

Extracellular matrix of the central nervous system: from neglect to challenge

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Abstract The basic concept, that specialized extracellular matrices rich in hyaluronan, chondroitin sulfate proteoglycans (aggrecan, versican, neurocan, brevican, phosphacan), link proteins and tenascins (Tn-R, Tn-C) can regulate cellular migration and axonal growth and thus, actively participate in the development and maturation of the nervous system, has in recent years gained rapidly expanding experimental support. The swift assembly and remodeling of these matrices have been associated with axonal guidance functions in the periphery and with the structural stabilization of myelinated fiber tracts and synaptic contacts in the maturing central nervous system. Particular interest has been focused on the putative role of chondroitin sulfate proteoglycans in suppressing central nervous system regeneration after lesions. The axon growth inhibitory properties of several of these chondroitin sulfate proteoglycans *in vitro*, and the partial recovery of structural plasticity in lesioned animals treated with chondroitin sulfate degrading enzymes *in vivo* have significantly contributed to the increased awareness of this long time neglected structure.

Keywords CSPG · Lectican · Hyalactan · ADAMTS · Hyaluronan synthase · Perineuronal net · Node of Ranvier · Knockout · Tenascin · Link protein · Central nervous system

Introduction

It was not until 1971 that the existence of an extracellular matrix (ECM) in the central nervous system (CNS) was generally acknowledged (Tani and Ametani 1971). Then a predominance of hyaluronan and chondroitin sulfate proteoglycans (CSPG) (Margolis et al. 1975) and the paucity of otherwise frequent ECM molecules, like fibronectin or collagens, have been described (Carbonetto 1984; Rutka et al. 1988; Sanes 1989). Today we know that this distinctive ECM is mainly composed of proteoglycans of the lectican/hyalactan-family and their binding partners, hyaluronan, link proteins and tenascins (Bandtlow and Zimmermann 2000; Novak and Kaye 2000; Rauch 1997, 2004; Ruoslahti 1996; Yamaguchi 2000). In the following, we will focus on the structure, expression and putative functions of this major matrix components that form this extraordinary extracellular meshwork. Not discussed in this review are less abundant, but nonetheless functionally important macromolecules of the nervous system, such as reelin, agrin and thrombospondins. For detailed information about these large glycoproteins, which are involved in the control of neuronal migration and establishment of synapses we would like to refer the reader to recent reviews and publications (Bezakova and Ruegg 2003; Christopherson et al. 2005; Herz and Chen 2006; Tissir and Goffinet 2003).

Structures and ligands of the major constituents of the neural ECM

Proteoglycans and hyaluronan

Proteoglycans (PGs) are glycoproteins carrying, in addition to variable numbers of N- and O-linked oligosaccharides, at

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least one glycosaminoglycan (GAG) side chain, which is covalently bound to a core protein. The GAGs themselves are long unbranched polymers of repetitive disaccharide units consisting of an uronic acid (glucuronic or iduronic) or galactose and an amino-sugar (*N*-acetylglucosamine or *N*-acetylgalactosamine). According to the combination of these sugars, the GAGs are subclassified into heparin/heparan-, keratan- or chondroitin/dermatan-sulfates (Bandtlow and Zimmermann 2000; Kjellén and Lindahl 1991; Prydz and Dalen 2000). The GAG chains are in general 20–200 disaccharide-repeats long. Whereas keratan sulfates are usually attached to the core proteins via short standard N- or O-glycan links to asparagine or serine/threonine, respectively, the binding of chondroitin/dermatan sulfate and of heparin/heparan sulfate chains is mediated by a serine residue and a specific linker tetrasaccharide composed of a xylose, two consecutive galactoses and a glucuronic acid molecule. Numerous modifications that include O- and N-sulfation and epimerization of glucuronic acid at the C5-position lead to a high structural variability and open up countless possibilities for the modulation of GAG-dependent interactions (Bulow and Hobert 2006; Kusche-Gullberg and Kjellen 2003). Moreover, the highly negatively charged GAGs attract and bind considerable amount of water and cations.

In contrast to the protein-bound GAGs, hyaluronan (also known as hyaluronic acid or hyaluronate) is incorporated into the extracellular matrix as a core protein-free glycosaminoglycan (Toole 2000, 2004). Hyaluronan is a very large linear polymer built of repetitive disaccharides units consisting of glucuronic acid and *N*-acetylglucosamine. This carbohydrate filament reaches a molecular mass of up to 10^7 Da and extends over lengths of 2–25 μm . Unlike the other GAGs, hyaluronan is not sulfated and the glucuronic acid units are not epimerized.

While the more complex core protein-bound GAGs are assembled and modified by a large set of glycosyl- and sulfotransferases, the structurally simpler hyaluronan is synthesized by a single enzyme, the hyaluronan synthase (HAS). Three vertebrate genes (*HAS1*, *HAS2*, and *HAS3*)

giving rise to three HAS isoenzymes have been identified (DeAngelis 1999; Itano and Kimata 2002; Spicer and McDonald 1998; Weigel et al. 1997). The HAS are rather unique, as they catalyze the incorporation of two different monosaccharides and are, in contrast to most other glycosyltransferases, localized at the inner surface of the cell membrane. From there the growing hyaluronan string directly extrudes into the pericellular space, while being still attached to the producing enzyme (Spicer and Tien 2004; Weigel et al. 1997).

The most prominent binding partners of hyaluronan in the nervous system are the extracellular matrix proteoglycans of the lectican family (also called hyalectans) (reviewed by Bandtlow and Zimmermann 2000; Iozzo and Murdoch 1996; Rauch 2004; Ruoslahti 1996; Yamaguchi 2000). In mammals, four distinct lectican genes encode brevican, neurocan, aggrecan and versican (Table 1; Fig. 1). Shared features among these large chondroitin sulfate proteoglycans are the highly homologous G1 and G3 domains, which appear in rotary shadowing electron microscopy as compact globular structures at either end of an extended, but flexible central region (Mörgelin et al. 1989; Retzler et al. 1996). This poorly sequence-conserved middle part carries most of the O- and N-linked oligosaccharides and all glycosaminoglycan side chains. It varies among the different family members in size and also in the number of carbohydrate substitutions. Occasionally, the GAGs might even be absent, like in a fraction of the part-time preteoglycan brevican, or they may be greatly diminished, as in CNS-derived aggrecan in relation to its cartilage counterpart. Apart from the variations within the carbohydrate moiety, alternative splicing also greatly adds to the structural diversity of lecticans. Four versican isoforms (V0, V1, V2 and V3) exist as a result of alternative usage of two giant exons encoding the central GAG- α and GAG- β domains (Dours-Zimmermann and Zimmermann 1994; Zako et al. 1995; Zimmermann and Ruoslahti 1989). Versican V0 contains both of these GAG-attachment modules, whereas versican V1 and V2 include only the GAG- β or GAG- α , respectively. Versican V3, the smallest lectican,

Table 1 Extracellular matrix CSPGs in the central nervous system

Name	Core size ^a		GAGs		Cellular origin	CNS-specific
	Calculated ^b	SDS-PAGE ^c	Type	Number		
Aggrecan	244	370	CS	?	Neurons/astrocytes	No
Versican V0	371	≈550	CS	17–23	Neurons/astrocytes?	No
Versican V1	263	≈500	CS	12–15	Astrocytes?	No
Versican V2	180	400	CS	5–8	Oligodendroglial lineage	Yes
Neurocan	141	245	CS	3	Astrocytes/neurons	Yes ^d
Brevican	97	145	CS	0–5	Glial cells/neurons	Yes
Phosphacan (-KS)	172	400	CS/(KS)	3–4	Glial cells/neurons	Yes

