



## NAMPT peptide (DAG-P1805)

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

### PRODUCT INFORMATION

<b>Antigen Description</b>	This gene encodes a protein that catalyzes the condensation of nicotinamide with 5-phosphoribosyl-1-pyrophosphate to yield nicotinamide mononucleotide, one step in the biosynthesis of nicotinamide adenine dinucleotide. The protein belongs to the nicotinic acid phosphoribosyltransferase (NAPRTase) family and is thought to be involved in many important biological processes, including metabolism, stress response and aging. This gene has a pseudogene on chromosome 10. [provided by RefSeq, Feb 2011]
<b>Specificity</b>	Expressed in large amounts in bone marrow, liver tissue, and muscle. Also present in heart, placenta, lung, and kidney tissues.
<b>Conjugate</b>	Unconjugated
<b>Sequence Similarities</b>	Belongs to the NAPRTase family.
<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="#">NAMPT nicotinamide phosphoribosyltransferase [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	NAMPT
<b>Synonyms</b>	NAMPT; nicotinamide phosphoribosyltransferase; VF; PBEF; PBEF1; VISFATIN; 1110035O14Rik; NAMPTase; pre-B cell-enhancing factor; pre-B-cell colony enhancing factor 1; pre-B-cell colony-enhancing factor 1;

<b>Entrez Gene ID</b>	<a href="#">10135</a>
<b>mRNA Refseq</b>	<a href="#">NM_005746.2</a>
<b>Protein Refseq</b>	<a href="#">NP_005737.1</a>
<b>UniProt ID</b>	P43490
<b>Chromosome Location</b>	7q22.3
<b>Pathway</b>	Adipogenesis, organism-specific biosystem; BMAL1:CLOCK/NPAS2 Activates Circadian Expression, organism-specific biosystem; Circadian Clock, organism-specific biosystem; Defective AMN causes hereditary megaloblastic anemia 1, organism-specific biosystem; Defective BTD causes biotinidase deficiency, organism-specific biosystem; Defective CD320 causes methylmalonic aciduria, organism-specific biosystem; Defective CUBN causes hereditary megaloblastic anemia 1, organism-specific biosystem; Defective G
<b>Function</b>	cytokine activity; nicotinamide phosphoribosyltransferase activity; nicotinate-nucleotide diphosphorylase (carboxylating) activity; protein binding;