

Review

Versican and the regulation of cell phenotype in disease[☆]Thomas N. Wight^{a,*}, Michael G. Kinsella^a, Stephen P. Evanko^a, Susan Potter-Perigo^a, Mervyn J. Merrilees^b^a Matrix Biology Program, Benaroya Research Institute at Virginia Mason, Seattle, WA, USA^b Department of Anatomy with Radiology, Faculty of Medical and Health Sciences, University of Auckland, Auckland, New Zealand

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ABSTRACT

Background: Versican is an extracellular matrix (ECM) proteoglycan that is present in the pericellular environment of most tissues and increases in many different diseases. Versican interacts with cells to influence the ability of cells to proliferate, migrate, adhere and assemble an ECM.

Scope of review: The structure of the versican molecule is briefly reviewed and studies highlighting those factors that promote versican synthesis and degradation and their impact on cell phenotype in disease are discussed. Particular attention is given to vascular disease, but other diseases where versican is important are covered as well, most notably different forms of cancers. Attention is given to mechanisms(s) by which versican influences cell behaviors through either direct or indirect processes. Versican produced by either stromal cells or myeloid cells can have a major impact influencing immunity and inflammation. Finally, studies controlling versican accumulation that either delay or inhibit the progression of disease will be highlighted.

Major conclusions: Versican is one component of the ECM that can influence the ability of cells to proliferate, migrate, adhere, and remodel the ECM. Targeting versican as a way to control cell phenotype offers a novel approach in the treatment of disease.

Significance: ECM molecules such as versican contribute to the structural integrity of tissues and interact with cells through direct and indirect means to regulate, in part, cellular events that form the basis of disease. This article is part of a Special Issue entitled Matrix-mediated cell behaviour and properties.

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1. Introduction

The extracellular matrix (ECM) is a reinforced composite of collagens and elastic fibers embedded in a viscoelastic gel of proteoglycans, hyaluronan (HA), and water, together with a wide variety and arrangement of assorted glycoproteins [1–4]. These molecules interact by entanglement and crosslinking to form a bioactive polymer which, in part, regulates the biomechanical properties of tissues and the phenotype of the cells that live in those tissues. This regulation involves molecular interactions that govern the attachment of cells to their ECM scaffolds through integrin and non-integrin receptors, detachment of cells from those scaffolds and molecular rearrangements in the ECM that allow cells to change shape during morphogenetic and remodeling events that occur in development and disease. The amount and composition of

the ECM is controlled by the co-ordinated and differential regulation of synthesis and turnover of each of the ECM components. It is becoming increasingly evident that individual components of the ECM can have dramatic effects on cell behavior.

Versican is a proteoglycan that is present in the ECM around cells and in the interstitial space between cells in most tissues [5–9]. In normal tissues, levels of versican are low but in disease versican increases dramatically, as is seen in the early intimal vascular lesions typical of developing human atherosclerosis (Fig. 1) (see reviews, [10,11]). In addition, versican accumulates in several types of cancers (see reviews [12–14]) and in many other diseases as well [10]. Versican is negatively charged due to its glycosaminoglycan (GAG) chains and attracts water, contributing to the viscoelasticity of the pericellular microenvironment [5,8]. In addition, versican interacts with a number of ECM components near the cell surface including HA, tenascin-R and -C, thrombospondin 1, fibronectin, and fibrillin [8,15–19] to create a mechanically active biopolymer around cells that influences the ability of cells to change shape, adhere, proliferate, migrate, and assemble other ECM components and survive. Versican, and ECM molecules that bind to versican, may modify the mechanical stiffness around cells contributing to alterations in mechanotransduction influencing cell behavior and phenotype [20–22]. Versican can also act as a reservoir for cytokines and growth factors to be released at various times, establishing fine control over cell activity and behavior [10,15,23,24]. All of these activities attributed to versican are diagrammatically depicted in Fig. 2 as a working

