

Fig. S1. Confirmation of the expression of differentially expressed miRs selected from next-generation sequencing technology by qRT-PCR. Data were presented as mean \pm SEM (n = 8) and were statistically analyzed using one-way ANOVA followed by Duncan's multiple-range test. Con, control; HFD, high-fat diet; HFD+Myr, high-fat diet with 100 mg/kg myricetin. **P* < 0.05 vs Con; #*P* < 0.05 vs HFD.

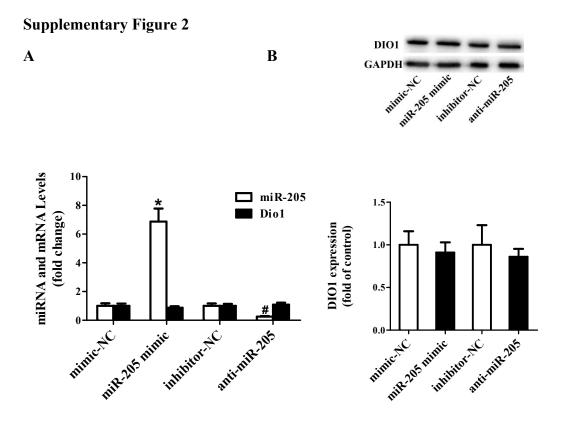


Fig. S2 miR-205 failed to regulate DIO1 expression in primary mouse hepatocytes. (A) miR-205 and *Dio1* mRNA levels, and (B) DIO1 protein levels were detected in primary hepatocytes transfected with miR-205 mimic or mimic negative control (mimic-NC) and the miR-205 inhibitor (anti-miR-205) or inhibitor negative control (inhibitor-NC). mRNA levels were detected in primary hepatocytes 24 h after transfection, while protein levels were detected by western blot 48 h after transfection. Mean \pm SEM shown were representative of at least three independent in *vitro* experiments, Student's *t*-test was used when comparing two groups. * *P* < 0.05 vs mimic-NC, #*P* < 0.05 vs inhibitor-NC.

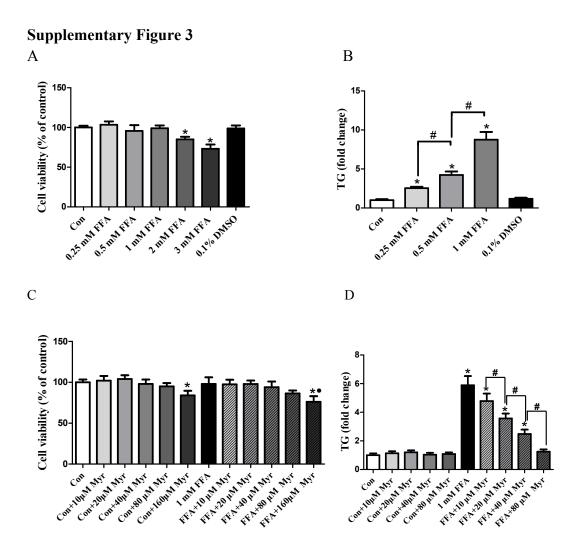


Fig.S3 Effects of FFA and myricetin on cell viability and lipid accumulation in primary mouse hepatocytes. Effects of FFA on cellular viability (as % of control) of hepatocytes as measured by MTT assay (A) and lipid accumulation (B). Effects of myricetin on cellular viability (as % of control) of 1 mM FFA treated hepatocytes as measured by MTT assay (C) and lipid deposition (D). All data were from three to five independent experiments performed in duplicate. Results were represented as mean \pm SEM and were statistically analyzed using Student's *t* test or one-way ANOVA followed by Duncan's multiple-range test.* *P* < 0.05 vs. Con group; #*P* < 0.05 between different concentrations of FFA-treated groups in (B) and between different concentrations of myricetin in 1 mM FFA-treated groups in (D); •*P* < 0.05 vs. 1mM FFA-treated group.