

*Division of Signal Transduction Therapy*

**Standard Operation Procedure**

**Preparation of UBE2G1**

<b><u>Enzyme description:-</u></b>	UBE2G1
<b><u>Clone number:-</u></b>	DU14054
<b><u>Source:-</u></b>	BL21 recombinant
<b><u>Tag:-</u></b>	cleaved from N-terminal GST-tag
<b><u>Purification method:-</u></b>	GSH-Sepharose
<b><u>Expression level:-</u></b>	10mg/L
<b><u>Calculated molecular mass:-</u></b>	
Monoisotopic	19908 Da
Average Mass	19920 Da
[cysteines reduced, methionines have not been oxidised]	
<b><u>Theoretical pI:-</u></b>	5.14
<b><u>Purity:-</u></b>	90%
<b><u>Enzyme storage buffer:-</u></b>	
50mM HEPES pH 7.5, 150mM NaCl, 10% glycerol, 1mM DTT	
<b><u>Storage temperature:-</u></b>	-80°C
<b><u>Assay:-</u></b>	
Loading with Ubiquitin and UBE1 in the presence of Mg-ATP	

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**Clone Data Sheet**

**UBE2G1**

<b><u>Protein</u></b>	UBE2G1
<b><u>Synonyms</u></b>	Ubc7, UBE2G
<b><u>Clone Number</u></b>	DU14054
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: P62253 DNA: NM_003342
<b><u>Tags</u></b>	N-terminal GST-tag (cleaved)
Aminoacid sequence of the expressed protein	<u>GPLGSMTELO</u> <b>SALLRRQLAELNKNPVEGFSAGLIDDNDLYRWEVLIIGPPD TLYEGGVFKAHLTFPKDYPLRPPKMKFITEIWHPNVDKNGDVCISILHEPGE DKYGYEKPEERWLP IHTVETIMISVISMLADPDGNSPANVDAAKEWREDRNG EFKRKVARCVRKSQETA</b> FE
Native sequence	in bold
Protease cleavage	Prescission protease site underlined
Cloning sites	BamH1 / Not1
<b><u>DNA sequence of insert</u></b>	GGATCCATGACGGAGCTGCAGTCGGCACTGCTACTGCGAAGACAGCTGGC AGAACTCAACAAAAATCCAGTGGAAAGCTTTTCTGCAGGTTTAATAGATG ACAATGATCTCTACCGATGGGAAGTCCTTATTATTGGCCCTCCAGATACA CTTTATGAAGGTGGTGTGTTTTTAAGGCTCATCTTACTTTCCAAAAGATTA TCCCCTCCGACCTCCTAAAATGAAATTCATTACAGAAATCTGGCACCCAA ATGTTGATAAAAAATGGTGTATGTGTGCATTTCTATTCCTCATGAGCCTGGG GAAGATAAGTATGGTTATGAAAAGCCAGAGGAACGCTGGCTCCCTATCCA CACTGTGGAAACCATCATGATTAGTGTGATTTCTATGCTGGCAGACCCTA ATGGAGACTCACCTGCTAATGTTGATGCTGCGAAAGAATGGAGGGAAGAT AGAAATGGAGAATTTAAAAGAAAAGTTGCCCGCTGTGTAAGAAAAAGCCA AGAGACTGCTTTTGAGTGAGCGGCCGC