Abbreviations: ECM, Extracellular matrix; HA, hyaluronan; GAG, glycosaminoglycan; CS, chondroitin sulfate; ASMCs, arterial smooth muscle cells; PDGF, platelet-derived growth factor; KO, knockout; miRNA, microRNA; UTR, untranslated region; CNS, central nervous system; MMPs, matrix metalloproteinases; PSGL-1, P-selectin glycoprotein-1; ADAMTS, A Disintegrin And Metalloproteinase with Thrombospondin Motifs; SNPs, single nucleotide polymorphisms; EGF, epidermal growth factor; TLR2, toll-like receptor 2; FAK, focal adhesion kinase; α_1 , inter-alpha-trypsin inhibitor; ER, endoplasmic reticulum

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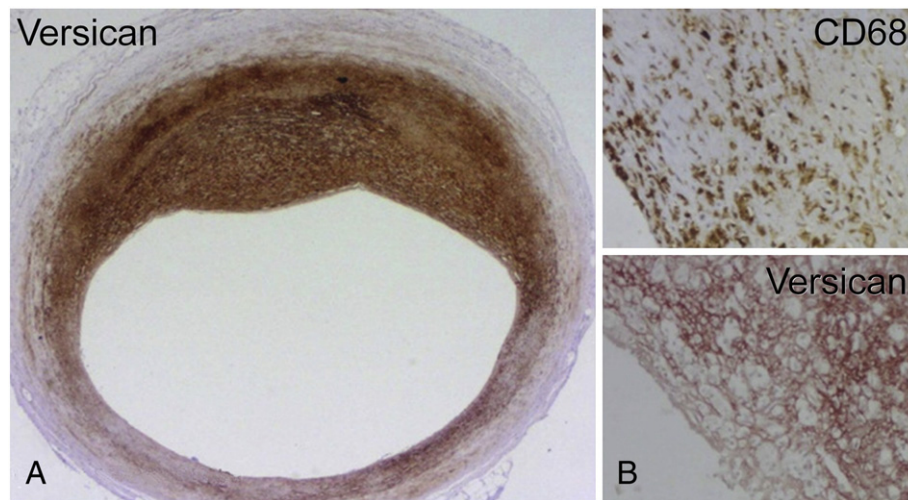


Fig. 1. A. A section through an early human coronary atherosclerotic lesion immunostained for versican illustrating marked accumulation of the ECM proteoglycan. B. Upper panel – These lesions frequently contain macrophages, as indicated by positive staining with antibody to CD68. Lower panel – Adjacent section immunostained for versican illustrating frequent co-localization of macrophage accumulation with versican. Sections were kindly provided by Drs. Frank Kolodgie and Renu Virmani, CV Path Institute, Inc., Gaithersburg, MD.

hypothesis and will be used in this review to discuss the importance of versican in influencing cell behavior in the pathogenesis of disease [5,11,25,26]. The focus will be primarily on vascular disease, but will highlight other diseases as well to show similarities for versican's action on the disease phenotype of cells. The fact that versican plays a central role in a number of diseases makes it a prime candidate for therapeutic intervention in the treatment of diseases [27–29].

The versican molecule exists in at least 4 different variants generated by alternative splicing of the two large internal exons that code for two chondroitin sulfate (CS) attachment domains in the core protein [5,6] (Fig. 3). These variants differ in the size of the core protein and the number of attached CS chains. While the V0 and V1 variants of versican bear CS chains and are the major forms that accumulate in most diseases, the V3 variant lacks CS chains and is not elevated in disease to any significant extent. While the V0 and V1 isoforms promote events associated with inflammation and disease, we [30–34] and others [35–38] have found that V3 acts as a dominant negative by reducing the CS-carrying isoforms of versican that accumulate in tissue, thus inhibiting the effects of V0/V1 on cell phenotype and the inflammatory response (see below). Such findings have led to the proposed use of V3 as a means to reduce

versican accumulation in tissues to inhibit the inflammatory response [27,39].

