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Product datasheet for SC206311

CDK2AP2 (NM_005851) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	CDK2AP2
Synonyms:	DOC-1R; p14
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_005851
Insert Size:	478 bp
Insert Sequence:	>SC206311 3'UTR clone of NM_005851 The sequence shown below is from the reference sequence of NM_005851. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GCAGAGACAGAGCGGAACGCCCGCACGTAACAGGAAGCGCCTCGGCCTCAGCGTCTGGACCTATCCGGC CACTGCAGAGCACCCGCTTCTCCCTGGCCTTCATCCCGAGTTGCACTAACCATCCTGGGCTTCCTGTCC TGTGTCCCTTGGTGGGTCCCCTCCAGGAACCAAGGAGTGGCCCTCCAGGTGGCAGCACTAAGGACACCC CCCCACAACAAGAGTTAGCAGCGAGGTCCCCATGAGTCCCACCCATGACCTGCCGACAGTGTTGCCCAC CGGAACTTTTGTGGCCCCTACCGCTCAGCCCTTCCCAGCACTTCTCCCACTTTGTCCCGAGCCTCCTTC TCCCCCAGCAGGGGCACAGGCCTGGCACCTCCCTGCCTTGTGTCCTGAGCCATAGTGACTCTTTTATCT GTGTGTCTTTTGCTAAATATGCCCTTTTTATATTAATAAAAGATGATTTGGAGTTGTGCTCTCA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
Restriction Sites:	SgfI-MluI
OTI Disclaimer:	Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).

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Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_005851.5
Summary:	This gene encodes a protein that interacts with cyclin-dependent kinase 2 associated protein 1. Pseudogenes associated with this gene are located on chromosomes 7 and 9. Alternatively spliced transcript variants have been observed for this gene. [provided by RefSeq, Dec 2012]
Locus ID:	10263
MW:	17.3