

Product datasheet for **SC309981**

ADAMTS10 (NM_030957) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ADAMTS10 (NM_030957) Human Untagged Clone
Tag:	Tag Free
Symbol:	ADAMTS10
Synonyms:	ADAM-TS10; ADAMTS-10; WMS; WMS1
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_030957 edited
GGCGCGGAGGCCCGGGCGCGCAGGCTCAAAGAAGAAGAAACCAAGGCCAGAGAG
GGAGGCCAGGTGCAGGGAGCAGCGAGGGAAGGATCCGTACAGGGGCCAACACTACTC
CACCAACCGAAGCCCCAAAAGGAGCCCGGTGATGCTGCGAAGGCTGTGAACAGGGGAGG
CGGCACGTGGGGGCTGCCGGCAGCCGGGCTGGGGAGAGACATGTGGACACGTGGCCTC
TATGGCTCCCGCTGCCAGATCCTCCGCTGGGCCCTCGCCTGGGGCTGGGCCTCATGTT
CGAGGTCACGCACGCCTCCGGTCTCAAGATGAGTTCTGTCCAGTCTGGAGAGCTATGA
GATCGCTTCCCCACCCGCTGGACCACAACGGGGCACTGCTGGCCTTCTCGCCACTCC
TCCCCGGAGGCAGCGCCGCGCACGGGGCCACAGCCAGTCCCCTCTTCTACAAAGT
GGCCTCGCCAGCACCCACTTCTGCTGAACCTGACCCGAGCTCCCGTCTACTGGCAGG
GCACGTCTCCGTGGAGTACTGGACACGGGAGGGCTGGCCTGGCAGAGGGCGGCCGGCC
CCACTGCCTCTACGCTGGTCACCTGCAGGGCCAGGCCAGCAGCTCCCATGTGGCCATCAG
CACCTGTGGAGGCTGCACGGCCTGATCGTGGCAGACGAGGAAGAGTACCTGATTGAGCC
CCTGCACGGTGGGCCCAAGGGTTCTCGGAGCCCGAGGAAAGTGGACCACATGTGGTGTA
CAAGCGTTCCTCTCTGCGTACCCCCACCTGGACACAGCCTGTGGAGTGAGAGATGAGAA
ACCGTGAAAGGGCGGCCATGGTGGCTGCGGACCTTGAAGCCACCGCCTGCCAGGCCCT
GGGGAATGAAACAGAGCGTGGCCAGCCAGGCTGAAGCGATCGGTGAGCCGAGAGCGCTA
CGTGGAGACCTGGTGGTGGCTGACAAGATGATGGTGGCCTATCACGGGCGCCGGGATGT
GGAGCAGTATGCTCTGGCCATCATGAACATTGTTGCCAACTTTTCCAGGACTCGAGTCT
GGGAAGCACCGTTAACATCCTCGTAACTCGCCTCATCCTGCTCACGGAGGACCAGCCAC
TCTGGAGATCACCCACATGCCGGAAAGTCCCTGGACAGCTTCTGTAAGTGGCAGAAATC
CATCGTGAACCACAGCGCCATGGCAATGCCATTCCAGAGAACGGTGTGGCTAACCATGA
CACAGCAGTGTCTATCACAGCTATGACATCTGCATCTACAAGAACAACCCTGCGGCAC
ACTAGGCCTGGCCCCGTGGCGGAATGTGTGAGCGGAGAGAAGCTGCAGCGTCAATGA
GGACATTGGCCTGGCCACAGCGTTCACCATTGCCACGAGATCGGGCACACATTCGGCAT
GAACCATGACGGCGTGGGAAACAGCTGTGGGGCCCGTGGTCAGGACCCAGCCAAGCTCAT
GGCTGCCACATTACCATGAAGACCAACCCATTCTGTGGTTCATCTGCACCCGTGACTA



[View online »](#)

CATCACCAGCTTTCTAGACTCGGGCCTGGGGCTCTGCCTGAACAACCGGCCCCAGACA
 GGACTTTGTGTACCCGACAGTGGCACCGGGCAAGCCTACGATGCAGATGAGCAATGCCG
