

Product datasheet for SC335955

TMPRSS4 (NM_001290094) Human Untagged Clone

Product data:

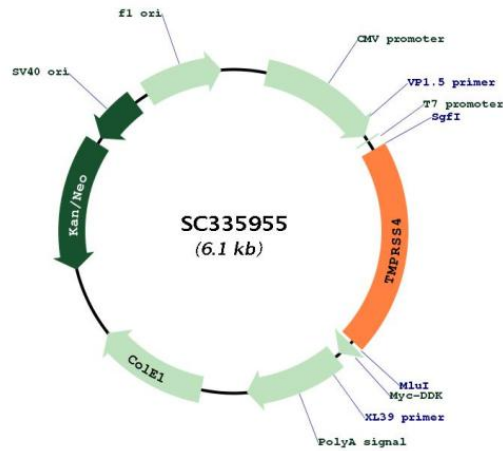
Product Type:	Expression Plasmids
Product Name:	TMPRSS4 (NM_001290094) Human Untagged Clone
Tag:	Tag Free
Symbol:	TMPRSS4
Synonyms:	CAP2; CAPH2; MT-SP2; TMPRSS3
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC335955 representing NM_001290094. Blue=Insert sequence Red=Cloning site Green=Tag(s)

```
GCTCGTTTAGTGAACCGTCAGAATTTGTAAACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGGAGACCTTCAGAAAGGTGGGGATCCCCATCATCATAGCACTACTGAGCCTGGCGAGTATCATCATT
GTGGTTGTCTCATCAAGGTGATTCTGGATAAATACTACTTCTCTGCGGGCAGCCTCTCCACTTCATC
CCGAGGAAGCAGCTGTGTACGGAGAGCTGGACTGTCCCTTGGGGGAGGACGAGGAGCACTGTGTCAAG
AGCTTCCCGAAGGGCCTGCAGTGGCAGTCCGCCTCTCCAAGGACCGATCCACACTGCAGGTGCTGGAC
TCGGCCACAGGGAAGTGTCTGCTGCTGTTTCGACAACCTCACAGAAGCTCTCGCTGAGACAGCCTGT
AGGCAGATGGGCTACAGCAGCAAACCCACTTTCAGAGCTGTGGAGATTGGCCAGACCAGGATCTGGAT
GTTGTTGAAATCACAGAAAACAGCCAGGAGCTTCGCATGCGGAACTCAAGTGGGCCCTGTCTCTCAGGC
TCCTGTCTCCCTGCACTGTCTTGCTGTGGGAAGAGCCTGAAGACCCCCCGTGTGGTGGTGGGGAG
GAGGCCTCTGTGGATTCTTGGCCTTGGCAGGTCAGCATCCAGTACGACAAACAGCACGTCTGTGGAGGG
AGCATCTGGACCCCACTGGGTCTCACGGCAGCCCACTGCTTCAGGAAACATACCGATGTGTTCAAC
TGGAAAGTGGGGCAGGCTCAGACAACTGGGCAGCTTCCCATCCCTGGCTGTGGCAAGATCATCATC
ATTGAATCAACCCATGTACCCCAAAGACAATGACATCGCCCTCATGAAGCTGCAGTCCCCTACT
TTCTCAGGCACAGTCAGGCCATCTGTCTGCCCTTCTTTGATGAGGAGCTCACTCCAGCCACCCCACTC
TGGATCATTGGATGGGGCTTACGAAGCAGAATGGAGGGAAGATGTCTGACATAGCTGCAGGCGTCA
GTCAGGTCATTGACAGCACACGGTGAATGCAGACGATGCGTACCAGGGGAAGTCAACGAGAAGATG
ATGTGTGCAGGCATCCCGAAGGGGTGTGGACACCTGCCAGGTTGACAGTGGTGGGCCCTGATGTAC
CAATCTGACCAGTGGCATGTGGTGGGCATCGTTAGTTGGGGCTATGGCTGCGGGGGCCCGAGCACCCCA
GGAGTATACACCAAGGTCTCAGCCTATCTCAACTGGATCTACAATGTCTGGAAGGCTGAGCTGTA
ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
```

Restriction Sites: Sgfl-Mlul



[View online »](#)

Plasmid Map:


ACCN: NM_001290094

Insert Size: 1239 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_001290094.1](#)

RefSeq Size: 3553 bp

RefSeq ORF: 1239 bp

Locus ID: 56649

Cytogenetics: 11q23.3

Protein Families: Druggable Genome, Protease, Transmembrane

MW: 45.3 kDa

Gene Summary:

This gene encodes a member of the serine protease family. Serine proteases are known to be involved in a variety of biological processes, whose malfunction often leads to human diseases and disorders. This gene was identified as a gene overexpressed in pancreatic carcinoma. The encoded protein is membrane bound with a N-terminal anchor sequence and a glycosylated extracellular region containing the serine protease domain. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]

Transcript Variant: This variant (6) uses an alternate splice junction at the end of a 5' exon compared to variant 1. The resulting isoform (6) is shorter at the N-terminus compared to isoform 1. 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.