The CS GAG chains attached to versican may vary in size and composition, depending upon the tissue type, species of origin, cell type, and culture conditions. For example, CS chains isolated from versican synthesized by non-human primate arterial smooth muscle cells (ASMCs) have a chondroitin-6-sulfate to chondroitin-4-sulfate ratio (6S:4S) of 2 which increases to approximately 4 upon platelet-derived growth factor (PDGF) stimulation and proliferation of the ASMCs [40,41]. The significance of such changes during the growth response is uncertain, although binding studies show that the 6S isoform binds with greater avidity to positively-charged proteins than the 4S isoform. Such stimulation also increases the length of the CS chains attached to versican, leading to an overall increase in the hydrodynamic size of the proteoglycan. Chain elongation increases the capacity of the versican molecule to bind lipoproteins, potentiating the capacity of versican to trap lipoproteins in the development of atherosclerosis [42–44]. Signaling pathways controlling transcription of the core protein in response to PDGF differ from the signaling pathways that regulate the post-translational processing of the CS chains attached to versican [40]. Other examples of post-translational modifications that can occur are in the degree of sulfation of the CS chains attached to versican. CS chains isolated from tumor tissue are frequently oversulfated [45] and such differences may be important in controlling key interactions of these molecules with ligands important in the progression of different types of cancers. An excellent comprehensive review of factors that control CS synthesis is now available [46].

Attempts to approach the functional aspects of versican's control over cell behavior by gene knockout (KO) in chick embryos have been frustrating as a result of the early mortality of versican null mice [47]. However, some recent success has been achieved obtaining a partial KO of the versican gene, resulting in the expression of versican lacking the A subdomain of G1, which, in turn, significantly decreases versican accumulation and impacts cell growth and migration [48,49].

2. Regulated versican synthesis and impact on cell phenotype

The synthesis and degradation of versican in many tissues is highly regulated during events associated with disease [50]. There are several transcription factor binding sites in the versican promoter in addition to a classic TATA box sequence (see reviews [51–53]). In addition to positive regulatory elements, versican synthesis is under microRNA (miRNA) control as well. For example, myocardin is a transcription factor that regulates ASMC-specific gene expression and stimulates

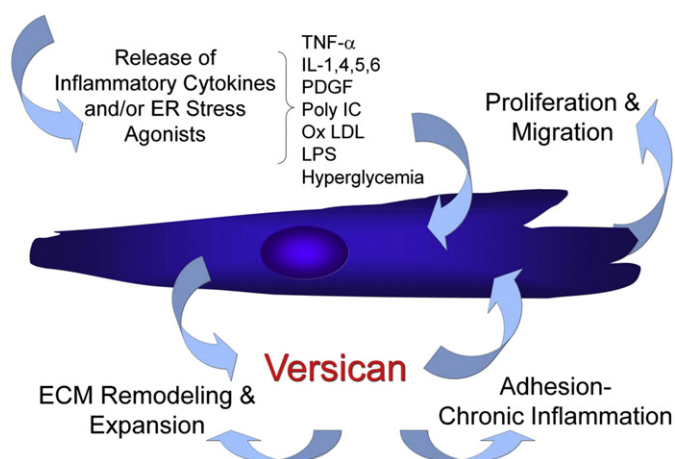


Fig. 2. A working model for the involvement of the ECM molecule versican influencing the behavior of cells during injury, remodeling, and inflammation in disease. The postulate follows that initial stimuli associated with disease initiation stimulate stromal cells or leukocytes to synthesize an ECM enriched in versican, which in turn influence these cells or other cells to proliferate, migrate, adhere, and remodel the ECM, promoting phenotypic changes associated with disease progression.

