Supplementary Table Legends

Supplementary Table S1a-c

All computationally identified DREs in the -5kb to + 2 kb region regardless of MS score. A) Human; B) Mouse and C) Rat Tables include gene name, gene abbreviation, LocusLink, RefSeq, DRE sequence, DRE location, MS Score and strand information.

Header definition:

Refseq: mRNA refseq ID from NCBI Locuslink: LocusLink ID from NCBI gene_abbr: Gene Name Abbreviation gene name: Official Gene Name

match ind: location of putative DRE relative to TSS, "-" indicates upstream of TSS and

"=" indicates downstream of TSS mat score: MS score of putative DRE

dre_seq: actual sequence of putative DRE with length of 19 bp match_str: positive or negative strand of identified DRE sequence

Supplementary Table S2a-c

All computationally identified DREs in the -1500bp to +1500 bp with MS score above the threshold. Tables include gene name, gene abbreviation, LocusLink, RefSeq, DRE sequence, DRE location, MS Score and strand information.

Header definition:

Refseq: mRNA refseq ID from NCBI Locuslink: LocusLink ID from NCBI gene_abbr: Gene Name Abbreviation gene name: Official Gene Name

match_ind: location of putative DRE relative to TSS, "-" indicates upstream of TSS and

"=" indicates downstream of TSS mat score: MS score of putative DRE

dre_seq: actual sequence of putative DRE with length of 19 bp match str: positive or negative strand of identified DRE sequence

Supplementary Table S4a-d

Orthologous genes with DREs:

- A) human-mouse orthologous genes both have DREs in the regions between -1500 and TSS;
- B) human-rat orthologous genes both have DREs in the regions between -1500 and TSS;
- C) mouse-rat orthologous genes both have DREs in the regions between -1500 and TSS.
- D) human-mouse-rat orthologous genes all have DREs in the regions between -1500 and TSS.

All detailed DRE information is available in table S1a-c.

Supplementary Table S5RTPCR Primers of genes which have positionally conserved DREs across human, mouse and rat.