

**SETD1A Blocking Peptide(Center)**  
**Synthetic peptide**  
**Catalog # BP19714c****Specification****SETD1A Blocking Peptide(Center) - Product Information**

Primary Accession  
Other Accession

[O15047](#)  
[NP\\_055527.1](#)

**SETD1A Blocking Peptide(Center) - Additional Information****Gene ID** 9739**Other Names**

Histone-lysine N-methyltransferase SETD1A, Lysine N-methyltransferase 2F, SET domain-containing protein 1A, hSET1A, Set1/Ash2 histone methyltransferase complex subunit SET1, SETD1A

**Target/Specificity**

The synthetic peptide sequence is selected from aa 942-956 of HUMAN SETD1A

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**SETD1A Blocking Peptide(Center) - Protein Information****Name** SETD1A**Function**

Histone methyltransferase that catalyzes methyl group transfer from S-adenosyl-L-methionine to the epsilon-amino group of 'Lys-4' of histone H3 (H3K4) via a non-processive mechanism (PubMed:<a href="http://www.uniprot.org/citations/25561738" target="\_blank">25561738</a>, PubMed:<a href="http://www.uniprot.org/citations/12670868" target="\_blank">12670868</a>). Part of chromatin remodeling machinery, forms H3K4me1, H3K4me2 and H3K4me3 methylation marks at active chromatin sites where transcription and DNA repair take place (PubMed:<a href="http://www.uniprot.org/citations/29937342" target="\_blank">29937342</a>, PubMed:<a href="http://www.uniprot.org/citations/31197650" target="\_blank">31197650</a>, PubMed:<a href="http://www.uniprot.org/citations/32346159" target="\_blank">32346159</a>). Responsible for H3K4me3 enriched promoters and transcriptional programming of inner mass stem cells and neuron progenitors during embryogenesis (By similarity) (PubMed:<a href="http://www.uniprot.org/citations/31197650" target="\_blank">31197650</a>). Required for

H3K4me1 mark at stalled replication forks. Mediates FANCD2-dependent nucleosome remodeling and RAD51 nucleofilaments stabilization at reversed forks, protecting them from nucleolytic degradation (PubMed:<a href="http://www.uniprot.org/citations/29937342" target="\_blank">29937342</a>, PubMed:<a href="http://www.uniprot.org/citations/32346159" target="\_blank">32346159</a>). Does not methylate 'Lys-4' of histone H3 if the neighboring 'Lys-9' residue is already methylated (PubMed:<a href="http://www.uniprot.org/citations/12670868" target="\_blank">12670868</a>). Binds RNAs involved in RNA processing and the DNA damage response (PubMed:<a href="http://www.uniprot.org/citations/38003223" target="\_blank">38003223</a>).

### **Cellular Location**

Nucleus speckle. Chromosome Cytoplasm. Note=Localizes to a largely non-overlapping set of euchromatic nuclear speckles with SETD1B, suggesting that SETD1A and SETD1B each bind to a unique set of target genes (PubMed:17355966). Predominantly nuclear (PubMed:38003223)

### **SETD1A Blocking Peptide(Center) - Protocols**

Provided below are standard protocols that you may find useful for product applications.

- [Blocking Peptides](#)

### **SETD1A Blocking Peptide(Center) - Images**

### **SETD1A Blocking Peptide(Center) - Background**

SET1A is a component of a histone methyltransferase (HMT) complex that produces mono-, di-, and trimethylated histone H3 at Lys4. The complex is the analog of the *S. cerevisiae* Set1/COMPASS complex (Lee and Skalnik, 2005 [PubMed 16253997]). Also see SET1B (MIM 611055).

### **SETD1A Blocking Peptide(Center) - References**

- Wu, M., et al. Mol. Cell. Biol. 28(24):7337-7344(2008)  
Nguyen, P., et al. Mol. Cell. Biol. 28(21):6720-6729(2008)  
Ansari, K.I., et al. Biochim. Biophys. Acta 1779(1):66-73(2008)  
Lee, J.H., et al. Mol. Cell. Biol. 28(2):609-618(2008)  
Cho, Y.W., et al. J. Biol. Chem. 282(28):20395-20406(2007)