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## **NEDD4: The founding member of a family of ubiquitin-protein ligases**

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## **Abstract**

Ubiquitination plays a crucial role in regulating proteins post-translationally. The focus of this review is on NEDD4, the founding member of the NEDD4 family of ubiquitin ligases that is evolutionarily conserved in eukaryotes. Many potential substrates of NEDD4 have been identified and NEDD4 has been shown to play a critical role in the regulation of a number of membrane receptors, endocytic machinery components and the tumour suppressor PTEN. In this review we will discuss the diverse pathways in which NEDD4 is involved, and the patho-physiological significance of this important ubiquitin ligase.

## **Keywords**

Ubiquitin; Ubiquitin protein ligases; NEDD4 family; IGF signaling; PTEN

## **1 Introduction**

Ubiquitination is a post-translational protein modification that is critical for a number of cellular processes. Ubiquitination involves the covalent attachment of the 8kDa protein ubiquitin to one or more lysine residues in the substrate protein to signal proteins for degradation, altered localisation, trafficking or function. Substrate proteins can be monoubiquitinated, multi-monoubiquitinated or poly-ubiquitinated, with the type of ubiquitination determining the fate of the protein. Ubiquitin itself has seven lysine residues, allowing for different ubiquitin linkage types; for example the well-studied K48-linkage typically targets proteins for proteasomal degradation (Hershko and Ciechanover, 1998) whereas K63 linkages are associated with protein trafficking and lysosomal degradation (Hicke and Dunn, 2003).

Ubiquitin is covalently attached to a protein substrate via an energy dependent three step process, involving an E1 ubiquitin activating enzyme, an E2 ubiquitin conjugating enzyme and an E3 ubiquitin protein ligase. The E3 ubiquitin ligase largely determines the substrate specificity of the system and in mammals there are several hundred ubiquitin protein ligases

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(Hershko and Ciechanover, 1998). These can be grouped into two main classes; the RING (Really Interesting New Gene) E3s which mediate the direct transfer of ubiquitin to the substrate (Deshaies and Joazeiro, 2009), and the HECT (Homologous to E6-AP C-Terminus) E3s which are involved in the transfer of activated ubiquitin from the E2 to the substrate by forming an intermediate complex with the C-terminus of the E3 (Rotin and Kumar, 2009). This review will focus on the HECT type ubiquitin ligase NEDD4, one of the first HECT E3 ligases discovered, and the founding member of the NEDD4 family of HECT ubiquitin ligases.

## **2 History of NEDD4 discovery**

The NEDD4 gene was cloned in 1992 as one of a number of murine Nedd (Neural precursor cell expressed developmentally down-regulated) genes differentially expressed in the central nervous system (Kumar et al., 1992). At the time of its cloning, the predicted protein had only one known domain – an N-terminal calcium/lipid-binding domain (C2 domain). The presence of three partial repeats of approximately 40 amino acids containing two conserved tryptophan residues in the middle part of the protein was also noted. These repeats, now known to occur in numerous proteins, are widely known as WW domains (Bork and Sudol, 1994). Subsequently, in the following year the C-terminal region of NEDD4 was found to be similar to human E6-AP, the papilloma virus oncoprotein E6-associated protein. E6-AP was the first discovered ubiquitin-protein ligase and it was shown to be involved in the E6 mediated ubiquitination of p53 (Scheffner et al., 1993). The C-terminus of E6-AP comprising the catalytic domain was named HECT (homologous to the E6-AP C-terminus) (Huibregtse et al., 1995). E6-AP became the founding member of the HECT type of E3 ubiquitin ligases, of which now there are 29 human members. In yeast the first HECT ligase was Rsp5p/Npi1p from *Saccharomyces cerevisiae*, which was originally discovered as a suppressor of mutations in the *SPT3* gene (Huibregtse et al., 1995). NEDD4 and similar proteins discovered subsequently became a family of HECT ligases, comprising 9 human proteins including NEDD4, NEDD4–2 (NEDD4L), ITCH, SMURF1, SMURF2, WWP1, WWP2, NEDL1 AND NEDL2 (Scheffner and Kumar, 2014). Rsp5 is also a member of the NEDD4 family, suggesting that these ligases are highly conserved.

## **3 NEDD4 orthologues and structure**

As mentioned above, NEDD4 is a highly evolutionarily conserved protein from yeast to man, and was initially cloned as a highly expressed gene in the early embryonic brain (Kumar et al., 1992; Kumar et al., 1997). There are 94 orthologues of NEDD4 in the NCBI database, all sharing the same modular structure consisting of an N-terminal C2 domain, 3–4 WW domains and a C-terminal catalytic HECT domain for ubiquitin protein ligation (Harvey and Kumar, 1999) (Figure 1A). The C2 domain is a calcium-dependent lipidbinding domain around 116 amino acids in length that targets proteins to phospholipid membranes (Dunn et al., 2004), and can also be involved in protein-protein interactions (Morrione et al., 1999; Plant et al., 2000). The C2 domain-mediated membrane translocation is required for some cellular functions of NEDD4 (Dunn et al., 2004). The WW domains are protein-protein interactions domains, usually around 40 amino acids in length, containing two conserved tryptophans (W) residues that are 21 amino acids apart (Bork and Sudol,

