



Supplementary Materials

Table S1. The primer pairs used in RT-qPCR.		
Genes	Primers	
FASN	5'-CCCCTGATGAAGAAGGATCA-3'	
	5'-ACTCCACAGGTGGGAACAAG-3'	
ACC1	5'-AGTGAGGATGGCAGCTCTGGA-3'	
	5'-TGAGATGTGGGCAGCATGAAC-3'	
SCD1	5'-GCAGGACGATATCTCTAGCT-3'	
	5'-GTCTCCAACTTATCTCCTCCATTC-3'	
SREBP-1c	5'-GGAGGGGTAGGGCCAACGGCCT-3'	
	5'-CATGTCTTCGAAAGTGCAATCC-3'	
LXRα	5'-GCCGAGTTTGCCTTGCTCA-3'	
	5'-TCCGGAGGCTCACCAGTTTC-3'	
ChREBP	5'-CAGCTGCGGGATGAGATTGA-3'	
	5'-AAACGCTGGTGTGTGATGGGTA-3'	
GLUT1	5'-CGGGCCAAGAGTGTGCTAAA-3'	
	5'-TGACGATACCGGAGCCAATG-3'	
GAPDH	5'-ATGAGAAGTATGACAACAGCCT-3'	
	5'-AGTCCTTCCACGATACCAAAGT-3'	



Figure S1. Effects of tanshinone IIA on Huh 7 cell viability. Huh 7 cells were treated with vehicle (0.1% DMSO) or tanshinone IIA (1,2,5,10 and 20 μ M) for 24 h. Cell viability was measured using an MTT assay. The data represent the mean ± SD of three independent experiments. ##*p* < 0.01 indicates significant differences compared to the vehicle-treated cells.



Figure S2. Effects of tanshinone IIA on FASN, ACC1 and SCD1 mRNA expression in Huh 7 cells. Huh 7 cells were treated with vehicle (0.1% DMSO) or tanshinone IIA (5 and 10 μ M) for 24 h. The mRNA expression of (**a**) FASN, (**b**) ACC1, and (**c**) SCD1 was measured by RT-qPCR analysis. The data represent the mean ± SD of three independent experiments. #*p* < 0.05 and ##*p* < 0.01 indicate significant differences compared to the vehicle-treated cells.



Figure S3. Effects of tanshinone IIA on ChREBP mRNA expression in HepG2 cells. HepG2 cells were treated with vehicle (0.1% DMSO) or tanshinone IIA (5 and 10 μ M) for 24 h. The mRNA expression of ChREBP was measured by RT-qPCR analysis. The data represent the mean ± SD of three independent experiments.



Figure S4. Effects of tanshinone IIA on LXR α -mediated transcriptional activity in HepG2 cells. HepG2 cells were pretreated with vehicle (0.1% DMSO) or tanshinone IIA (10 μ M) for 1 h followed by treatment with agonist T0901317 (1 μ M) for 24 h. The mRNA levels of (**a**) ACC1, (**b**) SCD1 and (**c**) GLUT1 were measured by RT-qPCR analysis. *p < 0.05 and **p < 0.01 indicate a significant difference compared to vehicle-treated group. #p < 0.05 and ##p < 0.01 indicate significant difference compared cells.