

Screening Libraries

Proteins

Inhibitors

Product Data Sheet



HIST1H2BM Protein, Mouse (His)

Cat. No.: HY-P72225

Synonyms: H2bc14; Hist1h2bm; Histone H2B type 1-M; H2B 291B

Species: Source: E. coli

Accession: P10854 (P2-K126)

Gene ID: 319186

Molecular Weight: Approximately 17.8 kDa

PROPERTIES

AA Seq	uence
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PEPTKSAPAP KKGSKKAVTK AOKKDGKKRK RSRKESYSVY VYKVLKQVHP DTGISSKAMG IMNSFVNDIF ERIAGEASRL AHYNKRSTIT SREIQTAVRL LLPGELAKHA VSEGTKAVTK

YTSSK

Lyophilized powder. **Appearance**

Formulation Lyophilized from a 0.2 μm solution of Tris-based buffer, 50% Glycerol.

Endotoxin Level <1 EU/µg, determined by LAL method.

Reconsititution It is not recommended to reconstitute to a concentration less than 100 $\mu g/mL$ in ddH₂O.

Storage & Stability Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.

Shipping Room temperature in continental US; may vary elsewhere.

DESCRIPTION

Background

HIST1H2BM protein functions as a core component of the nucleosome, a crucial architectural unit that envelops and compacts DNA into chromatin, thereby limiting DNA accessibility to cellular machineries dependent on DNA templates. Playing a pivotal role in transcription regulation, DNA repair, DNA replication, and the maintenance of chromosomal stability, histones contribute significantly to cellular processes. The regulation of DNA accessibility involves a sophisticated system of post-translational modifications, collectively known as the histone code, and dynamic nucleosome remodeling. The nucleosome structure consists of a histone octamer, including two molecules each of H2A, H2B, H3, and H4, assembled into one H3-H4 heterotetramer and two H2A-H2B heterodimers. This octamer efficiently wraps approximately 147 base pairs of DNA, reflecting its central role in chromatin organization and genomic function.

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