# **UBE2E2** (UbcH8) [GST-tagged]

E2 - Ubiquitin or ISG15 Conjugating Enzyme

Alternate Names: UbcH8

Cat. No. 62-0087-020

Lot. No. 1839

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). UBE2E2 is a member of the E2 ubiquitin-conjugating enzyme family and cloning of the human gene was first described by Kimura et al. (1997). The Ubc domain of UBE2E2 shares over 90% identity with human UBE2E1, mouse UbcM2, and Drosophila UbcD2 (Kimura et al., 1997). UBE2E2 has been shown to ubiquitylate the E3 ligase E6AP by binding to its HECT domain (Kumar et al., 1997). A yeast two hybrid screen identified two UBE2E2 binding proteins, UbcH7-Associated Protein (H7-AP1) and Human Homologue of Drosophila ARladne (HHARI); both of these proteins are characterized by the presence of a RING finger and In Between RING finger (IBR) domains (Moynihan et al., 1999). Studies using deletion mutants of UBE2E2 and two point mutants - ARA54 and C220S and RNF8 C403S, have demonstrated that ARA54 and RNF8 ring finger proteins interact with the Ubc domain of UBE2E2 (Ito et al., 2001). UBE2E2 binds directly to the BRCA1 RING motif of the human heterodimeric RING E3 ligase complex BRCA1-BARD1 and is active in causing autoubiquityla-

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## **Physical Characteristics**

20 µg

-70°C

Species: human

Quantity:

Storage:

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: ~49 kDa

Purity: >90% by InstantBlue™ SDS-PAGE

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

aliquot as required

#### **Protein Sequence:**

MSPILGYWKIKGLVQPTRLLLEYLEEKYEEH
LYERDEGDKWRNKKFELGLEFPNLPYYIDGD
VKLTQSMAIIRYIADKHNMLGGCPKERAEISMLE
GAVLDIRYGVSRIAYSKDFETLKVDFLSKLPEM
LKMFEDRLCHKTYLNGDHVTHPDFMLYDALDV
VLYMDPMCLDAFPKLVCFKKRIEAIPQIDKY
LKSSKYIAWPLQGWQATFGGGDHPPKSDLEV
LFQGPLGSMSTEAQRVDDSPSTSGGSSDGDQRES
VQQEPEREQVQPKKKEGKISSKTAAKLST
SAKRIQKELAEITLDPPPNCSAGPKGDNIYE
WRSTILGPPGSVYEGGVFFLDITFSPDYPFKP
PKVTFRTRIYHCNINSQGVICLDILKDNWSPAL
TISKVLLSICSLLTDCNPADPLVGSIATQYMT
NRAEHDRMARQWTKRYAT

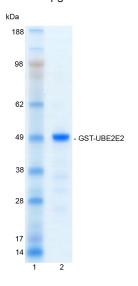
Tag (**bold text**): N-terminal GST
Protease cleavage site: PreScission™ (<u>LEVLFQ▼GP</u>)
UBE2E2 (regular text): Start **bold italics** (amino acid residues 1-201)

Accession number: NP 689866.1

## **Quality Assurance**

#### **Purity:**

4-12% gradient SDS-PAGE InstantBlue™ staining Lane 1: MW markers Lane 2: 1 μg GST-UBE2E2



#### **Protein Identification:**

Confirmed by mass spectrometry.

### **E2-Ubiquitin Thioester Loading Assay:**

The activity of GST-UBE2E2 was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the GST-UBE2E2 E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and GST-UBE2E2 enzymes in the presence of ubiquitin and ATP at  $30\,^{\circ}\text{C}$  was compared at two time points,  $T_0$  and  $T_{10}$  minutes. Sensitivity of the ubiquitin/GST-UBE2E2 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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tion *in vitro* (Christensen *et al.*, 2007). UBE2E2 has also been shown to bind the ubiquitin-protein ligase Parkin via its C-terminal ring-finger domain, resulting in ubiquitylation of the synaptic vesicle associated protein CDCrel-1 (Zhang *et al.*, 2000).

#### References:

Christensen DE, Brzovic PS, Klevit RE (2007) E2-BRCA1 RING interactions dictate synthesis of mono- or specific polyubiquitin chain linkages. *Nat Struct Mol Biol* **14**, 941-8.

Ito K, Adachi S, Iwakami R, Yasuda H, Muto Y, Seki N, Okano Y (2001) N-Terminally extended human ubiquitin-conjugating enzymes (E2s) mediate the ubiquitination of RING-finger proteins, ARAS4 and RNF8. *Eur J Biochem* **268**, 2725-32.

Kimura M, Hattori T, Matsuda Y, Yoshioka T, Sumi N, Umeda Y, Nakashima S, Okano Y (1997) cDNA cloning, characterization, and chromosome mapping of UBE2E2 encoding a human ubiquitin-conjugating E2 enzyme. Cytogenet Cell Genet 78, 107-11.

Kumar S, Kao WH, Howley PM (1997) Physical interaction between specific E2 and Hect E3 enzymes determines functional cooperativity. *J Biol Chem* **272**, 13548-54.

Moynihan TP, Ardley HC, Nuber U, Rose SA, Jones PF, Markham AF, Scheffner M, Robinson PA (1999) The ubiquitinconjugating enzymes UbcH7 and UbcH8 interact with RING finger/IBR motif-containing domains of HHARI and H7-AP1. *J Biol Chem* **274**, 30963-8.

Zhang Y, Gao J, Chung KK, Huang H, Dawson VL, Dawson TM (2000) Parkin functions as an E2-dependent ubiquitin-protein ligase and promotes the degradation of the synaptic vesicle-associated protein, CDCrel-1. *Proc Natl Acad Sci U S A* **97**, 13354-9.



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 International:
 +44 (0) 1382 381147 (9AM-5PM UTC)

 US/Canada:
 +1-617-245-0020 (9AM-5PM UTC)

 Email:
 tech.support@ubiquigent.com

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