Modulation of glucose metabolism by a natural compound from Chloranthus

japonicus via activation of AMP-activated protein kinase

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Supporting Information:

Figure S1. The survival analysis of shizukaol F treated C2C12 cells and primary hepatocytes. (A) Viability of C2C12 cells treated with shizukaol F at indicated concentrations for different time-points was analyzed by MTT assay. Result was normalized to DMSO group which is set as 100%. Error bars represent s.d. from three independent experiments. (B) Viability of primary liver cells treated with shizukaol F at indicated concentrations for different time-points was analyzed by MTT assay. Result was normalized to that of DMSO treated cells which is set as 100%. Error bars represent s.d. from three independent experiments. *, P<0.05; **, P<0.01 compared to corresponding controls (one way ANOVA).



Figure S2. C2C12 differentiation status detected by western. C2C12 cells differentiation was stimulated by 2% horse serum (6 days) and then cells were photographed by fluo-microscopy (A). Two differentiated protein marker, myoD and myogenein, were detected by immunoblotting (B).



Figure S3. Shizukaol F increases AMPKa phosphorylation in C2C12 cells and primary hepatocyte. C2C12 cells were treated with shizukaol F at indicated concentrations, the phosphorylation levels of AMPKa (A), GLUT-4 translocation (B) and phosphorylated ACC were quantified (C). *, P<0.05; **, P<0.01(one way ANOVA test), n=3 independent biological replicate experiments. (B) The phosphorylation level of AMPKa was quantified after treatment of shizukaol F in primary hepatocytes. **, P<0.01(one way ANOVA test), n=3 independent biological replicate experiments.



Figure S4. AMPKa phosphorylation activity affects shizukaol F's function. C2C12 myotubes were pretreated with 20 μ M AMPK inhibitor compound C, and followed by the treatment of 1 μ M shizukaol F, the phosphorylation levels of AMPKa (A) and GLUT-4 translocation (B) were quantified. Primary hepatocytes were pretreated with 10 μ M compound C, and followed by the incubation of 1 μ M shizukaol F. Then the phosphorylation levels of AMPKa was quantified (C). PEPCK and G6Pase level were detected by q-PCR (D, E). **, P<0.01(two-tailed Student t-test). n \geq 3 independent biological replicate experiments.



Figure S5. Shizukaol F regulates glucose metabolism AMPKa via phosphorylation C2C12 myotubes infected activity. by lenti-virus of shRNA-AMPKa1 were treated with 1 µM shizukaol F. AMPKa1 relative ratio (A), AMPKa phosphorylation and GLUT-4 translocation were quantified (B). Primary hepatocytes were transfected with shRNA-AMPKa1 lenti-virus or a negative control. Cells were incubation with 1 µM of shizukaol F for 24 h. AMPKa1 relative ratio and AMPKa phosphorylation were calculated (C). PEPCK and G6Pase level were detected by q-PCR (D, E). **, P<0.01(two-tailed Student t-test). $n \ge 3$ independent biological replicate experiments.



Figure S6. Analysis of respiration in intact C2C12 cells or in mitochondria isolated from C57BL/6J mice. (A) Metformin was set as control in cellular AMP/ATP ratio detection. (B) Rosiglitazone was set as control in cellular respiration analysis. By analyzing the ADP-stimulated respiration in the presence of complex I (glutamate + malate) or complex II (succinate) substrates in isolated mitochondria from HepG2 cells, rosiglitazone (C) was used as specific inhibitor for complex I. **, P<0.01(one way ANOVA), n=3 independent biological replicate experiments.

