



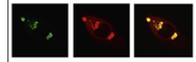
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Review

The role of the NG2 proteoglycan in OPC and CNS network function

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ABSTRACT

In the normal mammalian CNS, the NG2 proteoglycan is expressed by oligodendrocyte precursor cells (OPC) but not by any other neural cell-type. NG2 is a type-1 membrane protein, exerting multiple roles in the CNS including intracellular signaling within the OPC, with effects on migration, cytoskeleton interaction and target gene regulation. It has been recently shown that the extracellular region of NG2, in addition to an adhesive function, acts as a soluble ECM component with the capacity to alter defined neuronal network properties. This region of NG2 is thus endowed with neuromodulatory properties. In order to generate biologically active fragments yielding these properties, the sequential cleavage of the NG2 protein by α - and γ -secretases occurs. The basal level of constitutive cleavage is stimulated by neuronal network activity. This processing leads to 4 major NG2 fragments which all have been associated with distinct biological functions. Here we summarize these functions, focusing on recent discoveries and their implications for the CNS.

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1. Origin and structure of the NG2 proteoglycan

The NG2 proteoglycan is coded by the CSPG4 gene and belongs to the protein family of chondroitin sulfate proteoglycans (CSPGs). NG2 was first discovered in 1977 in a screen for neuronal and glial antigens in the rat (Stallcup, 1977). Homologs have been reported so far from mouse (Niehaus et al., 1999; Schneider et al., 2001; Stegmuller et al., 2002), human (Pluschke et al., 1996) and drosophila (Estrada et al., 2007; Schnorrer et al., 2007). Oligodendrocyte precursor cells (OPC) express the NG2 protein, while it is absent from other neural cells such as neurons (Clarke et al., 2012; Karam et al., 2008), astrocytes (Huang et al., 2014; Zhu et al., 2008) and resident microglia (Moransard et al., 2011). Within the oligodendrocyte lineage NG2 is down-regulated with ongoing differentiation of OPC into myelinating mature oligodendrocytes (De Biase et al., 2010; Kukley et al., 2010; Nishiyama et al., 2009). OPC are thus the exclusive source of NG2 amongst neural cells. OPC make up a stable cell population at all stages of development in gray and white matter, in the adult mammalian brain they make up around 5% of total neural cells (Dawson et al., 2003; Gallo et al., 2008). In addition to expression within the normal CNS, NG2 is expressed by populations of glioma cells, including the most aggressive glioblastoma (Al-Mayhany et al., 2011; Chekenya et al., 2008; Persson et al., 2010). Interestingly, OPC have been identified as constituting the cells of origin for gliomas (Liu et al., 2011) and NG2 seems to play an important role in likelihood for an OPC to become a tumor cell (Sugiarto et al., 2011). Subpopulations of pericytes, cells of the vascular system, also express NG2 (Ozerdem et al., 2002; You et al., 2014).

NG2 is a type-1 membrane protein with a protein core of 252 kD molecular weight (MW), the glycosylated form that is extractable from *in vivo* neural samples has a MW of 300 kD and exhibits a broad band pattern around this size on SDS gels, typical for a glycoprotein containing complex sugars. Glycosylation of the extracellular part yields one confirmed glycosaminoglycan (GAG-) chain (O-glycosylation) (Stallcup and Dahlin-Huppe, 2001), with several predicted N-glycosylation sites (Nishiyama et al., 1991). The extracellular part of 2200 amino acids (aa), makes up over 95% of the protein, while the transmembrane (TM) domain comprises 21 aa and the intracellular part 77 aa. Two N-terminal Laminin Neurexin Sex-Hormone Binding Globulin (LNS) domains represent conserved protein domains which recently have been shown to exhibit neuromodulatory properties (Sakry et al., 2014). These NG2 LNS domains are conserved throughout species from humans to drosophila (Rudenko et al., 2001). In the ensuing text we focus on summarizing functions of the intracellular part of NG2 including the binding partners, as well as novel functions of the extracellular part of NG2. We first provide an overview of the latest discoveries related to proteolytic processing of NG2, as this is essential to yield the biological functions of the protein.

