

Product datasheet for SC332019

TEX264 (NM 001243727) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: TEX264 (NM_001243727) Human Untagged Clone

Tag: Tag Free
Symbol: TEX264
Synonyms: ZSIG11

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC332019 representing NM_001243727.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

CCCACTGCCCCTGAGAAGGGCAAGGAGTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM 001243727

Insert Size: 720 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).

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).



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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: <u>NM 001243727.2</u>

 RefSeq Size:
 1181 bp

 RefSeq ORF:
 720 bp

 Locus ID:
 51368

 UniProt ID:
 Q9Y6I9

 Cytogenetics:
 3p21.2

Protein Families: Secreted Protein, Transmembrane

MW: 26 kDa

Gene Summary: 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]

Transcript Variant: This variant (6) differs in the 5' UTR and lacks an exon in the coding region, but maintains the reading frame, compared to variant 1. The encoded isoform (2) is shorter than isoform 1.