

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of UBE2H

<u>Enzyme description:-</u>	UBE2H
<u>Clone number:-</u>	DU32149
<u>Source:-</u>	BL21 recombinant
<u>Tag:-</u>	cleaved from N-terminal His ₆ -tag
<u>Purification method:-</u>	Ni ⁺⁺ -NTA-Sepharose, protease treatment
<u>Expression level:-</u>	2.5mg/L
<u>Calculated molecular mass:-</u>	
Monoisotopic	20940 Da
Average Mass	20952 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	4.47
<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-UBE2H

<u>Protein</u>	UBE2H
<u>Synonyms</u>	Ubc8, E2-20K, UbcH, UbcH2
<u>Clone Number</u>	DU32149
<u>Species</u>	Human
<u>Accession Number</u>	Protein: NP_003335, DNA: NM_003344
<u>Tags</u>	N-terminal His ₆ tag
Aminoacid sequence of the expressed protein	<u>GP</u> GS MSSPSPGKR MDTDVV KLIES KHEVTILGGLNEFVV KFYGPQGT PYEG GVWKVRVDLPDKY PFKSP SIGFMNKIFHPNIDEAS GTVC LDVINQ TWTALYD LTNIFESFLPQ LLAYPNPIDPLNGDAA MYLHRP EYKQKIKEYIQKYATEE ALKEQE EGTGDSSSESSMSDFSEDEA QDMEL
Native sequence	in bold
Protease cleavage	Prescission Protease site underlined
Cloning sites	BamH1 / NotI
<u>DNA sequence of insert</u>	GGATCCATGTCATCTCCCAGTCCGGGCAAGAGGCGGATGGACACGGACGT GGTCAAGCTCATCGAGAGTAAACATGAGGTTACGATCCTGGGAGGACTTA ATGAATTTGTAGTGAAGTTTATGGACCACAAGGAACACCATATGAAGGC GGAGTATGGAAAGTTAGAGTGGACCTACCTGATAAATACCCTTTCAAATC TCCATCTATAGGATTCATGAATAAAATTTTCCATCCCAACATTGATGAAG CGTCAGGAAGTGTGTGTCTAGATGTAATTAATCAAACCTTGGACAGCTCTC TATGATCTTACCAATATATTTGAGTCCTTCCTGCCTCAGTTATTGGCCTA TCCTAACCCCATAGATCCTCTCAATGGTGACGCTGCAGCCATGTACCTCC ACCGACCAGAAGAATACAAGCAGAAAATTAAAGAGTACATCCAGAAATAC GCCACGGAGGAGGCGCTGAAAGAACAGGAAGAGGGTACCGGGGACAGCTC ATCGGAGAGCTCTATGTCTGACTTTTCCGAAGATGAGGCCAGGATATGG AGTTGTAGGCGGCCGC