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Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

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Datasheet

RPL35A (Human) Recombinant Protein (P02)

Catalog Number: H00006165-P02

Regulation Status: For research use only (RUO)

Product Description: Human RPL35A full-length ORF (NP_000987.2, 1 a.a. - 110 a.a.) recombinant protein with GST-tag at N-terminal.

Sequence:

MSGRLWSKAIFAGYKRGLRNQREHTALLKIEGVYARD
ETEFYLGKRCAYVYKAKNNTVTPGGKPNKTRVIWGKV
TRAHGNSGMVRAKFRSNLPAKAIGHRIRVMLYPSRI

Host: Wheat Germ (in vitro)

Theoretical MW (kDa): 38.9

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: 6165

Gene Symbol: RPL35A

Gene Alias: DBA5

Gene Summary: Ribosomes, the organelles that catalyze protein synthesis, consist of a small 40S subunit and a large 60S subunit. Together these subunits are composed of 4 RNA species and approximately 80 structurally distinct proteins. This gene

encodes a ribosomal protein that is a component of the 60S subunit. The protein belongs to the L35AE family of ribosomal proteins. It is located in the cytoplasm. The rat protein has been shown to bind to both initiator and elongator tRNAs, and thus, it is located at the P site, or P and A sites, of the ribosome. Although this gene was originally mapped to chromosome 18, it has been established that it is located at 3q29-qter. Transcript variants utilizing alternative transcription initiation sites and alternative polyA signals exist. As is typical for genes encoding ribosomal proteins, there are multiple processed pseudogenes of this gene dispersed through the genome. [provided by RefSeq]