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Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

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Datasheet

RNPC3 (Human) Recombinant Protein (P01)

RNP

Catalog Number: H00055599-P01**Regulation Status:** For research use only (RUO)**Product Description:** Human RNPC3 full-length ORF (AAH10697, 1 a.a. - 261 a.a.) recombinant protein with GST-tag at N-terminal.**Sequence:**

MNKLMEANLQPKRPKTIKQRHVRKKRKKIKDMLNTPLC
PSHSSLHPVLLPSDVFDQPPVGNKRIEFHISTDMPAA
FKKDLEKEQNCEEKNHDLPEVDASNIGFGKIFPKPN
LDITEEIKEDSDEMPSECISRRELEKGRISREEMETLSV
FRSYEPGEPNCRIYVKNLAKHVQEKDLKYIFGRYVDFS
SETQRIMFDIRLMKEGRMKGQAFIGLPNEKAAAKALKE
ANGYVLFKGPMVVQFARSARPKQDPKEGKRKC

Host: Wheat Germ (in vitro)**Theoretical MW (kDa):** 54.45**Interspecies Antigen Sequence:** Mouse (87); Rat (87)**Applications:** AP, Array, ELISA, WB-Re

(See our web site product page for detailed applications information)

Protocols: See our web site at<http://www.abnova.com/support/protocols.asp> or product page for detailed protocols**Preparation Method:** [in vitro wheat germ expression system](#)**Purification:** Glutathione Sepharose 4 Fast Flow**Storage Buffer:** 50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.**Storage Instruction:** Store at -80°C. Aliquot to avoid repeated freezing and thawing.**Entrez GeneID:** 55599**Gene Symbol:** RNPC3**Gene Alias:** FLJ20008, FLJ25070, KIAA183, RBM40,

Gene Summary: Two types of spliceosomes catalyze splicing of pre-mRNAs. The major U2-type spliceosome is found in all eukaryotes and removes U2-type introns, which represent more than 99% of pre-mRNA introns. The minor U12-type spliceosome is found in some eukaryotes and removes U12-type introns, which are rare and have distinct splice consensus signals. The U12-type spliceosome consists of several small nuclear RNAs and associated proteins. This gene encodes a 65K protein that is a component of the U12-type spliceosome. This protein contains two RNA recognition motifs (RRMs), suggesting that it may contact one of the small nuclear RNAs of the minor spliceosome. [provided by RefSeq]