# UBE2I (Ubc9) [GST-tagged] E2 - SUMO Conjugating Enzyme

The enzymes of the SUMOylation pathway

play a pivotal role in a number of cellular processes including nuclear transport,

signal transduction, stress responses and

cell cycle progression. Covalent modifica-

tion of proteins by small ubiquitin-related modifiers (SUMOs) may modulate their

stability and subcellular compartmen-

talisation. Three classes of enzymes are

involved in the process of SUMOylation;

an activating enzyme (E1), conjugating

enzyme (E2) and protein ligases (E3s).

UBE2I is a member of the E2 conjugating

enzyme family and cloning of the human gene was first described by Wang et al.

(1996). The human UBE2I cDNA contains

7 exons sharing 56% and 100% identity

with the yeast and mouse homologues

(Nacerddine et al., 2005; Shi et al., 2000; Wang et al., 1996). The candidate tumor suppressor gene Fragile Histidine Triad

(FHIT) located on 3p14.2 is deleted in many types of human cancer. UBE2I binds

to FHIT and this interaction is thought to

be involved in the degradation of S and M

phase cyclins and cell cycle control. Pro-

liferating Cell Nuclear Antigen (PCNA) a

DNA polymerase sliding clamp involved

in DNA synthesis and repair is a substrate

for UBE2I. SUMOylation of PCNA is me-

diated by UBE2I and occurs on a specific

lysine residue - K146 - which may also

be modified by ubiquitin (Hoege et al.,

2002). Crystallography has revealed that

UBE2I forms part of a 4 protein complex

consisting of a NUP358/RANBP2 E3 ligase

domain, and SUMO1 conjugated to the

carboxy-terminal domain of RANGAP1. A model for the complex has been proposed

in which NUP358/RANBP2 acts as an E3 by binding both SUMO and UBE2I to position the SUMO-E2-thioester in an optimal orientation to enhance conjugation

(Reverter and Lima, 2005). SUMOylation

Alternate Names: P18, SUMO-1 protein ligase, UBC9, Ubiquitin conjugating enzyme UbcE2A, Ubiquitin like protein SUMO-1 conjugating enzyme

| Cat. No. | 62-0065-100 |
|----------|-------------|
| Lot. No. | 1426        |
|          |             |

Quantity: 100 µg Storage: -70°C

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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: ~45 kDa

Purity: >98% by InstantBlue<sup>™</sup> SDS-PAGE

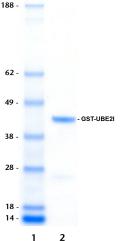
Stability/Storage: 12 months at -70°C; aliquot as required

# **Quality Assurance**

#### **Purity:**

4-12% gradient SDS-PAGE InstantBlue<sup>™</sup> staining lane 1: MW markers lane 2: 1 µg GST-UBE2I





**Protein Sequence:** 

**MSPILGYWKIKGLVQPTRLLLEYLEEKYEEH** LYERDEGDKWRNKKFELGLEFPNLPYYIDGD VKLTQSMAIIRYIADKHNMLGGCPKER AEISMLEGAVLDIRYGVSRIAYSKDFETLKVD FLSKLPEMLKMFEDRLCHKTYLNGDHVTHP DFMLYDALDVVLYMDPMCLDAFPKLVCFK KRIEAIPQIDKYLKSSKYIAWPLQGWQAT FGGGDHPPKSDLEVLFQGPLGSMSGIALSR LAQERKAWRKDHPFGFVAVPTKNPDGTMN LMNWECAIPGKKGTPWEGGLFKLRMLFKD DYPSSPPKCKFEPPLFHPNVYPSGTVCLSILEED KDWRPAITIKQILLGIQELLNEPNIQDPAQAEA YTIYCQNRVEYEKRVRAQAKKFAPS

Tag (bold text): N-terminal glutathione-S-transferase (GST) Protease cleavage site: PreScission™ (LEVLFQ▼GP) UBE2I (regular text): Start bold italics (amino acid residues 1-158) Accession number: NP 003336

#### **Protein Identification:**

Confirmed by mass spectrometry.

### SUMO-E2 Thioester Loading Assay:

The activity of GST-UBE2I was validated by loading E1 SAE1/SAE2 activated SUMO onto the active cysteine of the GST-UBE2I E2 enzyme via a transthiolation reaction. Incubation of the SAE1/SAE2 and GST-UBE2I enzymes in the presence of SUMO and ATP at 30°C was compared at two time points, T<sub>0</sub> and T<sub>10</sub> minutes. Sensitivity of the SUMO/ GST-UBE2I 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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#### UBE2I (Ubc9) [GST-tagged] E2 - SUMO Conjugating Enzyme Alternate Names: P18, SUMO-1 protein ligase, UBC9, Ubiquitin conjugating enzyme UbcE2A, Ubiquitin like protein SUMO-1 conjugating enzyme 62 0065 100 100 ..... Cat. No Quantitu

| Cal. NO. | 02-0003-100 | Quantity: | του μα |
|----------|-------------|-----------|--------|
| Lot. No. | 1426        | Storage:  | -70°C  |
|          |             |           |        |



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**CERTIFICATE OF ANALYSIS - Page 2 of 2** 

# Background

#### Continued from page 1

of Amyloid Precursor Protein (APP) was reported to be associated with decreased levels of beta amyloid (Abeta) aggregates, suggesting a role in the pathogenesis of Alzheimer's Disease (AD). An investigation into single nucleotide polymorphisms (SNPs) in the UBE2I gene have shown an association between this and the risk of late onset AD (Ahn et al., 2009).

#### **References:**

Ahn K, Song JH, Kim DK, Park MH, Jo SA, Koh YH (2009) Ubc9 gene polymorphisms and late-onset Alzheimer's disease in the Korean population: a genetic association study. *Neurosci Lett* 465, 272-5.

Hoege C, Pfander B, Moldovan GL, Pyrowolakis G, Jentsch S (2002) RAD6-dependent DNA repair is linked to modification of PCNA by ubiquitin and SUMO. *Nature* **419**, 135-41.

Nacerddine K, Lehembre F, Bhaumik M, Artus J, Cohen-Tan-noudji M, Babinet C, Pandolfi PP, Dejean A (2005) The SUMO pathway is essential for nuclear integrity and chromosome segregation in mice. Dev Cell 9, 769-79.

Reverter D, Lima CD (2005) Insights into E3 ligase activity revealed by a SUMO-RanGAP1-Ubc9-Nup358 complex. *Nature* **435**, 687-92.

Shi Y. Zou M. Farid NR. Paterson MC (2000) Association of FHIT (fragile histidine triad), a candidate tumour suppressor gene with the ubiquitin-conjugating enzyme hUBC9. Biochem / 352 Pt 2, 443-8.

Wang ZY, Qiu QQ, Seufert W, Taguchi T, Testa JR, Whitmore SA, Callen DF, Welsh D, Shenk T, Deuel TF (1996) Molecular cloning of the cDNA and chromosome localization of the gene for human ubiquitin-conjugating enzyme 9. J Biol Chem 271, 24811-6.



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