

Product datasheet

Recombinant Human U2AF35/U2AF1 protein (denatured) ab156324

1 Image

Description

Product name	Recombinant Human U2AF35/U2AF1 protein (denatured)
Purity	> 90 % SDS-PAGE.
Expression system	Escherichia coli
Accession	Q01081-2
Protein length	Full length protein
Animal free	No
Nature	Recombinant
Species	Human
Sequence	<p>MGSSHHHHHH SSGLVPRGSH MGSMAYELAS IFGTEKDKVN CSFYFKIGAC RHGDRC SRLH NKPTFSQTL IQNIYRNPQN SAQTADGSHC AVSDVEMQEH YDEFFEEVFT EMEEKYGEVE EMNVCDNLGD HLVGNVYVKF RREEDAEKAV IDLNNRWFNG QPIHAELSPV TDFREACCRQ YEMGECTRGG FCNFMHLKPI SRELRRRELYG RRRKKHRSRS RSRERRRSR DRGRGGGGGG GGGGGGRERD RRRSRDRERS GRF</p>
Predicted molecular weight	30 kDa including tags
Amino acids	1 to 240
Tags	His tag N-Terminus

Specifications

Our [Abpromise guarantee](#) covers the use of **ab156324** 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
Form	Liquid
Additional notes	This product was previously labelled as U2AF35

Preparation and Storage

Stability and Storage

Shipped at 4°C. Store at +4°C short term (1-2 weeks). Upon delivery aliquot. Store at -20°C or -80°C. Avoid freeze / thaw cycle.

pH: 8.00

Constituents: 2.4% Urea, 0.32% Tris HCl, 10% Glycerol (glycerin, glycerine)

General Info

Function

Plays a critical role in both constitutive and enhancer-dependent splicing by mediating protein-protein interactions and protein-RNA interactions required for accurate 3'-splice site selection. Recruits U2 snRNP to the branch point. Directly mediates interactions between U2AF2 and proteins bound to the enhancers and thus may function as a bridge between U2AF2 and the enhancer complex to recruit it to the adjacent intron.

Sequence similarities

Belongs to the splicing factor SR family.

Contains 2 C3H1-type zinc fingers.

Contains 1 RRM (RNA recognition motif) domain.

Domain

The C-terminal SR-rich domain is required for interactions with SR proteins and the splicing regulators TRA and TRA2, and the N-terminal domain is required for formation of the U2AF1/U2AF2 heterodimer.

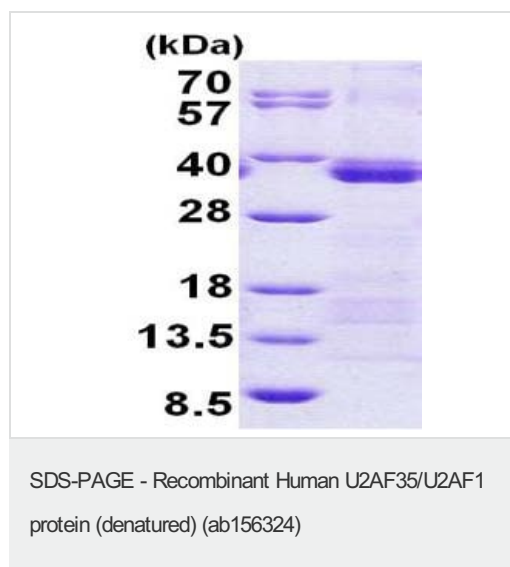
Post-translational modifications

Phosphorylated upon DNA damage, probably by ATM or ATR.

Cellular localization

Nucleus. Nucleus speckle.

Images



15% SDS-PAGE analysis of ab156324 (3µg).

Please note: All products are "FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES"

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