^a kDa

^b Mature polypeptide

^c Core glycoprotein after glycosaminoglycan removal

^d Minor expression in peripheral nervous system

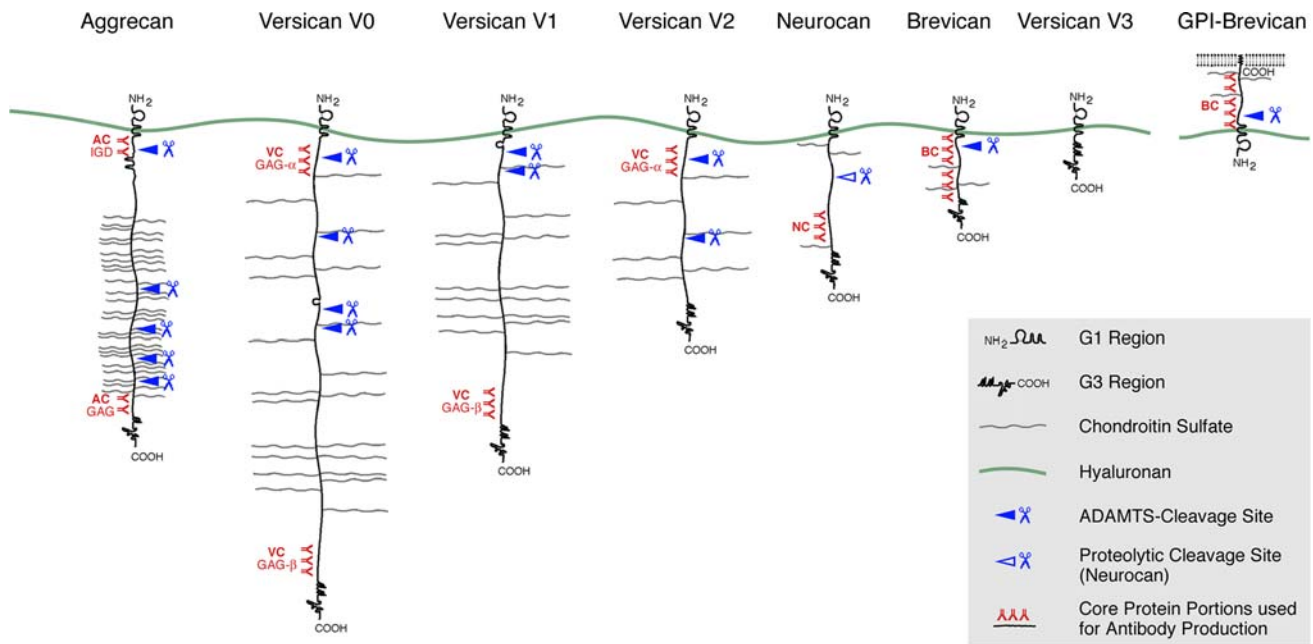


Fig. 1 Structural models of the chondroitin sulfate proteoglycans of the lectican family. ADAMTS cleavage sites and binding regions of the polyclonal antibodies used in immunohistochemical stainings displayed in the following figures are indicated

lacks both of these alternatively spliced elements and consequently carries no glycosaminoglycans.

In contrast to these largely diversified GAG-binding regions, the modules that form the globular G1 and G3 structures display little variability. For instance, all N-terminal G1 regions of the different lecticans include an immunoglobulin (Ig)-like loop and two link-protein-like tandem repeats that are involved in the binding of hyaluronan and link proteins (LeBaron et al. 1992; Mörgelin et al. 1989; Neame et al. 1987; Rauch et al. 2004). The G3 globule at the other end of the core protein contains a C-type lectin-like element, which is flanked by one or two EGF-repeats and a Sushi (SCR/CCP) domain, respectively. The entire G3 region is only absent in a GPI-anchored brevicant arising from alternative transcription termination (Seidenbecher et al. 1995). Recombinantly expressed G3 domains and/or the C-type lectin elements alone bind in vitro simple carbohydrates and heparin or heparan sulfate (Ujita et al. 1994), fibulin-1 and -2 (Aspberg et al. 1999; Olin et al. 2001), fibrillin-1 (Isogai et al. 2002), sulfoglycolipids (Miura et al. 1999) and the tenascins, Tn-C and Tn-R (Aspberg et al. 1995, 1997; Day and Prestwich 2002; Rauch et al. 1997). Of note, the majority of the G1- and G3-ligands are recognized by all lecticans, albeit differences in the affinities exist. Completing the structural models of the lecticans and their splice-variants by including the singular GAG-attachment regions between the largely homologous globular ends makes evident that this proteoglycan family forms an almost perfect array of functionally closely related but differently sized modular proteins (Fig. 1).

Besides the large aggregating proteoglycans of the lectican family, phosphacan, a secreted CSPG-isoform of the receptor-type protein-tyrosine phosphatase β (RPTP β), plays a prominent role in the brain ECM (Barnea et al. 1994; Maurel et al. 1994; Shitara et al. 1994). Like the other alternative splice products of the RPTP β gene, it is composed of a carbonic anhydrase domain, a fibronectin type III repeat and a spacer element, but lacks the transmembrane domain and the cytoplasmic tyrosine phosphatase modules. An additional element, present in phosphacan and the larger phosphatase variant, carries three to four chondroitin sulfate chains and sporadically also a few keratan sulfates (Rauch et al. 1991).

Phosphacan/RPTP β binds in a calcium-dependent manner to the ECM glycoproteins tenascin-R and tenascin-C (Milev et al. 1997; Xiao et al. 1997). It furthermore interacts in vitro with various cell adhesion molecules of the Ig-superfamily (IgCAMs) including N-CAM, Ng-CAM, axonin-1 (TAG-1) and contactin (F3/F11) (Milev et al. 1994, 1996; Peles et al. 1995), and it binds to the extracellular portion of voltage-gated sodium channels (Ratcliffe et al. 2000). The interaction with contactin and the sodium channels seems to be mediated by the carbonic anhydrase (CAH) domain of RPTP β /phosphacan (Peles et al. 1995; Ratcliffe et al. 2000).

Link proteins

The interaction between hyaluronan and lecticans is reinforced by small link proteins, collectively denominated

HAPLNs (hyaluronan and proteoglycan binding link proteins) (Spicer et al. 2003). This family of ancillary glycoproteins consists of four members, including the classical cartilage link protein (HAPLN1 = Crt11) (Neame and Barry 1993). Since its discovery several decades ago, HAPLN1 Crt11 has also been detected in brain, where it joins two other link proteins, the CNS-restricted HAPLN2/Bral1 and HPLN4/Bral2 (Bekku et al. 2003; Hirakawa et al. 2000). Only HAPLN3/Lp3 seems to be absent from the brain parenchyma, however, being expressed by the smooth muscle cells of larger blood vessels (Ogawa et al. 2004; Spicer et al. 2003).

The structure of the link proteins strongly resembles the globular G1 regions of lecticans, as they are all built-up of an Ig-fold and a hyaluronan-binding tandem repeat. There is some evidence that the interaction with lecticans may either be mediated by the Ig-fold (aggrecan) or by the tandem repeat of the link protein (versican). This functional relationship between HAPLNs and lecticans is also reflected in their chromosomal location and genomic organization. Intriguingly, each of the link protein genes is paired up with one of the lectican genes: *HAPLN1* with *VCAN*, *HAPLN2* with *BCAN*, *HAPLN3* with *AGC* and *HAPLN4* with *NCAN* (Spicer et al. 2003). Despite this particularity, the expression and binding preferences of link proteins and lecticans seem to be uncoupled from their genomic setup. For instance, HAPLN1/Crt11 is co-expressed and interacts with aggrecan in cartilage and possibly in brain, while versican, V0 and V1, and HAPLN3/Lp3 are both synthesized in smooth muscle tissues and may bind to each other at these locations (Ogawa et al. 2004). Alternatively, the same versican isoforms or neurocan may team up with HAPLN1/Crt11 to form ternary complexes with hyaluronan during brain development (Hirakawa et al. 2000; Matsumoto et al. 2003; Rauch et al. 2004; Seyfried et al. 2005; Shi et al. 2004). Finally, there are strong indications that pairs of HAPLN2/Bral1 and versican V2 (Oohashi et al. 2002) and of HAPLN4/Bral2 and brevican associate in mature brain tissues with hyaluronan (Bekku et al. 2003).