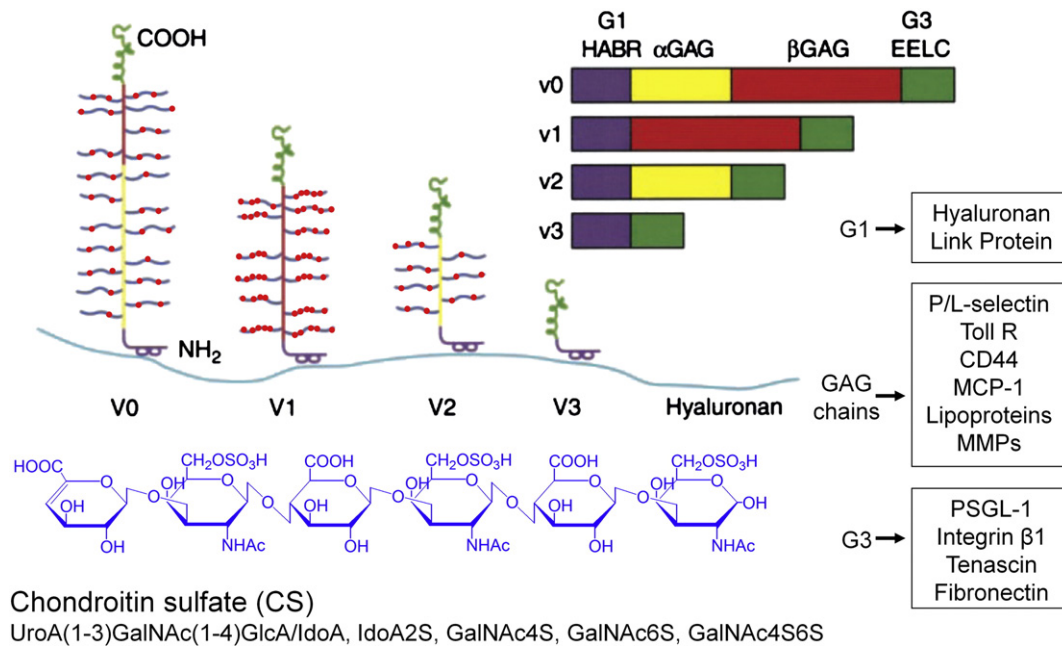


Fig. 3. A model of the different isoforms generated by alternative splicing of the mRNA transcript for versican. All isoforms interact with hyaluronan, forming different sized aggregates. Different colors denote specific domains in the gene and in the protein and carbohydrate product. Purple = hyaluronan binding region (HABR); yellow = α GAG exon and protein product; red = β GAG exon and protein product; green = two epidermal growth factor repeats (EE), a lectin-binding domain (L) and a complement regulatory region (C). Bottom panel shows monosaccharides linked together to form the CS GAG with red dots denoting charged residues. Versican interacts through specific domains in the molecule with a number of other molecules found on the surface of cells and within the ECM, some of these are depicted in the boxes to the right. These interactions form complexes which can also influence cell phenotypes. (Modified from Wight [5], and used with permission from Elsevier).

miRNA-143 expression. This miRNA binds to the 3' untranslated region (UTR) of the versican gene and suppresses versican expression affecting ASMC migration [54]. These results indicate that part of the ASMC differentiation program involves decreased versican production. The 3' UTR of the versican gene has other regulatory functions as well. For example, Burton Yang's group has recently demonstrated that the versican 3' UTR antagonizes other miRNAs [55] and most recently they have shown that this UTR can target miRNAs involved in regulating protein expression in growth control and facilitating tumor formation [56,57].

