

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 mmiR–144–3p (5'–UACAGUAUAGAUGAUGUACU–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 \pm 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