

ADR1

-1000 ATGCCTGGCTAATTTGTATTTAGTAGAGACGGAGTTCACCAAGGTTGCCAGGCTGGTCTTGA^{AC}
USF-1 GATA-1 GATA-2
ACCTGATCTCAGGTGATCCGCCATCTGGCTCTCAAAGTGCT^{GGGATT}ACAGGTGCGAGCCATCG
GCCTGGCTAGAGGTGTACATTTAACAGAACATTCAAAGGAGGTG^{GGGATC}ATGGACACT
TCCATGCTACAGCATTAATCTCCTAAGAATAAGGATAAC^{ACTCCCAC}ATACCATGACACTCTGTCACAC
CTAAAAAAATITACATTATTCCAGAACATCATCTAAATCTCCAGTCCTGCTTACATGTCCCCATTGTC
CCCAAAACATCTTTATAGATT^TTTTAAATTTGTTAAATGCCATATCCAATCGATATGGCAATCAA
C/EBP
TGCAAATCCATATTGC^{CATTGGT}ATG^TTCTCTTAGTCTTGCATAAGGGGGCCTCTT^{AGGATGC}
STRE
AAAATCTTATCATCTCTTCCACTGGGACTTGGCTGAAAATCAGGAGTGGCTGGAACACG
CCCATTTACTGTTGGTTGCAGGTTGGAGGGTACTACAGAACATCCCTCTGGAGAGGGC
CGTGAGCCTGGTTGCCCTAGACTGAGTGCCTGGCAGAGCTTCCATGTGTGCAGTGGAAAGA
AGCCCAGATCAGTCCAAAGGCCTAACCCCCACTCCCAGACCC^TACCTACTCTCAGAGAAATAGGC
TCCCTACCC^TGAACCC^AAGAACAGCTGTACCTCATCCCCAGGGACC^CAGCACCC^TCTGGCCTAT
AP-1 NF-1
CCCCAAAG^{AGTCACCC}CTGGTCTTAGGTAGTAGGTGGAGCTGAGGGATAAT^{GGCCCAAGGCCAAGA}
USF-1
GTTGATCCTCCAAC^TTGTT^CAGTGATCCAGCTT^CATAT^{CAGGT}ATCAGGACAACCAGGCCAATCT
SP1 +1
-43 GA^{TAGGGGCGGTGT}TTATAAAAGGCCACTCAC^TAGAGCCA^TGGCATTG^TCCCAGTCTGTTCC
CTTCTGGCCACAGAGCTTCTCCTGGC

Supplemental Figure 1. Mapping of putative transcription factor binding sites on human *CYP1A2* upstream sequence.

CYP1A2 mRNA:	
hCYP1A2 mRNA Left primer 5'-GTTCAAGCACAGCAAGAAGG-3'	
hCYP1A2 mRNA Right primer 5'-CTCAGGCTTGGTCACAAGGT-3'	
GAPDH mRNA:	
Left Primer 5'-TGGAAATCCC ^C ATCACC ^A TCT-3';	
Right Primer 5'-TTCACACCC ^C ATGACGAACAT-3'	
CHIP Primers:	
Left primer 5'-GCT CTT CCT CAT GTG TGC AGT G-3'	
Right primer 5'-GTC CTG ATC ACC TGA TAT GAA G-3'	
Southern Blot Probe:	
NF-1 5'-TAA TGG CCC AAG GCC AAG AGT TGA-3'	
USF-1 5'-TTT CAT ATC AGG TGA TCA GGA C-3'	

Supplemental Table 1. Primer sequences used for RNA/DNA manipulation.