Tenascins

Apart from hyaluronan, lecticans and link proteins, tenascins represent the fourth class of molecules that form the basic constituents of the brain ECM. Tenascins (Tn) are very large multimeric glycoproteins that are well conserved among vertebrates (reviewed by Chiquet-Ehrismann and Chiquet 2003; Hsia and Schwarzbauer 2005; Joester and Faissner 2001; Jones and Jones 2000). In mammals, the family consists of four members namely tenascin-C, -R, -X and -W (-N). The macromolecular structures of the different tenascin monomers are highly alike as they follow the same modular arrangement. They are built of an amino-terminal

cysteine-rich oligomerization region composed of three to four α -helical heptad repeats, EGF-like elements, fibronectin type III-repeats (FN III) and a carboxyl-terminal fibrinogen-like globular domain.

The heptad domains allow an N-terminal association of the individual subunits that primarily form homotrimers. In tenascin-C and tenascin-W, an additional cysteine residue in this region permits these trimers to further assemble into large hexameric structures, the so-called hexabrachions. Although this cysteine is also present in tenascin-R, only trimers of this ECM molecule have been observed so far.

While the number of EGF-like repeats varies only between the different tenascins, some of the FN III domains are also subjected to alternative splicing. Since different combinations of these variable FNIII repeats are possible, several isoforms of the individual tenascins exist (Jones and Jones 2000). For instance, the tenascin-C monomers bear 14.5 EGF-repeats plus 17 FN III repeats, 9 of which can be alternative spliced. As a result, up to 27 different tenascin-C transcripts may be expressed during mouse brain development (Joester and Faissner 1999), each subunit comprising molecular weights in the range of 180–300 kDa. In contrast, tenascin-R, which includes 4.5 EGF and 9 FN III repeats, gives rise only to two splice variants of 180 and 160 kDa per subunit. This difference depends on the presence or absence of a supplementary FN III module between the FN III-repeats 5 and 6 (Fuss et al. 1993). Alternative splicing seems also to generate different isoforms of Tn-X (Ikuta et al. 1998) and Tn-W/Tn-N, the latter representing variants originating from the same gene (Neidhardt et al. 2003; Scherberich et al. 2004). Because the neural expression of Tn-W/Tn-N is currently still rather controversial (Neidhardt et al. 2003; Scherberich et al. 2004) and because Tn-X does not reach significant levels in the central nervous system (Matsumoto et al. 1994), we will focus in the following only onto the two other tenascin-family members.

Both, tenascin-C and tenascin-R bind to a wealth of extracellular matrix and cell surface ligands (Jones and Jones 2000). These interactions are mainly mediated by the FN III modules. The majority of the cellular receptors for tenascin-C and tenascin-R belong to the integrins, to the cell surface heparan sulfate proteoglycans (syndecans, glypicans) or to the cell adhesion molecules of the immunoglobulin superfamily (Ig-CAMs contactin/F11/F3, axonin/TAG-1 and neurofascin). Other cell surface binding partners are annexin II and the receptor protein tyrosine phosphatase (RPTP- ζ/β). Albeit of uncertain physiological relevance, a low affinity of the EGF-like repeats of Tn-C for EGF-receptors has also been reported (Swindle et al. 2001). Among the major ECM binding partners of these tenascins are fibronectin, phosphacan and particularly lecticans.

Supramolecular assembly

The ability of the brain ECM-components to selectively aggregate, leads to the establishment of large, relatively loose and flexible meshworks where hyaluronan acts as backbone. The current concept of supramolecular organization and assembly is mainly based on affinity measurements *in vitro* (mentioned before) and on recent rotary shadowing electron micrographs displaying the corresponding large multi-molecular complexes (Lundell et al. 2004). In this model, filamentous hyaluronan-molecules extruding from the neuronal cell membrane associate first with the G1 domains of distinct lecticans and link proteins that stabilize the complex. At the other end of the lectican core protein, the G3 domains engage in an interaction with one of the arms of the tenascin oligomers. This way tenascins may cross-link up to three (Tn-R) or up to six lecticans (Tn-C) and consequently bridge neighboring hyaluronan/lectican/link protein-complexes to complete the extracellular network. Supposing that the different components are proportionally expressed, the mesh size of this structure will be determined by the length of the tenascin arm, the distance between the lectican G1/link protein-aggregates along the hyaluronan molecules and most importantly by the dimension of the lectican core protein. Hence, the network would become roughly twice as dense by replacing for instance the V0 isoform of versican by the V2 variant and switching from Tn-C to Tn-R. Along with the change of lectican expression, the proportion of negatively charged carbohydrates and consequently the amounts of attracted water and cations may alter the gel-like properties of the extracellular matrix that occupies the intercellular spaces within the central nervous system. Indeed, the estimated change of the extracellular volume from roughly 40% in the developing brain to about 20% in adults (Nicholson and Sykova 1998) may at least partly be linked to the general tendency to express smaller lectican variants in the course of tissue maturation.

Expression and distribution

Gaining a clear insight into the expression and distribution of ECM components in development and adulthood has not been an easy task as especially antibodies widely employed in the earlier immunohistochemical studies were often not monospecific. Since *ex vivo* isolates of the highly glycosylated molecules were frequently used for immunization, several monoclonal antibodies have recognized carbohydrate epitopes, which could be found in glycans of several glycoproteins and/or were sometimes only present in sub-fractions of a particular ECM component (Lander et al. 1997; Matthews et al. 2002; Yamagata et al. 1993; Zako et al.

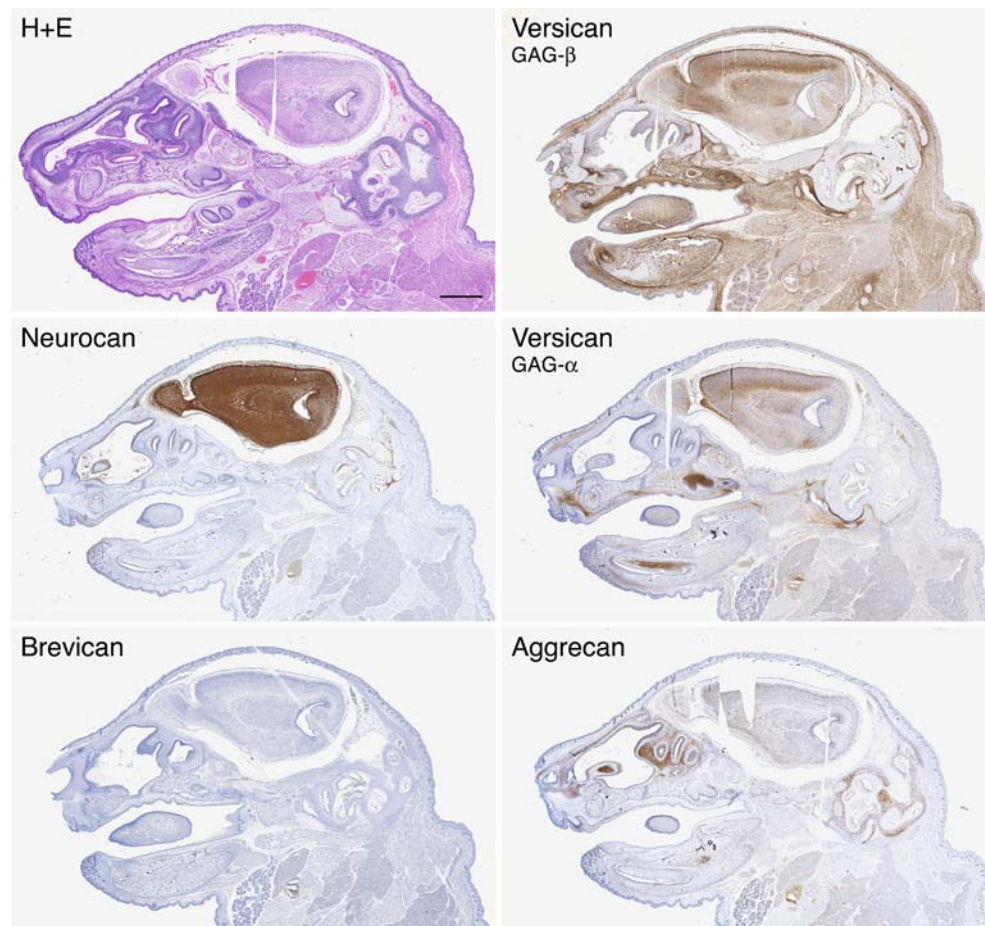
2002). Another cause for antibody cross-reactivity has also been the high protein sequence conservation, as in the G1 and G3 domains of the lecticans, and the consequent epitope sharing (Bignami et al. 1993; Perides et al. 1993; Yamada et al. 1997b). With the increasing availability of cDNAs and the advent of recombinant expression systems in bacteria, however, antibody production could be directed against more selective sites in little or non-glycosylated regions of the proteins (Milev et al. 1998b; Schmalfeldt et al. 1998; Zimmermann et al. 1994) (Fig. 1). Consequently, a set of highly specific and sensitive polyclonal antibodies with well-defined immunoreactivity has revealed a more consistent picture of ECM expression.

