# Transcriptional regulatory network: conserved-overrepresented oligos for promoter regions only

- for microRNA genes and their association to known transcription factors binding sites.

- Column I is the overrepresented DNA oligo S labeling the set. Columns 1 ... n are the Ensembl gene id

- of the microRNA in whose promoter region the oligo S is overrepresented (p < 0.1%). Only oligos surviving

- the conserved-overrepresentation test (BY < 0.1) are retained. For each oligo, we report also its

- association with known transcription factor binding sites, where possible. We divided the dataset according

- motif length, from 6 to 9 nt in this study. In this dataset are included only microRNA genes.

# At the end of each section, a validation with known TFBSs is reported, if possible.

A plain text ASCII version of this file is available at: http://personalpages.to.infn.it/~cora/circuits/index.html
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