 CTTTCAGCATGGAGTCAAATCGCGTCAGTGTAATACGGGGAGGTCTGCAGCGAGCTGTG
 GTGTCTGAGCAAGAGCAACCGGTGCATACCAACAGCATCCCGGCCGCCGAGGGCACGCT
 GTGCCAGACGCACACCATCGACAAGGGGTGGTGTACAAACGGGTCTGTGTCCCTTTGG
 GTCGCGCCAGAGGGTGTGGACGGAGCCTGGGGCCGTGGACTCCATGGGGCGACTGCAG
 CCGGACCTGTGGCGCGCGCTGTCTTCTAGCCGTCAGTGCAGACAGCCCAAGGCAAC
 CATCGGGGGCAAGTACTGTCTGGGTGAGAGAAGCGCGCACCGCTCCTGCAACACGGATGA
 CTGTCCCTGGCTCCAGGACTTCAGAGAAGTGCAGTGTCTGAATTTGACAGCATCCC
 TTTCCGTGGGAAATTCTACAAGTGGAAAACGTACCGGGGAGGGGGCGTGAAGGCCTGCTC
 GCTCACGTGCCTAGCGGAAGGCTTCAACTTCTACACGGAGAGGGCGGCAGCCGTGGTGA
 CGGGACACCTGCCGTCCAGACACGGTGGACATTTGCGTCAGTGGCGAATGCAAGCACGT
 GGGCTGCGACCGAGTCTGGCTCCGACCTGCGGGAGGACAAGTCCGAGTGTGTGGCGG
 TGACGGCAGTGCCTGCGAGACCATCGAGGGCGTCTTACGCCAGCCTCACCTGGGGCCGG
 GTACGAGGATGTGTCTGGATTCCAAAGGCTCCGTCCACATCTTCATCCAGGATCTGAA
 CCTCTCTCAGTCACTTGGCCCTGAAGGGAGACCAGGAGTCCCTGCTGCTGGAGGGGCT
 GCCYGGGACCCCCAGCCCCACCGTCTGCCTCTAGCTGGGACCACCTTTCAACTGCGACA
 GGGGCCAGACCAGGTCCAGAGCCTCGAAGCCCTGGGACCGATTAATGCATCTCTCATCGT
 CATGGTGTGGCCCGGACCGAGCTGCCTGCCCTCCGCTACCGCTTCAATGCCCCCATCGC
 CCGTGACTCGTGCCTCCCTACTCCTGGCACTATGCGCCCTGGACCAAGTGTCTGGCCCA
 GTGTGCAGGCGGTAGCCAGGTGCAGGCGGTGGAGTGCCGAACCAAGTGGACAGTCCGC
 GGTGCGCCCCACTACTGCAGTGCACAGCAAGCTGCCAAAAGGCAGCGCGCTGCAA
 CACGGAGCCTTGCCCTCCAGACTGGGTTGTAGGGAAGTGGTCTGCTGCAGCCGACGGA
 CGATGCAGGCGTGCAGCGCGCTCGGTCTGTGCCAGCGCCGCGTCTCTGCCGCGGAGGA
 GAAGGCGTGGACGACAGCGCATGCCCGCAGCCCGCCACCTGTACTGGAGGCCTGCCA
 CGGCCCACTTGCCCTCCGGAGTGGGCGGCCCTCGACTGGTCTGAGTGCACCCCAAGCTG
 CGGGCCGGGCTCCGCCACCGCGTGGTCTTTGCAAGAGCGCAGACCACTGCGCCACGCT
 GCCCCCGGCGCACTGCTCACCCCGCCCAAGCCACCGGCCACCATGCGCTGCAACTTGCG
 CCGCTGCCCCCGGCCGCTGGGTGGTGGCGAGTGGGGTGAAGTCTGTCACAGTGGG
 CGTCGGGCAGCGGAGCGCTCGGTGCGCTGCACCAGCCACACGGGCCAGGCGTCGCACGA
 GTGCACGAGGCCCTGCGGCCGCCACACGCAGCAGTGTGAGGCCAAGTGCAGACGCC
 AACCCCGGGGACGGCCCTGAAGAGTGAAGGATGTGAACAAGGTGCGCTACTGCCCCCT
 GGTGCTCAAATTTCAATTCTGCAGCCGAGCCTACTTCCGCCAGATGTGCTGCAAAACCTG
 CCAGGGCCACTAGGGGGCGCGCGCACCCGGAGCCACAGCTGGCGGGGTCTCCGCCGCCA
