

## 32-4767: Recombinant Human Sin3A-Associated Protein, 18kDa

**Alternative Name :** Histone deacetylase complex subunit SAP18, Sin3-associated polypeptide 18 kDa, Sin3-associated polypeptide p18, Cell growth-inhibiting protein 38, 2HOR0202, SAP18, GIG38.

### Description

Source : Escherichia Coli. SAP18 Human Recombinant fused with 20 amino acid His tag at N-terminus produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 173 amino acids (20- 172 a.a.) and having a molecular mass of 19.7 kDa. The SAP18 is purified by proprietary chromatographic techniques. SAP18 is a component of the histone deacetylase complex which has a significant role in the regulation of eukaryotic gene expression. SAP18 directly interacts with SIN3 and enhances SIN3-mediated transcriptional repression once bound to the promoter. SAP18 plays an important part in gene-specific recruitment of the HDAC complex by a number of transcription factors including Gli, GAGA, and Bicoid. Histone acetylation is significant in the regulation of eukaryotic gene expression. Histone acetylation and deacetylation is catalyzed by multisubunit complexes. SAP18 is a component of the histone deacetylase complex, which includes SIN3, SAP30, HDAC1, HDAC2, RbAp46, RbAp48, and other polypeptides. SAP18 is a component of a splicing-dependent multiprotein EJC (exon junction complex) placed at splice junction on mRNAs.

### Product Info

**Amount :** 20 µg  
**Purification :** Greater than 90.0% as determined by SDS-PAGE.  
**Content :** The SAP18 solution contains 20mM Tris-HCl buffer (pH8.0), 0.1M NaCl, 30% glycerol, 1mM DTT.  
**Storage condition :** Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid multiple freeze-thaw cycles.  
**Amino Acid :** MGSSHHHHHH SSGLVPRGSH MAVESRVTE EIKKEPEKPI DREKTCPLLL RVFTTNNGRH HRMDEFSGRN VPSSELQIYT WMDATLKELT SLVKEVYPEA RKKGTHFNFA IVFTDVKRPY YRVKEIGSTM SGRKGTDDSM TLQSQKFQIG DYLDIAITPP NRAPPSGRM RPY.

