Supplemental Figure 1. Hypoxia induces dynamic changes in the expression profile of HIF2A (EPAS1) in HUVECs cells. The mRNA levels were monitored in qRT-PCR experiments. The results from 3 independent experiments (n=32) are plotted normalized to 18S rRNA levels and expressed as a fold-change over the normoxic control. Error bars represent standard deviation. Significant (p<0.05) changes are marked with an "*".

Direct link to Supplemental Table 1. Genes related that were induced in HUVECs following miR-429 reduction. The significant induction was described by log₂ fold change above 2 and q (p value corrected for false discovery rate) value below 0.01.