2. Cleavage of NG2

Soluble forms of the NG2 ectodomain extractable from tissue with salt solutions (290 kD and 275 kD) were initially reported

in 1995 (Nishiyama et al., 1995) and confirmed by others (Deepa et al., 2006; Morgenstern et al., 2003). We recently showed that the full-length (FL) NG2 protein (300 kD) is processed sequentially by the α -secretase ADAM10 and subsequently by the γ -secretase complex Fig. 1. Both these enzymes are expressed by OPC. α -secretase cleavage leads to an ectodomain of around 290 kD which can be released from the cell and a matching c-terminal fragment (CTF) of 12 kD. The membrane-bound CTF can be further processed by the γ -secretase complex releasing the intracellular part from the membrane stump: this released intracellular domain is termed the ICD (8.5 kD) (Sakry et al., 2014). Similar cleavage cascades are known for other type-1 membrane proteins such as Notch, N-Cadherin, Neuroligin, L1 or APP, involving the same or similar proteases. The NG2 FL, the ectodomain, the CTF and the ICD are the major forms of NG2 known to date.

A new focus was given to the biological relevance of these cleavage events when it was shown that α -secretase mediated cleavage of neuronal surface proteins such as neuroligin-1 and N-cadherin was stimulated by neuronal activity (Malinverno et al., 2010; Suzuki et al., 2012). We have recently shown that NG2 cleavage by α - and γ -secretases occurs constitutively and can be increased by neuronal activity acting on OPC (Sakry et al., 2014). This is especially interesting since OPC are unique glia entering into synaptic contact with the neuronal network in all major areas of the brain (Bergles et al., 2000; Jabs et al., 2005; Kukley et al., 2008; Mangin and Gallo, 2011). This implies that NG2 cleavage-dependent functions can be modulated by the neuronal network (Sakry et al., 2014). An important unresolved question for all such cleavage events is the mechanism of activation of the α -secretase (Sonderegger and Matsumoto-Miyai, 2014).

OPC are likely to play multiple complex roles in different types of CNS lesion, which are at present incompletely understood. It has been established that together with microglia OPC constitute the cellular response of resident CNS cells within the first few days after injury (Buffo et al., 2005; Dimou et al., 2008). The response of OPC to lesions involves increased proliferation, migration and differentiation (Simon et al., 2011) and has largely been interpreted as a response of the oligodendrocyte lineage to damage of myelin; little is known about their role in the glial scar (Buss et al., 2009; Honsa et al., 2012; Vadivelu et al., 2015). Furthermore OPC have been associated with injury-dependent increase of NG2 cleavage from the OPC (Jones et al., 2002; Levine, 1994; Morgenstern et al., 2003). Some studies report identified microglia with NG2 surface staining which is sometimes purported to be lesion-dependent expression of NG2 by microglia (Jones et al., 2002; Zhu et al., 2012), but may in fact be bound OPC-derived NG2 ectodomain. A detailed lineage-tracing study reported that CNS resident macrophages (microglia) do not express NG2 while invading macrophages from the blood can be NG2-positive (Moransard et al., 2011).

3. Intracellular functions of the NG2 proteoglycan

The intracellular part of NG2 (77 aa) can be cleaved by γ -secretase activity yielding the NG2 ICD (Sakry et al., 2014).

