

Product datasheet for **TP300008L**

TEX264 (NM_015926) Human Recombinant Protein

Product data:

Product Type:	Recombinant Proteins
Description:	Recombinant protein of human testis expressed 264 (TEX264), transcript variant 1, 1 mg
Species:	Human
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>RC200008 protein sequence Red =Cloning site Green =Tags(s)
	<p>MSDLLLLGLIGGLTLLLLLTLAFAGYSGLLAGVEVSAGSPPIRNVTVAYKFHMGlyGETGRLFTESCSI SPKLRSAVYYDNPMMVPPDKCRCVAVGSILSEGEESPSPELIDLYQKFGFKVFSFPAPSHVVTATFPYTT ILSIWLATRRVHPALDITYIKERKLCAYPRLEIYQEDQIHFMCLARQGDYVPEMKETEWKWRGLVEAID TQVDGTGADTMSDTSSVSLEVSPGSRETSATLSPGASSRGWDDGDTRSEHSYSESGASGSSFEELDLEG EGPLGESRLDPGTEPLGTTKWLWEPTAPEKKGKE</p> <p>TRTRPLEQKLISEEDLAANDILDYKDDDDKV</p>
Tag:	C-Myc/DDK
Predicted MW:	34 kDa
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Preparation:	Recombinant protein was captured through anti-DDK affinity column followed by conventional chromatography steps.
Note:	For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.
Storage:	Store at -80°C.
Stability:	Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.
RefSeq:	<u>NP_057010</u>
Locus ID:	51368



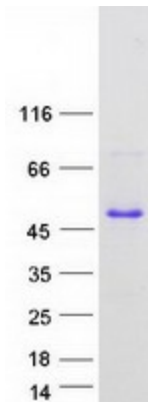
[View online »](#)

UniProt ID: [Q9Y6I9](#)
RefSeq Size: 1403
Cytogenetics: 3p21.2
RefSeq ORF: 939
Synonyms: ZSIG11

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]

Protein Families: Secreted Protein, Transmembrane

Product images:



Coomassie blue staining of purified TEX264 protein (Cat# [TP300008]). The protein was produced from HEK293T cells transfected with TEX264 cDNA clone (Cat# [RC200008]) using MegaTran 2.0 (Cat# [TT210002]).