

Product datasheet for SC310399

CROP (LUC7L3) (NM_016424) Human Untagged Clone

Product data:

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Product Type:	Expression Plasmids
Product Name:	CROP (LUC7L3) (NM_016424) Human Untagged Clone
Tag:	Tag Free
Symbol:	CROP
Synonyms:	CRA; CREAP-1; CROP; hLuc7A; LUC7A; OA48-18
Vector:	pCMV6 series
Fully Sequenced ORF:	>NCBI ORF sequence for NM_016424, the custom clone sequence may differ by one or more nucleotides ATGATTTCGGCCGCCAGTGTTGGATGAGTTAATGGGCCGGGACCGAAACCTAGCCCCG GACGAGAAGCGCCGCAGCTGCGGTGGGACCACGAGAGCGTTTGTAAATATTATCTCTGT GGTTTTGTCCTGCGGAATTGTTCACAAATACAGTTGGACTTGGTCGTGGAAAAA ATTCATGATGAGAAAACTACGAAAACAGTATGAGAAGAGGCTCTCGTTTCATGAAAGTTGGC TATGAGAGAGATTTTTTGCGATACTTACAGACTTACTTGCAGAAGTAGAACGTAGGACC AGGCGAGGCCATGCTCGTTTGGCATTATCTCAAAACCAGCAGTTGTCAGAAGATGAGAGCCCAGGGCCGCTGGC CCAACAGGCCAATGACTTAGGGTCTGAAGGAAAGTAGAAGAGCCCAGGGGCGCGGCGC CCAACAGGCAAAAATGAAGAGAAGAGAGAGAGCTGCTAAGGCCCACAGCGGATGATGAAA TTAGTTGAGCAATTAAAAGGAAGAGAGAGACTGCTAAGGCCCACACGGGGATGATGAAA TTAGTTGAGCAATTAAAAGAAAAACTTAGAGATTGGAAGTCGACACACGCGGACGATGAAAA TTAGTTGAGCAATTAAAAGAAAAGGAGAGACACTGCTAAGGTCCACAACGGCGATGATGAAA TTAGTTGAGCAATTAAAAGAAAAGGAGAGACACTGCTAAGGTCCACACACGGGCATATGAA AGCTTTGCTGCCCGGGTGAGTGACCATTTGATGGGAAAACAACACACAC
Restriction Sites:	Please inquire
ACCN:	NM_016424



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	ROP (LUC7L3) (NM_016424) Human Untagged Clone – SC310399
OTI Disclaimer:	Due to the inherent nature of this plasmid, standard methods to replicate additional amoun of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amount of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <u>custsupport@origene.com</u> or by calling 301.340.3188 option 3 for pricing and delivery.
	The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ throug naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <u>More info</u>
OTI Annotation:	This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water)
Reconstitution M	 1. Centrifuge at 5,000xg for 5min. 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. 3. Close the tube and incubate for 10 minutes at room temperature. 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liqu at the bottom. 5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date o shipping when stored at -20°C.
RefSeq:	<u>NM 016424.3, NP 057508.2</u>
RefSeq Size:	3477 bp
RefSeq ORF:	1299 bp
Locus ID:	51747
UniProt ID:	<u>095232</u>
Cytogenetics:	17q21.33
Domains:	DUF259
Protein Families:	Stem cell - Pluripotency

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Gene 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]
Transcript Variant: This variant (1) represents the longer transcript. Sequence Note: This
RefSeq record was created from transcript and genomic sequence data to make the sequence
consistent with the reference genome assembly. The genomic coordinates used for the
transcript record were based on transcript alignments.

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