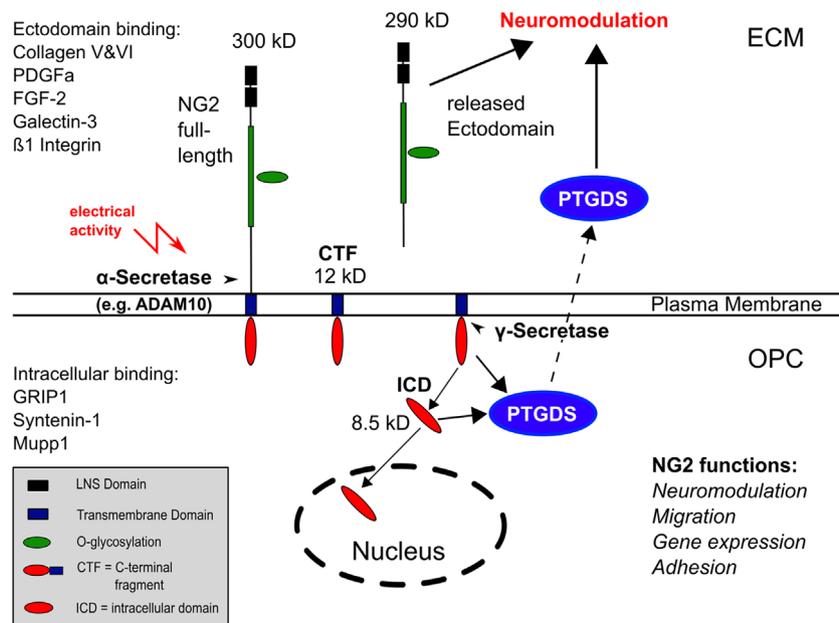


Fig. 1 – The role of the OPC-expressed NG2 protein in the CNS. The type-1 membrane protein NG2 (300 kD) is located at the OPC surface. Sequential cleavage of NG2 by α - and γ -secretases, leads to a released ectodomain (around 290 kD) and membrane bound c-terminal fragment (CTF, 12 kD). The CTF can then be further processed by the γ -secretase releasing the so called ICD (intracellular domain of 8.5 kD). Activity of the α -secretase ADAM10 on NG2 can be increased by electrical activity of the neuronal network via the neuron–glia synapse. The CTF and ICD modulate expression of the secreted enzyme PTGDS, which has reported neuromodulatory functions and has recently been shown to be expressed by OPC in an NG2-dependent manner. The LNS domains of the NG2 ectodomain alter defined glutamatergic synaptic properties of neuron–neuron synapses.

This region of the protein contains two phosphorylation sites, Thr-2256 and 2314, the first is a target for PKC α and the second for ERK (Makagiansar et al., 2007). The C-terminus contains a PSD95/DiscsLarge/Zho1 (PDZ) binding motif (QYWV). Binding partners are to date the thirteen PDZ domain containing scaffolding protein Mupp1 (Barritt et al., 2000), Syntenin, a cytoplasmic adapter whose interaction with NG2 is important for migration of OPC (Chatterjee et al., 2008) and the synaptic protein GRIP1 which binds the AMPAR subunit GluR2 together with NG2 on OPC (Stegmuller et al., 2003). We have observed that NG2 also binds to the serine protease OMI/HtrA2, localized in the intermembrane space in mitochondria. This molecule plays a role in apoptosis induction and we suggest that sequestration of the protease may help protect OPC from apoptosis-inducing stress. (Maus et al., unpublished results).

Recently a complex signal pathway resulting in directed migration of OPC in response to gradients of FGF or PDGF-aa has been reported (Biname et al., 2013). In this study FGF-dependent polarization of OPC was altered in cortical lesions of mice lacking the NG2 protein (NG2-KO mice, (Karram et al., 2008)). The underlying cellular signal cascade within the OPC has been shown to involve Rac/Rho dependent phosphorylation of the intracellular region of NG2, reviewed by Biname (2014).

The NG2 CTF and ICD influence target gene expression in OPC. The enzyme PTGDS catalyzes the conversion of prostaglandin H2 to the neuromodulatory form prostaglandin D2. PTGDS has been shown to be expressed and released by mature oligodendrocytes and the meninges (Taniike et al., 2002) and recently shown to be additionally expressed by OPC (Sakry

