abcam

Product datasheet

Recombinant Human SF2 protein ab82026

製品の詳細

製品名 Recombinant Human SF2 protein

生理活性 1 unit equals 1 nanogram of purified protein.

精製度 > 95 % SDS-PAGE.

purified by affinity column

発現系 Escherichia coli

タンパク質長 Full length protein

Animal free No.

由来 Recombinant

生物種 Human

特性

Our **Abpromise guarantee** covers the use of **ab82026** in the following tested applications.

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

アプリケーション Gel Supershift Assays

SDS-PAGE

EMSA

製品の状態 Liquid

備考 1 unit equals 1 nanogram of purified protein.

前処理および保存

保存方法および安定性 Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 7.90

Constituents: 0.55% Ammonium sulphate, 0.0077% DTT, 0.476% HEPES, 0.00584% EDTA,

20% Glycerol (glycerin, glycerine)

関連情報

機能 Plays a role in preventing exon skipping, ensuring the accuracy of splicing and regulating

alternative splicing. Interacts with other spliceosomal components, via the RS domains, to form a

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bridge between the 5'- and 3'-splice site binding components, U1 snRNP and U2AF. Can stimulate binding of U1 snRNP to a 5'-splice site-containing pre-mRNA. Binds to purine-rich RNA sequences, either the octamer, 5'-RGAAGAAC-3' (r=A or G) or the decamers,

AGGACAGAGC/AGGACGAAGC. Binds preferentially to the 5'-CGAGGCG-3' motif in vitro. Three copies of the octamer constitute a powerful splicing enhancer in vitro, the ASF/SF2 splicing enhancer (ASE) which can specifically activate ASE-dependent splicing. Isoform ASF-2 and isoform ASF-3 act as splicing repressors.

配列類似性 Belongs to the splicing factor SR family.

Contains 2 RRM (RNA recognition motif) domains.

The RRM 2 domain plays an important role in governing both the binding mode and the phosphorylation mechanism of the RS domain by SRPK1. RS domain and RRM 2 are uniquely positioned to initiate a highly directional (C-terminus to N-terminus) phosphorylation reaction in which the RS domain slides through an extended electronegative channel separating the docking groove of SRPK1 and the active site. RRM 2 binds toward the periphery of the active site and guides the directional phosphorylation mechanism. Both the RS domain and an RRM domain are required for nucleocytoplasmic shuttling.

Phosphorylated by CLK1, CLK2, CLK3 and CLK4. Phosphorylated by SRPK1 at multiple serines in its RS domain via a directional (C-terminal to N-terminal) and a dual-track mechanism incorporating both processive phosphorylation (in which the kinase stays attached to the substrate after each round of phosphorylation) and distributive phosphorylation steps (in which the kinase and substrate dissociate after each phosphorylation event). The RS domain of SRSF1 binds to a docking groove in the large lobe of the kinase domain of SRPK1 and this induces certain structural changes in SRPK1 and/or RRM 2 domain of SRSF1, allowing RRM 2 to bind the kinase and initiate phosphorylation. The cycles continue for several phosphorylation steps in a processive manner (steps 1-8) until the last few phosphorylation steps (approximately steps 9-12). During that time, a mechanical stress induces the unfolding of the beta-4 motif in RRM 2, which

which facilitates SRSF1 dissociation after phosphorylation is completed. Arg-97 is dimethylated, probably to asymmetric dimethylarginine.

Cytoplasm. Nucleus speckle. In nuclear speckles. Shuttles between the nucleus and the

then docks at the docking groove of SRPK1. This also signals RRM 2 to begin to dissociate,

cytoplasm.

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