1994). The WW domains interact with proline rich PPxY (PY) motifs and can also interact with phospho-serine/threonine residues in substrates (Sudol et al., 1995). The number of WW domains can vary between NEDD4 family members, and also between species i.e. the NEDD4 protein in human, chicken and Xenopus contains four WW domains, whereas mouse, zebrafish, *Drosophila* and yeast contains three WW domains (Yang and Kumar, 2010) (Figure 1B). The HECT domain is a highly conserved domain that comprises around 350 amino acids, and contains a conserved cysteine residue that forms an intermediate thioester bond with the activated ubiquitin accepted from an E2, before catalysing the ubiquitination of a lysine in the substrate protein (Rotin and Kumar, 2009). There are a number of E2s that are able to transfer ubiquitin to NEDD4, including Ubc4, UbcH5B, UbcH5C, UbcH6 and UbcH7 (Anan et al., 1998; Fotia et al., 2006).

The human NEDD4 gene is located on chromosome 15q21.3 and comprises 30 exons (HGNC:7727) shown to encode a ~120 KDa protein. There are five NEDD4 protein variants in the NCBI database, all of which share 100% identity from the first WW domain through to the end of the protein, only varying in the N-terminal region which includes the C2 domain. Recently it was reported that there is a 75kDa NEDD4 isoform found exclusively in myotonic dystrophy type 2 muscle in addition to full length NEDD4 (Screen et al., 2014). NEDD4 protein is localised to the cytoplasm, mainly in the perinuclear region and cytoplasmic periphery of human cultured cells (Anan et al., 1998). NEDD4 is also found in exosomes when recruited by the NEDD4 family interacting protein Ndfip1 (Putz et al., 2008).

## **4 NEDD4 binding partners and targets**

A number of in vitro binding studies and proteomic approaches have been used to identify potential NEDD4 substrates (see below for a summary and Table 1 for a list of interacting proteins).

#### **4.1 Ion channels and membrane transporters**

The epithelial sodium channel (ENaC) is a transmembrane ion channel that contains a PY motif in its cytoplasmic tail. Using yeast two hybrid studies, rat NEDD4 was shown to bind to the PY motif in ENaC via its WW domains (Staub et al., 1996). Furthermore, in response to increased intracellular sodium, NEDD4 binds and ubiquitinates ENaC to mediate the down-regulation of sodium channel activity (Dinudom et al., 1998). However, in vivo studies indicate that NEDD4–2 is the main NEDD4 family member that is responsible for ENaC regulation (Kamynina et al., 2001; Fotia et al., 2003; Boase et al., 2011; Goel et al., 2015).

A number of ion channels found in the brain are subject to NEDD4 regulation. Alpha-Amino-3-hydroxy-5-methyl-isoxazole-4-propionic acid receptors (AMPARs) are ion channels that mediate excitatory synaptic transmission in the brain. In rat neurons, NEDD4 was shown to ubiquitinate the AMPAR GLUR1 subunit, leading to lower levels of AMPAR at the cell surface, and a decrease in synaptic transmission (Lin et al., 2011). The voltagegated calcium ion channel  $Ca<sub>v</sub>1.2$  is regulated by NEDD4, inducing down-regulation of cell surface expression at the plasma membrane to affect whole cell  $Ca<sub>v</sub>$  currents, and also directing  $Ca<sub>v</sub>1.2$  sorting from the trans-golgi network to endosomes (Rougier et al., 2011).

Voltage gated sodium channels 1.2 and 1.7 (Na<sub>v</sub>1.2 and 1.7) contain PY motifs that interact with the WW domains of NEDD4 to result in down-regulation of the channel and reduced sodium channel activity in Xenopus oocytes (Fotia et al., 2004).

#### **4.2 Growth factor signalling**

In the epidermal growth factor signalling pathway, there are a number of components that are regulated by NEDD4. The epidermal growth factor receptor (EGFR) family of tyrosine kinase members, HER3 and HER4, both undergo NEDD4-mediated ubiquitination and degradation to down-regulate signalling (Zeng et al., 2009; Huang et al., 2014). Activated Cdc42-associated tyrosine kinase (ACK) is an ubiquitin-binding protein involved in EGFR regulation. The PY motif of ACK binds the WW3 domain of NEDD4, resulting in ubiquitination and lysosomal degradation of ACK and EGFR in response to EGF stimulation (Lin et al., 2010). Epidermal growth factor receptor substrate 15 (EPS15) is a protein in the EGFR pathway that contains a ubiquitin interacting motif (UIM), and has been shown to interact with ubiquitinated NEDD4 to promote mono-ubiquitination of itself in a process termed coupled monoubiquitination (Woelk et al., 2006).

