eRF1 (B-11): sc-365686



The Power to Question

BACKGROUND

Translation is carried out by the ribosome and several associated protein factors through three consecutive steps: initiation, elongation and termination. Termination of protein synthesis takes place when the ribosomal A site is occupied simultaneously by one of three stop codons and by a class 1 translation termination factor. In eukaryotes, this termination factor is the eukaryotic release factor 1 (eRF1), a protein that promotes hydrolysis of the last peptidyl-tRNA on the ribosome. eRF1 activity is stimulated by the association with the GTP-binding protein eRF3. eRF1 forms a quaternary complex with eRF3, GTP and the ribosome. This complex performs a dual role, where, in the "GTP state", it controls the positioning of eRF1 toward the stop codon and peptidyl-tRNA, and, in the "GDP state", it promotes the release of the eRFs from the ribosome. eRF1 contains a highly conserved Asn-Ile-Lys-Ser (NIKS) tetrapeptide, which is essential for the interaction of eRF1 with the ribosome. The gene encoding human eRF1 maps to chromosome 5q31.2.

REFERENCES

- Frolova, L., et al. 1996. Eukaryotic polypeptide chain release factor eRF3 is an eRF1 and ribosome dependent guanosine triphosphatase. RNA 2: 334-341.
- Le Goff, X., et al. 1997. Overexpression of human release factor 1 alone has an antisuppressor effect in human cells. Mol. Cell. Biol. 17: 3164-3172.

CHROMOSOMAL LOCATION

Genetic locus: ETF1 (human) mapping to 5q31.2; Etf1 (mouse) mapping to 18 B1.

SOURCE

eRF1 (B-11) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 411-437 at the C-terminus of eRF1 of human origin.

PRODUCT

Each vial contains 200 $\mu g \ lgG_{2a}$ kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

eRF1 (B-11) is available conjugated to agarose (sc-365686 AC), 500 μ g/0.25 ml agarose in 1 ml, for IP; to HRP (sc-365686 HRP), 200 μ g/ml, for WB, IHC(P) and ELISA; to either phycoerythrin (sc-365686 PE), fluorescein (sc-365686 FITC), Alexa Fluor* 488 (sc-365686 AF488), Alexa Fluor* 546 (sc-365686 AF546), Alexa Fluor* 594 (sc-365686 AF594) or Alexa Fluor* 647 (sc-365686 AF647), 200 μ g/ml, for WB (RGB), IF, IHC(P) and FCM; and to either Alexa Fluor* 680 (sc-365686 AF680) or Alexa Fluor* 790 (sc-365686 AF790), 200 μ g/ml, for Near-Infrared (NIR) WB, IF and FCM.

Blocking peptide available for competition studies, sc-365686 P, (100 μ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

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STORAGE

Store at 4° C, **DO NOT FREEZE**. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

APPLICATIONS

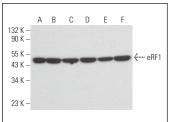
eRF1 (B-11) is recommended for detection of eRF1 of mouse, rat and human origin by Western Blotting (starting dilution 1:100, dilution range 1:100-1:1000), immunoprecipitation [1-2 μ g per 100-500 μ g of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

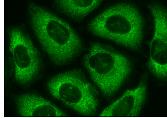
Suitable for use as control antibody for eRF1 siRNA (h): sc-37871, eRF1 siRNA (m): sc-37872, eRF1 shRNA Plasmid (h): sc-37871-SH, eRF1 shRNA Plasmid (m): sc-37872-SH, eRF1 shRNA (h) Lentiviral Particles: sc-37871-V and eRF1 shRNA (m) Lentiviral Particles: sc-37872-V.

Molecular Weight of eRF1: 50 kDa.

Positive Controls: SK-BR-3 cell lysate: sc-2218, MDA-MB-231 cell lysate: sc-2232 or MCF7 whole cell lysate: sc-2206.

DATA





eRF1 (B-11): sc-365686. Western blot analysis of eRF1 expression in MCF7 (**A**), MDA-MB-231 (**B**), SK-BR-3 (**C**), MH-S (**D**), NTERA-2 cl.D1 (**E**) and COLO 205 (**F**) whole

eRF1 (B-11): sc-365686. Immunofluorescence staining of methanol-fixed HeLa cells showing cytoplasmic localization.

SELECT PRODUCT CITATIONS

- Ortega, J.A., et al. 2020. Nucleocytoplasmic proteomic analysis uncovers eRF1 and nonsense-mediated decay as modifiers of ALS/FTD C9orf72 toxicity. Neuron 106: 90-107.e13.
- Sharma, J., et al. 2021. A small molecule that induces translational readthrough of CFTR nonsense mutations by eRF1 depletion. Nat. Commun. 12: 4358.
- Oltion, K., et al. 2023. An E3 ligase network engages GCN1 to promote the degradation of translation factors on stalled ribosomes. Cell 186: 346-362 e17
- 4. Müller, M.B.D., et al. 2023. Mechanisms of readthrough mitigation reveal principles of GCN1-mediated translational quality control. Cell 186: 3227-3244.e20.
- Gurzeler, L.A., et al. 2023. Drug-induced eRF1 degradation promotes readthrough and reveals a new branch of ribosome quality control. Cell Rep. 42: 113056.

RESEARCH USE

For research use only, not for use in diagnostic procedures.