

Description of Additional Supplementary Files

Supplementary Data 1. 109 significantly dysregulated circRNAs

Supplementary Data 2. Mass spectrometry identification of proteins pulled down by circNDUFB2 sense probe

Supplementary Data 3. RNA-seq analysis of the gene expression profile affected by circNDUFB2 overexpression

Supplementary Data 4. Sequences of primer, siRNA and RNA pull down probe used in this study