

Supplementary Materials: Ochratoxin A Induces Steatosis via PPAR γ -CD36 Axis

Qian-Wen Zheng, Xu-Fen Ding, Hui-Jun Cao, Qian-Zhi Ni, Bing Zhu, Ning Ma, Feng-Kun Zhang, Yi-Kang Wang, Sheng Xu, Tian-Wei Chen, Ji Xia, Xiao-Song Qiu, Dian-Zhen Yu, Dong Xie and Jing-Jing Li

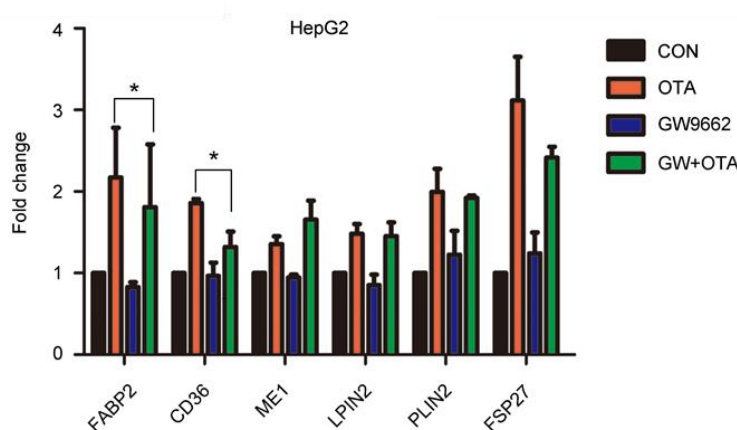


Figure S1. OTA affects PPAR γ signaling. Fold change of gene mRNA expression by qPCR in HepG2 cells under indicated treatment ($n = 6$ biological replicates). Data shown as the mean \pm S.E.M. * $P < 0.05$.

