# **DCNL2** [6His-tagged]

E3 Ligase

Alternate Names: C13orf17 protein, FLJ10704, FLJ20092, DCUN1D2

**Cat. No. 63-2004-025** Quantity: 25 μg **Lot. No. 30161** Storage: -70°C

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**CERTIFICATE OF ANALYSIS Page 1 of 2** 

### **Background**

The enzymes of the NEDDylation pathway play a pivotal role in the activation of the largest class of ubiquitin E3 ligases called Cullin-RING-Ligases (CRLs). Akin to ubiquitylation three classes of enzymes are involved in the process of mammalian NEDDylation; E1 activating enzyme (APP-BP1/ UBA3 heterodimer), E2 conjugating enzymes (UBE2M or UBE2F) and E3 ligases the defective in Cul neddylation 1 domain-containing proteins (DCUN1D1-5) (Meyer-Schaller et al., 2009; Huang et al., 2011). There are 5 human DCUN1D1-5 proteins are also named defective in Cul neddylation 1 like proteins (DCNL1-5) (Meyer-Schaller et al., 2009). Cloning of DCNL2 was first described by Kurz et al. (2005) and Lamesch et al. (2007). The DCNLs have distinct amino-terminal domains, but share a conserved C-terminal potentiating neddylation (PONY) domain (Kurz et al., 2008). It has been determined that the interaction between the DCNLs and Cul1 occurs through the PONY domain and the Winged Helix DNA binding domain (WHB) respectively (Kurz et al., 2008; Scott et al., 2011). Pairwise analysis of 30 combinations of the five DCNL PONY domains and six cullin WHB subdomains by isothermal titration calorimetry have all shown interaction albeit with differing affinities (Monda et al., 2013).

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## **Physical Characteristics**

Species: human

Source: E. coli

Quantity: 25 μg

Concentration: 0.5 mg/ml

Formulation: 50 mM HEPES pH 7.5,

150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

Molecular Weight: ~33.3 kDa

Purity: >98% by InstantBlue™ SDS-PAGE

Stability/Storage: 12 months at -70°C;

aliquot as required

#### **Protein Sequence:**

MGSSHHHHHHSSGLEVLFQGPGSPEFPGVDMH
KLKSSQKDKVRQFMACTQAGERTAIYCLTQNE
WRLDEATDSFFQNPDSLHRESMRNAVDKK
KLERLYGRYKDPQDENKIGVDGIQQFCD
DLSLDPASISVLVIAWKFRAATQCEFSRKE
FLDGMTELGCDSMEKLKALLPRLEQELKDTAK
FKDFYQFTFTFAKNPGQKGLDLEMAVAYWKLV
LSGRFKFLDLWNTFLMEHHKRSIPRDTWNLLL
DFGNMIADDMSNYDEEGAWPVLIDDFVEYAR

PVVTGGKRSLF

Tag (bold text): N-terminal His

Protease cleavage site: PreScission™ (<u>LEVLFQ▼GP</u>)
DCNL2 (regular text): Start **bold italics** (amino acid

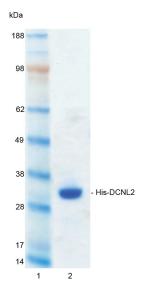
residues 1-259)

Accession number: NP\_001014305.1

# **Quality Assurance**

#### **Purity:**

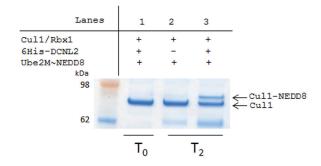
4-12% gradient SDS-PAGE InstantBlue™ staining Lane 1: MW markers Lane 2: 1 μg His-DCNL2



#### **Protein Identification:**

Confirmed by mass spectrometry.

E3 Ligase Assay: The activity of His-DCNL2 was validated through its ability to enhance the neddylation of Cul1/Rbx1 acting as a substrate in the presence of the thioester-loaded His-Ube2M~NEDD8. Incubation of Cul1/Rbx1 and thioester loaded His-Ube2M~NEDD8 in the presence or absence of His-DCNL2 at  $4\,^{\circ}\mathrm{C}$  was compared at two time points  $\mathrm{T_0}$  and  $\mathrm{T_2}$  minutes. Increased neddylation of the Cul1 subunit in the presence of His-DCNL2 was demonstrated.





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Lot-specific COA version tracker: v1.0.0

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### **Background**

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#### References:

Huang G, Kaufman A J, Ramanathan Y, Singh B (2011) SCCRO (DCUN1D1) promotes nuclear translocation and assembly of the neddylation E3 complex, J Biol Chem 286, 10297-10304.

Kurz T, Chou YC, Willems AR, Meyer-Schaller N, Hecht ML, Tyers M, Peter M, Sicheri F. (2008) Dcn1 functions as a scaffold-type E3 ligase for cullin neddylation, *Mol Cell* **29**, 23-35.

Kurz T, Ozlü N, Rudolf F, O'Rourke SM, Luke B, Hofmann K, Hyman AA, Bowerman B, Peter M. (2005) The conserved protein DCN-1/Dcn1p is required for cullin neddylation in C. elegans and S. cerevisiae, Nature 435, 1257-1261.

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Meyer-Schaller N, Chou YC, Sumara I, Martin DD, Kurz T, Katheder N, Hofmann K, Berthiaume LG, Sicheri F, Peter M. (2009) The human Dcn1-like protein DCNL3 promotes Cul3 neddylation at membranes, *Proc Natl Acad Sci U S A* **106**, 12365-12370.

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Scott D.C. Monda JK Bennett F.I. Harner JW Schulman B.A. (2011) N-terminal acetylation acts as an avidity enhancer within an interconnected multiprotein complex, Science 334, 674-678.



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