CellScanner

Import Data

Train Model

Run Prediction

Coculture Options for Prediction

01-t12_d50_wc_btriA-D7,01-t12_d50_wc_btriB-D8,01-t12_d50_wc_btriC-D9,01-t12_d50_wc_btriD-D10,01-t12_d50_wc_btriE-D11,01-t12_d50_wc_btriF-D12

Choose the Channels that will be used as x, y, z axis for the 3D plot:

X Axis: | SSC-H

Y Axis: | FSC-H

Z Axis: | FITC-H

☑ Apply filtering on the predictions based on their uncertainty scores.

Threshold for uncertainty filtering: | 0,95

☑ Apply line gating

Staining inactive cells (e.g. PI): | PI-H | > | 3.1

Staining all cells (e.g. SYBR/DAPI): | FITC-A | > | 4.5

Add extra stain

Predict