UBE2B (HR6B) [untagged]

E2 – Ubiquitin Conjugating Enzyme

Alternate Names: HHR6B, HR6B, RAD6B, Ubiquitin carrier protein B, Ubiquitin protein ligase B

Cat. No. 62-0004-020 Quantity: **Lot. No. 1456** Storage:

FOR RESEARCH USE ONLY NOT FOR USE IN HUMANS



CERTIFICATE OF ANALYSIS Page 1 of 2

Background

The enzymes of the ubiquitylation pathway play a pivotal role in a number of cellular processes including regulated and targeted proteasomal degradation of substrate proteins. Three classes of enzymes are involved in the process of ubiquitylation; activating enzymes (E1s), conjugating enzymes (E2s) and protein ligases (E3s). UBE2B is a member of the E2 ubiquitin-conjugating enzyme family and cloning of the human gene was first described by Koken et al. (1991). UBE2B shares 70% identity with its veast homologue but lacks the acidic C-terminal domain. The ring finger proteins RAD5 and RAD18 interact with UBE2B and other members of the RAD6 pathway (Notenboom et al., 2007; Ulrich and Jentsch, 2000). In complex UBE2B and RAD18 trigger replication fork stalling at DNA damage sites during the post replicative repair process (Tsuji et al., 2008). Null mutations of the UBE2B gene in mice are associated with structural abnormalities in sperm and SNP analysis of human UBE2B variants has provided evidence for association of this gene with male infertility (Escalier et al., 2003; Suryavathi et al., 2008).

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Physical Characteristics

20 μg -70°C

Species: human

Source: E. coli expression

Quantity: 20 µg

Concentration: 1 mg/ml

Formulation: 50 mM HEPES pH 7.5, 150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

Molecular Weight: ~17 kDa

Purity: >98% by InstantBlue™ SDS-PAGE

Stability/Storage: 12 months at -70°C;

aliquot as required

Protein Sequence:

GPLGS STPARRRLMRDFKRLQEDPPVGVS
GAPSENNIMQWNAVIFGPEGTPFEDGT
FKLVIEFSEEYPNKPPTVRFLSKMFHPNVY
ADGSICLDILQNRWSPTYDVSSILTSIQSLL
DEPNPNSPANSQAAQLYQENKREYEKRV
SAIVEQSWNDS

The residues <u>underlined</u> remain after cleavage and removal of the purification tag.

UBE2B (regular text): Start **bold italics** (amino acid residues

2-152)

Accession number: NP_003328

Quality Assurance

Purity:

4-12% gradient SDS-PAGE InstantBlue™ staining Lane 1: MW markers Lane 2: 1 µg UBE2B



Protein Identification:

Confirmed by mass spectrometry

E2-Ubiquitin Thioester Loading Assay:

The activity of UBE2B was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the UBE2B E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and UBE2B enzymes in the presence of ubiquitin and ATP at 30°C was compared at two time points, T_0 and T_{10} minutes. Sensitivity of the ubiquitin/UBE2B thioester bond to the reducing agent DTT was confirmed.



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Lot-specific COA version tracker: v1.0.0

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CERTIFICATE OF ANALYSIS Page 2 of 2

Background

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References:

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Tsuji Y, Watanabe K, Araki K, Shinohara M, Yamagata Y, Tsurimoto T, Hanaoka F, Yamamura K, Yamaizumi M, Tateishi S (2008) Recognition of forked and single-stranded DNA structures by human RAD18 complexed with RAD6B protein triggers its recruitment to stalled replication forks. *Genes Cells* 13, 343-54.

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