Α

Mouse Anxa2 Gene size: 1374 nt QGRS found: 6

GCCCAGCCGG	GCTGCTCCGC	TTCAAGGGAG	GCTCTCAGCG	ATACGTGCCC	GGCCCAGCTT	TTTTTTTCTT	CAAAAATGTCT	ACTGTCCACG	AAATCCTGTG	100
CAAGCTCAGC	CTGGAGGGTG	ATCATTCTAC	ACCCCCAAGT	GCCTACGGGT	CAGTCAAACC	CTACACCAAC	TTCGATGCTG	AGAGGGATGC	TCTGAACATT	200
GAGACAGCAG	TCAAGACCAA	AGGAGTGGAT	GAGGTCACCA	TTGTCAACAT	CCTGACAAAC	CGCAGCAATG	TGCAGAGGCA	GGACATTGCC	TTCGCCTATC	300
AGAGAAGGAC	CAAAAAGGAG	CTCCCGTCAG	CGCTGAAGTC	AGCCTTATCT	GGCCACCTGG	AGAC <mark>GG</mark> TGAT	TTT <mark>GG</mark> GCCTA	TTGAAGACAC	CTGCCCAGTA	400
TGATGCTTCG	GAACTAAAAG	CTTCCATGAA	G <mark>gg</mark> cct gggg	ACTGACGA <mark>GG</mark>	ACTCCCTCAT	TGAGATCATC	TGCTCACGAA	CCAACCAGGA	GCTGCAAGAG	500
ATCAACAGAG	TGTACAAGGA	AATGTACAAG	ACTGATCTGG	AGAAGGACAT	CATCTCTGAC	ACATCTGGAG	ACTTCCGAAA	GCTGATGGTC	GCCCTTGCAA	600
AGGGCAGACG	AGCAGAGGAT	GGCTCAGTTA	TTGACTACGA	GCTGATTGAC	CAGGATGCCC	GGGAGCTCTA	TGATGCC <mark>GGG</mark>	GTGAAGA <mark>GG</mark> A	AA <mark>GG</mark> AACCGA	700
CGTCCCCAAG	TGGATCAGCA	TCATGACTGA	GCGCAGTGTG	TGCCACCTCC	AGAAAGTGTT	CGAAAGGTAC	AAGAGCTACA	GCCCTTATGA	CATGCTGGAG	800
AGCATCAAGA	AAGA <mark>GG</mark> TCAA	A <mark>gggg</mark> acct <mark>g</mark>	GA GAACGCCT	TCCTGAACCT	GGTCCAGTGC	ATCCAGAACA	AGCCCCTGTA	CTTCGCTGAC	CGGCTGTACG	900
ACTCCATGAA	G <mark>GGCAAGGGG</mark>	ACTCGAGACA	A GG TCCTGAT	TAGAATCATG	GTCTCTCGCA	GTGAAGTGGA	CATGCTGAAA	ATCAGATCTG	AATTCAAGAG	1000
GAAATATGGC	AAGTCCCTGT	ACTACTACAT	CCAGCAAGAC	ACCAAGGGTG	ACTACCAGAA	GGCACTGCTG	TACCTGTG1 <mark>G</mark>	G TGGGGATGA	CTGAAGC GCT	1100
CAGCACAGTG	GATCACCCAG	AAGTGGCTCT	ACCTGTGCCC	CAACCTGGCG	TTCTAGAGAC	TTCGCTCTCC	ACTAATGGAC	CCCTGAGCTC	CTCCCTGTGA	1200
GGATGATGAC	AGGGCTGCCG	ACCCTTTCCC	CATCTTAGCT	GCCCTTGCCT	GGCTTTCTCC	TCATTCTCTC	CTTTATGCCA	AAGAAGTGAA	CATTCCAGGG	1300
AGTGGGGCGT	AGCGTCTGTG	ACATGAGACA	CTTCCTCTTA	TGTACTGTGT	CGTGAATAAA	CCGTTTTTAC	TTTA 1374			