Fibroblast growth factor receptor 1 (FGFR1) has important roles in regulating cellular proliferation and differentiation. FGFR1 contains a novel site (VL\*\*\*PSR) that binds the C2 and WW3 domain of NEDD4, leading to the ubiquitination and down-regulation of FGFR1 at the cell surface, and attenuation of downstream signalling (Persaud et al., 2011). The vascular endothelial growth factor receptor-2 (VEGF-R2) interacts with NEDD4 leading to the degradation of VEGF-R2, although this degradation is not due to NEDD4-mediated ubiquitination. Grb10 acts as a positive regulator of VEGF-R2 signalling by interacting with NEDD4 to inhibit the NEDD4-mediated degradation of VEGF-R2 (Murdaca et al., 2004). The beta 2 androgen receptor (β2-AR) interacts with NEDD4 via the adaptor protein betaarrestin 2 in vitro, leading to the ubiquitination and degradation of activated β2-AR in the lysosome (Shenoy et al., 2008).

#### **4.3 Tumor suppressors**

Beclin 1 is a tumor suppressor involved in the PI3 kinase pathway. The PY motif of Beclin1 interacts with NEDD4 WW domains, leading to polyubiquitination and subsequent downregulation of Beclin1 (Platta et al., 2012). Interestingly, the polyubiquitination involved ubiquitin-K11 linkages to promote Beclin1 down-regulation, rather than the typical ubiquitin-K48 linkages. Large tumor suppressor kinase 1 (LATS1) is a serine/threonine kinase that is a negative regulator of YAP1 in the Hippo signalling pathway, and directly binds NEDD4 leading to its ubiquitination and subsequent degradation, implicating NEDD4 as an additional regulator of the Hippo pathway (Salah et al., 2013). NEDD4 has also been shown to interact with the tumor suppressor PTEN (phosphatase and tensin homologue depleted on chromosome ten) (Wang et al., 2007), as discussed further in 5.2.

#### **4.4 Endocytic regulation proteins**

 $\gamma$ 2-adaptin is a member of the heterotetrameric clathrin adaptor complex family that contains a UIM, which binds to the ubiquitinated C2 domain of NEDD4 to facilitate its own ubiquitination (Rost et al., 2008). Furthermore, the high affinity of  $\gamma$ 2-adaptin for the C2

domain of NEDD4 allows not only mono-ubiquitination of  $\gamma$ 2-adaptin, but also multi and poly-ubiquitination chains. Another UIM-containing protein, Hgs (a mammalian homologue of a yeast vacuolar sorting adaptor), interacts with NEDD4 to target substrate proteins such as the EGFR receptor for ubiquitination to induce endocytosis (Katz et al., 2002). This UIM is often found in endocytic adaptors, indicating that ubiquitination, and NEDD4, could play a general role in vesicular trafficking network.

Lysosomal associated protein transmembrane 5 (LAPTM5) is a lysosomal transmembrane protein that is expressed in hematopoietic cells, and resides in the late endosome/lysosome. The PY motifs of LAPTM5 bind the WW domains of NEDD4 to control the sorting of Laptm5 from the golgi to the lysosome, but this does not require the ubiquitination of LAPTM5 (Pak et al., 2006). In addition, The NEDD4-LAPTM5 complex recruits GGA3 (Golgi-localised, γ-ear-containing, ADP ribosylation factor binding protein), a protein involved in regulating cargo trafficking, and NEDD4 ubiquitinates GGA3 (Pak et al., 2006).

#### **4.5 NEDD4 and viral budding**

A number of viruses contain the PY late budding domain expressed within their matrix proteins, mediating interactions with the WW domains of NEDD4 to facilitate viral budding. These include the matrix proteins of Ebola virus (Harty et al., 2000), Rous sarcoma virus (Kikonyogo et al., 2001), Mason-Pfizer monkey virus (Gottwein et al., 2003), the murine leukemia virus (Segura-Morales et al., 2005), the Marburg virus (Urata and Yasuda, 2010) and Human Immunodeficiency virus (HIV) (Sette et al., 2010). NEDD4 also interacts with components of the ESCRT machinery required for viral budding. For example, the vacuolar protein sorting protein Alix recruits NEDD4 to HIV-1 Gag protein to facilitate HIV-1 release via a mechanism that involves Alix ubiquitination (Sette et al., 2010). Finally, NEDD4 binds and ubiquitinates the latent membrane protein 2A (LMP2A) of the Epstein-Bar virus to modulate B-cell signal transduction (Ikeda et al., 2000).

## **4.6 Other NEDD4 targets**

Alpha-synuclein is an abundant protein in the human brain that leads to the neurodegenerative disorder Parkinson's disease when it is misfolded and accumulates in the brain. NEDD4 recognises the c-terminus of α-synuclein and attaches K63-type ubiquitin chains, resulting in lysosomal degradation (Tofaris et al., 2011).

The WW domains of NEDD4 bind to the PY motif of the gap junction protein connexin43  $(Cx43)$  (Leykauf et al., 2006). This results in the mono-ubiquitination of  $Cx43$  and its interaction with Eps15, an adaptor protein involved in endocytosis, leading to the internalisation of Cx43 from the plasma membrane (Girao et al., 2009). Recently it was shown that the interaction between NEDD4 and Cx43 is enhanced in rat astrocytes after lipopolysaccharide stimulation (Liao et al., 2013). NEDD4 also mediates the ubiquitination of the adaptor protein dishevelled-1, and associates with Rho-GTPase Rac1 to stimulate the maturation of epithelial cell-cell contacts (Nethe et al., 2012).

