



Figure S1 Hepatic deletion of SIRT1 significantly affects hepatic cholestasis and FXR/RXR activation pathways.

Total RNA from 6 pairs of control and SIRT1 LKO mice were analyzed using Agilent Whole Genome Mouse 4x44 multiplex format oligo arrays, following the Agilent 1-color microarray-based gene expression analysis protocol. The significantly altered gene probes ($p < 0.05$) in SIRT1 LKO mice were then analyzed by Ingenuity Pathway Analysis (IPA software) to identify significantly affected canonical pathways. The -log values of the P values in IPA analysis were displayed against each pathway.