There is considerably less known about whether the synthesis of the different isoforms of versican is differentially regulated in disease. The V1 isoform is the most abundant in most adult tissues and seems to be the isoform that is most highly regulated in events associated with repair and remodeling in injury and disease, with the exception of diseases of the nervous system (see below). The V0 isoform appears to be more prominent during embryonic development [58] and less abundant in adult tissues [59]. The V2 isoform is mostly found in the central nervous system (CNS) [60], but has been found in other tissues as well. A number of studies show mRNA expression of V3 in a variety of tissues and synthesized by a variety of cells [59], but few studies have identified native V3 protein in the tissue and cells. However, V3 has been identified when overexpressed in different experimental conditions. For example, forced expression of V3 in a chondrogenic cell line disrupts the deposition and organization of V0/V1 isoforms and inhibits mesenchymal condensation and chondrogenesis [35]. Kern and colleagues [36] transduced mouse embryonic cardiomyocytes with V3 and noticed a marked reduction in proliferation of the cardiomyocytes and a significant increase in myocardial cell–cell association. Furthermore, injection of an adenovirus that contained the V3 gene into a developing mouse heart led to an increase in the outflow track myocardium and at least a two-fold increase in the compact layer of the ventricular myocardium. Notably, this study found that when only the G1 domain of versican was expressed, opposite effects were seen [36]. Given that V3 may resist degradation, these findings suggest that V3 may have dominant effects on phenotype when compared to degraded forms of versican. We have found that forced V3 expression in ASMCs and fibroblasts creates

dramatic effects on cell phenotypes [30–34] (see below). Recently, a new versican isoform, V4, consisting of the G1 domain, the first 398 amino acids of the β -GAG region and the G3 domain has been found to be upregulated in human breast cancer lesions [61]. Other isoforms potentially may exist, such as a V5 isoform, consisting of essentially only the G1 domain, found by new gene discovery techniques and listed as a reference sequence for mouse versican in Entrez Gene.

Proteoglycans are also synthesized by leukocytes and may play a role in the inflammatory response [62], also see reviews [23,39]. A number of studies have identified versican as a gene in monocytes which is upregulated in a number of pro-inflammatory states (see reviews, [10,23,39]). Versican synthesis is increased when monocytes differentiate into macrophages along with other proteins involved in the inflammatory response [63] (Fig. 4). Versican has been identified as a gene that is differentially expressed in M1 macrophages as opposed to M2, but it is not clear if versican is critical for M1 macrophage differentiation [64,65]. Versican expressed by macrophages binds to matrix metalloproteinases (MMPs) such as MMP9 [66] and chemokines such as CCL2 [67,68], suggesting a role for versican in determining macrophage phenotype in inflammation. In addition, macrophages in tumors express versican and can regulate mesenchymal to epithelial transitions and metastasis through effects on tumor cell proliferation, as recently shown [69,70].

3. Regulated versican degradation and impact on cell phenotype

The degradation of versican is associated with several tissue remodeling and inflammatory events in disease. Once bound to the versican-containing ECM, leukocytes may degrade the ECM to generate pro-inflammatory fragments that further drive the inflammatory response [71–75]. For example, the G3 domain of versican interacts with P-selectin glycoprotein-1 (PSGL-1) on the surface of macrophages to promote macrophage adhesion and aggregation [76].

Versican is degraded by a number of different proteases, including several MMPs, plasmin, and at least 5 members of the ADAMTS (A Disintegrin And Metalloproteinase with Thrombospondin

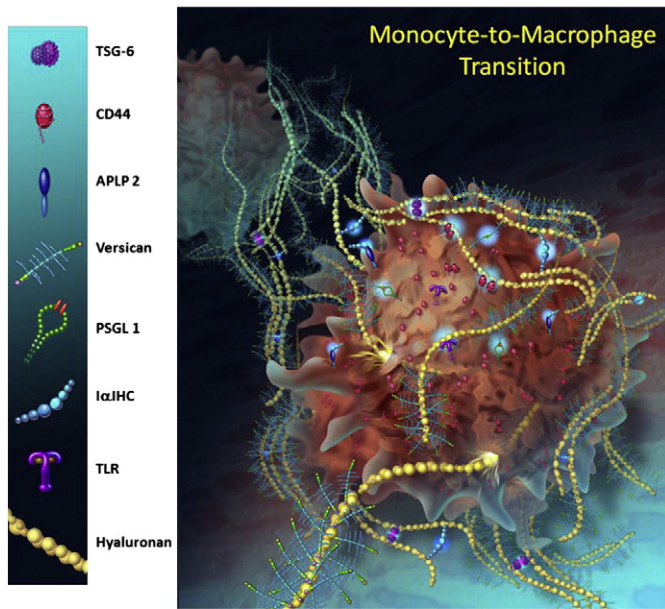


Fig. 4. A model to depict some of the ECM molecules expressed by monocytes, including versican, as they differentiate into macrophages. Illustration by Kate Sweeney, University of Washington; based on data presented in Chang et al., 2012, [63].

