Supporting Information

Hepatic loss of *Lis1* induces fatty liver and accelerates liver tumorigenesis in mice Xiaoling Li, Liansheng Liu, Ailing Wu, Jinqiu Lu, Qingzhe Wu, Mujun Zhao, Hai Song

Supporting Information includes:

Figure S1. Immunohistochemistry staining on control and Lis1 KO liver consecutive sections at 10 months with anti-LIS1 and CK19 antibodies.

Figure S2. Figure S2. Gene expression profile analysis of *Lis1* KO livers and human NAFLD samples.

Figure S3. Localization of COPI and COPII vesicles is altered in *LIS1* knockdown Hela cells.

Table S1. Differentially expressed genes in Lis1 KO livers comparing with controls.

Table S2. The sequences of qPCR primers for mouse genes.

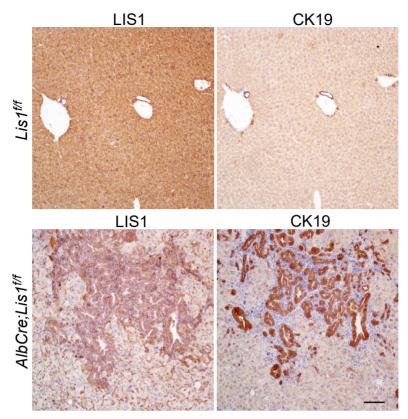


Figure S1. Immunohistochemistry staining on control and Lis1 KO liver consecutive sections at 10 months with anti-LIS1 and CK19 antibodies.

LIS1 is expressed in cholangiocytes and macrophages, but absent in hepatocytes in Lis1 KO livers, indicating a specific deletion of Lis1 in hepatocytes using AlbCre driver. Hyperplasia of bile duct was often seen in the older Lis1 KO mice. The scale bar represents 100 μ m.

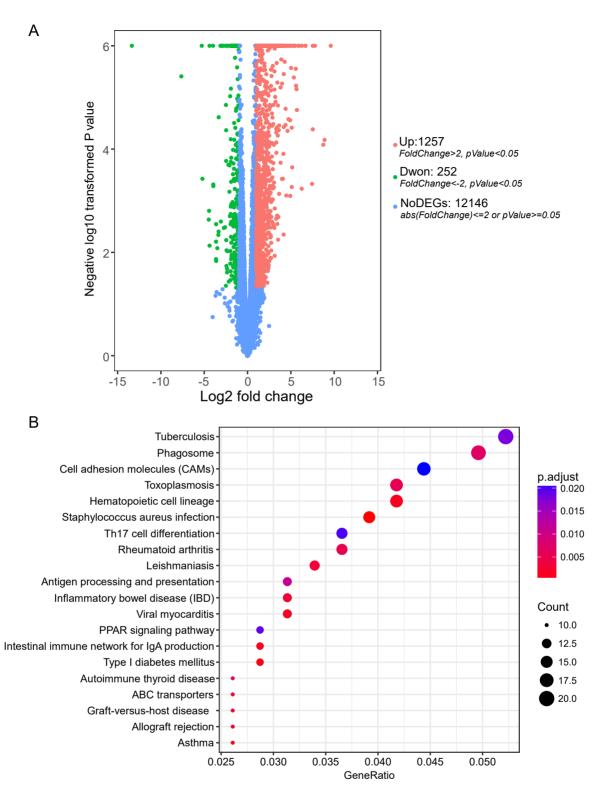


Figure S2. Gene expression profile analysis of *Lis1* KO livers and human NAFLD samples (A) Volcano analysis of gene expression in *Lis1* KO livers using RNA-Seq. (B) Top pathways that are enriched in differentially expressed genes in human NAFLD samples from public data set GSE48452.

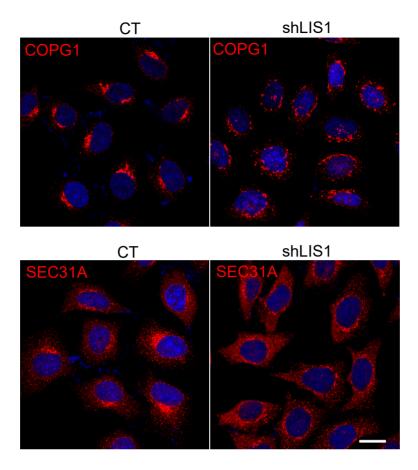


Figure S3. Localization of COPI and COPII vesicles is altered in *LIS1* knockdown Hela cells. Immunofluorescent staining of Hela cells transduced with *shLIS1* lentivirus with anti-COPG1 and SEC31A antibodies. Localization of COPG1 and SEC31A was dispersed in *LIS1* knockdown Hela cells.