

Product datasheet for **SC108811**

CTAGE5 (MIA2) (NM_005930) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	CTAGE5 (MIA2) (NM_005930) Human Untagged Clone
Tag:	Tag Free
Symbol:	CTAGE5
Synonyms:	CTAGE5; MEA6; MGEA; MGEA6; MGEA11; TALI
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL5</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >NCBI ORF sequence for NM_005930, the custom clone sequence may differ by one or more nucleotides

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ATGGAGGAGCCCGGGTTACCCCTCAACCGTATTTGGGGCTGCTCCTGGAGGAGCTACGCAGGGTTGTGG
CAGCACTGCCTGAAGGTATGAGACCAGATTCTAATCTTTATGGTTTTCCATGGGAATGGTGATATGTGC
AGCTGTTGTTGGATTTTTGCTGTTCTCTTTTTTTTGGGAGAAGTTTTAGATCGGTTAGGAGTCGGCTT
TATGTGGGACGAGAGAAAAAGCTTGCTCTAATGCTTTCTGGACTAATTGAAGAAAAAAGTAAACTACTTG
AAAAATTTAGCCTTGTTCAAAAAGAGTATGAAGGCTATGAAGTAGAGTCATCTTTAAAGGATGCCAGCTT
TGAGAAGGAGGCAACAGAAAGCACAAGTTTGGAGGCAACCTGTGAAAAGCTGAACAGGTCCAATTCTGAA
CTTGAGGATGAAATACTGTCTAGAAAAAGAGTTAAAAGAAGAGAAATCCAAACATTCTGAACAAGATG
AATTGATGGCGGATATTTCAAAAAGGATACAGTCTCTAGAAGATGAGTCAAAATCCCTCAAATCACAAAGT
AGCTGAAGCCAAAATGACCTTCAAGATATTTCAAATGAATGAAGAACGACTGAAGATAGCAATAAAAGAT
GCTTTGAATGAAAATTCTCAACTTCAGGAAAGCCAGAAACAGCTTTTGCAAGAAGCTGAAGTATGAAAAG
AACAAAGTGAAGTAAATAAACAGAAAGTAACATTTGAAGACTCCAAAGTACATGCAGAACAAAGTTCT
AAATGATAAAGAAAGTCACATCAAGACTCTGACTGAACGCTTGTTAAAGATGAAAGATTGGGCTGCATG
CTTGAGAGAAGACATAACGGATGATGATAACTTGGAAATAGAAATGAACAGTGAATCGGAAAAATGGTGCTT
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AACCTTAGAAGGAGAAAGAAACCAATTTATATTCAGTTGCTGAAGTTGATAAAAACAAAGGAGAGCTT
ACAGAGCATATTAATACTTCAGACTGAACAAGCATCTTTGCAGTCAGAAAACACACATTTTGAAGATG
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CAGGAAATTAACAGTAGAGGAAAATTATCGGTTAGAGAAAAGAAGGAAACTTTCTAAAGTAGATGAAAAG
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TCGGAATGCTGAAAGAAACCTCAATGATTTAAGGAAAGAAAATGCTCACACAGACAAAAATTAAGTAA
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GGCCAGGGAATCCTCTGGACCATCAGATTACCAATGAAAGAGGAGAATCAAGCTGTGATAGGTTAACCG
ATCCTCATAGGGCTCCCTGCACTGGGTCTCTGTACCTCCATGGGACCAGGACCGTAGGATGATGTT
TCCTCCGCCAGGACAATCATATCCTGATTACAGCCCTTCCACAAAGGCAAGACAGATTTGTTCTAAT
TCTGGTAGACTGTCTGGACCAGCAGAAGTAAAAATATGCCTTCTTTGGATAAAATGGATGGGT
CAATGCCTTCAGAAATGGAATCCAGTAGAAATGATACCAAGATGATCTTGGTAATTTAAATGTGCCTGA
TTCATCTCTCCCTGCTGAAAATGAAGCCACTGGCCCTGGCTTTGTTTCTCCACCTCTTGCTCCAATCAGA
GGTCCATTGTTCCAGTGGATGCAAGAGGCCAATCTTGAGAAGAGGACCTCCTTTCCCCCACCTCCTC
CAGGAGCCATGTTGGAGCTTCTCGAGATTATTTCCACCAGGGGATTTCCAGGTCCACCACCTGCTCC
ATTTGCAATGAGAAATGTCTATCCACCGAGGGGTTTTCTCTTACCTTCCCCAAGACCTGGATTTTTC
CCCCACCCACATTCTGAAGGTAGAAGTGAGTCCCCTCAGGTTTGATTCCACCTTCAAATGAGCCTG
CTACTGAACATCCAGAACCACAGCAAGAAACCTGA
    
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5' Read Nucleotide Sequence:

