

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of UBE2B

<u>Enzyme description:-</u>	UBE2B
<u>Clone number:-</u>	DU20017
<u>Source:-</u>	BL21 recombinant
<u>Tag:-</u>	cleaved from N-terminal His ₆ -tag
<u>Purification method:-</u>	Ni ⁺⁺ -NTA-Sepharose; protease treatment, SEC
<u>Expression level:-</u>	6mg/L
<u>Calculated molecular mass:-</u>	
Monoisotopic	17599 Da
Average Mass	17610 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	4.81
<u>Purity:-</u>	90%
<u>Enzyme storage buffer:-</u>	
50mM HEPES pH 7.5, 150mM NaCl, 10% glycerol, 1mM DTT	
<u>Storage temperature:-</u>	-80°C
<u>Assay:-</u>	
Loading with Ubiquitin and UBE1 in the presence of Mg-ATP	

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Clone Data Sheet

His-UBE2B

<u>Protein</u>	UBE2B
<u>Synonyms</u>	ubiquitin-conjugating enzyme E2B, Rad 6 homolog
<u>Clone Number</u>	DU20017
<u>Species</u>	Human
<u>Accession Number</u>	Protein: NP_003328 DNA: NM_003337
<u>Tags</u>	N-terminal His ₆ tag, cleaved
Aminoacid sequence of the expressed protein	<u>G</u> PGSMSTPARRRLMRDFKRLQEDPPVGVSGAPSENNIMQWNAVIFGPEGTPF EDGTFKLVIEFSEEYPNKPPTVRFLSKMFHPNVYADGSICLDILQNRWSPTY DVSSILTSIQSLLEPNPNPANSQAAQLYQENKREYEKRVSAIVEQSWNDS
Native sequence	
Protease cleavage	Prescission site underlined
Cloning sites	BamH1 / NotI
<u>DNA sequence of insert</u>	<u>GGATCC</u> ATGTCGACCCCGGCCCGGAGGAGGCTCATGCGGGATTTCAAGCG GTTACAAGAGGACCCACCTGTGGGTGTCAGTGGCGCACCATCTGAAAACA ACATCATGCAGTGGAATGCAGTTATATTTGGACCAGAAGGGACACCTTTT GAAGATGGTACTTTTAAACTAGTAATAGAATTTTCTGAAGAATATCCAAA TAAACCACCAACTGTTAGGTTTTTATCCAAAATGTTTCATCCAAATGTGT ATGCTGATGGTAGCATATGTTTAGATATCCTTCAGAATCGATGGAGTCCA ACATATGATGTATCTTCTATCTTAACATCAATTCAGTCTCTGCTGGATGA ACCGAATCCTAACAGTCCAGCCAATAGCCAGGCAGCACAGCTTTATCAGG AAAACAAACGAGAATATGAGAAAAGAGTTTCGGCCATTGTTGAACAAAGC TGGAATGATTCATAAGCGGCCGC