File Name: Supplementary Data 1  
**Description:** Distribution of Robust Rank Aggregation (RRA) scores in genome-wide negative CRISPR/Cas9 screens (GeCKOV2 library and Brunello library);

File Name: Supplementary Data 2  
**Description:** Targeted metabonomics unprocessed data determined by LC-MS in HeLa cells expressing inducible Tet-on LbNOX, EcSTH or EcSTH-LbNOX cultured with Dox (0.1 µg/ml) for 24 h.