Notch is an importing regulator of cell differentiation and proliferation, and with Deltex has been shown to interact with NEDD4 resulting in the ubiquitination of Notch and Deltex,

promoting their endocytosis and turnover (Sakata et al., 2004). In Drosophila, Ndfip1 acts as an adaptor protein to promote NEDD4 interaction with Notch (Dalton et al., 2011).

Spy1A is a constitutively expressed cyclin-like protein required for progression through the G1/Sphase of the cell cycle that is ubiquitinated by NEDD4 to be degraded in a cell cycle manner (Al Sorkhy et al., 2009). RNA polymerase II was shown to be ubiquitinated by NEDD4 in response to UV-induced DNA damage in human cells (Anindya et al., 2007).

The calcium-dependent cysteine protease calpain 2 undergoes NEDD4-mediated ubiquitination and degradation in response to Brucella bacterial infection in macrophages (Cui et al., 2014). The cysteinyl aspartate protease 9 (caspase-9) is directly ubiquitinated by NEDD4 in the Patched complex, which is required for Patched-mediated apoptosis (Fombonne et al., 2012). Sensitive to apoptosis gene (SAG) is an anti-apoptotic cellular survival protein that has been shown to promote cell proliferation and protect cancer cells from apoptosis. SAG was initially identified as a potential NEDD4 substrate in a proteomics study (Persaud et al., 2009), and was validated as a target by Zhou et al (2014). The RING domain of SAG interacts with NEDD4 in an atypical fashion via the HECT domain, leading to the ubiquitination and subsequent degradation of SAG (Zhou et al., 2014). NEDD4 levels are inversely correlated with protein levels of SAG in human lung cancer cell lines (Zhou et al., 2014).

## **5 Physiological functions of NEDD4**

NEDD4 is widely expressed in mammalian tissues. To investigate the physiological functions of NEDD4, a number of studies have focussed on  $NEDD4^{-/-}$  mice that lack NEDD4 expression. The first described  $NEDD4^{-/-}$  mice were neonatal lethal, with delayed embryonic development and reduced growth and body weight due (Cao et al., 2008). Consistent with this,  $NEDD4^{-/-}$  murine embryonic fibroblasts (MEFs) display reduced mitogenic activity (Cao et al., 2008). *NEDD4<sup>-/-</sup>* embryos have reduced skeletal muscle size, decreased motor neuron and axon numbers and abnormal neuromuscular junction structure and function (Liu et al., 2009).  $NEDD4^{-/-}$  embryos also have severe craniofacial defects, and are deficient in cranial neural crest cells (Wiszniak et al., 2013). Another independently generated NEDD4<sup>-/-</sup> mouse line also showed embryonic lethality, and NEDD4<sup>-/-</sup> embryos have pronounced heart defects, vasculature abnormalities and impaired dendrite development (Fouladkou et al., 2010; Kawabe et al., 2010). Recently, a conditional skeletal muscle-specific *NEDD4<sup>-/-</sup>* mouse demonstrated an important role for NEDD4 in mediating denervation-induced skeletal muscle atrophy (Nagpal et al., 2012). Due to the lethality of  $NEDD4^{-/-}$  mice, studies have been performed in  $NEDD4$  heterozygous mice, which are viable and live to maturity. NEDD4 heterozygous mice have normal motor function, but show significant age-dependent gait abnormalities (Camera et al., 2014).

#### **5.1 NEDD4 and IGF signalling**

One of the most pronounced phenotypes of the  $NEDD4^{-/-}$  mice is reduced growth, attributed to decreased insulin like growth factor-1 (IGF-1) and insulin signalling, highlighting the important role of NEDD4 in the regulation of cell proliferation, survival, differentiation and cell motility (Cao et al., 2008). Previous in vitro work showed that

NEDD4 bound to the adaptor protein Grb10 that is known to negatively regulate IGF1 signalling, to form a bridge between NEDD4 and the Insulin-like growth factor 1 receptor (IGF-1R) (Morrione et al., 1999). This NEDD4/Grb10 complex regulates the ubiquitination and stability of the IGR-1R (Vecchione et al., 2003) by mediating the multi-ubiquitination of IGF-1R, leading to receptor internalisation (Monami et al., 2008). These results suggest that NEDD4 acts as a negative regulator of IGF-1R signalling, and are inconsistent with the growth deficiency seen in the  $NEDD4^{-/-}$  mice, as described below.

 $NEDD4^{-/-}$  MEFs show reduced cell surface expression of both the Insulin Receptor (IR) and the Insulin-like growth factor 1 receptor (IGF-1R) (Cao et al., 2008). Upon ligand binding, IGF-1R undergoes auto-phosphorylation, leading to phosphorylation of substrates such as insulin receptor substrate (IRS) and the activation of the phosphoinositide 3-kinase (PI3K) and mitogen-activated protein kinase (MAPK) signalling pathways.  $NEDD4^{-/-}$ MEFS showed reduced IGF-1 induced tyrosine phosphorylation of IGF-1R and IRS-1, as well as reduced activation of Akt (PI3K pathway) and ERK (MAPK Pathway), although the total abundance of these proteins was not altered (Cao et al., 2008). Recently it was shown that defective IGF signalling in  $NEDD4^{-/-}$  MEFs, including the phosphorylation of IRS1 and Akt, could be rescued by ablation of the phosphatase PTEN (Shi et al., 2014). The role of NEDD4 in this growth hormone signalling was specific for insulin and IGF-1, and not observed with epidermal growth factor (EGF) or serum signalling (Fan et al., 2013; Shi et al., 2014). Moreover, IGF-1 signalling stimulates NEDD4 K63-type poly-ubiquitination of pAKT at the plasma membrane, without altering total AKT ubiquitination, to promote pAKT nuclear trafficking (Fan et al., 2013). The in vivo data above indicate that IGF-1 and insulin signalling require NEDD4 function (see Figure 2).

