

Supplementary Materials:

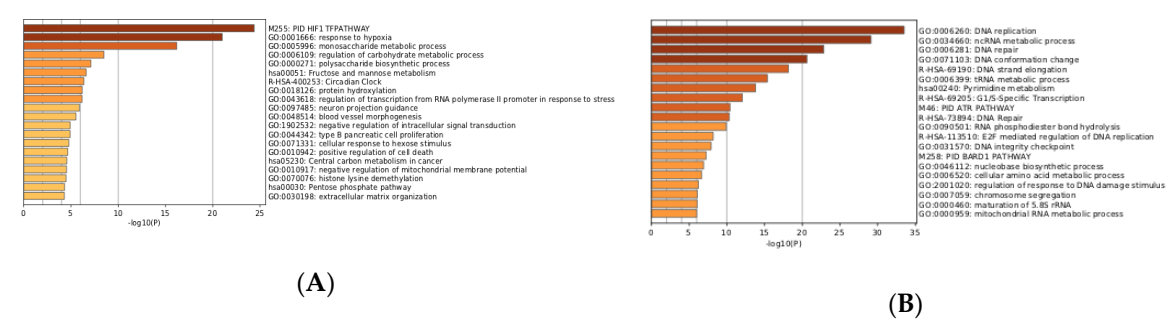


Figure S1: Gene set enrichment analysis of genes up- and downregulated in hypoxic transcriptomes. (A) Gene set enrichment analysis of genes with HN-score \Rightarrow 32; 374 genes (around 1% of all genes). (B) Gene set enrichment analysis of genes with HN-score \Leftarrow -34; 324 genes (around 1% of all genes)