



# Human ADAMTS5 peptide (DAG-P1346)

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

## PRODUCT INFORMATION

<b>Antigen Description</b>	This gene encodes a member of the ADAMTS (a disintegrin and metalloproteinase with thrombospondin motifs) protein family. Members of the family 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 enzyme encoded by this gene contains two C-terminal TS motifs and functions as aggrecanase to cleave aggrecan, a major proteoglycan of cartilage. [provided by RefSeq, Jul 2008]
<b>Specificity</b>	Expressed at low level in placenta primarily but also detected in heart and brain, cervix, uterus, bladder, esophagus, rib cartilage, chondroblastoma, fibrous tissue and a joint capsule from an arthritic patient.
<b>Conjugate</b>	Unconjugated
<b>Sequence Similarities</b>	Contains 1 disintegrin domain.Contains 1 peptidase M12B domain.Contains 2 TSP type-1 domains.
<b>Format</b>	Liquid
<b>Preservative</b>	None
<b>Storage</b>	Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw cycles. Information available upon request.

## GENE INFORMATION

<b>Gene Name</b>	<a href="#">ADAMTS5 ADAM metalloproteinase with thrombospondin type 1 motif, 5 [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	ADAMTS5

<b>Synonyms</b>	ADAMTS5; ADAM metalloproteinase with thrombospondin type 1 motif, 5; ADMP-2; ADAM-TS5; ADAMTS-5; ADAMTS11; ADAM-TS 5; ADAMTS-11; ADAM-TS 11; A disintegrin and metalloproteinase with thrombospondin motifs 5; aggrecanase-2; a disintegrin and metalloproteinase with thrombospondin motifs 11; a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2);
<b>Entrez Gene ID</b>	<a href="#">11096</a>
<b>mRNA Refseq</b>	<a href="#">NM_007038.3</a>
<b>Protein Refseq</b>	<a href="#">NP_008969.2</a>
<b>UniProt ID</b>	Q9UNA0
<b>Chromosome Location</b>	21q21.3
<b>Pathway</b>	Degradation of the extracellular matrix, organism-specific biosystem; Degradation of the extracellular matrix, organism-specific biosystem; Endochondral Ossification, organism-specific biosystem; Extracellular matrix organization, organism-specific biosystem; Extracellular matrix organization, organism-specific biosystem;
<b>Function</b>	integrin binding; metalloendopeptidase activity; metalloproteinase activity; protein binding; zinc ion binding;