 GCCCTGCAGCGGGCCGGCCAGAGGGGGCCCCGGGGGGCGGGAAGTGGGAGGGAAGGGTG
 AGACGGAGCCGGAAGTTATTTATTGGGAACCCCTGCAGGGCCCTGGTGGGGGATGGAG
 AGGGGTGGCTATCCCCCAGAGCCCTCTTACAGATCCGCCCTTCCAGTTCACATAGT
 GAGACCCCTTCCAGGGTGGGTCTGGGGAGGGGACAGCTGTTACCCCAAGCACCCTTTG
 AACCCACCCTTAGGGAGTCCCCAACACTCCCCCTCTTTGATCGGAATATGTACTGTG
 AAGAGTAGGGGTGGGAGGTGGTCTGCCGGTCCCTGCCCCAGCACTGCCCTACCCCT
 CCACTCCGAGCTGGGGGGAATCTGTGTCTGCTATGGGAGGGGGGGGTAGCACCACTC
 CCTGACACCTCCCTGACCTGACTAGGGGGAGGATCCACCCCAACCTCTGCCCTGCCCG
 CCCCAGGGGACCCCGACATCCAGGCCACCCCTCATGGTGTACAGACCCTGCCCTGGG
 GCCCACACTCTGCCAGGAAGCCCTACATCAATAAAGTTCTGTCTTGTGTAGATTTCT
 ACGAAAAAAAAAAAAAAAAAAAA

5' Read Nucleotide Sequence:	<p>>OriGene 5' read for NM_030957 unedited</p> <pre> NGTTCAGTTAAATTTGTATACGACTCACTATAGGCGGCCGCGNAATTCGCACGAGGGCGC GGAGGCCCCGGGCGCGGCCGAGGAGCCCGGTGATGCTGCGAAGCTGTGAACAGGGGAGGC GGCACTGTGGGGGCTGCCGGCAGCCGGGGCTGGGAGAGACATGTGGACACGTGGCCTCT ATGGTCCCAGCTGCCAGATCCTCCGCTGGGCCCTCGCCTGGGGCTGGCCTCATGTTT GAGGTACGCACGCCTTCCGGTCTCAAGATGAGTTCCTGTCCAGTCTGGAGAGCTATGAG ATCGCCTTCCCCACCCCGTGGACCACAACGGGGCACTGCTGGCCTTCTCGCCACCTCCT CCCCGGAGGCAGCGCCGCGGCACGGGGCCACAGCCGAGTCCCGCCTTCTTACAAAGTG GCCTCGCCAGCACCCTTCTGCTGAACCTGACCCGAGCTCCCGTCTACTGGCAGGG CACGTCTCCGTGGAGTACTGGACACGGGAGGGCCTGGCCTGGCAGAGGGCGGCCGGCC CACTGCCTCTACGCTGGTACCTGCAGGGCCAGGCCAGCAGCTCCCATGTGGCCATCAGC ACCTGTGGAGGCCTGCACGGCCTGATCGTGGCAGACGAGGAAGAGTACCTGATTGAGCCC CTGCACGGTGGGCCAAGGGTTCTCGGAGCCCGAAGAAAGTGGACCACATGTGGTGTAC AAGCGTTCCTCTCTGCGTACCCCCACCTGGACACAGCCTGTGGAGTGAGAGATGAGAAA CCGTNGGAAGGGCGCCATGGTGGCTGCCGAACCTTGAAGCCACCGCCTGCCAGGCCCC TGGGGGAATGAAAC </pre>
3' Read Nucleotide Sequence:	<p>>OriGene 3' read for NM_030957 unedited</p> <pre> CCCGTGTGGCTGGTGCAGCGCACNGACGCTGCCGCTGCCGNANCCGCACTGTGCAGNA CACTACCCCACTCGCCAGCCACCCAGCGGGCCGGGGGCAAGCGCGCAAGTTGCAGCGC ATGGTGGCCGGTGGCTTGGCGGGGGTGGAGAGTGCGCCGGGGGAGCGTGGCGCAGTGG TCTGCGCTCTTGCAAAGGACCACGCGGTGGCGGAGGCCCGGCCCGCAGCTGGGGGTGCAC TCAGACCAGTCGAGGGCCGCCACTCCGGAGGGCAAGTGGGGCCGTGGCAGGCCTCCAGT