Tenascins C and R, phosphacan, most of the link proteins and all lecticans, including alternative splice-variants, are at some stage present in the central nervous system (Bandtlow and Zimmermann 2000; Jones and Jones 2000; Rauch 2004; Yamaguchi 2000). While Tn-R, phosphacan, HAPLN2/Bral1, HAPLN4/Bral2, brevican, neurocan and versican V2 are uniquely expressed in brain and spinal cord, Tn-C and the versican isoforms V0, V1 and V3 are also found in many mesenchymal tissues during embryogenesis and later in association with remodeling processes induced by, e.g., injury, neovascularization, inflammation or neoplasia. Likewise, aggrecan and HAPLN1/Crtl1, which participate in the establishment of some specialized brain ECMs, are primarily components of developing and mature cartilage. Finally, hyaluronan, which interacts with all the lecticans, displays the widest tissue distribution, being (also outside the nervous system) most abundantly produced by morphogenetically active zones during development and adult remodeling processes (Toole 2001, 2004).

Juvenile matrix type

During late embryonic and early postnatal phases of mammalian development, a juvenile type of extracellular matrix is initially formed in the CNS (Fig. 2). It mostly consists of hyaluronan, neurocan, versican V0, versican V1, tenascin-C and HAPLN1/Crtl1 (Milev et al. 1998b; Rauch 2004). In addition, prominent CNS-expression of aggrecan has been reported in chick embryos (Schwartz and Domowicz 2004), while only very small amounts have been detected in prenatal rodent brains (Matthews et al. 2002; Milev et al. 1998c). Cellular origin of neurocan and the large versican variants are neurons (Engel et al. 1996; Yamagata and Sanes 2005), whereas aggrecan appears to be mainly expressed by cells of astroglial lineage at this early time point (Domowicz et al. 2008). The transient deposition of neurocan and versican V0/V1, peaks in rats shortly after birth (Fig. 3) (Meyer-Puttlitz et al. 1995; Milev et al. 1998a). Preferential accumulations are the marginal zone and the subplate of the

Fig. 2 Immunohistochemical analyzes of the lectican distribution in the head region of E18.5 mouse embryos. Note the strong brain-specific staining of antibodies against neurocan. The antibodies against versican detect at this time point mainly the versican splice-variants V0 (GAG- α and GAG- β reactive) and V1 (only GAG- β reactive). These versicans are deposited in the CNS and in most of the mesenchymal tissues of the embryo. Aggrecan is practically absent from the brain, but is strongly expressed in cartilaginous tissues. No brevican staining can be detected at this embryonic stage. Sagittal sections. Bar 100 μ m



neocortex, parts of the hypothalamus, the amygdala and the developing hippocampus and dentate gyrus (Meyer-Puttlitz et al. 1996; Miller et al. 1995; Popp et al. 2003). Early postnatally, neurocan and the versican isoforms V0 and V1 are also found in the presumptive white matter and in the internal granule cell layer of the cerebellum (Meyer-Puttlitz et al. 1996; Popp et al. 2003). Starting from the second week after birth they are down-regulated and finally only persist in a few specialized locations in the adult CNS.

The time course of this early lectican expression is paralleled by their ligands, hyaluronan and the HAPLN1/Crt11 link protein. In fact, the content of hyaluronan peaks in rodent brain shortly after birth, being subsequently reduced to one-fourth of its maximal level in the adult tissue (Margolis et al. 1975). In the developing cerebellum, hyaluronan is mostly present in the granular cell layer and prospective white matter (Ripellino et al. 1988), where it seems to be organized into extracellular arrays of fiber-like structures (Baier et al. 2007). There is evidence that glial cells synthesize hyaluronan in the early phase of the CNS formation (Deyst and Toole 1995), the hyaluronan synthase responsible for its production at this early stage has yet to be identified. Similar to hyaluronan, HAPLN1/Crt11, which may

stabilize the interaction of hyaluronan with neurocan, versican and aggrecan, is up to about postnatal day 10 increasingly expressed and then gradually diminishes in the maturing nervous tissue (Hirakawa et al. 2000).

A relatively early CNS expression is also observed for the glia-derived ECM glycoprotein Tn-C. It first accumulates around the fibrous processes of radial and Bergmann glial cells that direct the migration of neuronal precursors during cortical and cerebellar development, respectively (Crossin et al. 1986; Prieto et al. 1990). In rodents, it is rather widely distributed shortly after birth and displays at least in the cerebral cortex, hippocampus and cerebellum partial overlaps with neurocan and versican V0/V1 depositions (Bartsch et al. 1992; Laywell and Steindler 1991; Meyer-Puttlitz et al. 1996; Popp et al. 2003). Particularly intriguing is the transient association of Tn-C with the glial boundary tissues surrounding the functional sets of neurons, like for instance the vibrissae-related barrel fields of the developing somatosensory cortex (Steindler et al. 1989). About 2–3 weeks after birth, Tn-C levels decrease continuously, maintaining only a significant expression level in the neurogenetically active areas of the adult brain that encompass the subependymal zone and the hippocampus

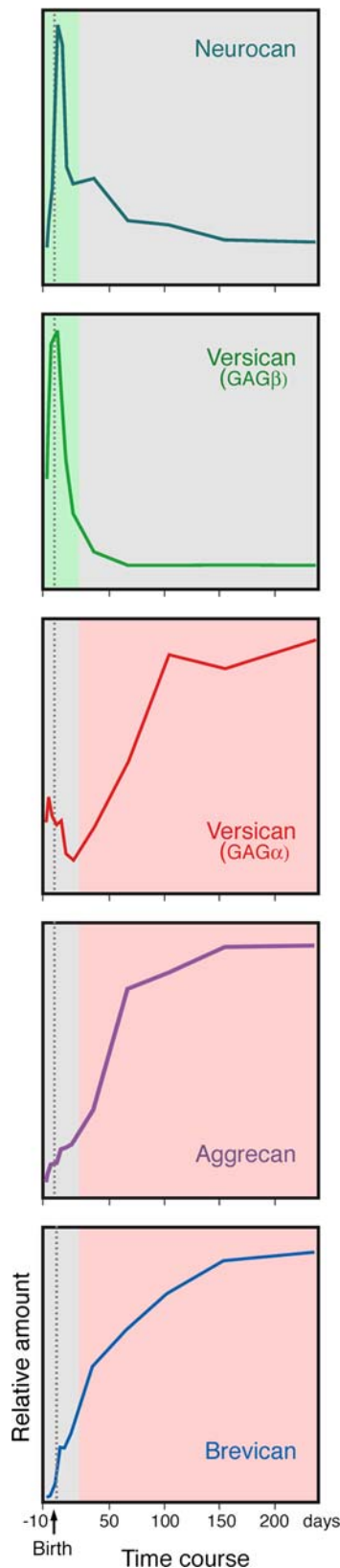


Fig. 3 Time course of lectican levels in extracts of embryonic and postnatal rat brains. Note the transition from the expression of the juvenile-type of matrix to its mature form around postnatal day 20. Diagrams adapted from (Milev et al. 1998b)

(Bartsch et al. 1992; Dorries and Schachner 1994; Gates et al. 1995; Thomas et al. 1996).