Motifs) family of proteases (see reviews [77,78]). The versican cleavage site for these ADAMTS enzymes is in the G1 domain of versican at the Glu441–Ala422 bond which generates a 70 kDa fragment that can be recognized by an antibody against the neopeptide sequence DPEAAE [79]. Altering the capacity of ADAMTS proteases to degrade versican leads to increases in versican in the pericellular matrix of fibroblasts and an increase in the myofibroblastic phenotype [80]. Such studies highlight a potentially significant role of versican in fibrosis. In addition, areas of increased ADAMTS-1 and -4 in early developing lesions of the vascular wall correlate with greater versican degradation and production of the amino terminal DPEAAE-containing versican fragment [79]. Alteration of blood flow in a baboon model of vascular graft repair promotes regression of vascular lesions by increasing versican degradation through ADAMTS activity [81]. The increase in cleaved versican correlates with regression of neointimal thickenings and loss of versican [81]. Interestingly, these changes in versican integrity also correlate with cell death in the regressed lesions [82]. It remains to be shown however, whether versican fragments promote ASMC death in this model. These findings are of interest because of studies demonstrating that cleaved versican regulates apoptosis during mammalian inter-digital web regression [83]. Of further interest is the finding that proteolytic cleavage products of versican are present in human plaques from endarterectomy segments, consistent with their generation in a pro-inflammatory microenvironment [84]. Notably, an MMP-12-derived versican fragment has been identified in plasma samples from patients with atherosclerotic heart disease, confirming versican degradation as part of the atherosclerotic response [85]. ADAMTS-1 mRNA transcript is also abundant in human aorta and increases as ASMCs migrate and proliferate *in vitro* [86]. A polymorphism in the ADAMTS-1 gene has been associated with an increase in cardiovascular disease in two separate studies [87,88]. Furthermore, high levels of ADAMTS-1 in brain tissues are associated with neurodegenerative diseases such as Down syndrome, Alzheimer's, and Pick's disease [89]. However, a direct causal link for versican in these diseases awaits further experimentation. On the other hand, single nucleotide polymorphisms (SNPs) and haplotype analyses of the versican gene in intracranial aneurysms revealed a strong association [90,91].

4. Versican and cell proliferation

Accumulation of versican occurs in tissues undergoing cellular proliferation. For example, mitogens such as PDGF upregulate versican expression in ASMCs as they are stimulated to divide [7,41,92,93]. Together with HA, versican contributes to the expansion of the pericellular ECM that is required for the proliferation of these cells to occur [7,8,92]. These complexes increase the viscoelastic nature of the pericellular matrix, creating a highly malleable extracellular environment influencing mechanotransduction and supporting a cell-shape change necessary for cell proliferation and migration to take place [8] (Fig. 5). Inhibiting the formation of this pericellular coat blocks the proliferation of ASMCs in response to PDGF [7,8]. Although PDGF stimulates the proliferation of ASMCs, TGF- β 1 which also stimulates versican synthesis [41] inhibits ASMC proliferation *in vitro*, suggesting that versican synthesis is not directly causatively linked to the proliferative phenotype. However, interference with versican synthesis in ASMCs, fibroblasts, and in some cancer cells inhibits their proliferation, suggesting that versican synthesis and accumulation is necessary, but not sufficient to cause changes in mitotic cell activity [12,14,31,34]. Thus, the versican–HA complex that surrounds cells serves as an important, but infrequently considered, mechanism for controlling cell shape and cell division.

