

## Product datasheet for SC207467

## SEZ6L2 (NM\_001114100) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	SEZ6L2 (NM_001114100) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	SEZ6L2
Synonyms:	BSRPA; PSK-1
ACCN:	NM_001114100
Insert Size:	570 bp
Insert Sequence:	<pre>&gt;SC207467 3'UTR clone of NM_001114100 The sequence shown below is from the reference sequence of NM_001114100. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG</pre>
	GGCAAGTTGGACGCCCGCAAGATTCCCACGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG         TAACAATTGGCAGAGCTCAGAATTCCAAGCGATCGCC         GATACGCGGGAGTATGAAGTTTCCATCTGAACCCCAAGACTACAGCTGCAGGACCCAGGACGCCCCTCC         CCTCCTCATTCGGGCAGAGGGGAAATACGGGACCCGGTCTCTGCCTCGCCTCGGCTGCCCCCCGGCCTGGCCCCCGGCGCGGCTG         TGTAAATAGTCTCCCTATCCCACGAGGGGGGCTTTGATGGCCCTGGAGATCCTACAGTAAATAAA
<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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	SEZ6L2 (NM_001114100) Human 3' UTR Clone – SC207467
RefSeq:	<u>NM 001114100.3</u>
Summary:	This gene encodes a seizure-related protein that is localized on the cell surface. The gene is located in a region of chromosome 16p11.2 that is thought to contain candidate genes for autism spectrum disorders (ASD), though there is no evidence directly implicating this gene in ASD. Increased expression of this gene has been found in lung cancers, and the protein is therefore considered to be a novel prognostic marker for lung cancer. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Aug 2011]
Locus ID:	26470
MW:	20.8

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