

Full-Length

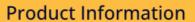
## EED (Human) Recombinant Protein (P01)

Catalog # H00008726-P01 Size 25 ug, 10 ug

## **Applications**



Specification	
Product Description	Human EED full-length ORF ( NP_003788.2, 1 a.a 441 a.a.) recombinant protein with GST-tag at N-terminal.
Sequence	MSEREVSTAPAGTDMPAAKKQKLSSDENSNPDLSGDENDDAVSIESGTNTERPDTPTNTPNAP GRKSWGKGKWKSKKCKYSFKCVNSLKEDHNQPLFGVQFNWHSKEGDPLVFATVGSNRVTLYE CHSQGEIRLLQSYVDADADENFYTCAWTYDSNTSHPLLAVAGSRGIIRIINPITMQCIKHYVGHGNAIN ELKFHPRDPNLLLSVSKDHALRLWNIQTDTLVAIFGGVEGHRDEVLSADYDLLGEKIMSCGMDHS LKLWRINSKRMMNAIKESYDYNPNKTNRPFISQKIHFPDFSTRDIHRNYVDCVRWLGDLILSKSCEN AIVCWKPGKMEDDIDKIKPSESNVTILGRFDYSQCDIWYMRFSMDFWQKMLALGNQVGKLYVWD LEVEDPHKAKCTTLTHHKCGAAIRQTSFSRDSSILIAVCDDASIWRWDRLR
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	76.6
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.





Note

Best use within three months from the date of receipt of this protein.

## Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — EED	
Entrez GenelD	<u>8726</u>
GeneBank Accession#	NM_003797.2
Protein Accession#	NP_003788.2
Gene Name	EED
Gene Alias	HEED, WAIT1
Gene Description	embryonic ectoderm development
Omim ID	605984
Gene Ontology	<u>Hyperlink</u>
Gene Summary	This gene encodes a member of the Polycomb-group (PcG) family. PcG family members form multimeric protein complexes, which are involved in maintaining the transcriptional repressive state of genes over successive cell generations. This protein interacts with enhancer of zeste 2, the cyto plasmic tail of integrin beta7, immunodeficiency virus type 1 (HIV-1) MA protein, and histone deac etylase proteins. This protein mediates repression of gene activity through histone deacetylation, and may act as a specific regulator of integrin function. Two transcript variants encoding distinct is oforms have been identified for this gene. [provided by RefSeq
Other Designations	WD protein associating with integrin cytoplasmic tails 1