

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of UBE2G2

<u>Enzyme description:-</u>	UBE2G2
<u>Clone number:-</u>	DU20174
<u>Source:-</u>	BL21 recombinant
<u>Tag:-</u>	cleaved from N-terminal His ₆ -tag
<u>Purification method:-</u>	Ni ⁺⁺ -NTA-Sepharose, Thrombin treatment
<u>Expression level:-</u>	3mg/L
<u>Calculated molecular mass:-</u>	
Monoisotopic	20083 Da
Average Mass	20096 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	4.67
<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 assay with Ubiquitin and UBE1 in the presence of Mg-ATP	

Division of Signal Transduction Therapy

Clone Data Sheet

UBE2G2

<u>Protein</u>	UBE2G2
<u>Synonyms</u>	
<u>Clone Number</u>	DU20174
<u>Species</u>	Human
<u>Accession Number</u>	Protein: NP_003334 DNA: NM_003343
<u>Tags</u>	N-terminal His ₆ tag (cleaved)
Aminoacid sequence of the expressed protein	<u>G</u>SHMASMTGGQ<u>Q</u>MGRGSAGTALKRLMAEYKQLTLNPPEGIVAGPMNEENFFE WEALIMGPEDTCF<u>E</u>FGVFPAILS<u>F</u>PLDYPLSPPKMRFTCEMFHPNIYPDGRV CISILHAPGDDPMGYESSAERWSPVQSVEKILLSVVSMLAEPNDESGANVDA SKMWRDDREQFYKIAKQIVQKSLGL
Native sequence	in bold
Protease cleavage	Thrombin site underlined
Cloning sites	BamH1 / Not1
<u>DNA sequence of insert</u>	GGATCCGCGGGGACCGCGCTCAAGAGGCTGATGGCCGAGTACAAACAATT AACACTGAATCCTCCGGAAGGAATTGTAGCAGGCCCCATGAATGAAGAGA ACTTTTTTGAATGGGAGGCATTGATCATGGGCCCAGAAGACACCTGCTTT GAGTTTGGTGTTCCTGCCATCCTGAGTTTCCCACTTGATTACCCGTT AAGTCCCCAAAGATGAGATTTACCTGTGAGATGTTTCATCCCAACATCT ACCCTGATGGGAGAGTCTGCATTTCCATCCTCCACGCGCCAGGCGATGAC CCCATGGGCTACGAGAGCAGCGCGGAGCGGTGGAGTCCTGTGCAGAGTGT GGAGAAGATCCTGCTGTCTGGTGGTGGAGTGGTGGAGTGGTGGAGTGGT AAAGTGGAGCTAACGTGGATGCGTCCAAAATGTGGCGCGATGACCGGGAG CAGTCTATAAGATTGCCAAGCAGATCGTCCAGAAGTCTCTGGGACTGTG AGCGGCCGC