S1 Fig. Sequence alignments of enhancers between different species and search for putative transcription factor binding sites.
The first panels show sequence alignments of zebrafish and mouse entire element A and core regions from elements B, C, E and F. The non-conserved element D sequence is also shown. The last panel shows a sequence alignment of zebrafish and koi carp element A. Conserved nucleotides are marked by a star, and characterized or putative binding sites for transcription factors are indicated: Krox20 (K20) in red, MafB in blue, vHnf1 in green, Hox/Pbx (HP) in purple, Meis in pink, and Sp in orange.