

## Product datasheet for SC322966

### TEX264 (NM\_001129884) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	TEX264 (NM_001129884) Human Untagged Clone
Tag:	Tag Free
Symbol:	TEX264
Synonyms:	ZSIG11
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC322966 representing NM_001129884. Blue=Insert sequence Red=Cloning site Green=Tag(s)

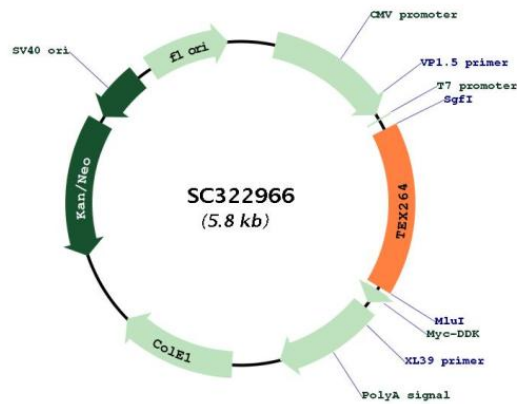
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GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGTCGGACCTGCTACTACTGGGCTGATTGGGGCCTGACTCTCTTACTGCTGCTGACGCTGCTGGCC
TTTGCCGGTACTCAGGGCTACTGGCTGGGGTGAAGTGAGTGCTGGGTACCCCCCATCCGCAACGTC
ACTGTGGCTACAAGTTCACATGGGGCTCTATGGTGAGACTGGGCGCTTTTCACTGAGAGCTGCAGC
ATCTCTCCCAAGCTCCGCTCCATCGCTGTCTACTATGACAACCCACATGGTCCCCCTGATAAGTGC
CGATGTGCCGTGGCAGCATCCTGAGTGAAGTGAGGAATCGCCCTCCCCTGAGCTCATCGACCTCTAC
CAGAAATTTGGCTCAAGGTGTTCTCCTTCCCGCACCCAGCCATGTGGTGACAGCCACCTTCCCCTAC
ACCACCATTCTGTCCATCTGGCTGGCTACCCGCGTGTCCATCCTGCCTTGGACACCTACATCAAGGAG
CGGAAGCTGTGTGCCTATCCTCGCTGGAGATCTACCAGGAAGACCAGATCCATTTTCATGTGCCACTG
GCACGGCAGGGAGACTTCTATGTGCCTGAGATGAAGGAGACAGAGTGAAATGGCGGGGGCTTGTGGAG
GCCATTGACACCCAGGTGGATGGCACAGGAGCTGACACAATGAGTGACACGAGTTCTGTAAGCTTGGAA
GTGAGCCCTGGCAGCCGGGAGACTTCAGCTGCCACACTGTACCTGGGGCGAGCAGCCGTGGCTGGGAT
GACGGTGACACCCGACGAGCAGCTACAGCGAGTCAGGTGCCAGCGGCTCCTCTTTTGGAGAGCTG
GACTTGGAGGGCGAGGGGCCCTTAGGGGAGTACGGCTGGACCTGGGACTGAGCCCTGGGACTACCC
AAGTGGCTTGGGAGCCCACTGCCCTGAGAAGGGCAAGGAGTAA
ACGCGTACGCGGCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
```

Restriction Sites: SgfI-MluI



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## Plasmid Map:



ACCN: NM\_001129884

Insert Size: 942 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:**

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 liquid at the bottom.
5. 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: [NM\\_001129884.2](#)

RefSeq Size: 1455 bp

RefSeq ORF: 942 bp

<b>Locus ID:</b>	51368
<b>UniProt ID:</b>	<a href="#">Q9Y6I9</a>
<b>Cytogenetics:</b>	3p21.2
<b>Protein Families:</b>	Secreted Protein, Transmembrane
<b>MW:</b>	34.2 kDa
<b>Gene Summary:</b>	<p>Major reticulophagy (also called ER-phagy) receptor that acts independently of other candidate reticulophagy receptors to remodel subdomains of the endoplasmic reticulum into autophagosomes upon nutrient stress, which then fuse with lysosomes for endoplasmic reticulum turnover (PubMed:31006538, PubMed:31006537). The ATG8-containing isolation membrane (IM) cradles a tubular segment of TEX264-positive ER near a three-way junction, allowing the formation of a synapse of 2 juxtaposed membranes with trans interaction between the TEX264 and ATG8 proteins (PubMed:31006537). Expansion of the IM would extend the capture of ER, possibly through a 'zipper-like' process involving continued trans TEX264-ATG8 interactions, until poorly understood mechanisms lead to the fission of relevant membranes and, ultimately, autophagosomal membrane closure (PubMed:31006537). [UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) differs in the 5' UTR, compared to variant 1. Variants 1, 2, 4, 5, and 7 encode the same isoform (1).</p>