Mature matrix type

Initiating about 2 weeks after birth, a major remodeling process takes place (Fig. 3), which replaces most of the juvenile type of matrix by its mature form (Rauch 2004). This exchange transforms the relatively loose embryonic and early postnatal ECM to a significantly firmer meshwork, which is subsequently maintained throughout adulthood. The large majority of the early matrix components are along the conversion substituted by a different, but homologous set of ECM proteins that include versican V2, aggrecan, brevican, phosphacan, tenascin-R and the brain link proteins HAPLN2/Bral1 and HAPLN4/Bral2 (Bekku et al. 2003; Hirakawa et al. 2000; Meyer-Puttlitz et al. 1995; Milev et al. 1998b; Pesheva et al. 1989).

In the adult CNS, various combinations of lecticans and link protein, which are generally associated with hyaluronan, Tn-R and phosphacan condense at some strategic locations (Fig. 4.). Versican V2 and the secreted brevican isoform emerge around the second and third week postpartum and subsequently evolve to the predominant constituents in the adult brain and spinal cord (Schmalfeldt et al. 1998; Yamaguchi 1996). Matrices that contain mainly versican V2 and some brevican are most prevalent in the white matter surrounding the myelinated fibers of all calibers (Ogawa et al. 2001; Schmalfeldt et al. 2000). Particularly dense accumulation of versican V2, HAPLN2/Bral1, Tn-R and phosphacan appear as ring-like structures around the CNS nodes of Ranvier (Dours-Zimmermann et al. unpublished; Melendez-Vasquez et al. 2005; Oohashi et al. 2002; Pesheva et al. 1989; Xiao et al. 1997). Origins of the Tn-R and versican V2 deposits in the white matter are primarily oligodendrocytes and their precursors (Asher et al. 2002; Pesheva et al. 1989; Schmalfeldt et al. 2000), while HAPLN2/Bral1 expression has been attributed to neurons (Oohashi et al. 2002) or oligodendrocytes (Carulli et al. 2006). Interestingly, the brevican expression shifts at the end of myelination from oligodendroglial to astrocytic lineage in the white matter (Ogawa et al. 2001). Starting from postnatal day 28, astrocytes seem to give rise to another particularly brevican-rich zone that forms after completion of the neuronal migration in the granular cell layer of the rat cerebellum. This more compact extracellular matrix ensheathes cerebellar glomeruli, in which the incoming mossy fibers contact the local neuronal processes (Yamada et al. 1997a).

The by far best studied extracellular matrix condensation in the adult central nervous system is the perineuronal net (PNN). This lattice-like structure engulfs the cell bodies, proximal dendrites and axon initial segments of specific

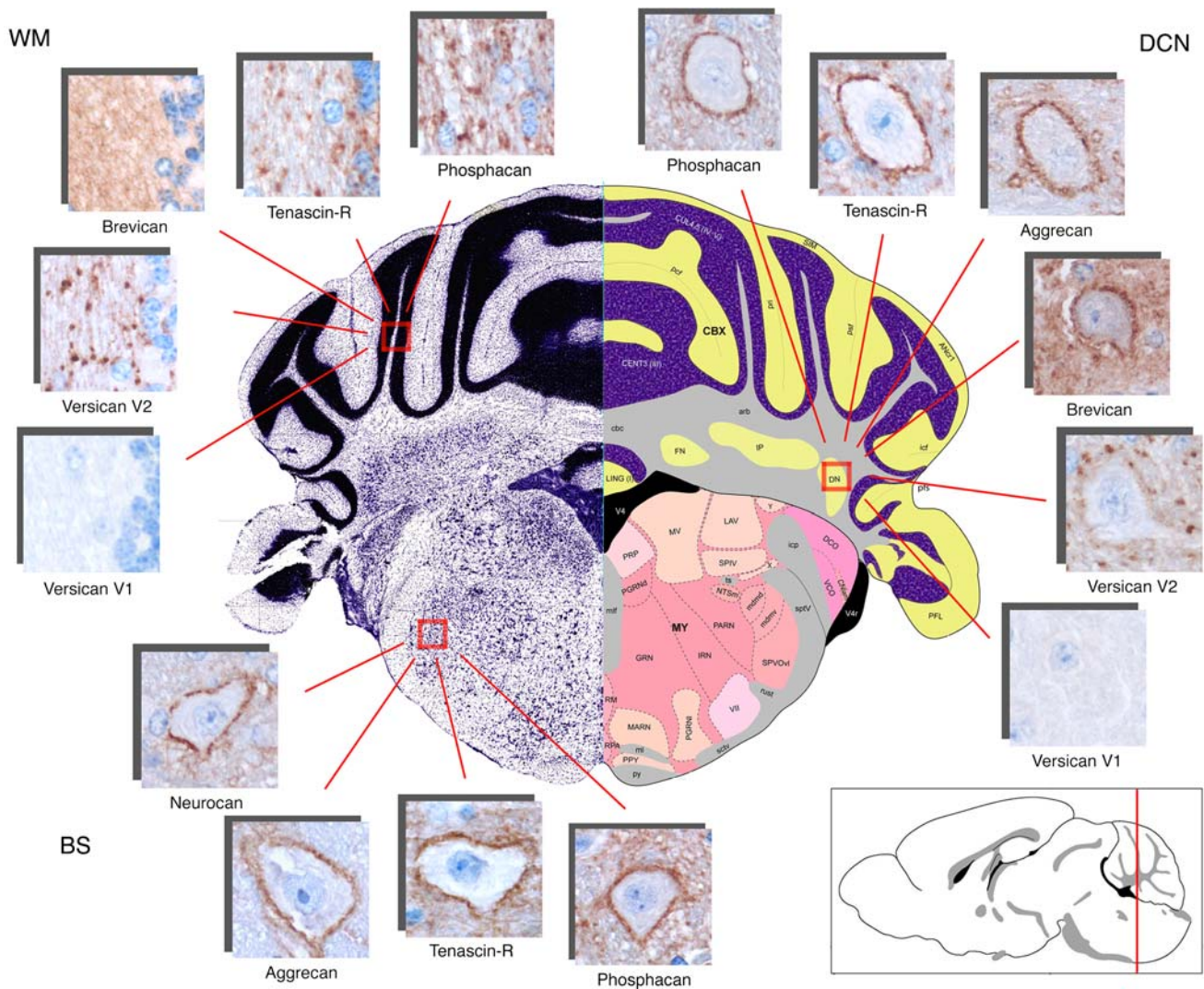


Fig. 4 Localization of lecticans and tenascins in coronal sections through adult cerebellum and medulla at Bregma -6.18 mm. Note the perineuronal net staining of aggrecan, tenascin-R and phosphacan in the deep cerebellar nuclei (DCN) and the brain stem (BS). Brevican and neurocan are only present in some of the PNNs. Versican V2, which stains exclusively with GAG- α -, but not with GAG- β -specific antibodies

is accumulating at the nodes of Ranvier in the white matter (WM), while brevican staining is rather diffused in the myelinated fiber tracts. Versican V0 and V1, which are recognized by the GAG- β -specific antibodies are absent from these brain regions in adult mice. Graphics modified from the Allen Brain Atlas (<http://www.brain-map.org>) (Lein et al. 2007)

subsets of neurons and embeds, with exception of the synaptic clefts, the presynaptic boutons that contact them (Bruckner et al. 1993, 2006; Celio and Blumcke 1994; Celio et al. 1998; Galtrey and Fawcett 2007; Murakami and Ohtsuka 2003; Rhodes and Fawcett 2004; Yamaguchi 2000). PNNs can be observed in many areas of the CNS including the cerebral cortex, the hippocampus, the thalamus, the cerebellum, the brain stem and the spinal cord. They most frequently surround parvalbumin-expressing GABAergic interneurons and certain cortical pyramidal neurons, as well as projection and large motor neurons of the brain stem and spinal cord. The formation of perineuronal nets occurs relatively late in postnatal development (in rodents 2–5 weeks after birth) and coincides with the end-

ing of the experience-dependent refinement of the synaptic network and the closure of the critical period, e.g., in the visual system and spinal cord (Guimaraes et al. 1990; Kalb and Hockfield 1988, 1990; Pizzorusso et al. 2002).

