# UBE2G2 (Ubc7) [untagged]

E2 – Ubiguitin Conjugating Enzyme

Alternate Names: EC 6.3.2.19, UBC7, Ubiquitin-conjugating enzyme E2G 2

Cat. No.	62-0030-100	Quantity:	100 µg
Lot. No.	30196	Storage:	-70°С
FOR RESEARCH USE ONLY		NOT FOR USE IN HUMANS	

The enzymes of the ubiquitylation

pathway play a pivotal role in a num-

ber of cellular processes including

regulated and targeted proteasomal

degradation of substrate proteins.

Three classes of enzymes are in-

volved in the process of ubiquitylation; activating enzymes (E1s), conjugating

enzymes (E2s) and protein ligases

(E3s). UBE2G2 is a member of the E2

conjugating enzyme family and cloning

of the human gene was first described by Katsanis and Fischer (1998). The

UBE2G2 gene is 2.9-kb and encodes a 165-amino-acid protein that shares

57% sequence identity with UBE2G1.

UBE2G2 is involved in protein deqradation, including a process known as Endoplasmic Reticulum-Associ-

ated Degradation (ERAD). Binding

of UBE2G2 to the E3 ligase GP78

via the G2BR domain significantly

increases the affinity of UBE2G2 for

GP78. The UBE2G2/GP78 interaction

results in the preassembly of Lys-48-

linked ubiguitin chains on the catalytic

cysteine of UBE2G2. Growth of the

polyubiquitin chain is mediated by an

aminolysis-based transfer reaction be-

tween two UBE2G2 proteins; a mech-

anism for transferring preassembled

ubiquitin chains from UBE2G2 to the

lysine residue in a substrate (Das et

al., 2009; Li et al., 2007). The E3 ligase HRD1 interacts with UBE2G2 to form

Lys-48-linked polyubiquitin chains on the substrate 3-Hydroxy-3-MethylGlu-

Background

# **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: ~19 kDa

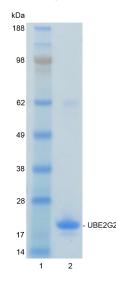
Purity: >90% by InstantBlue™ SDS-PAGE

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

## **Quality Assurance**

## **Purity:**

4-12% gradient SDS-PAGE InstantBlue<sup>™</sup> staining Lane 1: MW markers Lane 2: 1 µg UBE2G2



## **Protein Sequence:**

UBIQUIGENT

**CERTIFICATE OF ANALYSIS Page 1 of 2** 

GPLGSAGTALKRLMAEYKQLTLNPPE GIVAGPMNEENFFEWEALIMGPEDTCFEF **GVFPAILSFPLDYPLSPPKMRFTCEMFHP** NIYPDGRVCISILHAPGDDPMGYESSAER WSPVQSVEKILLSVVSMLAEPNDESGAN VDASKMWRDDREQFYKIAKQIVQKSLGL

The residues underlined remain after cleavage and removal of the purification tag UBE2G2 (regular text): Start bold italics (amino acid residues 2-165) Accession number: NP 003334

## **Protein Identification:**

Confirmed by mass spectrometry.

## E2-Ubiquitin Thioester Loading Assay:

The activity of UBE2G2 was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the UBE2G2 E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and UBE2G2 enzymes in the presence of ubiquitin and ATP at 30°C was compared at two time points, T<sub>0</sub> and T<sub>10</sub> minutes. Sensitivity of the ubiquitin/UBE2G2 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

Cat. No.

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#### Continued from page 1

taryl-coenzyme A Reductase (HMGR) targeting it for degradation (Kikkert *et al.*, 2004). A C4HC3 RING fingercontaining ubiquitin ligase of the endoplasmic reticulum - TEB4 - catalyses Lys-48-linked polyubiquitylation employing UBE2G2 *in vitro* (Hassink *et al.*, 2005).

## **References:**

Das R, Mariano J, Tsai YC, Kalathur RC, Kostova Z, Li J, Tarasov SG, McFeeters RL, Altieri AS, Ji X, Byrd RA, Weissman AM. (2009) Allosteric activation of E2-RING finger-mediated ubiquitylation by a structurally defined specific E2-binding region of gr78. *Mol Cell* **34**, 674-85.

Hassink G, Kikkert M, Voorden SV, Lee SJ, Spaapen R, Laar TV, Coleman CS, Bartee E, Fruh K, Chau V, Wiertz V. (2005) TEB4 is a C4HC3 RING finger-containing ubiquitin ligase of the endoplasmic reticulum. *Biochem J* **388**, 647-55.

Katsanis N, Fisher EM (1998) Identification, expression, and chromosomal localization of ubiquitin conjugating enzyme 7 (UBE2G2), a human homologue of the Saccharomyces cerevisiae ubc7 gene. *Genomics* **51**, 128-31.

Kikkert M, Doolman R, Dai M, Avner R, Hassink G, Voorden SV, Thanedar S, Roitelman J, Chau V, Wiertz E. (2004) Human HRD1 is an E3 ubiquitin ligase involved in degradation of proteins from the endoplasmic reticulum. *J Biol Chem* **279**, 3525-34.

Li W, Tu D, Brunger AT, Ye Y (2007) A ubiquitin ligase transfers preformed polyubiquitin chains from a conjugating enzyme to a substrate. *Nature* **446**, 333-7.



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