

Hsa_FAM214B promoter

V\$NANOG.01 V\$TCD11.01
AAGGCCTACGCCTTTGAAAGGGTTAAATGGCAGGGCAGGGTGTGCTTTG 50
TTCCGGATGCGGAAACTTTCCAATTTACCGTCCCGTCCCACAACGAAAC
V\$LEF1.02

V\$PPAR_RXR.01 V\$SMARCA3.01
GACTCCGGCAGCTGTTCGGGTTAAGGAGAAGGGCGGAGCGACCTTGGGTAA 100
CTGAGGCCGTCGACAGCCCAATTCCTCTTCCCGCCTCGCTGGAACCCATT
V\$PTX1.01

V\$ZNF692.01 V\$WT1.02
GCCTACCCCCTTATGGACCAATAGGCGTGTGCCTCTTGGAGGGGGCGTGC 150
CGGATGGGGGAATACCTGGTTATCCGCACACGGAGAACCCTCCCCGCACG

V\$SALL4.01 V\$PAX3.03
CTCGCACTTGAGGTCGCGGGGAGGCTGGCTTCCCTATGGACCAATGAGCAT 200
GAGCGTGAACTCCAGCGCCCCTCCGACCGAAGGATACCTGGTTACTCGTA

GGTCGCTTTCCAGGGAGGGCGTAGCCTGAGGGGGGCGTGGCGGCCGTGGC 250
CCAGCGAAAGGTCCCTCCCGCATCGGACTCCCCCGCACCGCCGGCACCG
V\$MZF1.02 V\$GATA1.01

V\$MARE_ARE.01 V\$CSRNP1.01
CGCCGCGGCGCGGGTGGTGGGGCACTGGGCGGGGCGGGCGGCCCATCAG 300
GCGGCGCCGCGCCACCACCCCGTGACCCGCCCCGCCC GCGGGGTAGTC

V\$STAT5B.01 V\$ZNF217.01
CTGGTAGGAAATGGCCTGTGAGTCAGAGGGGCAGTGGAGTCGGCGGGCGCG 350
GACCATCCTTTACCGGACACTCAGTCTCCCCGTACCTCAGCCGCCGCGC

GAGCCGGGGCGGCCCGGGAGGGACCGTCGCAGGGAGGGGCGCGGCAGC 400
CTCGGC CCGCCGGGGGCCCTCCCTG GCAGCGTCCCTCCCCGCGCCGTTCG

V\$BRN2.03

V\$GRHL1.01

V\$PPARG.02

V\$E2F.01

GGCTGGGGGGCTGCGGGCGCCAGGGCCCGCTAGTGTTCACACGAAGCTG 450
CCGACCCCCCGACGCCCGCGGTCCCGGGCGATCACAACGTGTGCTTCGAC

AACAGCGAAGAGCCTAGCAGAGCAGGGCTCAGCCTGAGCCACCCTTTGTC 500
TTGTCGCTTCTCGGATCGTCTCGTCCCGAGTCGGACTCGGTGGGAAACAG