



## Human SETD2 peptide (DAG-P0742)

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

### PRODUCT INFORMATION

<b>Antigen Description</b>	Huntingtons disease (HD), a neurodegenerative disorder characterized by loss of striatal neurons, is caused by an expansion of a polyglutamine tract in the HD protein huntingtin. This gene encodes a protein belonging to a class of huntingtin interacting proteins characterized by WW motifs. This protein is a histone methyltransferase that is specific for lysine-36 of histone H3, and methylation of this residue is associated with active chromatin. This protein also contains a novel transcriptional activation domain and has been found associated with hyperphosphorylated RNA polymerase II. [provided by RefSeq, Aug 2008]
<b>Specificity</b>	Ubiquitously expressed.
<b>Conjugate</b>	Unconjugated
<b>Sequence Similarities</b>	Belongs to the histone-lysine methyltransferase family. SET2 subfamily. Contains 1 AWS domain. Contains 1 post-SET domain. Contains 1 SET domain. Contains 1 WW domain.
<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="#">SETD2 SET domain containing 2 [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	SETD2
<b>Synonyms</b>	SETD2; SET domain containing 2; HYPB; SET2; HIF-1; HIP-1; KMT3A; HBP231; HSPC069; p231HBP; histone-lysine N-methyltransferase SETD2; huntingtin yeast partner B; lysine N-methyltransferase 3A; huntingtin interacting protein 1; huntingtin-interacting protein B;

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<b>Entrez Gene ID</b>	<a href="#">29072</a>
<b>mRNA Refseq</b>	<a href="#">NM_014159.6</a>
<b>Protein Refseq</b>	<a href="#">NP_054878.5</a>
<b>UniProt ID</b>	Q9BYW2
<b>Chromosome Location</b>	3p21.31
<b>Pathway</b>	Lysine degradation, organism-specific biosystem; Lysine degradation, conserved biosystem;
<b>Function</b>	histone-lysine N-methyltransferase activity; histone-lysine N-methyltransferase activity; protein binding;

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