UBE2D2 (UbcH5b) [untagged]

E2 - Ubiquitin Conjugating Enzyme

Alternate Names: E2(17)KB 2, EC 6.3.2.19, PUBC1, UBC4, UBC4/5 homolog of, UbcH5B, Ubiquitin conjugating enzyme UbcH5B, Ubiquitin-conjugating enzyme E2 D2 transcript variant 1, Ubiquitin-conjugating enzyme E2-17 kDa 2

 Cat. No.
 62-0012-100
 Quantity:
 100 μg

 Lot. No.
 30030
 Storage:
 -70°C

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). UBE2D2 is a member of the E2 ubiquitin-conjugating enzyme family and cloning of the human gene was first described by Jensen et al. (1995). UBE2D2 shares 95% and 79% sequence identity with the *Dros*ophila and S. cerevisiae homologues respectively. UBE2D2 can conjugate ubiquitin to targets in an E6AP dependent manner. UBE2D2 forms part of a ubiquitin E3 ligase complex with Cullin1, Skp1, Roc1 and BTRC (Kim et al., 2005). This complex has been shown to mediate activation of NFkB and promote degradation of phosphorylated IκBα. Co-immunoprecipitation has shown that the Shigella flexneri effector protein (OspG) interacts with UBE2D2 (Kim et al., 2005). Glial cell missing homolog 1 (GCM1) is an important transcription factor regulating placental cell fusion and it has been shown that UBE2D2 is required for Skp1-Cullin-F-Box (SCF) E3 ligase mediated ubiquitylation of GCM1 (Chiang et al., 2008). UBE2D2 also supports mdm2 mediated ubiquitylation of p53 (Saville et al., 2004).

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Physical Characteristics

Species: human

Source: E. coli expression

Quantity: 100 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: >95% by InstantBlue™ SDS-PAGE

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

aliquot as required

Protein Sequence:

GSHMASMTGGQQMGRGSALKRIHKELNDLARD PPAQCSAGPVGDDMFHWQATIMGPNDSPYQG GVFFLTIHFPTDYPFKPPKVAFTTRIYHPNIN SNGSICLDILRSQWSPALTISKVLLSICSLL CDPNPDDPLVPEIARIYKTDREKYNRIARE WTQKYAM

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

of the purification tag.

UBE2D2 (regular text): Start **bold italics** (amino acid

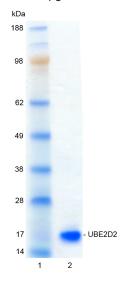
residues 2-147)

Accession number: NP_003330

Quality Assurance

Purity:

4-12% gradient SDS-PAGE InstantBlue™ staining Lane 1: MW markers Lane 2: 1 µg UBE2D2



Protein Identification:

Confirmed by mass spectrometry.

E2-Ubiquitin Thioester Loading Assay:

The activity of UBE2D2 was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the UBE2D2 E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and UBE2D2 enzymes in the presence of ubiquitin and ATP at 30°C was compared at two time points, T_0 and T_{10} minutes. The sensitivity of this ubiquitin/UBE2D2 thioester bond to the reducing agent DTT was demonstrated.



Dundee, Scotland, UK

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CERTIFICATE OF ANALYSIS Page 2 of 2

Background

Continued from page 1

References:

Chiang MH, Chen LF, Chen H (2008) Ubiquitin-conjugating enzyme UBE2D2 is responsible for FBXW2 (F-box and WD repeat domain containing 2)-mediated human GCM1 (glial cell missing homolog 1) ubiquitination and degradation. *Biol Reprod* **79**, 914-20.

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.

Kim DW, Lenzen G, Page AL, Legrain P, Sansonetti PJ, Parsot C (2005) The Shigella flexneri effector OspG interferes with innate immune responses by targeting ubiquitin-conjugating enzymes. *Proc Natl Acad Sci USA* 102, 14046-51.

Saville MK, Sparks A, Xirodimas DP, Wardrop J, Stevenson LF, Bourdon JC, Woods YL, Lane DP (2004) Regulation of p53 by the ubiquitin-conjugating enzymes UbcH5B/C in vivo. *J Biol Chem* **279**, 42169-81.



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