Another mechanism by which versican could influence proliferation is by acting as a mitogen itself, by binding to growth factor receptors via epidermal growth factor (EGF) sequences in the G3 domain of the molecule [15]. For example, expression of G3 mini-genes in NIH/3T3 cells enhances cell proliferation, and the effect can be blocked by deletion of the EGF domains in the G3 construct [94]. This same construct exerts a dominant negative effect on cell proliferation through inhibiting the binding of G3 to the cell surface, via the lectin domain in G3 [15,95]. The concentration of versican associated with the cell surface appears to be a critical factor, and loss of versican from the cell surface is associated with decreased cell proliferation. Maximal growth-promoting activity is achieved in NIH/3T3 cells and chondrocytes with both G1 and G3 mini-gene constructs, supporting the concept that versican regulates proliferation by binding directly to a growth factor receptor and by interfering with cell adhesion [94,96]. Work in NIH/3T3 cells *in vitro* suggests that V1 and V2 isoforms may have opposing activities. For example, the V1 isoform enhances the proliferation of NIH/3T3 cells and protects these cells from apoptosis, while V2 decreases their proliferation and has no activity on apoptosis [97]. V2 has recently been shown

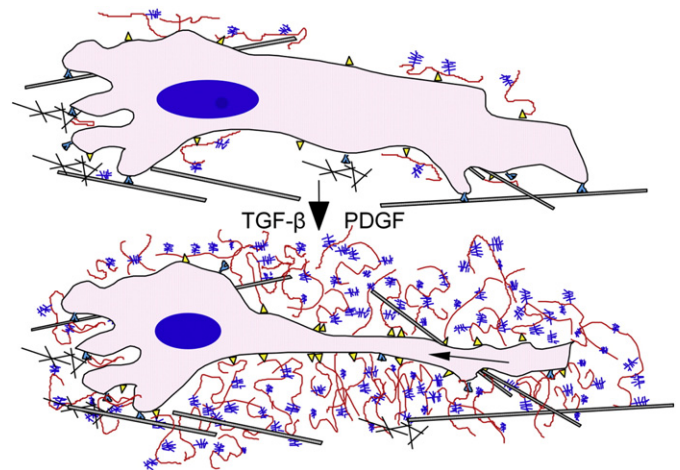


Fig. 5. ECM transitions required for cell proliferation and migration. In order for cells to change shape during division and migration, they must modify their pericellular environment by first degrading the existing ECM and replacing it with components that allow the cell to change shape. Two ECM molecules that are produced during these events and allow this to happen are hyaluronan (red) and versican (blue). These changes lead to expansion of the pericellular matrix and changes in the mechanical properties of the ECM that influence cell phenotype. [Modified from Wight et al., 2011, Am J Physiol Gastrointest Liver Physiol.]

to enhance angiogenesis of endothelial cells by slowing cell proliferation and enhancing fibronectin synthesis [98]. Additional recent studies indicate that the V3 isoform may regulate cell proliferation as well. The capacity of V3 to promote a specific phenotype opposite to what is seen with V0/V1 is manifested when ASMCs are transduced with V3 [32]. Forced expression of V3 causes decreases in cell growth [32]. Interestingly, suppression of V0/V1 synthesis had the same effect as V3 expression on ASMC and skin fibroblast proliferation [30,31].