In vitro NEDD4 has been shown to bind to Grb10, a negative regulator of IGF-1 signalling (Morrione et al., 1999). *NEDD4<sup>-/-</sup>* MEFs showed increased levels of the Grb10, and simultaneous loss of NEDD4 and Grb10 restores the cell-surface levels and signalling of IGF-1R, indicating that NEDD4 acts to oppose the function of the negative regulator Grb10 (Cao et al., 2008). Furthermore *NEDD4<sup>-/-*</sup> lethality is partially rescued by the maternal inheritance of a disrupted Grb10 allele (Cao et al., 2008). The role of NEDD4 in IGF-R1 signalling may be context specific. It was recently reported that during neurodegeneration in mice, NEDD4 was upregulated in cultured neurons and this led to the ubiquitination and proteasomal degradation of IGF-IRβ receptor (Kwak et al., 2012). Other ubiquitin ligases such as Mdm2 and c-Cbl have been shown to ubiquitinate IGF-1R leading to receptor endocytosis (Sehat et al., 2008) and Mdm2 has recently been identified as substrate of NEDD4 (Xu et al., 2014). Clearly, the exact role of NEDD4 on the regulation of IGR-1R is yet to be fully elucidated.

#### **5.2 NEDD4 and PTEN**

PTEN is a tumour suppressor protein that functions as a phosphatase for PIP3 (phosphatidylinositol-3,4,5-tri-phosphate), which is an important secondary messenger of the PI3K/AKT/mTOR pathway that regulates cell survival, proliferation and differentiation. PTEN was identified as a direct target of NEDD4 in vitro, resulting in PTEN ubiquitination and subsequent proteasomal degradation (Wang et al., 2007). Mono-ubiquitination of PTEN

by NEDD4 was shown to lead to PTEN nuclear import in both murine and human cells, thereby protecting PTEN from cytoplasmic degradation (Trotman et al., 2007). Thus NEDD4 can act as a proto-oncogene to degrade PTEN, or in a tumor-suppressive role to shuttle PTEN into the nucleus to protect PTEN, as even from the nucleus PTEN is still able to antagonize the AKT pathway and cause apoptosis (Trotman et al., 2007).

The role of NEDD4 in the regulation of PTEN in vivo however remains somewhat controversial. Increased NEDD4 expression leads to increased PTEN degradation in various human cancer cell lines overexpressing K-ras or treated with EGF (Zeng et al., 2014). In human melanoma cells, in the presence of indole-3-carbinol (I3C), stabilization of PTEN results from disrupting NEDD4 binding, thereby preventing the ubiquitination and proteasomal degradation of PTEN (Aronchik et al., 2014). Similarly, Rak functions as a tumour suppressor by phosphorylating PTEN to block NEDD4 binding, therefore stabilizing PTEN (Yim et al., 2009). NEDD4 co-localises with PTEN within sensory neurons in vivo and *in vitro*, and also within regenerating axons at an injury site (Christie et al., 2012). NEDD4 RNAi knockdown supports the role of NEDD4 mediated PTEN degradation in axon growth in Xenopus retinal ganglion cells (Drinjakovic et al., 2010) and in rat dorsal root ganglion cells (Christie et al., 2012). However, PTEN protein levels are equivalent in wildtype and  $NEDD4^{-/-}$  MEFs (Cao et al., 2008; Fouladkou et al., 2008), and PTEN stability and nuclear import are not affected in  $NEDD4^{-/-}$  MEFs (Fouladkou et al., 2008). In conditional neuronal-specific *NEDD4<sup>-/-</sup>* mice axon growth was affected by the lack of NEDD4, without changes in the activity, ubiquitination levels, or localisation of PTEN (Hsia et al., 2014). PTEN has been shown to interact with other E3 ligases (XIAP, WWP2, CHIP), so perhaps the interaction between NEDD4 and PTEN is context specific. Indeed, NEDD4 function in PTEN regulation may primarily occur in pathological contexts such as cancer or cellular stress.

