

Product datasheet for **SC333190**

TSPEAR (NM_001272037) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TSPEAR (NM_001272037) Human Untagged Clone
Tag:	Tag Free
Symbol:	TSPEAR
Synonyms:	C21orf29; DFNB98; ECTD14; TSP-EAR
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC333190 representing NM_001272037. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGAGCTTCCCAGCATCCAGGATTTTCTCCAGTGTGACCTCTTCCCTGAAGAATTTTCCATCGTCGTA
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GCCGGCGCTGGCAGACCCGAGTGTCTTCCGACGCCCCGCCCTGGTGGATGGCCGCTGGCACACACTG
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CAGGGCTACGAGGGCTTCGTGGCGGTGCACAGCCTCCCCACCGTCGGCTGCAGGGACTGGGAGGCCTTC
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AGGACACGCTGA
  
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Restriction Sites:	Sgfl-Mlul
ACCN:	NM_001272037
Insert Size:	1806 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).
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_001272037.1</u>
RefSeq Size:	4042 bp
RefSeq ORF:	1806 bp
Locus ID:	54084
UniProt ID:	<u>Q8WU66</u>
Cytogenetics:	21q22.3
MW:	67.9 kDa
Gene Summary:	This gene encodes a protein that contains a N-terminal thrombospondin-type laminin G domain and several tandem arranged epilepsy-associated repeats (EARs). A mutation in this gene is the cause of autosomal recessive deafness-98. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Dec 2012] Transcript Variant: This variant (2) contains an additional exon in the 5' region that causes translation initiation from a downstream AUG compared to variant 1. The resulting isoform (2) has a shorter N-terminus compared to isoform 1.