

Product datasheet

Recombinant Human 4E-BP2 protein ab104667

[1 References](#) [1 Image](#)

Description

Product name	Recombinant Human 4E-BP2 protein
Purity	> 85 % SDS-PAGE. purified by using anion-exchange chromatography (DEAE sepharose resin) and gel-filtration chromatography (Sephacryl S-200) with 20mM Tris pH 7.5, 2mM EDTA.
Expression system	Escherichia coli
Accession	<u>Q13542</u>
Protein length	Full length protein
Animal free	No
Nature	Recombinant
Species	Human
Sequence	MGSSHHHHHHSSGLVPRGSHMSSSAGSGHQPSQSRAI PTRTVAISDAAQL PHDYCTTPGGTLFSTTPGGTRIYDRKFLLDRRNSPMAQTP PCHLPNIPG VTSPGTLIEDSKVEVNNLNNLNNHDRKHAVGDDAQFEMDI
Predicted molecular weight	15 kDa including tags
Amino acids	1 to 120
Tags	His tag N-Terminus

Specifications

Our **Abpromise guarantee** covers the use of **ab104667** in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Applications	SDS-PAGE Mass Spectrometry
Mass spectrometry	MALDI-TOF
Form	Liquid

Preparation and Storage

Stability and Storage	Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw
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cycles.

pH: 8.00

Constituents: 0.0154% DTT, 0.316% Tris HCl, 10% Glycerol (glycerin, glycerine), 0.58% Sodium chloride

General Info

Function

Repressor of translation initiation involved in synaptic plasticity, learning and memory formation (By similarity). Regulates EIF4E activity by preventing its assembly into the eIF4F complex: hypophosphorylated form of EIF4EBP2 competes with EIF4G1/EIF4G3 and strongly binds to EIF4E, leading to repress translation. In contrast, hyperphosphorylated form dissociates from EIF4E, allowing interaction between EIF4G1/EIF4G3 and EIF4E, leading to initiation of translation (PubMed:25533957). EIF4EBP2 is enriched in brain and acts as a regulator of synapse activity and neuronal stem cell renewal via its ability to repress translation initiation (By similarity). Mediates the regulation of protein translation by hormones, growth factors and other stimuli that signal through the MAP kinase and mTORC1 pathways.

Sequence similarities

Belongs to the eIF4E-binding protein family.

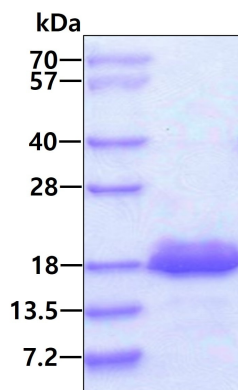
Domain

The TOS motif mediates interaction with RPTOR, leading to promote phosphorylation by mTORC1 complex. Intrinsically disordered protein that undergoes folding upon phosphorylation (PubMed:25533957). Hypophosphorylated form interacts strongly with EIF4E using (1) the YXXXXLPhi motif, that undergoes a disorder-to-helix transition upon binding and (2) the secondary EIF4E binding sites (residues 78-82) (PubMed:24207126, PubMed:25533957). Phosphorylation at Thr-37 and Thr-46 induces folding of region encompassing residues from Pro-18 to Arg-62 of into a four-stranded beta-domain that sequesters the helical YXXXXLPhi motif into a buried beta-strand, blocking accessibility to EIF4E. Protein phosphorylated at Thr-37 and Thr-46 is however unstable and subsequent phosphorylation at Ser-65, Thr-70 and Ser-83 is required to stabilize the fold, decreasing affinity for EIF4E by a factor of 4000 (PubMed:24207126, PubMed:25533957).

Post-translational modifications

Phosphorylation at Thr-37, Thr-46, Ser-65, Thr-70 and Ser-83 is mediated by MTOR and corresponds to the hyperphosphorylated form: it abolishes binding to EIF4E by inducing folding of intrinsically disordered regions (PubMed:24207126, PubMed:25533957). First phosphorylated at Thr-37 and Thr-46 by MTOR, inducing folding of region encompassing residues from Pro-18 to Arg-62 of into a four-stranded beta-domain that sequesters the helical YXXXXLPhi motif into a partly buried beta-strand, blocking accessibility to EIF4E. Protein phosphorylated at Thr-37 and Thr-46 is however unstable and subsequent phosphorylation at Ser-65, Thr-70 and Ser-83 is required to stabilize the fold, decreasing affinity for EIF4E by a factor of 4000 (PubMed:24207126, PubMed:25533957). Phosphorylated in response to insulin, EGF and PDGF. Deamidated at Asn-99 and Asn-102 to aspartate (Asp) in brain. Deamidation promotes interaction with RPTOR, subsequent phosphorylation by mTORC1 and increased translation, leading to impair kinetics of excitatory synaptic transmission. Deamidation takes place during postnatal development, when the PI3K-Akt-mTOR signaling is reduced, suggesting it acts as a compensatory mechanism to promote translation despite attenuated PI3K-Akt-mTOR signaling in neuron development. Deamidation converts Asn residues into a mixture of Asp and isoaspartate; interactions with PCMT1 is required to prevent isoaspartate accumulation and convert isoaspartate to Asp.

Images



SDS-PAGE of ab104667 (3 μ g) under reducing condition and visualized by coomassie blue stain.

SDS-PAGE - Recombinant Human 4E-BP2 protein
(ab104667)

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