

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of His-SUMO1 1-97 (mature full length)

<u>Enzyme description:-</u>	SUMO1 1-97
<u>Clone number:-</u>	DU32080
<u>Source:-</u>	human recombinant
<u>Tag:-</u>	His-
<u>Purification method:-</u>	Ni ²⁺ -NTA-Sepharose
<u>Expression system:-</u>	E.coli

Calculated molecular mass:-

Monoisotopic	13568 Da
Average Mass	13577 Da
[cysteines reduced, methionines have not been oxidised]	

Theoretical pI:- 6.09

Purity:- 90%

Enzyme storage buffer:-

10mM Ammonium bicarbonate

Storage temperature:- -80°C

Assay:-

Loading of UBE2I.

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Clone Data Sheet

His-SUMO 1-97

<u>Protein</u>	SUMO1 1-97
<u>Synonyms</u>	SUMO1 (SMT3 suppressor of mif two 3 homolog 1 isoform a)
<u>Clone Number</u>	DU32080
<u>Species</u>	Human
<u>Accession Number</u>	Protein: NP_003343
<u>Tags</u>	His-
<u>Aminoacid sequence of the expressed protein</u>	MGSSHHHHHSSGLEVLFOGPGSMSDQEAKPSTEDLGDKKEGEYIKL KVIGQDSSEIHFVKVMTTHLKKLKESYCORQGVPMNSLRFLFEGQRI ADNHTPKELGMEEEDVIEVYQEQTGG
<u>Native sequence</u>	The native sequence is a little longer and normally processed to this mature form by SENP1.
<u>Protease cleavage</u>	Precession protease site underlined
<u>Cloning sites</u>	BamHI / NotI
<u>DNA sequence of insert</u>	GGATCCATGTCTGACCAGGAGGCAAACCTTCAACTGAGGACTTGGGGGA TAAGAAGGAAGGTGAATATATTAACTCAAAGTCATTGGACAGGATAGCA GTGAGATTCACCTCAAAGTGAAAATGACAACACATCTCAAGAACTCAA GAATCATACTGTCAAAGACAGGGTGTTCCAATGAATTCACCTCAGGTTTCT CTTTGAGGGTCAGAGAATTGCTGATAATCATACTCCAAAAGAACTGGGAA TGGAGGAAGAAGATGTGATTGAAGTTTATCAGGAACAAACGGGGGGTTAA TCAACAGTTTAGGCCGCCGC