

Product datasheet for SC317714

SEPTIN7 (NM_001011553) Human Untagged Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	Expression Plasmids
Product Name:	SEPTIN7 (NM_001011553) Human Untagged Clone
Tag:	Tag Free
Symbol:	SEPTIN7
Synonyms:	CDC3; CDC10; NBLA02942; SEPT7; SEPT7A
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC317714 representing NM_001011553. Blue=Insert sequence Red=Cloning site Green=Tag(s)
	GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG GATCCGGTACCGAGGAGATCTGCCGCGCGCGCGCACGCC ATGTCGGTCAGTGCGAGATCCGCTGCTGCTGCTGCTGAGGAGAGGAGCGTCAACAGCAGCACCATGGCTCAACAG AAGAACCTTGAAGGCTATGTGGGATTTGCCAATCTCCCAAATCAAGTATACAGAAAATCGGTGAAGAGA GGTTTTGAATTCACGCTTATGGTAGTGGGTGAATCTGGATTGGGAAAGTCGACATTAATCAACTCATTA TTCCTCACAGATTTGTATTCTCCAGAGTATCCAGGTGCTTCACTAGAATTAAAAAGACTGTACAGGTG GAACAATCCAAAGTTTTAATCAAAGAAGGTGGTGTTCAGTTGCTGCTCACAATAGTTGATACCACAGGA TTTGGAGATGCAGTGGATAATAGTAATTGCTGGCAGCCTGTTATCGACTACATTGATAGTAAATTTGAG GACTACCTAAATGCAGAATCACGAGTGAACAGACGTCAGATGCGACTACAATGGAAGATGCAGTTGTTTA TACTTCATTGCTCCTCAGGACATGGACTTAAACCATTGGATATTGAGGTTATTGAAGCGTTTGCATGAA AAAGTGAATATCATCCCACTTATTGCCAAAGCAGACCACACCCACACAGAGGAATGCCAACAGTTTAAA AAAGTGAATATCATCCCACTTATTGCCAAAGCAGACCACACTCCACACCAGAGGAATGCCAACAGTTAAA AAAGTGAATATCATCCCACTTATTGCCAAAGCAGACCACTCCACACCAGAGGAATGCCAACAGTTTAAA AAACAGATAATGAAAGAAATCCAAGGACACATAAAATTAAAATATACGAATTTCCAGAAACAGATGATGAA AAACAGATAATGGCAAAAGGGTCCAGAGGAAGCAGTTCCTTGGCTGGGTAGTAATACGAACTATCA ATTGAAGTTATTAGCACAAAGGGTCAGAGGAAGGCAGTTACCTCTGGCGTGGTAGGTA

View online »

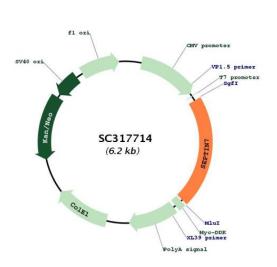
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Plasmid Map:

SgfI-Mlul



ACCN:	NM_001011553
Insert Size:	1311 bp
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
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 Method:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM 001011553.3</u>
RefSeq Size:	4377 bp
RefSeq ORF:	1311 bp
Locus ID:	989
Cytogenetics:	7p14.2

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MW:	50.6 kDa
Gene Summary:	This gene encodes a protein that is highly similar to the CDC10 protein of Saccharomyces cerevisiae. The protein also shares similarity with Diff 6 of Drosophila and with H5 of mouse. Each of these similar proteins, including the yeast CDC10, contains a GTP-binding motif. The yeast CDC10 protein is a structural component of the 10 nm filament which lies inside the cytoplasmic membrane and is essential for cytokinesis. This human protein functions in gliomagenesis and in the suppression of glioma cell growth, and it is required for the association of centromere-associated protein E with the kinetochore. Alternative splicing results in multiple transcript variants. Several related pseudogenes have been identified on chromosomes 5, 7, 9, 10, 11, 14, 17 and 19. [provided by RefSeq, Jul 2011] Transcript Variant: This variant (2) lacks an in-frame segment in the 5' coding region, compared to variant 1, resulting in an isoform (2) that is 1 aa shorter than isoform 1. 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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