

## Product datasheet for **RC233517**

### TEX264 (NM\_001243727) Human Tagged ORF Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** TEX264 (NM\_001243727) Human Tagged ORF Clone  
**Tag:** Myc-DDK  
**Symbol:** TEX264  
**Synonyms:** ZSIG11  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**ORF Nucleotide Sequence:** >RC233517 representing NM\_001243727  
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGCC**

ATGTCGGACCTGCTACTACTGGGCCTGATTGGGGCCTGACTCTCTTACTGCTGCTGACGCTGCTGGCCT  
TTGCCGGTACTCAGGGCTACTGGCTGGGGTGGAAAGTGAGTGCTGGGTACCCCCATCCGCAACGTCAC  
TGTGGCCTACAAGTTCACATGGGGCTCTATGGTGAGACTGGGCGGCTTTTCACTGAGAGCTGCAGCATC  
TCTCCAAGCTCCGCTCCATCGTGTCTACTATGACAACCCACATGGAGCGGAAGCTGTGTGCCTATC  
CTCGGCTGGAGATCTACCAGGAAGACCAGATCCATTTTCATGTGCCCACTGGCACGGCAGGGAGACTTCTA  
TGTGCCTGAGATGAAGGAGACAGAGTGGAAATGGCGGGGCTTGTGGAGGCCATTGACACCCAGGTGGAT  
GGCACAGGAGCTGACACAATGAGTGACACGAGTTCTGTAAGCTTGGAAAGTGAGCCCTGGCAGCCGGGAGA  
CTTCAGCTGCCCACTGTACCTGGGGCGAGCAGCCGTGGCTGGGATGACGGTGACACCCGAGCGAGCA  
CAGCTACAGCGAGTCAGGTGCCAGCGGCTCCTCTTTTGAGGAGCTGGACTTGGAGGGCGAGGGGCCCTTA  
GGGGAGTCACGGCTGGACCCTGGGACTGAGCCCTGGGGACTACCAAGTGGCTCTGGGAGCCCACTGCC  
CTGAGAAGGGCAAGGAG

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA



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**Protein Sequence:** >RC233517 representing NM\_001243727  
Red=Cloning site Green=Tags(s)

MSDLLLLGLIGLLTLLLLLTLAFAGYSGLLAGVEVSAGSPPIRNVTVAYKFMGLYGETGRLFTESCSI  
 SPKLRSIAVYYDNPHERKLCAYPRLEIYQEDQIHFMCLARQGFYVPEMKETEWKWRGLVEAIDTQVD  
 GTGADTMSDTSSVSLEVSPGSRSETSATLSPGASSRGWDDGDTRSEHSYSESGASGSSFEELDLEGEPL  
 GESRLDPGTEPLGTTKWLWEPTAPEKGKE

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Restriction Sites:** SgfI-MluI

**Cloning Scheme:**

Cloning sites used for ORF Shuttling:



**ACCN:** NM\_001243727

**ORF Size:** 717 bp

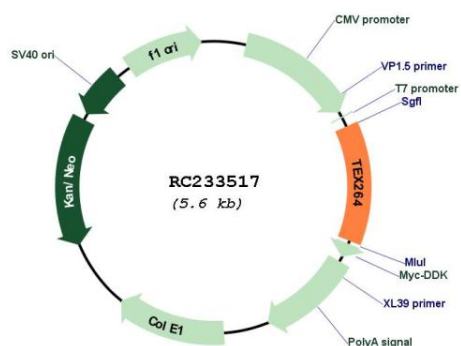
**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**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).

<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_001243727.2</a> , <a href="#">NP_001230656.1</a>
<b>RefSeq Size:</b>	1181 bp
<b>RefSeq ORF:</b>	720 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>	26.4 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>

## Product images:



Circular map for RC233517