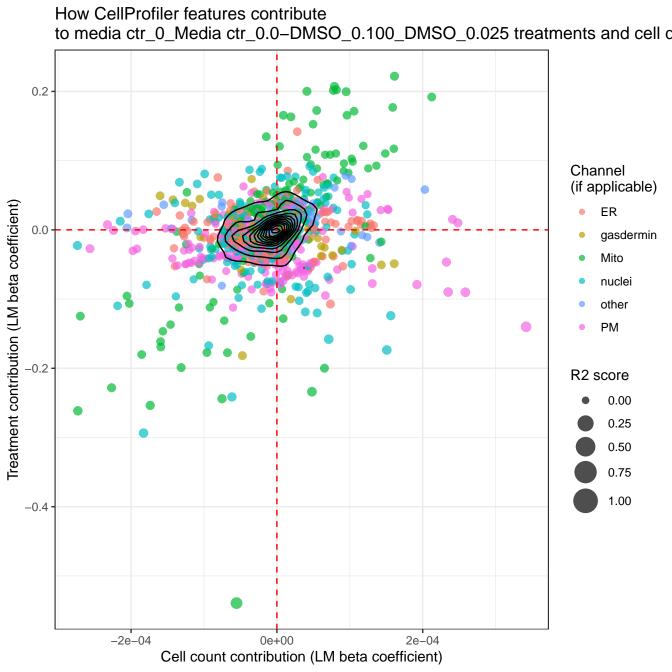
How CellProfiler features contribute to Disulfiram\_2.500\_DMSO\_0.025-DMSO\_0.100\_DMSO\_0.025 treatments and 0.5 Channel (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito 0.0 nuclei other PM R2 score 0.00 0.25 0.50 -0.5 0.75 1.00 -1.0-0.00050-0.000250.00000 0.00025 0.00050 0.00075 Cell count contribution (LM beta coefficient)



How CellProfiler features contribute to Disulfiram\_0.100\_DMSO\_0.025-DMSO\_0.100\_DMSO\_0.025 treatments and 0.50 Channel 0.25 -(if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei 0.00 other PM R2 score 0.00 -0.25 **-**0.25 0.50 0.75 1.00 -0.50 --2e-04 0e+00 2e-04 4e-04 -4e-04Cell count contribution (LM beta coefficient)







How CellProfiler features contribute to DMSO\_0.100\_Z-VAD-FMK\_100.0-DMSO\_0.100\_DMSO\_0.025 treatments 0.50 Channel (if applicable) Treatment contribution (LM beta coefficient) 0.25 ER gasdermin Mito nuclei other PM 0.00 R2 score 0.00 0.25 0.50 -0.25 0.75 1.00 -0.50 **-**0e+00 5e-04 Cell count contribution (LM beta coefficient)

How CellProfiler features contribute to LPS\_10.000\_Z-VAD-FMK\_100.0-DMSO\_0.100\_DMSO\_0.025 treatments a Channel 0.25 (if applicable) Treatment contribution (LM beta coefficient) ER gasdermin Mito nuclei other 0.00 PM R2 score 0.00 0.25 0.50 -0.25 0.75 1.00 -0.50-4e-04 0e+00 4e-04 Cell count contribution (LM beta coefficient)