This specialized matrix consists of hyaluronan, different lecticans, the link proteins HAPLN1/Crtl1 and HAPLN4/Bral2, Tn-R, and phosphacan (Asher et al. 1995; Bekku et al. 2003; Bruckner et al. 2000; Carulli et al. 2006; Haunso et al. 1999; Maeda et al. 1995). The perineuronally accumulating hyaluronan is synthesized by HAS2 and HAS3 expressed in the net-carrying neurons, while HAS1 is generally absent from the CNS (Carulli et al. 2006, 2007). During the PNN-formation, HAS3 may slightly precede HAS2. The expression of both of these hyaluronan

synthases is subsequently attenuated, but pertains throughout adulthood.

Among the lectican family members, aggrecan is the most common perineuronal net component. Although present in all PNNs, it may contribute to a certain sub-specialization by varying its glycan structures between distinctive sets of neuronal coats (Matthews et al. 2002). Brevican and neurocan display in many CNS regions an expression pattern similar to aggrecan (Fig. 4) and there is indeed some evidence that they co-localize within the same perineuronal structure (Carulli et al. 2006; Galtrey et al. 2008). At least hyaluronan, aggrecan and Tn-R seem to be rather uniformly distributed within a single net covering the cell body, proximal dendrites and the axon initial segment (AIS) irrespective of the presence of synapses (Bruckner et al. 2006). In contrast, a certain preference of brevican deposition around the AIS has been observed in neuronal cell cultures (Hedstrom et al. 2007; John et al. 2006). Using a monoclonal antibody (12C5) against the hyaluronan-binding G1 domain of versican, immunohistological examinations of brain and spinal cord sections have also suggested that the fourth lectican member is present in PNNs (Carulli et al. 2006; Deepa et al. 2006). Surprisingly, none of our antibodies against the two GAG-attachment regions (Fig. 1) has displayed a similar perineuronal net staining (Fig. 4). It is therefore conceivable that the 12C5 antibody, in fact does not detect intact versican V2 in the PNNs, but rather a proteolytic remnant of the early postnatal versican V0 and V1 expressions that correspond to the versican fragment, formerly named as glial hyaluronate-binding protein (GHAP) or hyaluronectin (Delpech and Halavent 1981; Perides et al. 1989). Also neurocan in the PNNs may at least partly be present in form of a hyaluronan-binding truncation product (Neurocan-*N*) (Deepa et al. 2006), which eventually has become trapped after the conversion of the juvenile matrix.

No matter whether the lecticans are intact or truncated, their interactions with hyaluronan seem to be either stabilized by HAPLN1/Crt11 or by HAPLN4/Bral2 (Bekku et al. 2003; Carulli et al. 2006, 2007). Both of these link proteins form an integral part of perineuronal nets. Whereas HAPLN1/Crt11 seems to bind to its classical partner aggrecan and probably also to neurocan or neurocan-*N*, there is some indication from knockout mice that brevican may be required for the PNN localization of HAPLN4/Bral2 (Bekku et al. 2003).

Most of the perineuronal net constituents seem to be expressed by the engulfed neurons themselves (Carulli et al. 2006). Nonetheless, contributions from surrounding astrocytes that extend cellular processes into the reticular structure cannot be excluded (Carulli et al. 2007). This concerns particularly brevican and to some extent also neurocan, which may be produced bilaterally by the ensheathed neurons and the contacting astrocytes. Co-culture experi-

ments of primary hippocampal neurons and glial cells indeed revealed that the perineuronal deposits of brevican are in vitro primarily astrocyte-derived (John et al. 2006), although perineuronal net-like structures develop apparently also in the virtual absence of glial cells (Miyata et al. 2005).

Lesion-associated reactive matrix

Once established, the composition of the mature type of extracellular matrix is rather stable with little or no turnover of their components. This changes, however, radically, when lesions to the adult central nervous system occur. Under these circumstances, the expression of various extracellular matrix molecules is highly up-regulated and major depositions are observed in and around the lesion site, in particular, in association with the glial scar tissue that forms (Bradbury and McMahon 2006; Galtrey and Fawcett 2007; Gonzenbach and Schwab 2008; Morgenstern et al. 2002; Zurn and Bandtlow 2006). The freshly produced ECM components may be secreted by reactive astrocytes, oligodendrocyte precursors, microglia/macrophages and eventually by meningeal cells. The lesion and consequent reactive processes induce a matrix accumulation that strongly resembles the juvenile-type of meshwork previously observed during early nervous system development. For instance, surgical incisions in the cerebral cortex or spinal cord provoke a relatively fast and transient up-regulation of Tn-C and neurocan (Asher et al. 2000; Haas et al. 1999; Jones et al. 2003; Laywell et al. 1992; Matsui et al. 2002; McKeon et al. 1999; Tang et al. 2003). Also the expression of versican V2, brevican and phosphacan appear to be affected. Yet, recent reports about alterations in the production of versican V2 (direction of regulation) and brevican (timing) are somewhat controversial (Asher et al. 2002; Jones et al. 2003; Tang 2003). There are nevertheless indications that the lecticans brevican, aggrecan and versican V2 follow the time course of phosphacan, whose content first diminishes during the acute phase and then increases in a late attempt to restore the mature type of matrix at the injury site (Lemons et al. 2001; Tang et al. 2003).

A similar post-traumatic upregulation of Tn-C and lecticans has also been observed after disruption of sensory axons at the dorsal root (Beggah et al. 2005; Pindzola et al. 1993). Importantly, this type of lesion near the PNS/CNS interface lacks a scar formation within the CNS, but still leads to a reactive gliosis in the dorsal root entry zone (DREZ) and Wallerian degeneration of the central axon branches, now separated from their cell bodies. Despite the absence of a glial scar tissue, elevated depositions of neurocan and versicans (also V1 isoform) are prominent in the DREZ suggesting that the remodeling process temporally

reactivates a juvenile matrix-like expression profile analogous to direct damages to the CNS.

Matrix turnover

The rapid switch from embryonic and early postnatal extracellular matrices to their mature form in the normal adult CNS (Rauch 2004), the fast disappearance of CSPGs from the predestined axonal pathways observed in the developing periphery (Landolt et al. 1995; Oakley and Tosney 1991), as well as the reactive changes following nervous system lesions (Galtrey and Fawcett 2007; Zurn and Bandtlow 2006) cannot solely be attributed to adjustments in the ECM expression patterns, but must also depend on highly active proteolytic processes (Agrawal et al. 2008; Ethell and Ethell 2007; Flannery 2006; Gottschall et al. 2005; Milward et al. 2007; Porter et al. 2005). The selective turnover of lecticans may in this context be essential for altering the cell migration and axon growth properties of many nervous tissues. Metalloendopeptidases of the ADAMTS family seem to be primarily responsible for the lectican catabolism (ADAMTS: a disintegrin and metalloproteinase with thrombospondin motifs). Conversely, matrix metalloproteases (MMPs) may only play a subordinate role in the proteoglycan degradation, but they could preferentially target the link proteins and tenascins of the central nervous system. In any case, the proteolysis is tightly controlled by the tissue inhibitors of metalloproteinases (TIMPs), which are physiological antagonists of both enzyme types, MMPs (TIMP-1 to -4) and ADAMTSs (TIMP-3).