ACAGGTGGGCGCGGCTGCCGGCATGCGCTGTCTCCAGCGCCTTCTCCTCCGCGGCAGAG ACGCGGCGCTGGCACACGACCGAGCGGCTGCGCACGCCTGCATCGCAGCTGCGGCTGCAG AGCGACCAGTTCCTACAACCCAGTCTGGAGGGCAAGGCTCCGTGTTGCAGGCGCGCTGC CTTTTGGGCAGCTTGTGTGGGCACTGCACTAGTGGGGGGCAGCCGGGAGCTGTCCAGC TGGTTGCGGCACTCCACCGCCTGCACCTGGCTACCGCCTGCACACTGGGCCGAGCACTTG GNTCAGGGCGCATAGTGCAGGAGTANGGGGGCAGCGAGTACGGGCGATGGGGGATTG AAGCCGTAGCGGAGGGCAAGCAGCTCGGTCCGGGCCAGCACCATGACGATGAGAGATGCA TTAATCGGTCCCAGGGCTTCCAGGCTCTGGACCTGGTCTGGCCCTGTGCGATTGAAAG GTGGTCCCCTANAGGCAAACGGTGGGGTGGGGGTCCCGGGCAGCCCTTCCACAGCAA GGACTCCTGGTCTCCCTCCAGGGCCAGTACTGAAAAAAGGTTN </pre>
Restriction Sites:	Please inquire
ACCN:	NM_030957
Insert Size:	4500 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).
OTI Annotation:	The open reading frame of this TrueClone was fully sequenced and found to be a perfect match to the protein associated to this reference.
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_030957.2 , NP_112219.2
RefSeq Size:	4237 bp
RefSeq ORF:	3312 bp
Locus ID:	81794
UniProt ID:	Q9H324
Cytogenetics:	19p13.2
Domains:	tsp_1, Reprolysin, Pep_M12B_propep
Protein Families:	Druggable Genome, Secreted Protein
Gene Summary:	<p>This gene belongs to the ADAMTS (a disintegrin and metalloproteinase domain with thrombospondin type-1 motifs) family of zinc-dependent proteases. ADAMTS proteases are complex secreted enzymes containing a prometalloprotease domain of the reprolysin type attached to an ancillary domain with a highly conserved structure that includes at least one thrombospondin type 1 repeat. They have been demonstrated to have important roles in connective tissue organization, coagulation, inflammation, arthritis, angiogenesis and cell migration. The product of this gene plays a major role in growth and in skin, lens, and heart development. It is also a candidate gene for autosomal recessive Weill-Marchesani syndrome. [provided by RefSeq, Jul 2008]</p> <p>Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1).</p>