# **UBA6** [6His-tagged]

E1 - Ubiquitin Activating Enzyme

Alternate Names: E1-L2, UBE1L2

Cat. No. 61-0002-050

Lot. No. 1367

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 proteosomal 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). UBA6 is a member of the E1 activating enzyme family and the human gene was first described by Jin et al. (2007). UBA6 shares 42% homology with UBE1 and contains all the structural elements required for E1 enzyme activity (Groettrup et al., 2008). UBA6 interacts with a number of E2 and E3 enzymes and has been shown to be involved with p53 ubiquitylation in vitro (Groettrup et al., 2008; Pelzer et al., 2007). UBA6 activates ubiquitin and the ubiquitin-like protein human leukocyte antigen F-Associated Transcript 10 (FAT10), both of which may serve as a signal for proteasomal degradation. FAT10, is encoded by the major histocompatibility (MHC) class I locus, and its expression is induced by tumor necrosis factor alpha (TNF $\alpha$ ) and interferon-gamma (IFNγ). FAT10 expression is significantly upregulated in hepatocellular carcinoma as well as in gastrointestinal and gynecological cancers (Lee et al., 2003), however its precise biochemical and cellular functions have yet to be determined. Although FAT10 forms covalent conjugates with cellular proteins through its C-terminal diglycine motif (Raasi et al., 2001), substrates remain to be identified. Knockdown of UBA6 results in decreased FAT10 conjugation, indicating that UBA6 is required to activate FAT10 and facilitate its conjugation (Chiu et al., 2007).

### References:

Chiu YH, Sun Q, Chen ZJ (2007) E1-L2 activates both ubiquitin and FAT10. Mol Cell 27, 1014-23.

Continued on page 2

## **Physical Characteristics**

50 µg

-70°C

Species: human

Quantity:

Storage:

Source: Sf21 insect cell-baculovirus expression

Quantity: 50 µg

Concentration: 0.5 mg/ml

Formulation: 50 mM HEPES pH 7.5, 150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

Molecular Weight: ~123 kDa

Purity: >98% by InstantBlue™ SDS-PAGE

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

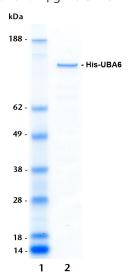
aliquot as required

# **Protein Sequence:** Please see page 2

# **Quality Assurance**

## **Purity:**

4-12% gradient SDS-PAGE InstantBlue™ staining lane 1: MW markers lane 2: 1 µg His-UBA6



#### **Protein Identification:**

Confirmed by mass spectrometry.

#### E1-Ubiquitin Thioester Loading Assay:

The activity of His-UBA6 was validated by loading ubiquitin onto the active cysteine of His-UBA6. Incubation of the His-UBA6 enzyme in the presence of ubiquitin and ATP at 30°C was compared at two time points, To and T<sub>10</sub> minutes. Sensitivity of the ubiquitin/His-UBA6 thioester bond to the reducing agent DTT was confirmed.



Dundee, Scotland, UK

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Lot-specific COA version tracker: v1.0.0

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CERTIFICATE OF ANALYSIS Page 2 of 2

## **Background**

#### Continued from page 1

Groettrup M, Pelzer C, Schmidtke G, Hofmann K (2008) Activating the ubiquitin family: UBA6 challenges the field. *Trends* Biochem Sci 33, 230-7.

Jin J, Li X, Gygi SP, Harper JW (2007) Dual E1 activation systems for ubiquitin differentially regulate E2 enzyme charging. *Nature* **447**, 1135-8.

Lee CG, Ren J, Cheong IS, Ban KH, Ooi LL, Yong Tan S, Kan A, Nuchprayoon I, Jin R, Lee KH, Choti M, Lee LA (2003) Expression of the FAT10 gene is highly upregulated in hepatocellular carcinoma and other gastrointestinal and gynecological cancers. Oncogene 22, 2592-603.

Pelzer C, Kassner I, Matentzoglu K, Singh RK, Wollscheid HP, Scheffner M, Schmidtke G, Groettrup M (2007) UBE1L2, a novel E1 enzyme specific for ubiquitin. *J Biol Chem* **282**, 23010-4.

Raasi S, Schmidtke G, Groettrup M (2001) The ubiquitin-like protein FAT10 forms covalent conjugates and induces apoptosis. J Biol Chem **276**, 35334-43.

# **Physical Characteristics**

Continued from page 1

**Protein Sequence:** 

**MSYYHHHHHHDYDIPTT**ENLYFQGAMGS GIQRPTSTSSLVAAA**M**EGSEPVAAHQGEEASC SSWGTGSTNKNLPIMSTASVEIDDALYSRQRYV LGDTAMQKMAKSHVFLSGMGGLGLEIAKN LVLAGIKAVTIHDTEKCQAWDLGTNFFLSED DVVNKRNRAEAVLKHIAELNPYVHVTSSSVPF NETTDLSFLDKYQCVVLTEMKLPLQKKINDFCR SQCPPIKFISADVHGIWSRLFCDFGDEFEVLDT TGEEPKEIFISNITQANPGIVTCLENHPHKLET GQFLTFREINGMTGLNGSIQQITVISPFSFSIGDT TELEPYLHGGIAVOVKTPKTVFFESLEROLKHPK CLIVDFSNPEAPLEIHTAMLALDQFQEKYSRKPN VGCQQDSEELLKLATSISETLEEKPDVNADIVH WLSWTAQGFLSPLAAAVGGVASQEVLKAVTG KFSPLCQWLYLEAADIVESLGKPECEEFLPRGDRY DALRACIGDTLCQKLQNLNIFLVGCGAIGCEM LKNFALLGVGTSKEKGMITVTDPDLIEKSNLN RQFLFRPHHIQKPKSYTAADATLKINSQIKIDAHL NKVCPTTETIYNDEFYTKQDVIITALDNVEARRYVD SRCLANLRPLLDSGTMGTKGHTEVIVPHLTESYN SHRDPPEEEIPFCTLKSFPAAIEHTIQWARDKFESSF SHKPSLFNKFWQTYSSAEEVLQKIQSGHSLEGCFQ VIKLLSRRPRNWSQCVELARLKFEKYFNHKALQLL **HCFPLDIRLKDGSLFWQSPKRPPSPIKFDLNEPL** HLSFLQNAAKLYATVYCIPFAEEDLSADALLNIL SEVKIQEFKPSNKVVQTDETARKPDHVPISSED ERNAIFQLEKAILSNEATKSDLQMAVLSFEKD DDHNGHIDFITAASNLRAKMYSIEPADRFKT KRIAGKIIPAIATTTATVSGLVALEMIKVTGGYP **FEAYKNCFLNLAIPIVVFTETTEVRKTKIRNGISFTI** WDRWTVHGKEDFTLLDFINAVKEKYGIEPTMV VQGVKMLYVPVMPGHAKRLKLTMHKLVKPTTEK KYVDLTVSFAPDIDGDEDLPGPPVRYYFSHDTD

Tag (bold text): N-terminal His

Protease cleavage site: TEV (ENLYFQ▼G)

UBA6 (regular text): Start bold italics (amino acid resi-

dues 1-1052)

Accession number: NP\_060697



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