UBE2D1 (UbcH5a) [untagged]

E2 – Ubiquitin Conjugating Enzyme

Alternate Names: E2(17)KB 1, EC 6.3.2.19, SFT, Stimulator of Fe transport, homolog of UBC4/5, UbcH5, UbcH5A, Ubiquitin protein ligase, Ubiquitin-conjugating enzyme E2-17 kDa 1, Ubiquitin-conjugating enzyme UbcH5A

Cat. No.	62-0010-020
Lot. No.	1458

Quantity: 20 µg Storage: -70°C

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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). UBE2D1 is a member of the E2 ubiquitin-conjugating enzyme family and cloning of the human gene was first described by Scheffner et al. (1994). UBE2D1 shares 89% sequence identity with its Drosophila homologue and mediates E6/UBE3A (E6AP)-induced ubiquitylation of p53 (Jensen et al., 1995; Scheffner et al., 1994). Ubiquitylation of the veast PTS1 import receptor (pex5p) has been demonstrated in an in vitro assay in the presence of the human UBE2D1 in combination with the ring domain of the yeast E3 ligase pex10p (Williams et al., 2008). Sequence encoding the stimulated Iron transport gene SFT overlaps with intron 7 and exon 6 of UBE2D1, and RT/PCR has shown significantly upregulated levels of UBE2D1 in livers of iron-overloaded patients with hereditary hemochromatosis (Gehrke et al., 2003).

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Physical Characteristics

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: ~18 kDa

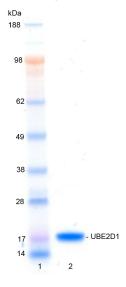
Purity: >98% by InstantBlue™ SDS-PAGE

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

Quality Assurance

Purity:

4-12% gradient SDS-PAGE InstantBlue[™] staining Lane 1: MW markers Lane 2: 1 µg UBE2D1



Protein Sequence:

GSHMASMTGGQQMGRGS**A**LKRIQKELSDLQRD PPAHCSAGPVGDDLFHWQATIMGPPDSAYQG **GVFFLTVHFPTDYPFKPPKIAFTTKIYHPNIN** SNGSICLDILRSQWSPALTVSKVLLSICSLL CDPNPDDPLVPDIAQIYKSDKEKYNRHARE WTQKYAM

The residues underlined remain after cleavage and removal of the purification tag UBE2D1 (regular text): Start bold italics (amino acid residues 2-147) Accession number: NP_003329

Protein Identification:

Confirmed by mass spectrometry.

E2-Ubiquitin Thioester Loading Assay:

The activity of UBE2D1 was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the UBE2D1 E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and UBE2D1 enzymes in the presence of ubiquitin and ATP at 30°C was compared at two time points, T₀ and T₁₀ minutes. Sensitivity of the ubiquitin/UBE2D1 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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Background

Continued from page 1

References:

Gehrke SG, Riedel HD, Herrmann T, Hadaschik B, Bents K, Veltkamp C, Stremmel W (2003) UbcH5A, a member of human E2 ubiquitin-conjugating enzymes, is closely related to SFT, a stimulator of iron transport, and is up-regulated in hereditary hemochromatosis. *Blood* **101**, 3288-93.

Jensen JP, Bates PW, Yang M, Vierstra RD, Weissman AM (1995) Identification of a family of closely related human ubiquitin conjugating enzymes. *J Biol Chem* **270**, 30408-14.

Scheffner M, Huibregtse JM, Howley PM (1994) Identification of a human ubiquitin-conjugating enzyme that mediates the E6-AP-dependent ubiquitination of p53. *Proc Natl Acad Sci USA* **91**, 8797-801.

Williams C, van den Berg M, Geers E, Distel B (2008) Pex10p functions as an E3 ligase for the Ubc4p-dependent ubiquitination of Pex5p. *Biochem Biophys Res Commun* **374**, 620-4.



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