

Supplementary Figure 4. p-values of enrichment for hexamers complementary to miR-124 in the UTRs of the miR-124 downregulated genes. All perfectly complementary hexamers are shown, as well as complementary hexamers (marked by an asterisk) that would form a G:U basepair and have a p-value < 0.05. Position of the hexamer with respect to the 5' end of the miRNA is given. p-values were calculated as in Supplementary Table 3, but no Bonferroni correction has been made.

