

Division of Signal Transduction Therapy

Standard Operation Procedure

Preparation of Cullin-2 / Rbx1

<u>Enzyme description:-</u>	Cullin-2 / Rbx1 dimer
<u>Clone number:-</u>	DU23263
<u>Source:-</u>	Recombinant
<u>Tag:-</u>	cleaved from Dac-tag
<u>Purification method:-</u>	Ampicillin Sepharose, SEC
<u>Expression level:-</u>	3mg/L

Calculated molecular mass:-

Monoisotopic	Cullin-2:	87127 Da
Average Mass	Cullin-2:	87181 Da
Monoisotopic	Rbx1:	12265 Da
Average Mass	Rbx1:	12273 Da

[cysteines reduced, methionines have not been oxidised]

<u>Theoretical pI:-</u>	Cullin2: 6.82	Rbx1: 7.00
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<u>Purity:-</u>	90%
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Enzyme storage buffer:-

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

<u>Storage temperature:-</u>	-80°C
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Assay:-

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Clone Data Sheet

Protein name

Protein Cullin-2 1-745 (full length)

Synonyms Cul2

Clone Number DU23263

Species Human

Accession Number Q13617

Tags cleaved from Dac-tag

Aminoacid sequence of
the purified and cleaved
Cullin2

GGSMSLKPRVVDFDETWNKLLTTIKAVVMLEYVERATWDRFSDIYAL
CVAYPEPLGERLYTETKIFLENHVRHLHKRVLESEEQVLVMYHRYWEE
YSKGADYMDCLYRYLNTQFIKKNKLTEADLOYGGVDMNEPLMEIGE
LALDMWRKLMVEPLQAILIRMLLREIKNDRGGEDPNQKVIHGVINSFV
HVEQYKKKFFPLKFYQEIFESPFLTETGEYYKQEASNLLQESNCSQYME
KVLGRLKDEEIRCRKYLHPSSYTKVIHECQORMVADHLQFLHAECHNI
IRQEKKNDMANMYVLLRAVSTGLPHMIQELQNHIDEGLRATSNLTQE
NMPTLFVESVLEVHGKVFQLintVLNGDQHFMSALDKALTSVVNYREP
KSVCKAPELLAKYCDNLLKKSAGMTENEVEDRLTSFITVFKYIDDKD
VFQKFYARMLAKRLIHGLSMSMDSEAMINKLKQACGYEFTSKLHRMY
TDM SVSADLNNKFNNFIKNQDTVIDLGISFQIYVLQAGAWPLTQAPSS
TFAIPQELEKSVQMFELFYSQHFSGRKLTLWHYLCTGEVKMNYLGKPY
VAMVTTYQMAVLLAFNNSETVSYKELQDSTQMKNEKELTKTIKSLLDVK
MINHDSEKEDIDAESSFLNMNFSKRTKFKITTSMQKDTPOEMEQR
SAVDEDRKMYLQAAIVRIMKARKVLRHNALIQEVISQSRARFNPSISM
IKKCIEVLIDKQYIERSQASADEYSYVA

