

Prediction of *cis*-regulatory elements controlling genes differentially expressed by retinal and choroidal vascular endothelial cells

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In Results, under the heading, “Comparison of *Cis*-Regulatory Motifs in Genes Differentially Expressed by Retinal and Choroidal Endothelial Cells”, we inadvertently omitted one of the two motif consensus core sequences corresponding to heat shock transcription factor 1 (HSF1), a transcription factor with *cis*-regulatory motifs that were more abundant in retinal endothelial promoter sequences. The relevant section of text should read: “For five transcription factors, motifs were significantly more abundant in retinal endothelial promoter sequences. Transcription factors and corresponding motif consensus core sequences (defined by IUPAC nucleotide code) [19] were: (1) glucocorticoid receptor (GCCR) ($p=0.015$), GTTCT; (2) high mobility group at hook-1 (HMGIY) ($p=0.022$), GGAAA ; (3) heat shock transcription factor 1 (HSF1) ($p=0.025$), AGAAY and TCTAG; (4) p53 ($p=0.025$), CATGY; and (5) vitamin D receptor (VDR) ($p=0.024$), GGGTS and TGAMC.”

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