

Product datasheet for **SC335967**

TMPRSS5 (NM_001288750) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TMPRSS5 (NM_001288750) Human Untagged Clone
Tag:	Tag Free
Symbol:	TMPRSS5
Synonyms:	SPINESIN
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC335967 representing NM_001288750. Blue=Insert sequence Red=Cloning site Green=Tag(s)

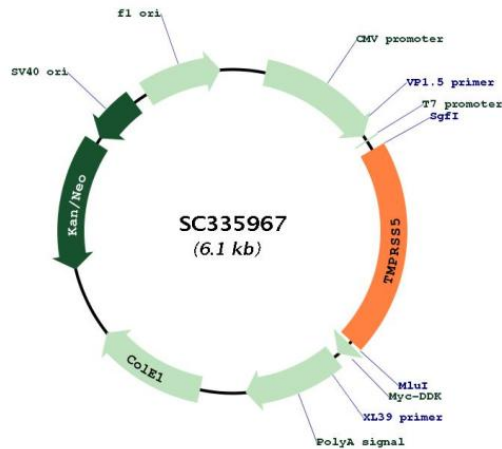
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GGTGTCTACGCCAAGGTAGCTGAGTTTCTGGACTGGATCCATGACACTGCTCAGGACTCCCTCCTTGA
ACGGTACGGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
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Restriction Sites: SgfI-MluI

Plasmid Map:



ACCN: NM_001288750

Insert Size: 1242 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: [NM_001288750.1](#)

RefSeq Size: 2157 bp

RefSeq ORF: 1242 bp

Locus ID: 80975

UniProt ID: [Q9H3S3](#)

Cytogenetics: 11q23.2

Protein Families: Druggable Genome, Protease, Transmembrane

MW: 44.7 kDa

Gene Summary: This gene encodes a protein that belongs to the serine protease family. Serine proteases are known to be involved in many physiological and pathological processes. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2014]
Transcript Variant: This variant (3) differs in its 5' UTR and initiates translation at a downstream start codon, compared to variant 1. The encoded isoform (3) has a shorter N-terminus, compared to isoform 1.