

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of Ubiquitin linear dimer

<u>Enzyme description:-</u>	Ubiquitin linear dimer
<u>Clone number:-</u>	DU20729
<u>Source:-</u>	BL21 recombinant
<u>Tag:-</u>	cleaved from GST-
<u>Purification method:-</u>	GSH-Sepharose, protease treatment, depletion over Q-sepharose, Source 15 S
<u>Expression level:-</u>	1.2mg/L
<u>Calculated molecular mass:-</u>	
Monoisotopic	17640 Da
Average Mass	17651 Da
[cysteines reduced, methionines have not been oxidised]	
<u>Theoretical pI:-</u>	7.7
<u>Purity:-</u>	90%
<u>Enzyme storage buffer:-</u>	
50 mM HEPES pH 7.5, 10% glycerol, 150mM NaCl, 1mM DTT	
<u>Storage temperature:-</u>	-80°C
<u>Assay:-</u>	
Protease treatment with USP2, USP5 or CYLD	

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Clone Data Sheet

Ubiquitin-dimer (linear)

<u>Protein</u>	Ubiquitin dimer (linear)
<u>Synonyms</u>	Ubiquitin-60S ribosomal protein L40; AltName: CEP52; Ubiquitin A-52 residue ribosomal protein fusion product 1
<u>Clone Number</u>	DU20729
<u>Species</u>	Human
<u>Accession Number</u>	Protein: P62987
<u>Tags</u>	cleaved from GST-Prescission
Aminoacid sequence of the purified protein	<u>G</u>PLGSAGMQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQ <u>Q</u>RLIFAGKQLEDGRTLSDYNIQKESTLHLVLRRLRGGMQIFVKTLTGK <u>T</u>ITLEVEPSDTIENVKAKIQDKEGIPPDQ<u>Q</u>RLIFAGKQLEDGRTLSD <u>Y</u>NIQKESTLHLVLRRLRGG
Native sequence	
Protease cleavage	Prescission Protease
Cloning sites	NaeI / SfoI

**DNA sequence of
insert**

GCCGGCATGCAGATCTTCGTGAAGACCCTGACTGGTAAGACCATCACTCT
CGAAGTGGAGCCGAGTGACACCATTGAGAATGTCAAGGCAAAGATCCAAG
ACAAGGAAGGCATCCCTCCTGACCAGCAGAGGTTGATCTTTGCTGGGAAA
CAGCTGGAAGATGGACGCACCCTGTCTGACTACAACATCCAGAAAGAGTC
CACCTGCACCTGGTCCTCCGTCTCAGAGGCGCC