UBE2Q1 (NICE-5) [6His-tagged]

E2 – Ubiquitin Conjugating Enzyme

Alternate Names: UBE2Q1, NICE5, PRO3094, GTAP

Cat. No.	62-0081-020
Lot. No.	1832

Quantity: 20 µg Storage: -70°C

FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS



CERTIFICATE OF ANALYSIS Page 1 of 1

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). UBE2Q1 is a member of the E2 conjugating enzyme family and cloning of the human gene was described by Marenholz et al. (2001). UBE2Q1 shares 50-75% sequence identity with its homologues in Mus musculus, Drosophila, C. elegans and Xenopus. Murine UBE2Q1 has a conserved sequence for ubiquitin binding shared by all the ubiquitin-conjugating enzymes, however its NH(2)-terminal domain appears critical for the binding and internalization of cell surface galactosyltransferase 1 (GalT1) in embryonic stem cells. UBE2Q1 regulates GalT1-associated, laminin-dependent embryonic cell adhesion and the formation of embryoid bodies (Wassler et al., 2008).

References:

Marenholz I, Zirra M, Fischer DF, Backendorf C, Ziegler A, Mischke D (2001) Identification of human epidermal differentiation complex (EDC)-encoded genes by subtractive hybridization of entire YACs to a gridded keratinocyte cDNA library. *Genome Res* **11**, 341-55.

Wassler MJ, Shur BD, Zhou W, Geng YJ (2008) Characterization of a novel ubiquitin-conjugating enzyme that regulates beta1,4-galactosyltransferase-1 in embryonic stem cells. *Stem Cells* 26, 2006-18.

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: ~50 kDa

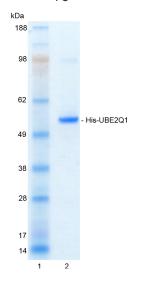
Purity: >98% by InstantBlue™ SDS-PAGE

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 His-UBE2Q1



Protein Sequence:

MHHHHHHSSGLVPRGSHMASMTGGQQMGRG SEFQQPQPQGQQQPGPGQQLGGQGAAPGAG GGPGGGPGPGPCLRRELKLLESIFHRGHER FRIASACLDELSCEFLLAGAGGAGAGAAPGPHLP PRGSVPGDPVRIHCNITESYPAVPPIWSVESD DPNLAAVLERLVDIKKGNTLLLQHLKRIISDL CKLYNLPQHPDVEMLDQPLPAEQCTQEDVSSED EDEEMPEDTEDLDHYEMKEEEPAEGKKSEDD GIGKENLAILEKIKKNQRQDYLNGAVSGSVQAT DRLMKELRDIYRSQSFKGGNYAVELVNDSLYD WNVKLLKVDQDSALHNDLQILKEKEGADFILL NFSFKDNFPFDPPFVRVVSPVLSGGYVLGG GAICMELLTKQGWSSAYSIESVIMQISATLVKG KARVQFGANKSQYSLTRAQQSYKSLVQIHEKNGW YTPPKEDG

Tag (**bold text**): N-terminal His Protease cleavage site: Thrombin (<u>LVPR▼GS</u>) UBE2Q1 (regular text): Start *bold italics* (amino acid residues 2-422). Accession number: NP_060052

Protein Identification:

Confirmed by mass spectrometry.

E2-Ubiquitin Thioester Loading Assay:

The activity of His-UBE2Q1 was validated by loading E1 UBE1 activated ubiquitin onto the active cysteine of the His-UBE2Q1 E2 enzyme via a transthiolation reaction. Incubation of the UBE1 and His-UBE2Q1 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/His-UBE2Q1 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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