>OriGene 5' read for NM_005930 unedited
 GGTGCAATATATTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGCACGAGGCCGG
 TTGCCGGGTGCGGATTCGGGTTCCGGACCGAAGGCTGTGTGTTCTCCGCCGTTTATTGTG
 GCCCCGACAGGCCGGGGTACTGTGGCGACCACGAGAGCAGCTTTGGCGCTATGGAGGAG
 CCCGGGGTTACCCCTCAACCGTATTTGGGGCTGCTCCTGGAGGAGCTACGCAGGGTTGTG
 GCAGACTGCCTGAAGGTATGAGACCAGATTCTAATCTTTATGGTTTTCCATGGGAATTG
 GTGATATGTGCAGCTGTTGTTGGATTTTTGCTGTTCTCTTTTTTTTTGTGGAGAAGTTTT
 AGATCGGTTAGGAGTCGGCTTTATGTGGACGAGAGAAAAAGCTTGTCTAATGCTTTCT
 GGACTAATTGAAGAAAAAGTAACTACTTGAAAAATTTAGCCTTGTTCAAAAAGAGTAT
 GAAGGCTATGAAGTAGAGTCATCTTTAAAGGATGCCAGCTTTGAGAAGGAGGCAACAGAA
 GCACAAAGTTTGGAGGCAACCTGTGAAAAGCTGAACAGGTCCAATTCTGAACTTGAGGAT
 GAAATACTCTGTCTAGAAAAAGAGTTAAAAGAAGAGAAATCCAAACATTCTGAACAAGAT
 GAATTGATGGCGGATATTTCAAAAAGGATACAGTCTCTAGAAGATGAGTCAAAATCCCTC
 ANATCACAAAGTAGCTGAAGCCAAAATGACCTTCAAGATATTTANATGAATGAAGAACGA
 CTGAAGATAGCCATAAAAGATGCTTTTGAATGAANATTCTCAACTCCAGGAAAGCCAGAA
 ACAGCTTTTTGCAGAAAGCTGAGTATGGAAAGAAACAGTGAGTGAACCTAATAAACAGAA
 GTTACATTTGAAGACTCCAAGTC

3' Read Nucleotide Sequence:

>OriGene 3' read for NM_005930 unedited
 ACCGCGGCCGCAATCTAGAGTCGAGTTTTTTTTTTTTTTTTTATAGAAAACTACTGAT
 TTTATTATTATCAATGCCAAACAATCTTTAAATCAACTATAATTTATTTAAGCATAAAG
 TTAATTTACATTTGTCCACAACCACAGTAAATACAGTCTTGGCATAAAGACACAGACT
 TCAGAATTTAAAGCCTCCCCACCTGAAAGATTACATATATAAACTCCCACTATTGTTAA
 TATAATAGGGGGATAATTCAAATTTGGAATAGTTTACCCGGTTAAAATAAAAAGGGGAA
 AATTTTTTCCCCCAAGAAGACCTTTTTTAAAAAATTTTTTTTTTTAAAAAAAACCT
 CCTCCGGGGGAAAAATTATAACCCCCCTTCCCCCCCCCGGGGGGGGGGAGGGGGG
 GGGGGTTTTTTTTTTTTTATAAAACGCGCTTCTTTTTTTTTGGAGGGGGGAGAGAGGA
 GCCCGCCCCCTTTTTTTTTTTTTTAAAAAAAAAAAAAAAAAGAAAAANCTCTCT
 TTGTGTGGGTGGGGGGGGCGGCTCTTCTTCTTCTTCTTGGGGGGGGGGGGGGGGA
 AAAAAAGAGGGGGGGCGCTCTTCTTCTTCTTCTTGGGGGGGGGGGGGGGGGGGGG
 GGGGGGAACACACACATCTTCTGGGGGGGGGGGGGGGAAACACACCACCCCTCTC
 CCCTTTTTTCTTTTTTTTTTTTTTTTTTTTTNNGGGGGGGGGGGGGGGGGGGGGGG
 CGGGGGGGCGCCCTCCCTTTCCCTTAAAATAATATAAAAATAATTATCCCCCCTC
 CTCCTCTCTCCCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGAGGNNANNTTATAN
 TTTTTATTNNTATTTCTTTCCCCCCCCCATATNCACAAAGAAAGAAAAAAAAG

Restriction Sites:

NotI-NotI

ACCN:

NM_005930

Insert Size:

3090 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:	<ol style="list-style-type: none">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_005930.2 , NP_005921.2
RefSeq Size:	3852 bp
RefSeq ORF:	2415 bp
Locus ID:	4253
UniProt ID:	Q96PC5
Cytogenetics:	14q21.1
Protein Families:	Transmembrane
Gene Summary:	<p>This gene encodes s receptor in the endoplasmic reticulum, which plays a role in the export of large pre-chylomicrons and pre-very low density lipoproteins (pre-VLDLs). Three major classes of transcripts are generated from this gene- melanoma inhibitory activity 2-specific transcripts, cTAGE family member 5-specific transcripts and transcripts that include exons from both these transcript species (TANGO1-like or TALI). Additionally, alternative splicing in these transcripts results in multiple transcript variants encoding multiple isoforms. [provided by RefSeq, Sep 2016]</p> <p>Transcript Variant: This variant (1) uses an alternate in-frame splice junction at the 3' end of an exon compared to variant 6. It encodes isoform 1, which lacks an internal segment compared to isoform 6.</p>