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-1921 TAGCTACTCA GGAAGCTGAG GCAGGAGAAT CACTTGAACC CGGGAGGTAG ACGTTGCCGT GAGCCGAGAT
        SRE3
-1851 CGCGCCACTG CACTCCAGCG TGGGTGACGG AGCGAGGCTC GGTCTCAAAA AAAAAATTCC ACTGTGAACC
-1781 ACTAAACAAA TTCATAGAAG AAAGGAACTA CCTTCCAAAC GTTTTCACGA TGATAAGAGT CATCTTGGCT
                                                                SRE2
-1711 AAATAAACTT CTGCCTTGCT AACTCACTTT GTAACATCCC CCCAAATTCT ACTCCACGAC TCCACTATAC
                                SRE1
-1641 CCAGGAAATC CTTTCCCTGA CCCCTCCAGG TGGAGAAACC ACTCTCTCAA CAAAAGATAC TCACTGAAGT
-1571 CCACAGCAGC TGAAATCTGC ATCTCGTAGG CTGGCTGCTC TTACTAGAGT GAGTTATAGG CTTCTCAAGA
-1501 GTTAAATTTG TTCCTAAATA TATGCCAGTG GCACGTGGTA TTGGGAATAG AGAATTTAAT TAAATATTAC
-1431 TAAATGTCAA TGAAATACAA GTGCAAAAAG AAAGTTGCTC TTGTGCAAAC TTAATGAATG ACTTTCGAGA
-1361 CAATCGACAA AGGAGAGTCA CAATCTTATT AGGCAAGGAC CGAAGGGATC CTAGACATTG TCCGGGAAAT
-1291 CGCAAAAATT CAGGCTTCTG TAATCAGATT GCTTCTCCGG TGAGCTTTAC GCTCTCCATG TATTTTTCAA
-1221 AAAAAATCCA ACATGAAAAT TCTAGAGGAG GCTTTATTGG GTGCGGTCTG TGCAAGAGAG ACGATGGGGA
-1151 ATGCCAATCG GAGAACCCAG ATTGAAAGGA AAGGTTTTTG GAACCCACAT CAAAAGACTG GCTGCTCGGA
-1081 AAGTTTTCTC GCTAGCGACA AAGGCTTGGA GGCTGGAAAA AGCGTCGTGG TCGCCACCTA CCGAGAAGCG
-1011 GCGGGTGGGG GGAGGATGGC CGTCGTGGCC CGAATCCCCA GTGGGTGGTA CTGGGTGGGG GGGCGGGCGG
-941  GAATCAAAAA ACCACTCAAT CTCGTGCTTT ACATAAACAT CTAAAAATCG TCGGCCTCGG GGTTGTAAAG
-871  TTTTGGAGTT TAACTTAGGT GAACAGTAAT GTCTCTCCAT TTTCAAAACC CACCGGCTAA TTCAAGTTGA
-801  GAAGCAAAAA ATAGTGGTAG GGGGAGGAGT TTGAGTTAAA TCGCATCGCG GTTGACAGAA CAAAACAGGA
-731  GCTACGAGTC AGGATTTCTG CTCCCACTCC CTTGCAAGTT TCCTTCCAAA TCCTGAGGGA GAGATAGAGG
-661  CGGAGGGTCT GGGGGAGGAA TTAAAACACA AGGTCTCCTC CCCTCTCGCC TGTCCGAACT TGGGGGCAAC
-591  GGTCTCTTTG GAGGCCAGCT TGGAAAACAC TAGAGAGGAG TTCTTTGTGT CCTGGAGGGT GGTGCAAGGC
-521  GGAAGGACTC CCCTTTTGCA GAGCAAGGAG GGCGACGGCC CAGCCCGGGG CTCCCCGGCT TTTGCAGCTT
-451  CCCTGCCTCG CAATTTGCAA GGAATGTGGG GTGCAAGAGC AAGTGGCGGA CTTGGACGGC GGCTGGCGCT
-381  CGGTGGGGAC GACGCCGGAG CCACCACCCG TCCAGATTTC CGTTTCTGCA CCCCGGCTCC CCTCCCCCAC
-311  CCGGGCGGCC GCGCAGGCCC CAGGTGCAGC CATGTTCTCG CTCCCACCAC CCTCGCCTCC ATTGACAGCC
-241  CTTGCTGGAG AGGGCCGGGC GCCGCGCTTG CCTCGTCCTC GGGTTCGGGG CTAGCCGGCG CGGTGCTTCC
-161  TGGGAGTTGT AGTCCACGAC CGCCGCCGCC GCGTTGACGG GGAACGCATC CGAAACTACA ACTCCCGCTG
-91   GGCGGCGCTG CGGATGCGCA GGCGCAACGC GCCTTCGAGG AACAAAAAAA AAAAAAAAAA AAAAAAAAAA
-21   AGAAAAAAGA AAAAAAAAAG GAGGAGGAGG AGGAGGGTGA GAGAGAAGCT GGGAGAGCAG AGAAAAGGGG
 50   CCACCGGTCG CCCCCCCGCT TCCCCGCACG CGCTCTCCAG CCGCGGCCGC CCGCCTGCCG CGGTCACCCC
                          ↳ +1
 120  GGCCTCTGCC TCTGTCCCCC AGTGATCGGA TCAAGGCGCT GAGCGAGGCC CTGCCTGCGG GGCGGCCATG
 190  CGGCGGTGAC AGGAG
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Supplementary Figure 1. Nucleotide sequence of the 5'-flanking region of the human TFII-I gene. The nucleotide sequence from -1922 to +204 is shown. Transcription state site was determined by using the Database of Transcriptional Start Sites (DBTSS) and is indicated by an arrow. Putative sterol-regulatory elements SRE1-SRE3 are in bold and underlined.