

Figure. MicroRNA array analysis of human lung cell line. The heatmap was generated from microRNA array with miRNA from normal lung cells without any treatment. This figure presented that array analysis has a intrinsic limitation to predict or explain some phenomena.

Supplementation

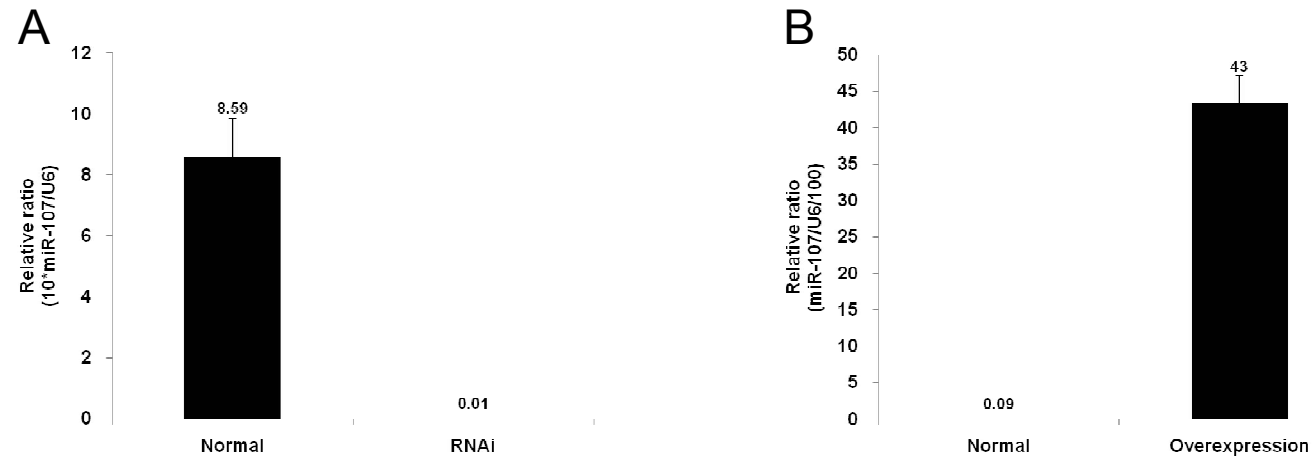


Figure. Quantitative RT-PCR analysis (n=3) after miR-107 was knocked down (A) and overexpressed (B) in murine hepatic cells. The data are expressed as an arbitrary unit and as the mean \pm SEM.