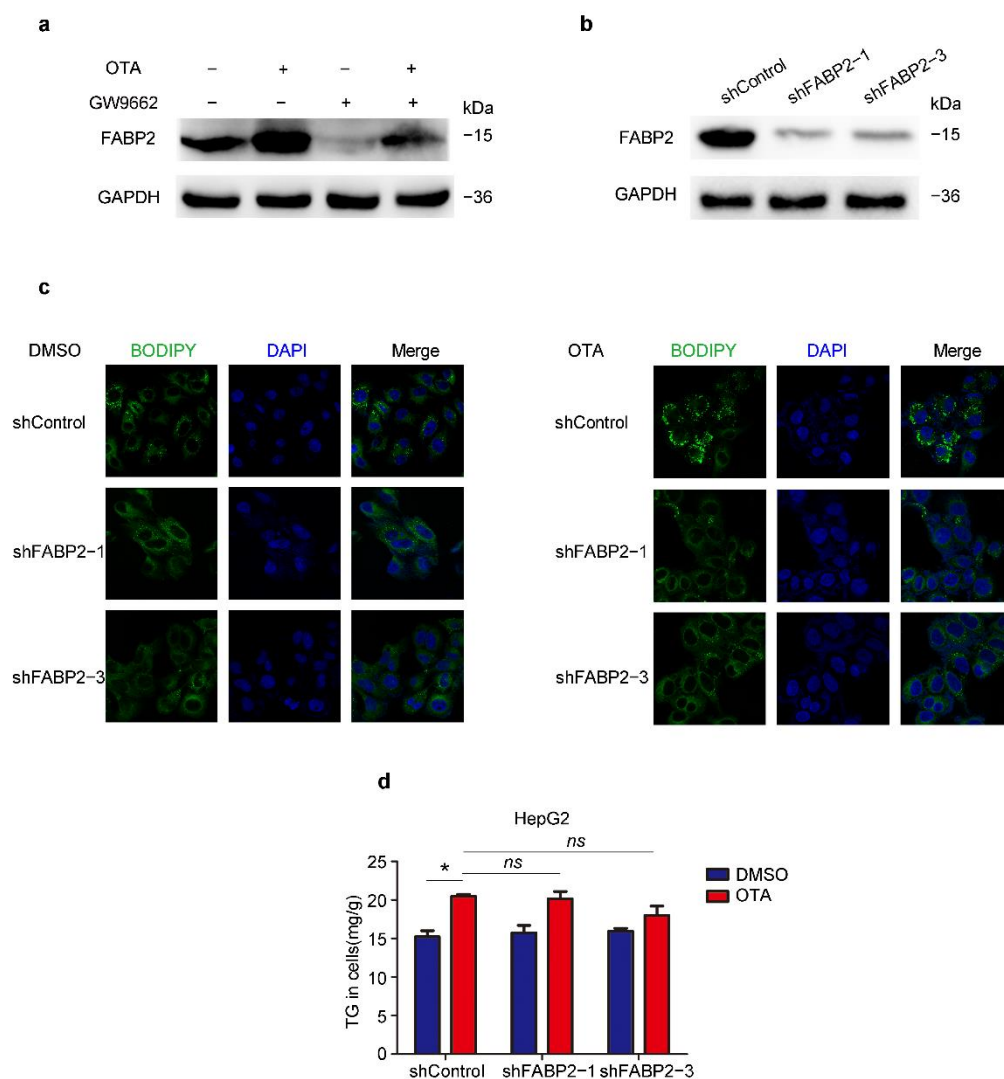


Figure S2. FABP2 is involved in the effect of OTA on lipid metabolism in liver cells. **(A)** Western blot analysis of FABP2 expression in HepG2 cells under indicated treatment. **(B)** Knockdown efficiency of FABP2 in HepG2 cells. **(C)** BODIPY staining of lipid droplets in control and FABP2-knockdown HepG2 cells treated with DMSO and OTA. **(D)** TG contents in control and FABP2-knockdown HepG2 cells treated with DMSO and OTA ($n = 6$ biological replicates). Data shown as the mean \pm S.E.M. * $P < 0.05$, *ns* means no significant difference.

Table S1. Primers of shRNA.

Primers	Sequence 5'→3'
CD36 shRNA2	F: CCGGGCCATAATCGACACATATAAACTCGAGTTTATATGTGTCGATTATGGCTTTTGG R: AATTCAAAAAGCCATAATCGACACATATAAACTCGAGTTTATATGTGTCGATTATGGC
CD36 shRNA3	F: CCGGACGGCTGCAGGTCAACCTATTCTCGAGAATAGGTTGACCTGCAGCCGTTTTTGG R: AATTCAAAAACGGCTGCAGGTCAACCTATTCTCGAGAATAGGTTGACCTGCAGCCGTT
FABP2 shRNA1	F: CCGGTGGAGCCTTGAGGGAAATAAACTCGAGTTTATTTCCTCAAGGCTCCATTTTGG R: AATTCAAAAATGGAGCCTTGAGGGAAATAAACTCGAGTTTATTTCCTCAAGGCTCCA
FABP2 shRNA3	F: CCGGCGAGAAATTATAGGTGATGAACCTCGAGTTCATCACCTATAATTTCTCGTTTTTGG R: AATTCAAAAACGAGAAATTATAGGTGATGAACCTCGAGTTCATCACCTATAATTTCTCG

Table S2. Primers for Real-Time PCR detection.

Gene	GenBank Accession Number	Sequence 5'→3'
Mouse Fabp2	14079	F: GCTGATTGCTGTCCGAGAGGTT R: AGCCTGGCATTAGCATGATGGA
Mouse Fads2	56473	F: GATGGCTGCAACATGACTATGG R: GCTGAGGCACCCTTAAGTGG
Mouse PPARg	19016	F: GGAAGACCACTCGCATTCTT R: GTAATCAGCAACCATGGGTCA
Mouse Me1	17436	F: GCCGGCTCTATCCTCCTTTG R: TTTGTATGCATCTTGACAACTCTT
Mouse Lpin2	64898	F: CAGAGTTCAGACGTTTCTCACAC R: GCTCCTTGATGCTCTTCTCCT
Mouse Plin2	11520	F: CTGTGTCTCCTCCGCTTATGTC R: GCAGAGGTCACGGTCTTCAC
Mouse Cd36	12491	F: ATGGGCTGTGATCGAACTG R: GTCTTCCCAATAAGCATGTCTCC
Mouse Fsp27	14311	F: ATGGACTACGCCATGAAGTCT R: CGGTGCTAACACGACAGGG
Mouse Siah2	20439	F: CCAATGCCGCCAGAAGTTAAG R: CAGGGAACAGAACTGCCGA
Mouse Gapdh	14433	F: AGGTCGGTGTGAACGGATTG R: TGTAAGACCATGTAGTTGAGGTCA
Human FABP2	2169	F: ATGGCGTTTGACAGCACTTG R: TCAGTTCCGCTGCTAGATTGTA
Human FADS2	9415	F: GACCACGGCAAGAACTCAAAG R: GAGGGTAGGAATCCAGCCATT
Human LPIN2	9663	F: TCTACAAGGGCATTAAACCAGGC R: AACGTGAAAAGGTGAACACTGA
Human PLIN2	123	F: TTGCAGTTGCCAATACCTATGC R: CCAGTCACAGTAGTCGTCACA
Human PPARG	5468	F: GGGATCAGCTCCGTGGATCT R: TGCACTTTGGTACTCTTGAAGTT
Human FSP27	63924	F: AAGTCCCTTAGCCTTCTCTACC R: CCTTCCTCACGCTTCGATCC
Human GAPDH	2597	F: CTGGGCTACACTGAGCACC R: AAGTGGTCGTGAGGGCAATG
Mouse/Human MKRN1	Human: 23608 Mouse: 54484	F: GAGCAGGTTTCAGAGGACTGG R: CACTCTCCCACTGCAGCATA
Human SIAH2	6478	F: CGCCAGAAGTTGAGCTGCT R: TGGTGGCATACTTACAGGGAA