

Product datasheet for SC207669

PRSS8 (NM_002773) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	PRSS8 (NM_002773) Human 3' UTR Clone
Symbol:	PRSS8
Synonyms:	CAP1; PROSTASIN
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_002773
Insert Size:	573 bp
Insert Sequence:	>SC207669 3' UTR clone of NM_002773 The sequence shown below is from the reference sequence of NM_002773. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site Blue=Stop Codon
	CAATTGGCAGAGCTCAGAATTCAAGCGATCGC
	ATGGCTCAGCGAGCAC TGA GCTGGCCCTACTTCCAGGATGGATGCATCACACTCAAGGACAGGAGCCTGG TCCTTCCCTGATGGCCTTTGGACCCAGGGCCTGACTTGAGCCACTCCTTCCT
	ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCG
Restriction Sites:	Sgfl-Mlul
OTI Disclaimer:	Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).



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Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
RefSeq:	<u>NM 002773.3</u>
Summary:	This gene encodes a member of the peptidase S1 or chymotrypsin family of serine proteases. The encoded preproprotein is proteolytically processed to generate light and heavy chains that associate via a disulfide bond to form the heterodimeric enzyme. This enzyme is highly expressed in prostate epithelia and is one of several proteolytic enzymes found in seminal fluid. This protease exhibits trypsin-like substrate specificity, cleaving protein substrates at the carboxyl terminus of lysine or arginine residues. The encoded protease partially mediates proteolytic activation of the epithelial sodium channel, a regulator of sodium balance, and may also play a role in epithelial barrier formation. [provided by RefSeq, Feb 2016]
Locus ID:	5652

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