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Product datasheet for SC202832

CSRP2 (NM_001321) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	CSRP2 (NM_001321) Human 3' UTR Clone
Symbol:	CSRP2
Synonyms:	CRP2; LMO5; SmLIM
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001321
Insert Size:	273 bp
Insert Sequence:	>SC202832 3'UTR clone of NM_001321 The sequence shown below is from the reference sequence of NM_001321. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GGAGCAGGGGCTCTTGTTCATGCCCAGTAAGATGTAAACCCTGAACTAAACATCACACACTGAGAATCT CTTCATAATCTAGGCACAGATAATCTTTAACACTAAACTACTGTGAAATTCTACCAGCATTAAGTACTG TATATCGCCCTGTACTTGGATAGGCTGGCTAACTCGTAGGAAGAGAGCACTGTATGGTATCCTTTTGCT TTATTCACCAGCATTTTGGGGGAACATTTCTTTTACATTTTAAATAAAACTTCAGCTTGATTTGGG 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).
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.
RefSeq:	NM_001321.3

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Summary: CSRP2 is a member of the CSRP family of genes, encoding a group of LIM domain proteins, which may be involved in regulatory processes important for development and cellular differentiation. CRP2 contains two copies of the cysteine-rich amino acid sequence motif (LIM) with putative zinc-binding activity, and may be involved in regulating ordered cell growth. Other genes in the family include CSRP1 and CSRP3. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2014]

Locus ID: 1466

MW: 10.3