

Product datasheet for **MC200096**

Tex264 (NM_011573) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Tex264 (NM_011573) Mouse Untagged Clone
Tag: Tag Free
Symbol: Tex264
Synonyms: TEG-264
Mammalian Cell Selection: Neomycin
Vector: PCMV6-Kan/Neo (PCMV6KN)
E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >BC002248 sequence for NM_011573
 CCCACGCGTCCGCCACGCGTCCGCCACGCGTCCGGGGAGGCGGTCTGACCTCTCGGCGGGATCCTG
 CCTCGGTGCTGGCGCGGAGTACTGGAGCCCTGCTGGACTGGATGCTATGAGTTGCTAAAAGTGAGCCTG
 ACTTTTTTCGCTGCCTTGGGGTGCTGAGAGGGAGTCCCAGAAAGAACCATGCCGGATCTCCTACTACTGGG
 CCTGATTGGGGCCTGACGCTGCTGTTGCTGCTGACGCTGCTGGCCTTTGCTGGTTATTACGACTGCTG
 ACTGGGGTGACAGTGAGCGCTGGATCACCCCAATCCGCAACATAACTGTGGCCTACAAGTTCCACGTGG
 GGTCTATGGTGACACTGGGCACCTTTTCACAGAGAGCTGCAGCATCTCTCCAAGCTCCGTTCCATCGC
 TGTCTACTATGACAACCCCATACGGTGCCTCCTGAGAAGTGCCGCTGTGCAGTCGGCAGCATCCTGAGT
 GAGGGGGAGGAGTCGCCTTCACCTGAGCTCATCCACCTCTATCAGAAATTTGGCTTCAAGATATTCTCCT
 TCCCAGCACCTAGCCATGTGGTCATAGCTACCTTCCCTTACACCACCCCATATCCATCTGGCTGGCTGC
 CCGCCGAGTCCATCCTGCCTTGGATACCTACATCAAGGAGCGGAAGCTGTGTGCTCACCTCGCCTGGAG
 ATCTACCAGCAAGACAAGATCCATTTTCATGTGCCCACTGGCAAGGCAAGGAGATTCTACGTGCCAGAGG
 TGAAGGAGACAGAGCGGAAATGCCGGGAGCTTGCAGGAGGCCACTGACACCCAGACGGATGGCACAGGAGC
 TGATACAAGTGATGCAAGTTCTGTGAGCCTGGATGTTTCGCCCTGGCAGCCGGGAGACTTCAGCCACCACA
 CTTTCTCCTGGGGCAGGCAACCGTGGCTGGGACGACGGTGACAACCGCAGCGAGCACAGCTACAGTGAAT
 CGGGTGCCAGTGGCTCGTCTTTGAGGAGCTGGACCTGGAGGGCGAGGGACCTTTGGGAGAACCCCGACT
 GAACCTGAAGCCAAGCTTCTGGGGCCCTCGGGAGCTCAGCACCCCTGAGAGGGGTGAGGAGTAATGA
 TAAGCCCTCAACCCTCCTGTGGTGCATTTGCTAAGGAAGTGAAGCAACTCTAGCCCTTCTTCTCCTTA
 ACCTGCTAGCTTGGCTGGGAGTACCAGGAGCCTGAGTTGTCTGCTCCAAGCCTAACCTCCTTCTCCT
 ACTGCCCTTTTGGCTCCAGGGCCGAAGGAGCCAGAGACTGTTATCTGCACCAGCCTCCTGGGCTGCCAA
 CTTTGCAGACTCTTATTGGAGCTTCCAGAACCCAGAATAAAGTGAATTACTTTCTTGTTCACCTGGAAA
 AAAAAAAAAAAAAAAAAAAAAA

Restriction Sites: RsrII-NotI
ACCN: NM_011573
Insert Size: 930 bp



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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).
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:	<ol style="list-style-type: none">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:	BC002248 , AAH02248
RefSeq Size:	1423 bp
RefSeq ORF:	930 bp
Locus ID:	21767
UniProt ID:	E9Q137
Cytogenetics:	9 F1
Gene Summary:	<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. 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. 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. Also involved in the repair of covalent DNA-protein cross-links (DPCs) during DNA synthesis: acts by bridging VCP/p97 to covalent DNA-protein cross-links (DPCs) and initiating resolution of DPCs by SPRTN.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (1) represents the longest transcript. Variants 1, 2, and 3 encode the same protein. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p>