et al., 2015). In OPC, the level of PTGDS expression is modulated by the NG2 CTF and ICD, as well as by a complete lack of NG2 (Sakry et al., 2015). Interestingly expression of high levels of the NG2 ICD in OPC after transfection results in localization of the ICD into the nucleus, suggesting a potential regulatory function in gene expression (Sakry et al., 2015). CNS PTGDS levels have been shown to be increased in lesions of the human demyelinating disease Multiple Sclerosis (MS) (Chabas et al., 2001) and neurological disorders (Marin-Mendez et al., 2012), in MS lesions PTGDS can be additionally expressed by astrocytes (Kagitani-Shimono et al., 2006). Taken together, OPC can thus be seen as an additional cellular source of PTGDS expression in the CNS and OPC contribute to the neuromodulatory functions of PTGDS in an NG2-dependent manner. Interestingly in the PNS, PTGDS is reported to be expressed by DRG neurons where the neurotrophin I ICD influences expression levels (Trimarco et al., 2014).

4. Extracellular functions of the NG2 proteoglycan

NG2 has been postulated as a cell adhesion molecule for quite some time. This has been strengthened by the observation that the extracellular region between the TM domain and the LNS domains binds to collagen V and VI and integrins (Fukushi et al., 2004; Tillet et al., 1997); thus NG2 functions as a cell-surface anchor within the extracellular matrix (ECM). Other family members of the chondroitin sulfate proteoglycan (CSPG) family such as brevican are secreted proteins with defined roles within the ECM (Deepa et al., 2006; Frischknecht

et al., 2014). ECM integrity is important for AMPAR signaling at synapses influencing lateral diffusion of the receptor and consequently synaptic signal propagation (Frischknecht et al., 2009).

Interestingly, the two NG2 N-terminal LNS domains (which so far lack a binding partner) are similar to the LNS domains of the neuronal synaptic adhesion proteins the neuexins (NX), here they are responsible for binding to the interaction partners the neuroligins (NL) (Aoto et al., 2013; Ichtchenko et al., 1995; Ichtchenko et al., 1996; Krueger et al., 2012) as well as LRRTMs, Calsyntenins, Cerebellin, dystroglycan, and neuexophilins (Ko et al., 2009; Linhoff et al., 2009; Petrenko et al., 1996; Pettem et al., 2013; Sugita et al., 2001; Uemura et al., 2010). NL-NX interaction is an essential for forming and maintaining chemical synapses of the neuronal network (Krueger et al., 2012). Mutations within the NX LNS domains have been related to severe synaptic phenotypes found in human patients with neural diseases such as autism spectrum disorder (ASD) (Sudhof, 2008). The sequence conservation of LNS domains between laminins and neuexins is around 20–25% together with high functional similarities of the 3-dimensional structure (Rudenko et al., 1999; Rudenko et al., 2001). The NG2 LNS domains are no exception here: they show highest sequence similarities of up to 26% with the LNS domains of neuexins.

As discussed above, the NG2 extracellular region can be released by the activity of the α -secretase ADAM10 into the ECM, and is thus regarded as a soluble 290 kD NG2 ectodomain (Sakry et al., 2014). This cleavage from the OPC surface occurs constitutively, is increased in an activity-dependent fashion and elevated in response to lesion. The question thus posed is: what is the biological function of the NG2 ectodomain in its soluble form as it can be extracted from the ECM? We recently showed that the two LNS domains of the NG2 ectodomain modulate synaptic AMPAR currents and kinetics of pyramidal neurons within the rodent somatosensory cortex, suggesting an influence on subunit composition of the AMPAR (Sakry et al., 2014). The AMPAR-phenotype observed in this neuronal population in mice lacking NG2 (NG2-KO mice) can be rescued by incubation of brain slices with a recombinant protein containing solely the two NG2 LNS domains. Furthermore, in NG2-KO mice reduced NMDAR-dependent LTP is observed within the same neuronal population as a consequence of the altered AMPAR and NMDAR currents. NG2-KO mice exhibit a postsynaptic phenotype which is regulated by the NG2 LNS domains, while presynaptic characteristics such as the paired-pulse ratio remain unchanged (Sakry et al., 2014). Thus the NG2 LNS domains regulate postsynaptic signaling and modulate glutamergic signaling of the neuronal network. Interestingly in NG2-KO mice behavioral differences compared to wild-type littermates have been observed in tests related to integration of sensory input. In particular, a test based on the response of the auditory system (acoustic startle response, paired-pulse-ratio) showed abnormalities in NG2-KO mice (Sakry et al., 2014) similar to those observed in patients with schizophrenia.