#### **5.3 NEDD4 in the neuron**

Given that NEDD4 was originally identified as a neuronally expressed, developmentally downregulated gene, it is not surprising that NEDD4 is highly expressed in the central nervous system and plays an important physiological role in neuronal development (recently reviewed in (Donovan and Poronnik, 2013)). Using both conventional and neuron-specific conditional *NEDD4<sup>-/-</sup>* mouse models, NEDD4 was identified as an important regulator of dendrite formation and arborisation in both hippocampal and cortical neurons (Kawabe et al., 2010). NEDD4 forms a complex with the Traf2 and NCK Interacting Kinase (TINK), and Ras-related protein 2A (Rap2A), a negative regulator of dendritogenesis, and this trimeric complex formation results in the NEDD4 dependent mono- and di-ubiqutination of Rap2A to reduce downstream signalling and promote dendrite growth and arborisation (Kawabe et al., 2010). NEDD4-mediated ubiquitination and subsequent down-regulation of PTEN in *Xenopus* root ganglion cell axons also promotes axon branching in vivo (Drinjakovic et al., 2010). Upon zinc-mediated damage of the central nervous system, NEDD4 ubiquitinates PTEN to mediate the neuronal response (Kwak et al., 2010). Similarly during the regenerative response to axonal injury, NEDD4 associates with PTEN in dorsal root ganglion neurons to promote nerve regeneration (Christie et al., 2012). Recently NEDD4-binding protein 3 (N4BP3) has also been implicated in axonal and dendritic

branching in developing neurons (Schmeisser et al., 2013). NEDD4 is also required for the normal formation and functioning of neuromuscular junctions, normal motor neuron and axon numbers and cranial neural crest (Liu et al., 2009; Wiszniak et al., 2013).

#### **5.4 NEDD4 and T-cell function**

The NEDD4 family E3s NEDD4 and Itch are both expressed in T-cells (Heissmeyer et al., 2004) and although in vitro data shows these ligases share a number of substrates involved in T-cell regulation, *Itch* and *NEDD4* knockout mouse models display unique phenotypes, suggesting discrete functions in vivo (Fang et al., 2002; Yang et al., 2008). Fetal liver chimeras that lack NEDD4 expression in cells of haemopoietic origin show that NEDD4 is not required for T-cell development, or for initial T-cell antigen receptor mediated activation events (Yang et al., 2008). However  $NEDD4^{-/-}$  mice have fewer effector T-cells, and in response to antigen immunization T-cells lacking NEDD4 proliferate poorly, and produce less interleukin 2, suggesting the role of NEDD4 is to convert naïve T-cells into activated Tcells. The hypo-responsiveness of  $NEDD4^{-/-}$  T-cells can be explained by the impaired ubiquitination and degradation of Cbl-b, a ubiquitin ligase that plays a critical role in T-cell activation and tolerance induction, as NEDD4 is required for the poly-ubiquitination of Cblb (Magnifico et al., 2003; Yang et al., 2008). Recently it was shown that Cbl-b inhibits T-cell activation by impeding the association of NEDD4 with PTEN in T-cells to suppress PTEN inactivation (Guo et al., 2012). In addition NEDD4 is not required for B-cells to become activated, but *NEDD4<sup>-/--</sup>* T-cells are unable to provide adequate help for B-cells to undergo immunoglobulin class switching (Yang et al., 2008).

#### **5.5 NEDD4 and cancer**

NEDD4 is frequently overexpressed in many different types of cancer (for reviews see (Chen and Matesic, 2007; Ye et al., 2014)). NEDD4 was first described as a proto-oncogene for its role in negatively regulating the tumor suppressor PTEN via ubiquitination in vitro (Wang et al., 2007). An inverse correlation between (increased) NEDD4 and (decreased) PTEN has been observed in many human cancer cell lines, including breast cancer MDA-MB-231 and prostate cancer DU145 cell lines (Liu et al., 2014). After overexpressing K-ras or EGF treatment, increased NEDD4 levels and PTEN degradation are observed in various type of human cancer cell lines including cervical adenocarcinoma HeLa, colorectal adenocarcinoma HT-29, gastric adenocarcinoma BGC-823 and hepatocellular carcinoma HepG2 (Zeng et al., 2014). Given the lack of NEDD4 regulation of PTEN in  $NEDD4^{-/-}$ mice, this led to the hypothesis that perhaps NEDD4-mediated PTEN degradation primarily occurs in cancer cells under certain oncogenic circumstances (Zeng et al., 2014). Furthermore, the CDK-4 binding partner p34 has been identified as an interactor of NEDD4 in cancer cells lines, and co-expression of p34 and NEDD4 is correlated with lowered PTEN levels in colon cancer tissues, suggesting that NEDD4 positively regulates tumorigenesis via the p34-dependent PTEN proteasomal degradation (Hong et al., 2014). Contrary to this, there is also evidence of NEDD4 overexpression in cancer that promotes cell growth independent of PTEN signalling, such as in human colorectal cancer lines HCT-15 and LoVo (Eide et al., 2013).

Decreased levels of NEDD4 can also be associated with cancer. NEDD4 directly ubiquitinates oncoproteins N-Myc in neuroblastoma and c-Myc in pancreatic cancer cells to target these Myc proteins for proteasomal degradation (Liu et al., 2013). The histone deacetylase SIRT2 enhances expression of N-Myc and c-Myc by directly binding to the NEDD4 promoter and repressing NEDD4 gene expression by deacetylating histone H4 lysine 16 (Liu et al., 2013). Importantly, NEDD4 gene expression could be reactivated by the addition of SIRT2 inhibitors, resulting in reduced N-Myc and c-Myc protein expression, and suppressing neuroblastoma and pancreatic cancer cell proliferation (Liu et al., 2013). An inverse relationship between protein levels of (low) NEDD4 and (high) HER3 (an EGFR tyrosine kinase member) is observed in ductal cells of prostate cancer tumors compared to surrounding tissues, and knockdown of NEDD4 in human prostate and breast cancer cell lines leads to increased HER3-mediated cell migration and proliferation in vitro, and xenoplant tumor growth in vivo (Huang et al., 2014).

