

PARN Protein, Human, Recombinant (His)

General Information

Synonyms:	Poly(A)-specific ribonuclease PARN;DAN;PARN;Deadenylation nuclease;Deadenylating nuclease;Polyadenylate-specific ribonuclease
Protein Construction:	1-639 aa
Species:	Human
Expression Host:	E. coli
Accession:	O95453
Molecular Weight:	77.5 kDa (predicted)
AA Sequence:	MEIIRSNFKSNLHKVYQAIIEADFFAIDGEGFSGISDGPSVSALTNGFDTPPEERYQKLKHKHSMDFLLFQFGLCTFK YDYTDSKYITKSFNFYVFPKPFNRSSPDVKFVCQSSSIDFLASQGFDFNKVFRNGIPYLNQEEERQLREQYDEK RSQANGAGALSYVSPNTSKCPVTIPEDQKKFIDQVVEKIEDLLQSEENKNLDLEPCTGFQRKLIYQTLQSWKYPK GIHVETLETEKKERYIVISKVDEEERKRREQQKHAKQEELNDAVGFSRVIHAIANSGKLVIGHNMLLDVMHTV HQFYCPLPADLSEFKEMTTCVFPRLLDTKLMASQPFKDIINNTSLAELEKRLKETPFNPPKVESAEGFPSYDTA SEQLHEAGYDAYITGLCFISMANYLGSFLSPPKIHVSARSKLIEPFNKLFMRVMDIPYLNLEGPDLQPKRDHV LHVTFPKEWKTSDLYQLFSAFGNIQISWIDDTSAFVLSQPEQVKIAVNTSKYAESYRIQTYAEYMGRKQEEKQ IKRKWTEDSWKEADSKRLNPQCIPYTLQNHYYRNNSTAPSTVGKRNLSPSQEEAGLEDGVSGEISDTELEQT DSCAEPLSEGRKKAKKLRMKKELSPAGSISKNSPATLFEVPDTW

QC Testing

Biological Activity:	Activity has not been tested. It is theoretically active, but we cannot guarantee it. If you require protein activity, we recommend choosing the eukaryotic expression version first.
Purity:	> 90% as determined by SDS-PAGE.
Endotoxin:	< 1.0 EU/μg of the protein as determined by the LAL method.
Formulation:	Tris-based buffer, 50% glycerol

Preparation and Storage

Reconstitution:

A Certificate of Analysis (CoA) containing reconstitution instructions is included with the products. Please refer to the CoA for detailed information.

Stability & Storage:

Lyophilized powders can be stably stored for over 12 months, while liquid products can be stored for 6-12 months at -80°C. For reconstituted protein solutions, the solution can be stored at -20°C to -80°C for at least 3 months. Please avoid multiple freeze-thaw cycles and store products in aliquots.

Actual storage temperature shall be subject to the COA.

Shipping:

In general, lyophilized powders are shipped with blue ice, while solutions are shipped with dry ice.

Protein 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 embryonic 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 premature stop codons. Also involved in degradation of inherently unstable mRNAs that contain AU-rich elements (AREs) in their 3'-UTR, possibly via its interaction with KHSRP. Probably mediates the removal of poly(A) tails of AREs mRNAs, which constitutes the first step of destabilization. Also able to recognize and trim poly(A) tails of microRNAs such as MIR21 and H/ACA box snoRNAs (small nucleolar RNAs) leading to microRNAs degradation or snoRNA increased stability.

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