

*Division of Signal Transduction Therapy*

**Standard Operation Procedure**

**Preparation of His-Ubiquitin 2-76 [K48R]**

**Enzyme description:-** Ubiquitin 2-76 [K48R]

**Clone number:-** DU8736

**Source:-** BL21 recombinant

**Tag:-** N-terminal His<sub>6</sub>-tag

**Purification method:-** Ni<sup>++</sup>-NTA-Sepharose

**Expression level:-** 8 mg/L

**Calculated molecular mass:-**

Monoisotopic 10909 Da

Average Mass 10916 Da

[cysteines reduced, methionines have not been oxidised]

**Theoretical pI:-** 7.17

**Purity:-** 90%

**Enzyme storage buffer:-**

50mM HEPES pH 7.5, 150mM NaCl, 10% glycerol, 1mM DTT

**Storage temperature:-** -80°C

**Assay:-**

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**Clone Data Sheet**

**His-Ubiquitin 2-76 [K48R]**

<b><u>Protein</u></b>	Ubiquitin 2-76 [K48R]
<b><u>Synonyms</u></b>	
<b><u>Clone Number</u></b>	DU8736
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: 62988
<b><u>Tags</u></b>	N-terminal His <sub>6</sub> tag
Aminoacid sequence of the expressed protein	MGSPHHHHHHSSGLEVLFOGPGS <b>QIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQRLIFAGRQLEDGRTLSDYNIQESTLHLVLR</b> RGG
Native sequence	in bold
Protease cleavage	Prescission protease site underlined
Cloning sites	BglIII/BamH1 XhoI
<b><u>DNA sequence of insert</u></b>	GGATCCCAGATCTTCGTGAAGACCCTGACTGGTAAGACCATCACTCTCGA AGTGGAGCCGAGTGACACCATTGAGAATGTCAAGGCAAAGATCCAAGACA AGGAAGGCATCCCTCCTGACCAGCAGAGGTTGATCTTTGCTGGGAGACAG CTGGAAGATGGACGCACCCTGTCTGACTACAACATCCAGAAAGAGTCCAC CCTGCACCTGGTCCCTCCGTCTCAGAGGTGGGTGATAACTCGACTCGAG