## **6 Regulation of NEDD4**

#### **6.1 NEDD4 auto-inhibition**

NEDD4 activity is in part regulated by auto-inhibition. In the absence of calcium, an autoinhibitory conformation of NEDD4 is formed by the C2 domain of NEDD4 binding to the HECT domain via intramolecular interactions, thereby inhibiting the enzymatic activity of NEDD4 (Wang et al., 2010). In the presence of calcium, the binding of the C2 domain to the HECT domain is disrupted and the C2 domain recruits NEDD4 to the lipid membrane promoting its ubiquitin ligase activity (Wang et al., 2010). Proteins or compounds that prohibit this auto-inhibitory conformation of NEDD4 serve as activators. The adaptor proteins Ndfip1 and Ndfip2 can stimulate NEDD4 activity by binding to the WW domains of NEDD4 via their PY motifs to release the auto-inhibitory conformation between the C2 and HECT domains of NEDD4 (Mund and Pelham, 2009). Recently, the c-Src kinase was demonstrated to phosphorylate NEDD4 at specific tyrosine residues Y43 (in the C2 domain) and Y585 (in the HECT domain), resulting in the disruption of the auto-inhibitory conformation of NEDD4 and consequently activating NEDD4 ubiquitin ligase activity (Persaud et al., 2014).

Conversely, stabilizing the C2 and HECT domain auto-inhibitory conformation would lead to an inhibition of NEDD4 activity, as has been proposed by in silico molecular modelling for the chemical indole-3-carbinol that binds to the HECT domain of NEDD4 (Aronchik et al., 2014). Increasing intracellular calcium can also act to release the auto-inhibitory conformation of NEDD4, for example gram-negative Brucella infection induces increases in NEDD4 activity in an intracellular calcium-dependent manner (Cui et al., 2014).

#### **6.2 Ndfip adaptors**

In addition to the activating roles of Ndfips in abrogating the auto-inhibitory conformation of NEDD4 (see above), these adaptors can facilitate substrate binding to target proteins that lack PY motifs (Foot et al., 2008; Foot et al., 2011). The Ndfip proteins contain 3 PY motifs with which they interact with the WW domains of NEDD4 (Shearwin-Whyatt et al., 2006).

Ndfip1 is also involved in exosomal secretion, and overexpression of Ndfip1 results in NEDD4 being recruited to the exosome (Howitt et al., 2009).

#### **6.3 Oxidative stress**

In the presence of neurotoxins that elicit oxidative stress in neurons, such as camptothecin, hydrogen peroxide and zinc, NEDD4 protein expression was up-regulated (Kwak et al., 2012). Reactive oxygen species (ROS) induces up-regulation of NEDD4 in primary rat cortical neurons after zinc treatment, and pre-treatment of neurons with antioxidants prevents the zinc-induced NEDD4 up-regulation. The oxidative stress induces NEDD4 transcriptional activation, with FOXM1B (Forkhead box protein M1B) identified as the key transcription factor mediating this ROS response (Kwak et al., 2012). The transcription factor FoxM1 is a member of the Forkhead box (Fox) family, and has been shown to bind directly to the endogenous human Nedd4 promoter at two FoxM1 binding sites to result in up-regulation of Nedd4 expression (Dai et al., 2010).

#### **6.4 Phosphorylation**

NEDD4 protein stability is governed by the  $SCF<sup>β-TRCP</sup>$  (Skp, Cullin, F-box containing complex, a multi-protein E3 ligase complex) (Liu et al., 2014). NEDD4 bound both Cullin1 and the F-box β-TRCP component of the SCF directly, and depletion of endogenous Cullin1 or β-TRCP led to the up-regulation of NEDD4 protein in human cells. Furthermore, Casein Kinase Iδ was identified as the kinase that phosphorylated NEDD4 at both S347 and S348 to trigger its interaction with β-TRCP and lead to the ubiquitination and degradation of NEDD4 (Liu et al., 2014). As mentioned above, NEDD4 phosphorylation by c-Src enhances its catalytic activity (Persaud et al., 2014).

#### **6.5 Other NEDD4 regulators**

The interferon inducible ISG15 is a small ubiquitin-like protein that negatively regulates NEDD4 by binding directly to NEDD4 to block its interaction with the conjugating E2 enzyme, thus preventing the ubiquitin transfer from the E2 to NEDD4 (Malakhova and Zhang, 2008). Another inhibitor of NEDD4, heclin (HECT ligase inhibitor) was recently identified that broadly inhibits HECT ligases in cells by binding to the HECT domain to cause a conformational change that allows the oxidation of the active Cys site, blocking formation of a Ub-E3 thioester bond (Mund et al., 2014) The p34 protein was recently identified as interacting with the WW1 domain of NEDD4 to enhance stability of NEDD4 by inhibiting NEDD4 auto-ubiquitination and subsequent proteasomal degradation (Hong et al., 2014). Similarly, both wild-type and catalytically inactive Cbl-b ligase inhibited NEDD4 auto-ubiquitination to unexpectedly enhances NEDD4 ubiquitination activity (Guo et al., 2012). These data suggest that Cbl-b regulates NEDD4 ubiquitin ligase activity independently of its RING finger domain.

