



## Human NMNAT1 peptide (DAG-P0982)

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

### PRODUCT INFORMATION

<b>Antigen Description</b>	This gene encodes an enzyme which catalyzes a key step in the biosynthesis of the coenzyme NAD. The encoded protein is one of several nicotinamide nucleotide adenyltransferases. Studies in Drosophila and mammalian neurons have shown the encoded protein can confer protection to damaged neurons. This protection requires enzymatic activity which increases NAD levels and activates a nuclear deacetylase which is the protective molecule. Pseudogenes of this gene are located on chromosomes 1, 3, 4, 14 and 15. [provided by RefSeq, Dec 2011]
<b>Specificity</b>	Widely expressed with highest levels in skeletal muscle, heart and kidney. Also expressed in the liver pancreas and placenta. Widely expressed throughout the brain.
<b>Conjugate</b>	Unconjugated
<b>Sequence Similarities</b>	Belongs to the eukaryotic NMN adenyltransferase 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="#">NMNAT1 nicotinamide nucleotide adenyltransferase 1 [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	NMNAT1
<b>Synonyms</b>	NMNAT1; nicotinamide nucleotide adenyltransferase 1; LCA9; NMNAT; PNAT1; nicotinamide mononucleotide adenyltransferase 1; NMN adenyltransferase 1; NaMN adenyltransferase 1; Lebers congenital amaurosis 9; pyridine nucleotide adenyltransferase 1; nicotinate-nucleotide adenyltransferase 1;

<b>Entrez Gene ID</b>	<a href="#">64802</a>
<b>mRNA Refseq</b>	<a href="#">NM_022787.3</a>
<b>Protein Refseq</b>	<a href="#">NP_073624.2</a>
<b>UniProt ID</b>	Q9HAN9
<b>Chromosome Location</b>	1p36.22
<b>Pathway</b>	Defective AMN causes hereditary megaloblastic anemia 1, organism-specific biosystem; Defective BTM 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 GIF causes intrinsic factor deficiency, organism-specific biosystem; Defective HLCS causes multiple carboxylase deficiency, organism-specific biosystem; Defective LMBRD1
<b>Function</b>	ATP binding; nicotinamide-nucleotide adenylyltransferase activity; nicotinamide-nucleotide adenylyltransferase activity; nicotinate-nucleotide adenylyltransferase activity; protein binding;