

Product datasheet for SC205155

CTSA (NM_001127695) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	CTSA (NM_001127695) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	CTSA
Synonyms:	GLB2; GSL; NGBE; PPCA; PPGB
ACCN:	NM_001127695
Insert Size:	413 bp
Insert Sequence:	>SC205155 3'UTR clone of NM_001127695 The sequence shown below is from the reference sequence of NM_001127695. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site
	GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA <mark>GCGATCGC</mark> C TCCCGCTTCCTGAACAAGCAGCCATACTGATGACCACAGCAACCAGCTCCACGGCCTGATGCAGCCCCT CCCAGCCTCTCCCGCTAGGAGAGTCCTCTTCTAAGCAAAGTGCCCCTGCAGGCCGGGGTTCTGCCGCCAG GACTGCCCCCTTCCCAGAGCCCTGTACATCCCAGACTGGGCCCAGGGTCTCCCATAGACAGCCTGGGGG CAAGTTAGCACTTTATTCCCGCAGCAGTTCCTGAATGGGGTGGCCTGGCCCCTTCTCTGCTTAAAGAAT GCCCTTTATGATGCACTGATTCCATCCCAGGAACCCAACAGAGCTCAGGACAGCCCACAGGGAGGTGGT GGACGGACTGTAATTGATAGATTAGGAATTAAATTGGGTACAGCTTCAAAAAAAA
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).
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 001127695.2</u>



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Summary:	This gene encodes a member of the peptidase S10 family of serine carboxypeptidases. Alternative splicing results in multiple transcript variants, at least one of which encodes a preproprotein that is proteolytically processed to generate two chains that comprise the heterodimeric active enzyme. This enzyme possesses deamidase, esterase and carboxypeptidase activities and acts as a scaffold in the lysosomal multienzyme complex. Mutations in this gene are associated with galactosialidosis. [provided by RefSeq, Nov 2015]
Locus ID:	5476
MW:	15.3

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