

Product datasheet for SC202456

ADAMTS13 (NM_139027) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Product Type:	3' UTR Clones
Product Name:	ADAMTS13 (NM_139027) Human 3' UTR Clone
Symbol:	ADAMTS13
Synonyms:	ADAM-TS13; ADAMTS-13; C9orf8; vWF-CP; VWFCP
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_139027
Insert Size:	236 bp
Insert Sequence:	<pre>>SC202456 3'UTR clone of NM_139027 The sequence shown below is from the reference sequence of NM_139027. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CAGTCCTGGAAGGGAAAGGAAGGAAGCAAGCTGAGGGTCATTGAACATTTGTTCCGTGTCTGGCCAGCCCTGG AGGGTTGACCCCTGGTCTCAGTGCTTTCCAATTCGAACATTTGTTCCGTGTCTGGCCAGCCCTGG AGGGTTGACCCCTGGTCTCAGTGCTTTCCAATTCGAACTTTTTCCAATCTTAGGTATCTACTTTAGAGT CTTCTCCAATGTCCAAAAGGCTAGGGGGTTGGAAGGTGGGGGACTCTGGAAAAGCAGCCCCATCTCCG GGTACCAATAAATAAAACATGCAGGCTGA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACC</pre>
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 139027.6</u>



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	ADAMTS13 (NM_139027) Human 3' UTR Clone – SC202456
Summary:	This gene encodes a member of a family of proteins containing several distinct regions, including a metalloproteinase domain, a disintegrin-like domain, and a thrombospondin type 1 (TS) motif. The enzyme encoded by this gene specifically cleaves von Willebrand Factor (vWF). Defects in this gene are associated with thrombotic thrombocytopenic purpura. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2013]
Locus ID:	11093
MW:	8.3

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