Human Anxa2 Gene size: 1471 nt QGRS found: 7

GCTCAGCATT	TGGGGGACGCT	CTCAGCTCTC	GGCGCACGGC	CCAGGGTGAA	AATGTTTGCC	ATTAAACTCA	CATGAAGTAG	GAAATATTTA	TATGGATACA	100
AAAGGCACCT	GCATGGGATA	ATGTCAAATT	TCATAGATAC	TGCTTTGTGC	TTCCTTCAAA	ATG TCTACTG	TTCACGAAAT	CCTGTGCAAG	CTCAGCTTGG	200
AGGGTGATCA	CTCTACACCC	CCAAGTGCAT	ATGGGTCTGT	CAAAGCCTAT	ACTAACTTTG	ATGCTGAGCG	GGATGCTTTG	AACATTGAAA	CAGCCATCAA	300
GACCAAAGGT	GTGGATGAGG	TCACCATTGT	CAACATTTTG	ACCAACCGCA	GCAATGCACA	GAGACAGGAT	ATTGCCTTCG	CCTACCAGAG	AAGGACCAAA	400
AAGGAACTTG	CATCAGCACT	GAAGTCAGCC	TTATCT <mark>GG</mark> CC	ACCT <u>GG</u> AGAC	GG TGATTTT G	G CCTATTGA	AGACACCTGC	TCAGTATGAC	GCTTCTGAGC	500
TAAAAGCTTC	CATGAA <mark>GGGG</mark>	CTG GG AACCG	ACGA GG ACTC	TCTCATTGAG	ATCATCTGCT	CCAGAACCAA	CCAGGAGCTG	CAGGAAATTA	ACAGAGTCTA	600
CAAGGAAATG	TACAAGACTG	ATCT <mark>GG</mark> AGAA	GG ACATTATT	TC GG ACACAT	CTGG TGACTT	CCGCAAGCTG	ATGGTTGCCC	T <mark>GG</mark> CAAAG <mark>GG</mark>	TAGAAGAGCA	700
GA <mark>GG</mark> AT <mark>GG</mark> CT	CTGTCATTGA	TTATGAACTG	ATTGACCAAG	ATGCTCGGGA	TCTCTATGAC	GCTGGAGTGA	AGAGGAAAGG	AACTGATGTT	CCCAAGTGGA	800
TCAGCATCAT	GACCGAGCGG	AGCGTGCCCC	ACCTCCAGAA	AGTATTTGAT	AGGTACAAGA	GTTACAGCCC	TTATGACATG	TTGGAAAGCA	TCA <mark>GG</mark> AAAGA	900
GG TTAAA <mark>GG</mark> A	GACCT GG AAA	ATGCTTTCCT	GAACCTGGTT	CAGTGCATTC	AGAACAAGCC	CCTGTATTTT	GCTGATCGGC	TGTATGACTC	CATGAA <mark>GG</mark> GC	1000
AA <mark>GGGG</mark> ACGC	GAGATAA <mark>GG</mark> T	CCTGATCAGA	ATCATGGTCT	CCCGCAGTGA	AGTGGACATG	TTGAAAATTA	GGTCTGAATT	CAAGAGAAAG	TACGGCAAGT	1100
CCCTGTACTA	TTATATCCAG	CAAGACACTA	AGGGCGACTA	CCAGAAAGCG	CTGCTGTACC	TGTGTGGTGG	AGATGAC TGA	AGCCCGACAC	GGCCTGAGCG	1200
TCCAGAAATG	GTGCTCACCA	TGCTTCCAGC	TAACAGGTCT	AGAAAACCAG	CTTGCGAATA	ACAGTCCCCG	TGGCCATCCC	TGTGAGGGTG	ACGTTAGCAT	1300
TACCCCCAAC	CTCATTTTAG	TTGCCTAAGC	ATTGCCTGGC	CTTCCTGTCT	AGTCTCTCCT	GTAAGCCAAA	GAAATGAACA	TTCCAAGGAG	TTGGAAGTGA	1400
AGTCTATGAT	GTGAAACACT	TTGCCTCCTG	TGTACTGTGT	CATAAACAGA	TGAATAAACT	GAATTTGTAC	A 1471			

В

Mouse Anxa2 QGRS sequences found

Position	Length	QGRS	G-score
351	25	GG CCACCT GG AGAC GG TGATTTT GG	18
432	19	GGCCTGGGGGACTGACGAGG	13
678	17	GGGGTGAAGAGGAAAGG	15
815	17	GGTCAAAGGGGGACCTGG	16
912	22	GGCAAGGGGACTCGAGACAAGG	10
1080	18	GG TGGGGATGACTGAA GG	12

human Anxa2 QGRS sequences found

Position	Length	QGRS	G-score
12	28	GGGG ACGCTCTCAGCTCTC GG CGCAC GG	6
437	25	GG CCACCT GG AGAC GG TGATTTT GG	18
517	20	GGGGCTGGGAACCGACGAGG	12
625	30	GG AGAA GG ACATTATTTC GG ACACATCT GG	15
682	27	GGCAAAGGGTAGAAGAGCAGAGGATGG	11
894	24	GGAAAGAGGTTAAAGGAGACCTGG	20
997	23	GG GCAA GGGG ACGCGAGATAA GG	10



Supplemental Figure 2



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NUG-MONTH 3UTR

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Supplemental Figure 4



Supplemental Figure 5