Various ADAMTS-enzymes cleave lecticans at a few specific sites *in vitro* and corresponding digestion products have been identified *in vivo* (Flannery 2006; Gottschall et al. 2005; Porter et al. 2005). The secreted ADAMTSs are close relatives of the cell membrane-bound ADAM metalloproteases. ADAMs and ADAMTSs are expressed as zymogens containing a prodomain and a zinc-binding catalytic element linked to a disintegrin-like motif. In ADAMTSs, several additional modules may control tissue localization and contribute to the substrate specificity. This C-terminal set of ancillary domains includes a central thrombospondin type I repeat, a cysteine-rich region, a spacer element and several supplementary thrombospondin motifs in most of the family members. The ADAMTS-zymogens are in the trans-Golgi or at the cell surface activated by N-terminal propeptide cleavage mediated by furin or furin-like convertases. Additional proteolytic or autocatalytic removal of a C-terminal portion has been demonstrated to modulate the enzymatic activity of some of these metalloproteases (Gao et al. 2002; Rodriguez-Manzanique et al. 2000; Zeng et al. 2006).

Five ADAMTS sites have been characterized in the aggrecan core protein (Caterson et al. 2000; Lemons et al.

2001), two to four in the proteoglycan isoforms of versican (V2–V0) (Jonsson-Rylander et al. 2005; Sandy et al. 2001; Westling et al. 2004) and one in brevican (Matthews et al. 2000; Yamada et al. 1995). Interestingly, in all of these lecticans one ADAMTS-site is located next to the globular G1-domain allowing the selective release of the large GAG-carrying portions from the complex with hyaluronan and link proteins. Finally, also neurocan, which is in adult tissues often present in form of two well-defined N- and C-terminal fragments, may be cleaved by an ADAMTS- or MMP-protease. Based on comparisons of known substrates, specific sites for by MMP-2- and ADAMTS-processing have been postulated. Whereas MMP-2 indeed cleaves neurocan *in vitro*, the susceptibility to ADAMTS-digestion still has to be confirmed in an experimental setting (Sandy et al. 2001; Turk et al. 2001).

Among the 19 ADAMTSs, seven exhibit cleavage capacity towards lecticans (ADAMTS-1, -4, -5, -8, -9, -15 and -20; ADAMTS-5 and -11 are identical). The classical aggrecanases are ADAMTS-4 and -5 (Abbaszade et al. 1999; Tortorella et al. 1999). Albeit less efficient, aggrecan-processing activities have also been observed of ADAMTS-1, -8, -9 and -15 (Collins-Racie et al. 2004; Flannery 2006; Kuno et al. 2000; Somerville et al. 2003). At least four ADAMTS-proteases including ADAMTS-1, -4, -9 and -20 recognize versican substrates *in vitro* (Sandy et al. 2001; Silver et al. 2008; Somerville et al. 2003; Westling et al. 2004), whereas only cleavage by ADAMTS-4 and -5 has been demonstrated for brevican (Matthews et al. 2000; Nakada et al. 2005; Nakamura et al. 2000).

Until now, only a few studies have explored the expression of distinct ADAMTSs during nervous system development and virtually no data are available for peri- and early postnatal phases that are particularly interesting regarding the remodeling of the neural ECM. From the limited insights one can currently conclude, that ADAMTS-1 and -9 are expressed relatively early in the embryonic CNS (Gunther et al. 2005; Jungers et al. 2005; Thai and Iruela-Arispe 2002), while ADAMTS-4 appears only in adult nervous tissues (Abbaszade et al. 1999; Jungers et al. 2005). Moderate to low expression of ADAMTS-4, for instance has been detected in pyramidal neurons of various cortical areas and in granule cells of the dentate gyrus in normal adult brain (Yuan et al. 2002).

This base-expression of ADAMTSs rises considerably in the presence of different types of lesions. After a kainate-induced excitotoxic insult, a transient up-regulation of ADAMTS-4 and ADAMTS-1 coincides with a marked increase of brevican proteolysis (Mayer et al. 2005; Yuan et al. 2002). A similar reaction is also observed in an experimental model of cerebral ischemia (Cross et al. 2006). Moreover, motor axon disruption induces the ADAMTS-1 production in the affected neurons (Sasaki et al. 2001) and

there is growing evidence that the same enzyme is also over-expressed in association with various neurodegenerative disorders (Miguel et al. 2005; Satoh et al. 2000). In addition, glioblastomas display elevated levels of brevican and its proteolytic fragments (Nutt et al. 2001) are associated with an increased expression of ADAMTS-5 and, although somewhat controversial, possibly also with ADAMTS-4 (Held-Feindt et al. 2006; Nakada et al. 2005). This close relationship between ECM remodeling and ADAMTS expression under traumatic and pathogenic conditions, provides an indirect indication for an analogous functional involvement of ADAMTSs in the matrix conversion of normal neural development.

Putative functions

The progress in unraveling the precise functions of the extracellular matrix in the central nervous system has for long been rather slow due to the high structural variability of its main constituents and the great complexity of the dynamic remodeling process involved in normal development and disease conditions. Albeit correlations of the distribution of specific ECM components with cellular processes like proliferation, migration, axonal growth or synapse formation have been very valuable for gaining insights into the putative functions *in vivo*, they may have been somewhat distorted by the fact that most antibodies used in these studies are unable to discriminate between intact and ADAMTS- or MMP-processed forms of a particular molecule. This distinction could be of considerable

importance as the specific cleavage may attenuate or even neutralize certain ECM functions without becoming apparent in the immunohistological stainings.

Moreover, partial overlapping rather than completely different functions of structurally related matrix proteins may be responsible for the unexpectedly mild phenotypes of several knockout mice, while the absence of potential key components has in contrast resulted in embryonic or perinatal lethality that prevents the study of their role in the mainly postnatal neural development (Table 2). Nevertheless, evidence for an extracellular matrix involvement in neural migration, axon guidance, plasticity restriction and fiber tract stabilization has in recent years come from various *in vitro* and *in vivo* observations.

Axon growth inhibition

Phosphacan and all lecticans of the central nervous system inhibit in their intact form axonal growth *in vitro* (Bandtlow and Zimmermann 2000; Yamaguchi 2000). This functional property appears to be mostly core protein-dependent in versican V0, V1 and V2 (Dutt, Stöckli and Zimmermann unpublished; Niederöst et al. 1999; Schmalfeldt et al. 2000;), neurocan (Margolis et al. 1996) and phosphacan (Maeda and Noda 1996), whereas no inhibition of brevican (Yamada et al. 1997a) and cartilage-derived aggrecan (Snow et al. 1990) has been detected after chondroitinase ABC digestion. Notably, we also observed two- to three-fold reductions, but never an abolition of the inhibitory capacity of versican V2 in stripe choice experiments after complete glycosaminoglycan-removal (Schmalfeldt et al.

Table 2 Available ECM-knockout mouse strains

Gene	Viability	CNS phenotype
Brevican (<i>Bcan</i>)	Normal	Reduced LTP
Neurocan (<i>Ncan</i>)	Normal	Reduced LTP
Versican (<i>Vcan/hdf</i> -strain)	Die at E10.5	–
Aggrecan (<i>Acan/cmd</i> -strain)	Die at birth	ND
RPTP- β /phosphacan (<i>Ptprz1</i>)	Normal	Enhanced LTP; impaired recovery from EAE (transmembrane variants?)
Tenascin-R (<i>Tnr</i>)	Normal	Aberrant PNNs; reduced conduction velocity; disturbed neuroblast migration in olfactory bulb; some behavioral abnormalities
Tenascin-C (<i>Tnc</i>)	Normal	Reduced LTP; mild behavioral abnormalities
HAPLN1/CrtI1 (<i>Hapln1</i>)	Die at birth	ND
HAS2 (<i>Has2</i>)	Die at E10.5	–
HAS3 (<i>Has3</i>)	Normal	ND

LTP Hippocampal long-term potentiation, EAE experimental autoimmune encephalomyelitis, PNN perineuronal net, ND not described

Brevican (Brakebusch et al. 2002); neurocan (Zhou et al. 2001); versican (Mjaatvedt et al. 1998); aggrecan (Watanabe et al. 1994); RPTP- β /phosphacan (Harroch et al. 2000, 2002; Niisato et al. 2005); TnR (Bruckner et al. 2000; Freitag et al. 2003; Haunso et al. 2000; Montag-Sallaz and Montag 2003; Saghatelian et al. 2004; Weber et al. 1999); TnC (Evers et al. 2002; Forsberg et al. 1996; Fukamauchi et al. 1996; Kiernan et al. 1999; Saga et al. 1992); HAPLN1/CrtI1 (Watanabe and Yamada 1999); HAS2 (Camenisch et al. 2000); HAS3 (Bai et al. 2005)

2000). This moderate decrease might be related to the partial collapse of the extended core protein structure after elimination of the highly negatively charged chondroitin sulfate side chains.

