

Enzyme catalysed Human Poly-ubiquitin (n2-7; K63-linked) Protein

Cat. No. UBA52-02H **Lot. No.** (See product label)

SPECIFICATION

Product Overview	Enzyme catalysed Human Poly-ubiquitin (n2-7; K63-linked) Protein
Species	Human
Source	Enzyme catalysed
Description	Ubiquitin is a highly conserved nuclear and cytoplasmic protein that has a major role in targeting cellular proteins for degradation by the 26S proteasome. It is also involved in the maintenance of chromatin structure, the regulation of gene expression, and the stress response. Ubiquitin is synthesized as a precursor protein consisting of either polyubiquitin chains or a single ubiquitin moiety fused to an unrelated protein. This gene encodes a fusion protein consisting of ubiquitin at the N terminus and ribosomal protein L40 at the C terminus, a C-terminal extension protein (CEP). Multiple processed pseudogenes derived from this gene are present in the genome.
Molecular Mass	various
AA Sequence	MQIFVKLTGKTTITLEVEPSDTIENVKAKIQDKEGIPPDQQRLIFAGKQLEDGRTLSDY NIQKESTLHLVLRRLRGG
Purity	>98% by InstantBlue SDS-PAGE
Stability	12 months at -70 centigrade; aliquot as required

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 Email: info@creative-biomart.com  Fax: 1-631-938-8127

 45-1 Ramsey Road, Shirley, NY 11967, USA

Storage	At -70 centigrade.
Concentration	5 mg/mL
Storage Buffer	50 mM HEPES pH 7.5, 150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

GENE INFORMATION

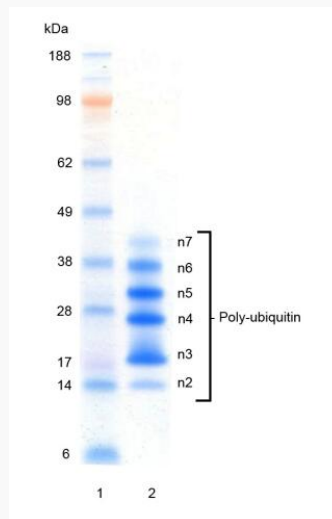
Gene Name	UBA52 ubiquitin A-52 residue ribosomal protein fusion product 1 [Homo sapiens (human)]
Official Symbol	UBA52
Synonyms	UBA52; ubiquitin A-52 residue ribosomal protein fusion product 1; L40; CEP52; RPL40; HUBCEP52; ubiquitin-60S ribosomal protein L40; ubiquitin carboxyl extension protein 52; ubiquitin-52 amino acid fusion protein; ubiquitin-CEP52
Gene ID	7311
mRNA Refseq	NM_003333
Protein Refseq	NP_003324
MIM	191321
UniProt ID	P62987

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Purity



4-12% gradient SDS-PAGE InstantBlue staining

Lane 1: MW markers

Lane 2: 4 µg Poly-ubiquitin (K63-linked)

Protein Sequence

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MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n2 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n3 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n4 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n5 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n6 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
n7 MQIFVKLTGKTTITLEVPSDTIENVKAKIQDKEGIPPDQQLIFAGKQLEDGRTLSDYNIQKESTLHLVLRGG
    
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