

PRODUCT DATASHEET

EZH2 Trimer Complex

(Enhancer of Zeste Homolog 2 in complex with EED & SUZ12)

CATALOG NO.: HMT-23-501 LOT NO.:

DESCRIPTION: Human recombinant EZH2 (residues 2-746; Genbank Accession # NM_001203247; MW = 88.6 kDa) in complex with human recombinants EED (2-441; NM_003797; 53.9 kDa) and SUZ12 (2-739; NM_015355; 86.3 kDa). Total complex MW is 228.7 kDa. All proteins are full-length (residue 2 through C-terminus) and co-expressed in an insect cell/baculovirus expression system. The EED subunit incorporates an N-terminal Strep and Flag-tag and all others include an N-terminal His-tag. Catalyzes the transfer of methyl groups from S-adenosyl-L-methionine (SAM) to the ε-amino function of protein L-lysine residues, specifically lysine-27 of histone H3 (H3K27). During development, Polycomb Repressive Complex 2 (PRC2) is the principal methyltransferase responsible for generating trimethylated histone H3 lysine-27 (H3K27me3), an epigenetic mark essential for programmed repression of gene expression 1-5. EZH2, which includes a SET methyltransferase domain, is the catalytic subunit of PRC2^{1,6}. The core of the catalytic complex includes EZH2, EED, SUZ12 and RbAp48, while addition of AEBP2 significantly enhances the methyltransferase activity of the complex (>3x)6. EZH2 is overexpressed in a wide range of human cancers and its overexpression can correlate with tumor progression, increased metastasis and poor prognosis (see review⁷). Depletion of EZH2 and/or other PRC2 components can inhibit growth or induce apoptosis in cancer cells⁶, 8-10. Consequently, EZH2 is considered a promising target for the development of anti-cancer therapies¹¹.

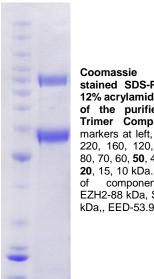
PURITY: >90% by SDS-PAGE.

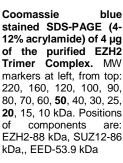
ASSAY CONDITIONS: RBC's EZH2 Trimer Complex displays histone methyltransferase activity at enzyme concentrations of 25.6 nM and above, 30°C, with HeLa oligo nucleosomes (0.05mg/mL), chicken core histones (0.05 mg/mL) 1µM rH3.3 histone or 0.5µM recombinant (H3-H4)₂ tetramer as TCA-precipitated counts in a scintillation/filter plate assay (Multiscreen FB, Topcount). Reaction conditions are: 50 mM Tris-HCl, pH 8.0, 50 mM NaCl, 1 mM EDTA, 1 mM DTT, 0.01% Brij30, with substrates at concentrations indicated above and [3H]-SAM.

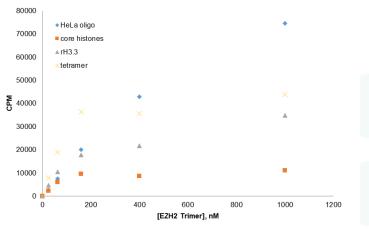
SUPPLIED AS: __ µM EZH2 Complex, as defined above, (__ µg/µl total protein) in 20 mM Tris-HCl, pH 8, 150 mM NaCl, 2 mM MgCl₂, 2 mM DTT, 20% glycerol (w/v), 0.01% NP-40 as determined by OD₂₈₀.

STORAGE: -70°C. Thaw quickly and store on ice before use. The remaining, unused, undiluted enzyme should be snap frozen, for example in a dry/ice ethanol bath or liquid nitrogen. Minimize freeze/thaws if possible, but very low volume aliquots (<5 µl) or storage of diluted enzyme is not recommended.

REFERENCES: 1) R. Cao et al. Science 2002 298 1039; 2) K. Plath et al. Science 2003 300 131; 3) J. Silva et al. Dev. Cell 2003 4 481; 4) S. Erhardt et al. Development 2003 130 4235; 5) R. Cao & Y. Zhang Curr. Opin. Genet. Dev. 2004 14 155; 6) R. Cao & Y. Zhang Mol. Cell 2004 15 57; 7) D.P.F. Tsang & A.S.L. Cheng J. Gastroenterol. Hepatol. 2011 26 19; 8) S. Varambally et al. Nature 2002 419 624; 9) A.P. Bracken EMBO J. 2003 22 5323; 10) J. Tan et al. Genes Dev. 2007 21 1050; 11) R.A. Copeland et al. Nature Rev. Drug Disc. 2009 8 724







Methyltransferase Activity of the EZH2 Trimer Complex. Methylation determined as TCA-precipitable counts in in a scintillation/filter plate assay. Reactions were 60 min., 30°C, 1 µM [3H]-SAM with 0.05mg/mL HeLa oligonucleosomes, 0.05mg/mL core histones, 1 µM histone H3 or 0.5µM tetramer.

This product is not intended for therapeutic or diagnostic use in animals or in humans.

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