

Errata

Identification of regulatory elements in the human adipose most abundant gene transcript-1 (apM-1) promoter: role of SP1/SP3 and TNF- α as regulatory pathways

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Table 1 was missing in this article.

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Table 1. Oligonucleotide sequences used for electrophoretic mobility shift assays

Oligonucleotide	Nucleotide Sequence
SP1 consensus	5'- AGATTCGATCGGGCGGGGCGAGC-3'
wild type prom -28/-80	5'-GTGTTGAGGCTGGGCCATCTCCTCCTCACTTCATTCTGACTGCAGTCTGTGGT-3'
mutated prom-mut -28/-80	5'- GTGTTGATTCTTTTCCATCTCCTCCTCACTTCCATTCTGACTGCAGTCTGTGGTTGCA-GTCTGTGT-3'
PPAR γ -593/-622	5'-AGGGCGTCCCTGACTTATTAGGGCAAATCTATTTATTTTATAGACCTCC-3'
C/EBP -157/-204	5'-GGGTGCAGGCTGTGGGCAACTGCCAGGGGCATGTGCCTGCCACCCGGCC-3'
C/EBP -222/-262	5'-AGTTGACTGAAGCTCCACTGGGCTTCCGAAACCCAAGCTGGG-3'
C/EBP -434/-382	5'-AGTTTCTCCGTCAGATGCAGGAAAAAATCTTGTTCCTGC-3'
C/EBP -753/-782	5'-AGTCACATTTTATGTAAGTTTTCTTTATATT-3'
C/EBP -883/-912	5'-AGATATGCTTAAGGAACTGGAAAAACAGGCTC-3'
AP1 -28/-80	5'AGTGTGAGGCTGGGCCATCTCCTCCTCACTTCCATTCTGACTGCAGTCTGTGGT-3'
AP1 -109/-156	5'-AGTCTGGCCCTCACTGAGTTGGCCAATGGGAAATGACAATTGTGAGGTGG-3'
AP1 -420/-466	5'-AGGGCGTCCCTGACTTATTAGGGCAAATCTATGGGAGGGGGAGACCTCC-3'
AP1 -483/-512	5'-AGCTCAGACTTAATCATGCACAGTGAGCAAGA-3'
USF-1 -167/-204	5'-AGGGTGCAGGCTGTGGGCAACTGCCAGGGGCATGTGCCTGCCACCCGGCC-3'
YY-1 -683/712	5'-AGCAGGGGCCTTTTAACTTGCCAAGCCCCAC-3'