

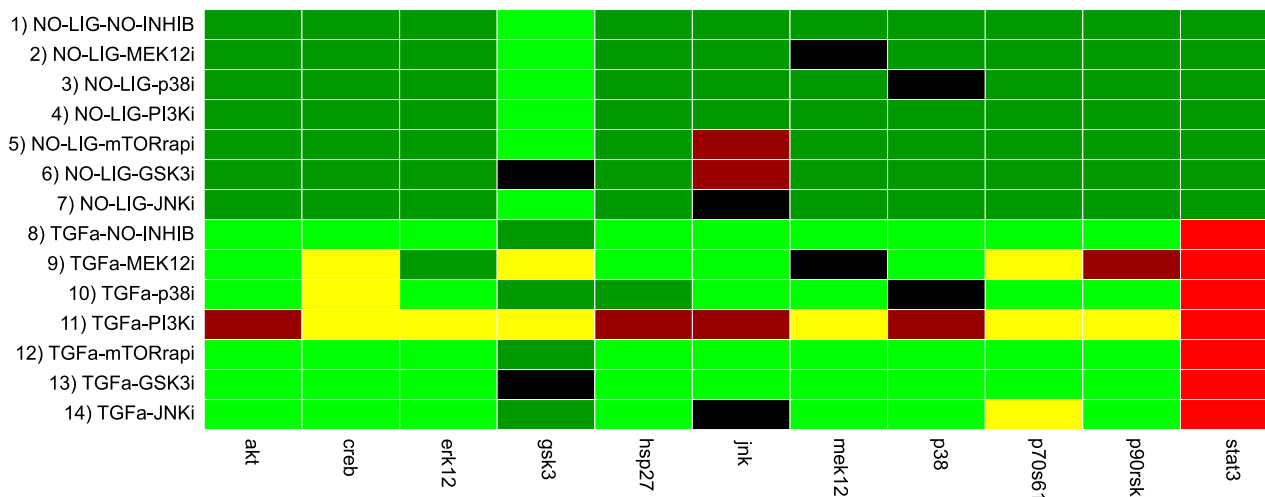
Figure S3. Comparison of the discretized data with predictions from model M2.

A Primary human hepatocytes (data from Alexopoulos *et al*, in preparation).

B HepG2 cells (the horizontal line separates the the first (top) from the second (bottom) dataset for HepG2 cells; see also text).

Each row represents one treatment and the readouts are shown in the columns. Light green: predicted correctly, "on"; dark green: predicted correctly, "off"; light red: predicted "on", measured "off"; dark red: predicted "off", measured "on"; yellow: state cannot be determined in logical steady state analysis; black: data points where the measured species is inhibited are not considered.

A



B