The axonal growth inhibition is likely dependent on the presence of a pericellular hyaluronan coat that covers many cell types and often includes variable amounts of lecticans, link proteins and tenascins (Evanko et al. 2007). Synthesis of hyaluronan alone promotes in various cell types the formation of plasma membrane protrusions (Kultti et al. 2006; Rilla et al. 2008). It is therefore conceivable that a coat containing exclusively hyaluronan has a similar effect in generating growth cone filopodia exploring the environment at the tip of growing axons. In situations, in which these filopodia encounter areas expressing lecticans, they may incorporate the CSPGs into the pericellular hyaluronan structure and consequently increase the hydration capacity and thus the thickness of their coat. Depending on the extent of this coat swelling, the contacts to the surface of neighboring cells or the extracellular matrix may be increasingly compromised by sterical hindrance. Accordingly, the advancing growth cone should slow down, turn away or retract from the lectican-containing zones. Furthermore, the expression level, core protein size and carbohydrate substitution of the encountered lecticans would control the coat dimensions and thus, modulate the inhibitory effect. In line with this hypothesis are the concentration-dependency of the versican inhibition (Schmalfeldt et al. 2000) and a certain tendency to accentuate the effect by using larger splice-variants *in vitro* (Dutt, Stöckli and Zimmermann, unpublished). In addition, neurons may regulate the coat size by the extent of the hyaluronan synthesis and/or by secreting ADAMTS-proteases, which would reduce the inhibitory activity through release of the GAG-carrying core protein portions from the pericellular structure.

While high-versican concentrations provoke retraction, low concentrations still allow a reduced growth, but promote enlargement of presynaptic varicosities in chick retinal axons *in vitro* (Yamagata and Sanes 2005). This finding is corroborated by RNA interference experiments *in vivo*, where depletion of versican causes a significant size reduction of the varicosities in the retinal arbors of the optic tectum suggesting that low-versican expression attenuates axonal growth and induces lamina-specific presynaptic maturation. Conversely, high expression of versican V0/V1 may also contribute to the formation of molecular barriers that block axon extension and may in addition, direct migratory neural crest cells in the developing peripheral nervous system (Dutt et al. 2006; Landolt et al. 1995; Oakley and Tosney 1991). Unfortunately, these suspected modulator functions of the V0/V1 isoforms cannot be verified in versican null mice (*hdf* strain), as they suffer from problems in heart segmentation and

consequently die early, at around embryonic day 10.5 (Mjaatvedt et al. 1998).

Although the highest matrix protein expression in the CNS during early neural development has been attributed to Tn-C and neurocan, their *in vivo* functions in the juvenile-type of matrix are still largely unknown. Single knockouts of neurocan and Tn-C showed no apparent anatomical abnormalities in the CNS (Steindler et al. 1995; Zhou et al. 2001). A potential compensation by up-regulation of the closely related brevican or Tn-R, respectively, has not been observed. Nonetheless, recent generation of a Tn-C, Tn-R, neurocan and brevican quadruple knockout suggested a partial replacement of the tenascins by fibulin-1 and -2, which are generally not expressed in the brain parenchyma, but may similarly cross-link the C-terminal domains of the remaining lecticans (Rauch et al. 2005). This complex may however, be less stable as suggested by the partly disturbed structure of the perineuronal nets in these quadruple and in the Tn-R single knockout mice (Bruckner et al. 2000; Haunso et al. 2000; Rauch et al. 2005; Weber et al. 1999).

Regulation of plasticity

There are increasing indications that the transition from the juvenile type of matrix to its mature form terminates the highly dynamic periods of cell and axonal migrations and restricts plasticity and regeneration in the adult central nervous system (Galtrey and Fawcett 2007; Rauch 2004). In white matter, specialized matrices rich in brevican and versican V2 contribute to axon growth inhibitory environment associated with CNS myelin (Niederöst et al. 1999) and possibly prevent abnormal axon branching by accumulating at the nodes of Ranvier. In addition, the establishment of perineuronal nets seems to stabilize the neuronal circuitry by suppressing the formation of new synaptic contacts (Galtrey and Fawcett 2007; Hockfield et al. 1990; Rauch 2004; Yamaguchi 2000). These speculations that are mostly based on the inhibitory properties of lecticans *in vitro*, have been further nourished by observations that injections of bacterial chondroitinase into the visual cortex of rats affect the integrity of perineuronal nets and restore ocular dominance plasticity even after closure of the critical period (Berardi et al. 2004; Hooks and Chen 2007; Pizzorusso et al. 2002, 2006). Similarly enhanced plasticity after chondroitinase treatment has been reported from experimental brain and spinal cord injuries (Barritt et al. 2006; Bradbury et al. 2002; Moon et al. 2001). In these studies, the digestion of chondroitin sulfates at the lesion sites resulted in increased axonal sprouting and some axonal re-growth across the normally non-permissive glial scar tissue, which expresses CSPGs abundantly. In addition, in adult brevican and neurocan double-knockout mice, the obstruction of the lectican-rich dorsal root entry

zone appears to be partly lifted, as a significant number of sensory axons cross back into the DREZ after proximal nerve disruption and subsequent growth stimulation through a late conditioning lesion of the peripheral branch (Quaglia et al. 2008). Because the suppression of a single myelin- or ECM-associated CNS-inhibitor typically leads to less than 10% robust re-growth of cut axons in all different experimental lesion systems tested (Bradbury and McMahon 2006; Gonzenbach and Schwab 2008; Zheng et al. 2006), such multimodal strategies may be required to achieve an improved neutralization of the inhibition and a more effective promotion of the regenerative response in the CNS.

Other putative functions

Apart from these putative roles connected to axon growth inhibition, the condensed matrices in the mature central nervous system may take part in axo-glial interactions and regulate the ion homeostasis required for the rapid and timely induction and propagation of action potentials at the axon initial segments (AIS) and at the nodes of Ranvier, respectively (Bruckner et al. 1993, 2006; Hedstrom and Rasband 2006; Poliak and Peles 2003; Salzer 2003). In this context it is interesting to note, that neurocan, brevican, RPTP- β /phosphacan and also Tn-C null-mice display certain alterations of hippocampal long-term potentiation (in RPTP- β knockout mice probably only linked to transmembrane variants), while a decreased axonal conductance has been measured in the optic nerve of Tn-R mutants (Brakbusch et al. 2002; Evers et al. 2002; Weber et al. 1999; Zhou et al. 2001).

It has also been suggested that proteoglycans in the extracellular meshwork surrounding myelin-free and thus, exposed nodal regions and axon initial segments may be implemented in neuroprotective functions (Miyata et al. 2007; Morawski et al. 2004).

Finally, abrogation of Tn-R expression impairs in adult knockout mice the detachment and radial migration of neuroblasts into the outer layers of the olfactory bulb. In contrast, ectopic expression reroutes the neuroblasts that originate from the subventricular zone of the lateral ventricles and first move tangentially in the rostral migratory stream. This suggests that Tn-R alone may also act as positive cue for adult neuroblast migration (Saghatelian et al. 2004).

Despite the panoply of putative functions of the extracellular matrix in the nervous system presented in the last years, the picture of its main roles in development and maturation has remained largely fragmented. It is to hope, that after a long period of neglect, the recently renewed research interests will boost the assembly of this fascinating puzzle.

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