

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of UBE2F

<u>Enzyme description:-</u>	UBE2F
<u>Clone number:-</u>	DU14051
<u>Source:-</u>	BL21 recombinant
<u>Tag:-</u>	cleaved from N-terminal GST-tag
<u>Purification method:-</u>	GSH-Sepharose, protease treatment
<u>Expression level:-</u>	3mg/L
<u>Calculated molecular mass:-</u>	
Monoisotopic	21474 Da
Average Mass	21487 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	6.7
<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 NEDD8 and NAE1/NAE2 in the presence of Mg-ATP	

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Clone Data Sheet

UBE2F

<u>Protein</u>	UBE2F
<u>Synonyms</u>	
<u>Clone Number</u>	DU14051
<u>Species</u>	Human
<u>Accession Number</u>	Protein: Q969M7 DNA: NM_080678
<u>Tags</u>	N-terminal GST-tag (cleaved)
Aminoacid sequence of the expressed protein	<u>GPLGSMLTLASKLKRDDGLKGSRTAATASDSTRRVSVRDKLLVKEVAELEAN LPCTCKVHFDPNKLHCFQLTVTPDEGYIQGGKQFETEVPDAYNMVPPKVK CLTKIWHPNITETGEICLSLLREHSIDGTGWAPTRTLKDVVWGLNSLFTDLL NFDDPLNIEAAEHHLRDKEDFRNKVDDYIKRYAR</u>
Native sequence	in bold
Protease cleavage	Prescission protease site underlined
Cloning sites	BamH1 / NotI
<u>DNA sequence of insert</u>	GGATCCATGCTAACGCTAGCAAGTAAACTGAAGCGTGACGATGGTCTCAA AGGGTCCCAGGACGGCAGCCACAGCGTCCGACTCGACTCGGAGGGTTTCTG TGAGAGACAAATTGCTTGTTAAAGAGGTTGCAGAACTTGAAGCTAATTTA CCTTGATACATGTAAAGTGCATTTTCCTGATCCAAACAAGCTTCATTTGTTT TCAGCTAACAGTAACCCAGATGAGGGTTACTACCAGGGTGGAAAATTTT AGTTTGAACTGAAGTTCCCGATGCGTACAACATGGTGCCTCCCAAAGTG AAATGCCTGACCAAGATCTGGCACCCCAACATCACAGAGACAGGGGAAAT ATGCTGAGTTTATTGAGAGAACATTC AATTGATGGCACTGGCTGGGCTC CCACAAGAACATTAAAGGATGTCGTTTGGGGATTAAACTCTTTGTTTACT GATCTTTTGAATTTGATGATCCACTGAATATTGAAGCTGCAGAACATCA TTTGCGGGACAAGGAGGACTTCCGGAATAAAGTGGATGACTACATCAAAC GTTATGCCAGATGAGCGGCCGC