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PARN Protein (AA 1-639, full length) (His tag)



Image



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Overview

Quantity:	100 μg
Target:	PARN
Protein Characteristics:	AA 1-639, full length
Origin:	Human
Source:	Yeast
Protein Type:	Recombinant
Purification tag / Conjugate:	This PARN protein is labelled with His tag.
Application:	SDS-PAGE (SDS)

Product Details

Sequence:

MEIIRSNFKS NLHKVYQAIE EADFFAIDGE FSGISDGPSV SALTNGFDTP EERYQKLKKH
SMDFLLFQFG LCTFKYDYTD SKYITKSFNF YVFPKPFNRS SPDVKFVCQS SSIDFLASQG
FDFNKVFRNG IPYLNQEEER QLREQYDEKR SQANGAGALS YVSPNTSKCP VTIPEDQKKF
IDQVVEKIED LLQSEENKNL DLEPCTGFQR KLIYQTLSWK YPKGIHVETL ETEKKERYIV
ISKVDEEERK RREQQKHAKE QEELNDAVGF SRVIHAIANS GKLVIGHNML LDVMHTVHQF
YCPLPADLSE FKEMTTCVFP RLLDTKLMAS TQPFKDIINN TSLAELEKRL KETPFNPPKV
ESAEGFPSYD TASEQLHEAG YDAYITGLCF ISMANYLGSF LSPPKIHVSA RSKLIEPFFN
KLFLMRVMDI PYLNLEGPDL QPKRDHVLHV TFPKEWKTSD LYQLFSAFGN IQISWIDDTS
AFVSLSQPEQ VKIAVNTSKY AESYRIQTYA EYMGRKQEEK QIKRKWTEDS WKEADSKRLN
PQCIPYTLQN HYYRNNSFTA PSTVGKRNLS PSQEEAGLED GVSGEISDTE LEQTDSCAEP
LSEGRKKAKK LKRMKKELSP AGSISKNSPA TLFEVPDTW

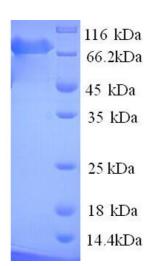
Purification: SDS-PAGE

Product Details > 90 % Purity: **Target Details** Target: **PARN** PARN (PARN Products) Alternative Name: Background: 3'-exoribonuclease that has a preference for poly(A) tails of mRNAs, thereby efficiently degrading poly(A) tails. Exonucleolytic degradation of the poly(A) tail is often the first step in the decay of eukaryotic mRNAs and is also used to silence certain maternal mRNAs translationally during oocyte maturation and early bryonic development. Interacts with both the 3'-end poly(A) tail and the 5'-end cap structure during degradation, the interaction with the cap structure being required for an efficient degradation of poly(A) tails. Involved in nonsense-mediated mRNA decay, a critical process of selective degradation of mRNAs that contain prature stop codons. Also involved in degradation of inherently unstable mRNAs that contain AU-rich elents (AREs) in their 3'-UTR, possibly via its interaction with KHSRP. Probably mediates the roval of poly(A) tails of AREs mRNAs, which constitutes the first step of destabilization. 75.4 kDa Molecular Weight: UniProt: 095453 **Application Details**

Application Notes:	Optimal working dilution should be determined by the investigator.
Restrictions:	For Research Use only

Handling

Format:	Liquid
Concentration:	0.1-2 mg/mL
Buffer:	20 mM Tris-HCl based buffer, pH 8.0
Storage:	-80 °C,4 °C,-20 °C
Storage Comment:	Store at -20°C, for extended storage, conserve at -20°C or -80°C. Repeated freezing and thawing is not recommended. Store working aliquots at 4°C for up to one week.



SDS-PAGE

Image 1.