

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of UBE2E1

<u>Enzyme description:-</u>	UBE2E1 1-193 (full length)
<u>Clone number:-</u>	DU12803
<u>Source:-</u>	human recombinant
<u>Tag:-</u>	cleaved from N-terminal His ₆ -tag
<u>Purification method:-</u>	Ni ⁺⁺ -NTA-Sepharose, protease treatment
<u>Expression system:-</u>	E.coli
<u>Calculated molecular mass:-</u>	
Monoisotopic	22858 Da
Average Mass	22872 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	9.07
<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	

Division of Signal Transduction Therapy

Clone Data Sheet

UBE2E1

<u>Protein</u>	UBE2E1 1-193 (full length)
<u>Synonyms</u>	UbcH6, E2E1
<u>Clone Number</u>	DU12803
<u>Species</u>	Human
<u>Accession Number</u>	Protein: NP_003332
<u>Tags</u>	N-terminal His ₆ tag
Aminoacid sequence of the expressed protein	<u>G</u> PGSPEFPGVDSKAAAMSDDDSRASTSSSSSSSSSNQOTEKETNTPKKKE SKVSMKNSKLLSTSAKRIQ KE LADITLDPPPNC SAGPKGDNIYWRST ILGPPGSVYEGGVFFLDITFTPEY PFKPPKVTFRTR RIYHCNINSQGVIC LDILKDNWSPALTI SKVLLS ICSL LLTDCNPADPLVGS IATQYMTNRAEH DRMARQWTKRYAT
Native sequence	in bold
Protease cleavage	Prescission Protease site underlined
Cloning sites	Not1
<u>DNA sequence of insert</u>	GCGGCCGCGATGTCGGATGACGATTTCGAGGGCCAGCACCAGCTCCTCCTC ATCTTCGTCTTCCAACCAGCAAACCGAGAAAGAAACAAACACCCCAAGA AGAAGGAGAGTAAAGTCAGCATGAGCAAAAACTCCAACTCCTCTCCACC AGCGCCAAGAGAAATTCAGAAGGAGCTGGCGGACATCACTTTAGACCCTCC ACCTAATTGCAGTGCTGGTCCCAAAGGCGATAACATCTATGAATGGAGAT CAACCATTCTAGGGCCTCCAGGATCCGTGTATGAGGGTGGTGTATTCTTT CTCGATATCACTTTTACACCAGAATATCCCTTCAAGCCTCCAAAGGTTAC ATTTGCGACAAGAATCTATCATTTGTAATATTAACAGTCAAGGTGTTATTT GCTTGGACATATTGAAAGATAATTGGAGTCCAGCACTAACCATTTCTAAA GTCTCCTTTCTATCTGCTCACTTCTTACAGACTGTAATCCTGCCGACCC CTTGGTGGGAAGTATTGCCACTCAGTATATGACCAACAGAGCAGAACATG ACAGAATGGCCAGACAGTGGACCAAGAGATACGCTACATAAGCGGCCGC