



## **ADAMTS16** peptide (DAG-P0132)

This product is for research use only and is not intended for diagnostic use.

## PRODUCT INFORMATION

Antigen Description	This gene encodes a member of the ADAMTS (a disintegrin and metalloproteinase with thrombospondin motifs) protein family. ADAMTS family members share several distinct protein modules, including a propeptide region, a metalloproteinase domain, a disintegrin-like domain,
	and a thrombospondin type 1 (TS) motif. Individual members of this family differ in the number of C-terminal TS motifs, and some have unique C-terminal domains. The protein encoded by this gene has high sequence similarity to the protein encoded by ADAMTS18, another family member. [provided by RefSeq, Jul 2008]
Purity	> 95 % by SDS-PAGE.

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Conjugate	Unconjugated
Applications	ELISA, WB
Format	Liquid
Buffer	Preservative: None Constituents: 0.001% Tween 20, 30mM HEPES, 2mM EDTA, 150mM Sodium chloride, pH 6.75
Preservative	None
Storage	Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw cycles. Preservative: None Constituents: 0.001% Tween 20, 30mM HEPES, 2mM EDTA, 150mM Sodium chloride, pH 6.75

## **GENE INFORMATION**

Gene Name	DAMTS16 ADAM metallopeptidase with thrombospondin type 1 motif, 16 [ Homo sapid	<u>ens</u>

(human)]

Official Symbol ADAMTS16

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Synonyms	ADAMTS16; ADAM metallopeptidase with thrombospondin type 1 motif, 16; ADAMTS16s; A disintegrin and metalloproteinase with thrombospondin motifs 16; ADAM-TS16; ADAMTS-16; ADAM-TS 16; a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 16;
Entrez Gene ID	170690
mRNA Refseq	NM 139056.2
Protein Refseq	NP 620687.2
UniProt ID	Q2XQZ0
Chromosome Location	5p15
Pathway	Degradation of the extracellular matrix, organism-specific biosystem; Degradation of the extracellular matrix, organism-specific biosystem; Extracellular matrix organization, organism-specific biosystem; Extracellular matrix organization, organism-specific biosystem;
Function	metalloendopeptidase activity; zinc ion binding;