

## Product datasheet for **SC203984**

### ULK2 (NM\_001142610) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	ULK2 (NM_001142610) Human 3' UTR Clone
Symbol:	ULK2
Synonyms:	ATG1B; Unc51.2
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001142610
Insert Size:	308 bp
Insert Sequence:	>SC203984 3'UTR clone of NM_001142610 The sequence shown below is from the reference sequence of NM_001142610. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b> =Stop Codon <b>Red</b> =Cloning site  GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA <b>GCGATCGCC</b> GCGCTCTGCCATAGCACCGCAACCGTGT <b>GA</b> GCAGCAGGCTCATCCCGTGGACCGGTGGTGGGAACGTGA GGAAGAGGGGAAGGAAGGAAGAGCTTTCCATTGGTGCTCCAATGTCTCCTGCTGGACCCATCTGCCT AGTGGAAGGCAGCAAAATTTCAAGAAACAGGTGAGGTTGAGCAGCTTGGTCAACCCCATGGGGCCTGG AGTTGGAGCTCAACAGCAATGGATTT <b>CAGAGACCACCCTGAAACTCCCAGTAAAAAAGACTTGGGAGAC</b> ATGTTAATAAACTCAAGCATTGATCGACCCA <b>ACGCGT</b> AAGCGGCCGCGCATCTAGATT <b>CGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA</b> CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
Restriction Sites:	Sgfl-Mlul
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:	<u><a href="#">NM_001142610.2</a></u>



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**Summary:**

This gene encodes a protein that is similar to a serine/threonine kinase in *C. elegans* which is involved in axonal elongation. The structure of this protein is similar to the *C. elegans* protein in that both proteins have an N-terminal kinase domain, a central proline/serine rich (PS) domain, and a C-terminal (C) domain. The gene is located within the Smith-Magenis syndrome region on chromosome 17. Alternatively spliced transcript variants encoding the same protein have been identified. [provided by RefSeq, Dec 2008]

**Locus ID:**

9706

**MW:**

11.1