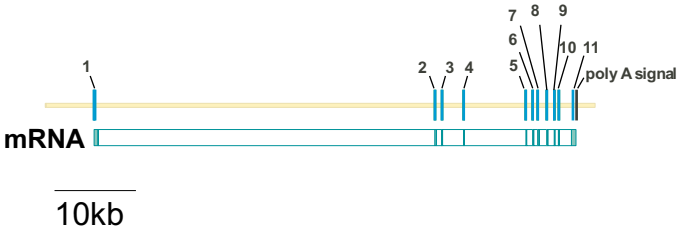


A

1 M P G L R R D R L L A L L L L G A L F S A D L Y F H L W P Q
ATGCCCGGGC TGC GCAGGGA CCGCCTGCTG GCCCTGCTGC TCCTGGGGCG GCTCTTCTCC GCCGACCTCT ACTTCCACCT CTGGCCGCAA
V Q R Q L R P G E R P A A C P C S G R A P S A S L H S A A A
91 GTGCAGCGCC AGTCCCGCC GGGCGAGCGC CCCGCCGCTT GCCCGTGCTC TGGCCGCGCC CCCTCCGCGT CTCTACACTC AGCCGCAGCC
S R D L G T A S H N F S G A L R P R V E H P S R G H P A P R S
181 TCCCGAGACC TTGGCACAGC CTCGCACAAC TTTTCAGGAG CTTTGCCCCG GGTGAGCAT CCCAGCCGCG GCCACCTGCG CCCGCGCTCT
K L Q A L F A H S L Y Q V L E D P P L L G P E D W L L A S Q
271 AAGCTGCGAG CCCTCTTCGC CCACTCGCTC TACCAAGTCC TGGAGGATCC GCCTCTCCTG GGACCCGAAG ACTGGCTCCT GGCCAGCCAG
E A L R Y Y R R K V A R W N R R H K I Y K E Q F N L T S L D
361 GAGGCGCTGC GGTATTACCG GAGGAAGGTG GCCCGCTGGA ACAGACGGCA CAAGATATAC AAAGAACAAT TCAACCTTAC CTCCTTGGAT
P P L Q F R P E A S W V Q F H L G I N S H G L Y S R S S L A
451 CCCCGCTCC AGTTTCGACC AGAGGCCAGC TGGGTCCAGT TCCACTTGGG GATCAACAGC CACGGGCTGT ACTCCAGGTC CAGCCTTGCC
I S K L L H D M R H F P T I S A D Y S Q D E K A L L G A C D
541 ATCAGCAAGC TCCTCCATGA CATGAGACAC TTCCCACCA TCAGTGCTGA TTACAGCCAG GATGAAAAAG CCTTGCTAGG GGCCTGTGAC
C S Q I V K P S G V H L K L V L R F S D F G K A M F K P M R
631 TGTCACAGA TTGTGAAACC CAGTGGGGTC CACCTGAAAT TGGTTCTGAG GTTCTCGGAC TTCGGGAAAG CCATGTTCAA ACCCATGAGA
Q Q R E E E T P E D F F Y F I D F Q R H N A E I A A F H L D
721 CAGCAGCGAG AGGAGGAGAC GCCTGAGGAC TTCTTCTATT TCATAGACTT TCAGAGACAC AATGCTGAGA TCGCAGCTTT CCACCTGGAC
R I L D F R R V P P T V G R L V N V T K E I L E V T K N E I
811 AGGATTCTGG ACTTCCGACG GGTACCGCCA ACAGTAGGGA GGCTAGTGAA TGTCAACCAAG GAAATCCTAG AGGTACACAA GAACGAGATC
L Q S V F F V S P A N N V C F F A K C P Y M C K T E Y A V C
901 CTACAGAGTG TGTTCTTTGT GTCTCCAGCT AACACGTTGT GCTTCTTCGC CAAGTGTCCG TACATGTGCA AGACCGAGTA CGCTGTCTGT
G N P H L L E G S L S A F L P S L N L A P R L S V P N P W I
991 GGCAACCCAC ACCTGCTGGA AGGCTCCCTC TCCGCCTTCC TGCCATCCCT CAACCTAGCC CACGAGTGT CTGTGCCCAA CCCCTGGATC
R S Y S L S G K E E W E L N P L Y C D T V K Q I Y P Y N S S
1081 CGTTCCTACT CGCTGTGAGG AAAAGAGGAG TGGGAGCTCA ACCCCCTTTA CTGTGACACA GTCAAGCAGA TCTACCCGTA CAACAGCAGC
N R L L G I I D M A V F D F L I G N M D R H H Y E M F T K F
1171 AACAGACTCC TCGGCATCAT TGACATGGCC GTCTTTGACT TCCTGATTGG AAATATGGAC CGACACCATT ATGAGATGTT CACCAAATTT
G D D G Y L I H L D N A R G F G R H S Q D E I S I L A P L A
1261 GGGGATGATG GATACTTAT TCATCTTGAC AATGCCAGAG GGTTCGGACG ACATTCCCG GATGAAATCT CTATCCTCGC CCCTCTAGCG
Q C C M I K R K T L L H L Q L L A Q A D Y R L S D V V M R E S
1351 CAGTGTTGCA TGATCAAAAG GAAAACACTC TTGCACCTTC AGTGCTGGC TCAGGCTGAC TACAGGCTCA GCGACGTGAT GCGGGAGTCA
L L L E D Q L S P V L T E P H L L A L D R R L Q I I L K T V E
1441 CTTCTAGAAG ACCAGCTCAG CCCTGTCCCTC ACAGAACCC ACCTCCTGGC CCTGGACCGC AGACTCCAGA TCATCCTGAA GACCGTGGAG
D C I E A H G E R R V I A E G S A Q R S A P D S G Q A N L T
1531 GACTGCATAG AGGCCACGG CGAGCGAAGA GTCATAGCCG AAGGGTCAGC ACAACGTTCC GCTCCAGACT CCGGCCAGGC TAATCTGACG
S *
1621 AGCTAAT

B



C

	Exon	Intron
1	532	44,511
2	185	731
3	51	2,800
4	79	8,149
5	93	819
6	116	528
7	181	1,026
8	110	882
9	82	488
10	60	1,595
11	618	