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Lieferung & Zahlungsart

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Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

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Datasheet

MRPL47 (Human) Recombinant Protein (P01)

Catalog Number: H00057129-P01

Regulation Status: For research use only (RUO)

Product Description: Human MRPL47 full-length ORF (NP_817125.1, 1 a.a. - 140 a.a.) recombinant protein with GST-tag at N-terminal.

Sequence:

MLLTLEQEAKRQRLPMPSPERLDKVVDSMDALDKVVQ
EREDALRLLQTGQERARPGAWRRDIFGRIIWHKFKQW
VIPWHLNKRYNRKRFFALPYVDHFLRLEREKRARIKAR
KENLERKKAKILLKKFPHLAEAQKSSLV

Host: Wheat Germ (in vitro)

Theoretical MW (kDa): 43.4

Interspecies Antigen Sequence: Mouse (80); Rat (80)

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: 57129

Gene Symbol: MRPL47

Gene Alias: CGI-204, MGC45403, NCM1

Gene Summary: Mammalian mitochondrial ribosomal proteins are encoded by nuclear genes and help in

protein synthesis within the mitochondrion. Mitochondrial ribosomes (mitoribosomes) consist of a small 28S subunit and a large 39S subunit. They have an estimated 75% protein to rRNA composition compared to prokaryotic ribosomes, where this ratio is reversed. Another difference between mammalian mitoribosomes and prokaryotic ribosomes is that the latter contain a 5S rRNA. Among different species, the proteins comprising the mitoribosome differ greatly in sequence, and sometimes in biochemical properties, which prevents easy recognition by sequence homology. This gene encodes a 39S subunit protein. This gene is immediately adjacent to the gene for BAF complex 53 kDa subunit protein a (BAF53a), in a tail-to-tail orientation. Two transcript variants encoding different protein isoforms have been identified. [provided by RefSeq]