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Datasheet

RPSA (Human) Recombinant Protein (Q01)

Catalog Number: H00003921-Q01

Regulation Status: For research use only (RUO)

Product Description: Human RPSA partial ORF (NP_002286, 196 a.a. - 295 a.a.) recombinant protein with GST-tag at N-terminal.

Sequence:

EVMPDLYFYRDPEEIEKEEQAAAEKAVTKEEFQGEWT APAPEFTATQPEVADWSEGVQVPSVPIQQFPTEDWS AQPATEDWSAAPTAQATEWVGATTDWS

Host: Wheat Germ (in vitro)

Theoretical MW (kDa): 36.74

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 GenelD: 3921

Gene Symbol: RPSA

Gene Alias: 37LRP, 67LR, LAMBR, LAMR1, LRP, p40

Gene Summary: Laminins, a family of extracellular matrix glycoproteins, are the major noncollagenous constituent of basement membranes. They have been implicated in a wide variety of biological processes including cell adhesion, differentiation, migration, signaling, neurite outgrowth and metastasis. Many of the effects of laminin are mediated through interactions with cell surface receptors. These receptors include members of the integrin family, as well as non-integrin laminin-binding proteins. This gene encodes а high-affinity, non-integrin family, laminin receptor 1. This receptor has been variously called 67 kD laminin receptor, 37 kD laminin receptor precursor (37LRP) and p40 ribosome-associated protein. The amino acid sequence of laminin receptor 1 is highly conserved through evolution, suggesting a key biological function. It has been observed that the level of the laminin receptor transcript is higher in colon carcinoma tissue and lung cancer cell line than their normal counterparts. Also, there is a correlation between the upregulation of this polypeptide in cancer cells and their invasive and metastatic phenotype. Multiple copies of this gene exist, however, most of them are pseudogenes thought to arisen from retropositional have events. Two alternatively spliced transcript variants encoding the same protein have been found for this gene. [provided by RefSeq]