

Key: GGACATTTT 5' UPSTREAM

ACTTTGTG 5' EXON UNTRANSLATED

ATGCTTGGC EXON PROTEIN CODING

GTGAGTGAT INTRON

CACGTG CONSENSUS E BOX

CATGTG NONCONSENSUS E BOX (BOTH ORIENTATIONS)

(C/G)(A/T)**GGGGGC**(A/T) MOTIF A (EGR1-LIKE REVERSE
COMPLEMENT)

GGGGTGGGG MOTIF B (EGR1-LIKE)

CCTCTGCCTCC MOTIF C (WT1-LIKE)

Adam8 :

GGACATTTTCTCAGTTGGGGTTCCTCTTCTAAAATGACTCCGGCATGTATGAAATTGACA
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Aim1:

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Alas1:

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Aldh1a1:

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Bin1:

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gdf3:

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Mdm1 :

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Rcn2 :

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Tyropb Truncated (2873 bp) :

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