The proto-oncogene Np63a negatively regulates NEDD4.  $p63^{-/-}$  MEFs display increased NEDD4 protein, and silencing of ΔNp63α in adult human keratinocytes led to increased transcript and protein levels of NEDD4 (Leonard et al., 2013)

Recently it was demonstrated that PTEN negatively regulates NEDD4 expression at the translational level in mouse hippocampal neurons by antagonising the PI3K pathway (Hsia et al., 2014). Another recent study showed Ras signalling up-regulates NEDD4 transcription (Zeng et al., 2014). It was observed in HEK293T cells that over-expressing K-Ras or treatment with EGF significantly increased endogenous protein levels of NEDD4 by enhancing transcription of NEDD4, indicating a negative feedback loop between Ras signalling and NEDD4 (Zeng et al., 2014).

## **7 Conclusions**

NEDD4 is the founding member of the NEDD4 family of ubiquitin protein ligases that function in the ubiquitin proteasome system. NEDD4 has many important cellular functions, as indicated by the high degree of evolutionary conservation observed across many species. NEDD4 has roles in regulating viral budding, IGF-1 signalling, in T-cell function and in PTEN signalling, although this is yet to be fully understood as there are currently a number of conflicting models. The role of NEDD4 in cancer suggests that NEDD4 could be a potential therapeutic target for the treatment of human cancer.

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#### **Figure 1. (A) Schematic structure of the NEDD4 protein**

Schematic of the modular structure of the human NEDD4 protein. The C2 calcium/ phospholipid binding domain mediates NEDD4 binding to membranes, and is also involved in substrate recognition. The WW domains are protein-protein interaction domains that bind to conserved PY motifs in substrates and regulatory proteins. The catalytic ubiquitin ligase domain binds the E2 conjugation enzyme and forms a thioester bond with ubiquitin before transferring ubiquitin to the substrate.

**(B) Phylogenetic relationship of NEDD4 proteins from various species**. NEDD4 sequences were obtained from the NCBI protein database as follows; *S. pombe*  (Schizosaccharomyces pombe Pub3; NP\_595793.1), *S. cerevisiae* (Saccharomyces cerevisiae Rsp5p; AAC03223.1), **mosquito** (Anopheles gambiae; XP\_003436401.1), *Drosophila* [Drosophila melanogaster; NP\_996116.1), *Xenopus* (Xenopus laevis; NP\_001084258.1), **zebrafish** (Danio rerio; NP\_001029358.1), **chicken** (Gallus gallus; XP\_413791.3), **mouse** (Mus musculus; NP\_035020.2), **rat** (Rattus norvegicus; NP\_037118.1), **monkey** (Macaca fascicularis; XP\_005559683.1), **chimpanzee** (Pan troglodytes; XP\_523083.3),and **human** (Homo sapiens; NP\_006145.2 ). Sequences were aligned using NCBI COBALT (Constraint Based Multiple Protein Alignment Tool) [http://](http://www.stva.ncbi.nlm.nih.gov/tools/cobalt/re_cobalt.cgi) [www.stva.ncbi.nlm.nih.gov/tools/cobalt/re\\_cobalt.cgi](http://www.stva.ncbi.nlm.nih.gov/tools/cobalt/re_cobalt.cgi) (Papadopoulos and Agarwala, 2007) and the minimum-evolution Phylogenetic Tree output displayed. The individual domains on the NEDD4 schematics were identified using NCBI Conserved Domain Database search <http://www.ncbi.nlm.nih.gov/Structure/bwrpsb/bwrpsb.cgi>(Marchler-Bauer et al., 2011) and are drawn roughly to scale. The scale bar indicates evolutionary distance (Grishin, 1995).



#### **Figure 2. The proposed role of NEDD4 in IGF-1R signaling**

Stimulation of the IGF-1R with IGF-1 ligand results in auto-phosphorylation of the receptor, leading to phosphorylation of insulin receptor substrate 1 (IRS1) and the activation of the PI3K pathway that ultimately results in cellular growth and proliferation. PTEN has recently been shown to be a protein phosphatase for IRS-1, and NEDD4 antagonizes this phosphatase activity (Shi et al., 2014). IGF-1 signalling stimulates the ubiquitination of pAKT by NEDD4 to promote the peri-nuclear trafficking of pAKT (Fan et al., 2013). The adaptor protein Grb10 negatively regulates IGF-1 signalling by binding the IGF-1R, and in vivo data indicates that NEDD4 down-regulates Grb10 expression (Cao et al., 2008). IGF-1R is also targeted by ubiquitin ligases such as MDM2 and c-Cbl for endocytosis and degradation.

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## Potential binding partners of NEDD4



