

*Division of Signal Transduction Therapy*

**Standard Operation Procedure**

**Preparation of DCUN1D1**

**Enzyme description:-** His-DCUN1D1

**Clone number:-** DU20631

**Source:-** Recombinant

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

**Purification method:-** Ni<sup>++</sup>-Sephrose

**Expression level:-** 5mg/L

**Calculated molecular mass:-**

Monoisotopic 32548 Da

Average Mass 32568 Da

[cysteines reduced, methionines have not been oxidised]

**Theoretical pI:-** 5.55

**Purity:-** 90%

**Enzyme storage buffer:-**

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

**Storage temperature:-** -80°C

**Assay:-**

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**Clone Data Sheet**

**Protein name DCUN1D1**

<b><u>Protein</u></b>	His-DCUN1D1 1-259 (full length)
<b><u>Synonyms</u></b>	DCUN1L1, RP42, SCCRO
<b><u>Clone Number</u></b>	DU20631
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein Q96GG9
<b><u>Tags</u></b>	N-terminal His <sub>6</sub>
Aminoacid sequence of the expressed protein	<b>MGSSHHHHHSSG<u>LEVL</u>FQGP<u>GS</u>MNKLKSSQKDKVRQFMIF<u>TQ</u>SSSEKTAV SCL<u>SQ</u>NDWKL<u>DV</u>ATDNFFQNP<u>EL</u>YIRESVKGSLDRKKLEQLYNRYKDPQD ENKIGIDGIQ<u>Q</u>F<u>C</u>DDLALDPASISVLI IAWKFRAATQCEFSKQEFMDGMT ELGCDSIEKLKAQIPKMEQELKEPGRFKDFYQFTFNFAKNPGQKGLDLEM AIAYWNLVNLGRFKFLDLWNKFLLEHHKRSIPKDTWNLLDFSTMIADDM SNYDEEGAWPVLI<u>DD</u>FVEFAR<u>PQ</u>IAGTKSTTV</b>
Native sequence	in bold
Protease cleavage	Prescission site underlined
Cloning sites	BamH1 Not1

<b><u>DNA sequence of insert</u></b>	<b>GGATCCATGAACAAGTTGAAATCATCGCAGAAGGATAAAGTTCGTCAGTT TATGATCTTCACACAATCTAGTGAAAAACAGCAGTAAGTTGTCTTTCTC AAAATGACTGGAAGTTAGATGTTGCAACAGATAATTTTTTCCAAAATCCT GAACTTTATATACGAGAGAGTGTAAAAGGATCATTGGACAGGAAGAAGTT AGAACAGCTGTACAATAGATACAAAGACCCCTCAAGATGAGAATAAAATTG GAATAGATGGCATAACAGCAGTCTGTGATGACCTGGCACTCGATCCAGCC AGCATTAGTGTGTTGATTATTGCATGGAAGTTCAGAGCAGCAACACAGTG CGAGTTCTCCAAACAGGAGTTCATGGATGGCATGACAGAATTAGGATGTG ACAGCATAGAAAACTAAAGGCCAGATAACCAAGATGGAACAAGAATTG AAAGAACCAGGACGATTTAAGGATTTTTTACCAGTTTACTTTTAATTTTGC AAAGAATCCAGGACAAAAAGGATTAGATCTAGAAATGGCCATTGCCTACT GGAACCTAGTGCTTAATGGAAGATTTAAATCTTAGACTTATGGAATAAA TTTTTGTTGGAACATCATAAACGATCAATACCAAAAGACACTTGGAATCT TCTTTTAGACTTCAGTACGATGATTGCAGATGACATGTCTAATTATGATG AGAAGGAGCATGGCCTGTTCTTATTGATGACTTTGTGGAATTTGCACGC CCTCAAATTGCTGGGACAAAAAGTACAACAGTGTAGGCGGCCGC</b>
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