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Product datasheet for SC207420

CROP (LUC7L3) (NM_006107) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	CROP (LUC7L3) (NM_006107) Human 3' UTR Clone
Symbol:	CROP
Synonyms:	CRA; CREAP-1; CROP; hLuc7A; LUC7A; OA48-18
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_006107
Insert Size:	575 bp
Insert Sequence:	>SC207420 3'UTR clone of NM_006107 The sequence shown below is from the reference sequence of NM_006107. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AAATCTGAAGGTGACACTCAGTCCAATTAAAACTGATCTGATAAGACCTCAGATCAGACAGAGGACTAC TGTTCGAAGATTTTTGGAAGAATACTGAGAACGGCATAAAGTGAAGATCGACATTTAAAAAATGAGGTG AAAGAAAGCTATAGTGGCATAGAAAAAGTATAAAGCTCAGTTAGTTTTTTTATTATTATTATTATTAAA AGTTAATTCAGGACTGATGTGACCTACCAGATTTCAGAACATGTGTTAATAGTATATATGCCACTGAAA ACTTAGGTCCTGTATCATACTTTTTTCTTTAAGACTTTTTAAGAAATATTACTTAAACATGTGGCTTGC TCAGTGTTTAATTGCAAGTTTTCAATCTTGGACTTTGAAAACAGGATTAAACGTTAGTATTCGTGTGAA TCAGACTAAGTGGGATTTCATTTTTACAACTCTGCTCTACTTAGCCTTTGGATTTAGAAGTAAAAATAA AGTATCTCTGACTTTCTGTTACAAAGTTGATTGTCTCTGTCATTGAAAAGTTTTAGTATTAATCTTTTT CTAATAAAGTTATTGACTCTGAA 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.
RefSeq:	NM_006107.4
Summary:	This gene encodes a protein with an N-terminal half that contains cysteine/histidine motifs and leucine zipper-like repeats, and the C-terminal half is rich in arginine and glutamate residues (RE domain) and arginine and serine residues (RS domain). This protein localizes with a speckled pattern in the nucleus, and could be involved in the formation of splicesome via the RE and RS domains. Two alternatively spliced transcript variants encoding the same protein have been found for this gene. [provided by RefSeq, Aug 2009]
Locus ID:	51747
MW:	22.2