Native sequence in bold

Protease cleavage TEV site underlined

Cloning sites BamH1 / NotI

**DNA sequence
of the Cullin 2
insert**

GGATCCATGTCTTTGAAACCAAGAGTAGTAGATTTTGATGAAACATGGAACAA
ACTTTTGACGACAATAAAAGCCGTGGTCATGTTGGAATACGTGCAAAGAGCAA
CATGGAATGACCGTTTCTCAGATATCTATGCTTTATGTGTGGCCTATCCTGAA
CCCCTTGGAGAAAGACTTTTATACAGAACTAAGATTTTTTTGGAAAATCATGT
TCGGCATTTGCATAAGAGAGTTTTGGAGTCAGAAGAACAAGTACTTGTATGT
ATCATAGGTACTGGGAAGAATACAGCAAGGGTGCAGACTATATGGACTGCTTA
TATAGGTATCTCAACACCCAGTTTATTA AAAAGAATAAATTAACAGAAGCGGA
CCTTCAGTATGGCTATGGTGGTGTAGATATGAATGAACCACTTATGGAAATAG
GAGAGCTAGCATTGGATATGTGGAGGAAATTGATGGTTGAACCACTTCAGGCC
ATCCTTATCCGAATGCTGCTCCGAGAAATCAAAAATGATCGTGGTGGAGAAGA
CCCAAACCAGAAAGTAATCCATGGGGTTATTAACCTCTTTGTTTCATGTTGAAC
AGTATAAGAAAAAATTCCTTAAAGTTTTATCAGGAAATTTTTGAGTCTCCC
TTTCTGACTGAAACAGGAGAGTATTACAAACAAGAAGCTTCAAATTTATTACA
AGAATCAAACCTGCTCACAGTATATGGAAAAGGTTCTAGGTAGATTA AAAAGATG
AAGAAATTCGATGTCGAAAATACCTACATCCAAGTTCATATACTAAGGTGATT
CATGAATGTCAACAACGAATGGTAGCAGACCACTTACAGTTTTTACATGCAGA
ATGTCATAATATAATTCGACAAGAGAAAAAAAATGACATGGCAAATATGTACG
TCTTACTCCGTGCTGTGTCCACTGGTTTACCTCATATGATTCAGGAGCTGCAA
AACCACATCCATGATGAGGGCCTTCGAGCAACCAGCAACCTTACTCAGGAAAA
CATGCCAACACTATTTGTGGAGTCAGTTTTGGAAGTGCATGGTAAATTTGTTT
AGCTTATCAACACTGTTTTGAATGGTGATCAGCATTTTATGAGTGCCTGGAT
AAGGCCCTTACGTCAGTTGTAAATTACAGAGAACC TAAGTCTGTTTGCAAAGC
ACCTGAACTGCTTGCTAAGTACTGTGACAACCTTACTGAAGAAGTCAGCGAAAAG
GGATGACAGAGAATGAAGTGGAAAGACAGGCTCACGAGCTTTCATCACAGTGTTT
AAATACATTGATGACAAGGACGCTTTTCAAAGTTCTACGCAAGAATGCTGGC
AAAACGTTTAATTCATGGGTATCCATGTCTATGGACTCTGAAGAAGCCATGA
TCAACAAATTAAGCAAGCCTGTGGTTATGAGTTTACCAGCAAGCTACATCGG
ATGTATACAGATATGAGTGTGAGCGCTGATCTCAACAATAAGTTCAACAATTT
TATCAAAAACCAAGACACAGTAATAGATTTGGGAATTAGTTTTCAAATATATG
TTCTACAGGCTGGTGCCTGCTTACTCAGGCTCCTTCATCTACGTTTGCA
ATTCCCAGGAATTAGAAAAAGTGTACAGATGTTTGAATTATTTTATAGCCA
ACATTTTCAGTGAAGGAACTTACATGGTTACATTATCTGTGTACAGGTGAAG
TTAAAATGAACTATTTGGGCAAACCATATGTAGCCATGGTTACAACATACCAA
ATGGCAGTTCTTCTTGCCTTTAACAACAGTGAAACTGTCAGTTATAAAGAGCT
TCAGGACAGCACTCAGATGAATGAAAAGGAACTGACAAAAACAATCAAATCAT
TACTTGATGTGAAAATGATTAACCATGATTCAGAAAAGGAAGATATTGATGCA
GAATCTTCGTTTTTCATTAATATGAACTTTTAGCAGTAAAAGAACA AAAATTTAA
AATTA CTACATCAATGCAGAAAGACACACCACAAGAAATGGAGCAGACTAGAA
GTGCAGTTGATGAGGACCGAAAATGTATCTCCAAGCTGCTATAGTTTCGTATC
ATGAAAGCACGAAAAGTGCTTCGGCACAATGCCCTTATTCAAGAGGTGATTAG
CCAGTCAAGAGCTAGGTTTAATCCCAGTATCAGCATGATTAAGAAGTGTATTG
AAGTTCTGATAGACAAACAATACATAGAACGCAGCCAGGCGTCGGCAGATGAA
TACAGCTACGTCGCGTGAGCGGCCGC

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Clone Data Sheet

Protein name

<u>Protein</u>	Rbx1 1-108 (full length)
<u>Synonyms</u>	RNF75, ROC1
<u>Clone Number</u>	DU23263
<u>Species</u>	Human
<u>Accession Number</u>	P62877
<u>Tags</u>	N/A
Aminoacid sequence of the purified Rbx1	MAAAMDVDTPSGTNSGAGKKRFEVKKWNAVALWAWDIIVVDNCAICRNH IMDLCIECQANQASATSEECTVAWGVCNHAFHFHCISRWLKTRQVCPL DNREWEFQKYGH
Native sequence	in bold
Protease cleavage	N/A
Cloning sites	Nhe1 / Kpn1
DNA sequence of the Rbx1 insert	gctagc ATGGCGGCAGCGATGGATGTGGATACCCGAGCGGCACCAAC AGCGGCGGGCAAGAAGCGCTTTGAAGTGAAAAAGTGGAAATGCAGTA GCCCTCTGGGCCTGGGATATTGTGGTTGATAACTGTGCCATCTGCAGG AACCACATTATGGATCTTTGCATAGAATGTCAAGCTAACCAGGCGTCC GCTACTTCAGAAGAGTGTACTGTTCGCATGGGGAGTCTGTAACCATGCT TTTCACTTCCACTGCATCTCTCGCTGGCTCAAAAACACGACAGGTGTGT CCATTGGACAACAGAGAGTGGGAATTCCAAAAGTATGGGCACTAG ggt acc