

Description: The mRNA levels of protein–coding genes *ACCA, DGAT2, FAS, PNPLA2,PPARG*, involved in glucose and lipid metabolism, were assessed. HepG2 cells were transfected with 50 nM of mirVana mimic *miR*–372–39 (5'–AAAGUGCUGCGACAUUUGAGCGU–3') along with a randomized sequence as control (Negative Control 1). Twenty–four hours post–transfection, cells were treated with oleic acid (0.5 mM) for 24 hours. Results are mean relative gene expression ± SEM (n = 3). *Abbreviations: NC* (negative control), *OA* (Oleic Acid), *ACCA* (Acetyl–CoA Carboxylase Alpha), *DGAT2* (Diacylglycerol O–Acyltransferase 2), *FAS* (Fatty Acid Synthase), *PNPLA2* (Patatin–like phospholipase domain–containing protein 2), *PPARG* (Peroxisome Proliferator Activated Receptor Gamma). *p*–value: \* *p* < 0.05.