

## Product datasheet for SC204542

## SEC13L1 (SEC13) (NM\_183352) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	SEC13L1 (SEC13) (NM_183352) Human 3' UTR Clone
Symbol:	SEC13L1
Synonyms:	D3S1231E; npp-20; SEC13L1; SEC13R
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_183352
Insert Size:	360 bp
Insert Sequence:	<pre>&gt;SC204542 3'UTR clone of NM_183352 The sequence shown below is from the reference sequence of NM_183352. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GTGACAGAGGGCCAGCAGAGACGAGCAGTGACAAGACAGGTGGGGCCTGGCTCCCCACCCGCCAGCTCCA GGACTGCCCCTTCCTGGGCCAACTAACCAGACAAGTGAGACAGGTGGGGCCTGGCTCCCCAACAGGAGTTATTTT CCCAGGAGGGCCAGCAGAGAGCAGTGACAACTGGGAAGAACTGCGAGATGCTCTGTAA TCTACTGTCCAGCTGAAAGCACTCATGTTACGAGGAAGAAACTACAAGTGATCGTACAATCTATTTTGG GTCATTTTTATGTACCTTTGGGTTCAGGCATTATTTGGGGGGGTTTTGTTTCCAAAGGAACTAAATAAA</pre>
<b>Restriction Sites:</b>	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.



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	SEC13L1 (SEC13) (NM_183352) Human 3' UTR Clone – SC204542
RefSeq:	<u>NM 183352.3</u>
Summary:	The protein encoded by this gene belongs to the SEC13 family of WD-repeat proteins. It is a constituent of the endoplasmic reticulum and the nuclear pore complex. It has similarity to the yeast SEC13 protein, which is required for vesicle biogenesis from endoplasmic reticulum during the transport of proteins. Multiple alternatively spliced transcript variants have been found. [provided by RefSeq, Oct 2008]
Locus ID:	6396
MW:	13.7

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