Subpopulations of pericytes express NG2. A contribution of NG2-expressing pericytes of the vascular system to the described ectodomain effects cannot be completely excluded but is likely to play if at all, a minor role. Pericytes ensheath the endothelial cells making up blood vessels. It is not known

so far if and how efficient pericytes can release protein factors through the other layers of the blood-brain-barrier into the ECM of the CNS. Cleavage of OPC NG2 has been shown in acute slices, primary OPC (pOPC) and an OPC cell-line (Sakry et al., 2014), our pOPC culture lacks pericytes (Sakry et al., 2015). Furthermore in slices and pOPC NG2 cleavage is increased by neuronal activity and glutamate (Sakry et al., 2014), as far as is known pericytes do not respond to network activity nor receive synaptic input from neurons.

Recent analysis of the response of the NG2 KO mice to lesions also suggests that the loss of NG2 has effects beyond the neuronal network and that cytokine production by neural and immune cells may also be influenced by the lack of NG2.

Even though we know very little at present about the exact molecular mechanism underlying the modulation of neuronal signaling by the two NG2 LNS domains, these results show that the release of NG2 ectodomain (including the LNS domains) is under the control of the neuronal network via activity-dependent cleavage in OPC, initiated at their so called “neuron–glia” synapses (Bergles et al., 2010; Mangin and Gallo, 2011; Sakry et al., 2011). In response, the neuronal network is regulated by the released NG2 ectodomain. These novel results have introduced a new paradigm: a feedback mechanism (neuron–glia–neuron) integrating OPC as a glial cell population within the neuronal network (Sakry et al., 2014) as originally hypothesized in (Sakry et al., 2011).

5. Summary

Recent studies focused on the NG2 protein have revealed a defined role of NG2 in brain homeostasis. This includes intracellular functions within the OPC including modulation of migration, target gene expression and AMPAR clustering. The first two functions of NG2 seem to be of particular importance in lesioned brain tissue. Cleavage of the full-length NG2 protein by α - and subsequently γ -secretase activity turns out to be a core aspect, as the cleavage fragments (ectodomain, CTF, ICD) exhibit defined roles within the CNS. The fact that increased α -secretase mediated NG2 cleavage is modulated by neuronal activity is especially interesting as it integrates NG2-expressing OPC into the neuronal network. The story becomes even more exciting, since the released NG2 ectodomain, specifically the two LNS domains, are able to modulate excitatory neuronal synapses; thus, introducing a neuron–OPC–neuron feedback loop within the CNS.

6. Outlook

Future studies are aimed at defining the biological roles of the NG2 intracellular cleavage fragments CTF and ICD within the OPC. These are likely to include a nuclear function of the NG2 ICD and defined roles of the CTF and ICD in target-gene regulation.

NG2-dependent expression of PTGDS by OPC especially in CNS diseases like MS, needs further investigation as well.

The neuromodulatory function of the two LNS domains of the NG2 ectodomain evoke questions as to the identity of possible neuronal binding partners and an explanation of the effects on AMPA receptor characteristics, as well as detailed further characterization of the involved neuronal networks. A role in excitatory/inhibitory homeostasis of the neuronal network has already become apparent as the LNS domains increase synaptic transmission at excitatory synapses on pyramidal neurons within the somatosensory system. Furthermore, since a lack of the NG2 LNS domains was linked to altered animal behavior similar to that found in human diseases with known excitatory/inhibitory misbalance such as schizophrenia, a link to other neurological disorders involving excitatory/inhibitory misbalance is very likely. It is also possible that NG2-expressing gliomas may influence the surrounding neuronal networks by the mechanisms described above.

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