PRODUCT INFORMATION



SET7/9 (human recombinant)

Item No. 10320

Overview and Properties

KMT7, SETD7, SETD7/9, SET Domain-Containing Protein 7/9 Synonyms:

Source: Active recombinant N-terminal His-tagged SET7/9 purified from E. coli

Amino Acids: 1-366 (full-length)

Uniprot No.: Q8WTS6 Molecular Weight: 43.3 kDa

-80°C (as supplied) Storage:

Stability: ≥6 months

Purity: batch specific (≥95% estimated by SDS-PAGE)

Supplied in: 50 mM HEPES, pH 7.2, 100 mM sodium chloride, and 20% glycerol

Protein

Concentration: batch specific mg/ml

batch specific U/ml. Determined using 50 μM human TAF10 peptide (amino acids Activity:

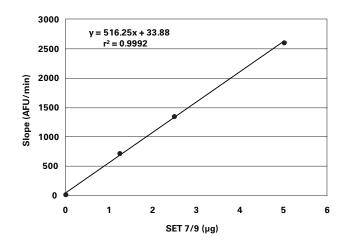
186-195) (Item No. 10228) (Ac)SKSKDRKYTL at 37°C using Cayman's Methyltransferase

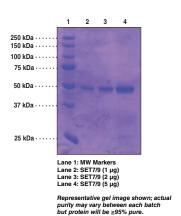
Colorometric Assay Kit (Item Number 700140).

batch specific nmoles/min/mg Specific Activity:

Information represents the product specifications. Batch specific analytical results are provided on each certificate of analysis.

Images





WARNING
THIS PRODUCT IS FOR RESEARCH ONLY - NOT FOR HUMAN OR VETERINARY DIAGNOSTIC OR THERAPEUTIC USE.

This material should be considered hazardous until further information becomes available. Do not ingest, inhale, get in eyes, on skin, or on clothing. Wash thoroughly after handling. Before use, the user must review the complete Safety Data Sheet, which has been sent via email to your institution.

WARRANTY AND LIMITATION OF REMEDY

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Description

Methylation of lysine can promote transcriptional activation or repression and is critical in regulating histone function. Lysine residues can be mono-, di-, or tri-methylated, and unlike most SET proteins, SET7/9 is exclusively a mono-methylase. SET7/9 methylates histone H3, tumor suppressor p53, and transcription factor TAF10. Recently, SET7/9 was shown to catalyze methylation of p53 in response to DNA damage thereby activating p53 for subsequent acetylation. SET7/9 is able to modulate p53 activity in a human cancer cell line, implying that it may play a significant role in human tumorigenesis.

References

- 1. Kurash, J.K., Lei, H., Shen, Q., et al. Methylation of p53 by Set7/9 mediates p53 acetylation and activity in vivo. Mol. Cell 29, 392-400 (2008).
- 2. Xiao, B., Jing, C., Wilson, J.R., et al. Structure and catalytic mechanism of the human histone methyltransferase SET7/9. *Nature* **421**, 652-656 (2003).
- 3. Couture, J.F., Collazo, E., Hauk, G., et al. Structural basis for the methylation site specificity of SET7/9. *Nat. Struct. Mol. Biol.* **13(2)**, 140-146 (2006).

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