Thus, versican expression is associated with a proliferative cell phenotype and is often found in tissues exhibiting elevated proliferation, such as in tumors (see reviews, [12,14]). Using a mouse model of spontaneous breast cancer, Gao and colleagues demonstrated that bone marrow-derived myeloid progenitor cells in the pre-metastatic mouse lung secrete versican, which promotes a mesenchymal to epithelial transition of tumor cells and subsequent metastasis [69,70]. Versican derived from myeloid cells in this model appears to promote tumor growth by enhancing tumor cell proliferation, possibly by blocking the TGF β -smad2/3 pathway. Earlier studies demonstrated that versican accumulation in Lewis lung carcinoma can interact with macrophage TLR2 to induce secretion of inflammatory cytokines, such as tumor necrosis factor- α (TNF α) and IL-1–6 [99,100] promoting tumor expansion and metastasis. It is of interest that highly sulfated CS GAG chains on versican may be critical to promote this activity [45,101]. Li and colleagues recently demonstrated that in co-culture experiments of macrophages and ovarian cancer cells, versican produced by the tumor cells promoted the production of an antimicrobial protein through interacting with toll-like receptor 2 (TLR2) on macrophages that, in turn, promoted ovarian tumor cell proliferation [102]. Evidence is accumulating that versican, either directly or indirectly, is functioning as a matrine or agonist in promoting tumor cell proliferation. Whether similar activity for versican exists during inflammation in non-cancerous tissue awaits further investigations. It may be that ligation of immune receptors, such as TLR2, by versican is responsible for the activation of multiple cell types and the induction of inflammatory cytokine secretion in many disease situations [100,103]. Along with its effects on the proliferation of cancer cells, versican enhances cell survival and apoptotic resistance of these cells [104,105], and protects them through inhibiting cytotoxic drug action and Fas-dependent programmed cell death [105].

There is also evidence that specific isoforms of versican may affect the proliferative activity of tumor cells. For example, Bassols and colleagues found that expression of V0/V1 isoforms of versican is increased in malignant melanoma, which contributes to the increased proliferation rate and decrease adhesion of the tumor cells [106,107]. Overexpression of V3 by melanoma tumor cells reverses this phenotype by decreasing proliferation and increasing adhesion [108]. The decrease in proliferation was accompanied by a decrease in the activation of ERK1/2 in response to EGF [38]. Recent studies by this group now show that V3 interferes with the CD44-EGFR/ErbB2 pathway in the regulation of cell proliferation and migration [109,110].

5. Versican and cell migration

Versican also affects the migration of a number of cells [111]. For example, PDGF not only promotes versican expression and ASMC proliferation, but can also affect ASMC migration primarily through inhibition of miRNA-143 and regulation of versican synthesis [54]. Versican is expressed also along neural crest pathways and influences neural cell migration [58,112]. A number of studies suggest that versican guides neural crest migration because cells do not enter the tissues that express versican [113–115]. Pax3 is a transcription factor associated with defective neural cell migration. Splotch mice are characterized by mutations in the Pax3 gene and exhibit neural crest-related abnormalities, including the failure of neural crest cells to colonize target tissues. However, neural crest cells derived from these mutant mice migrate as controls *in vitro*, so it has been suggested that the defect may not reside in the

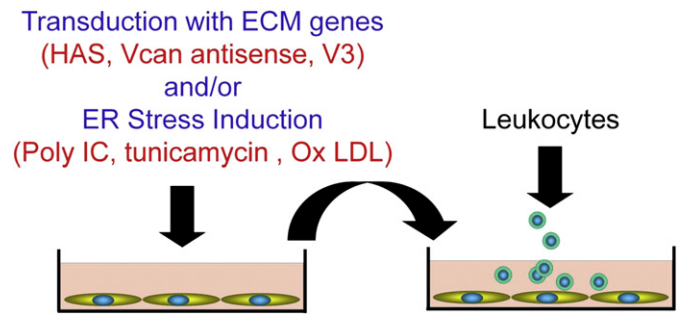


Fig. 6. An *in vitro* protocol to examine the generation of ECMs that bind leukocytes. Stromal cells such as ASMCs, lung fibroblasts, or synoviocytes are transduced with matrix genes or exposed to agents that promote ER stress, and the ECM generated and assessed for ability to support leukocyte adhesion. HAS = hyaluronan synthase; Vcan = versican; V3 = versican isoform V3; Ox LDL = oxidized low density lipoprotein.

neural crest cells themselves, but rather in the ECM environment through which they migrate. Indeed, earlier studies [113] demonstrated that versican was markedly overexpressed in Splotch mutants in neural crest cell migration pathways, suggesting that versican may be responsible for defective cell migration in this species. Other studies show that overexpression of Pax3 in a medulloblastoma cell line causes upregulation of the V2 splice variant of versican and a downregulation of the V3

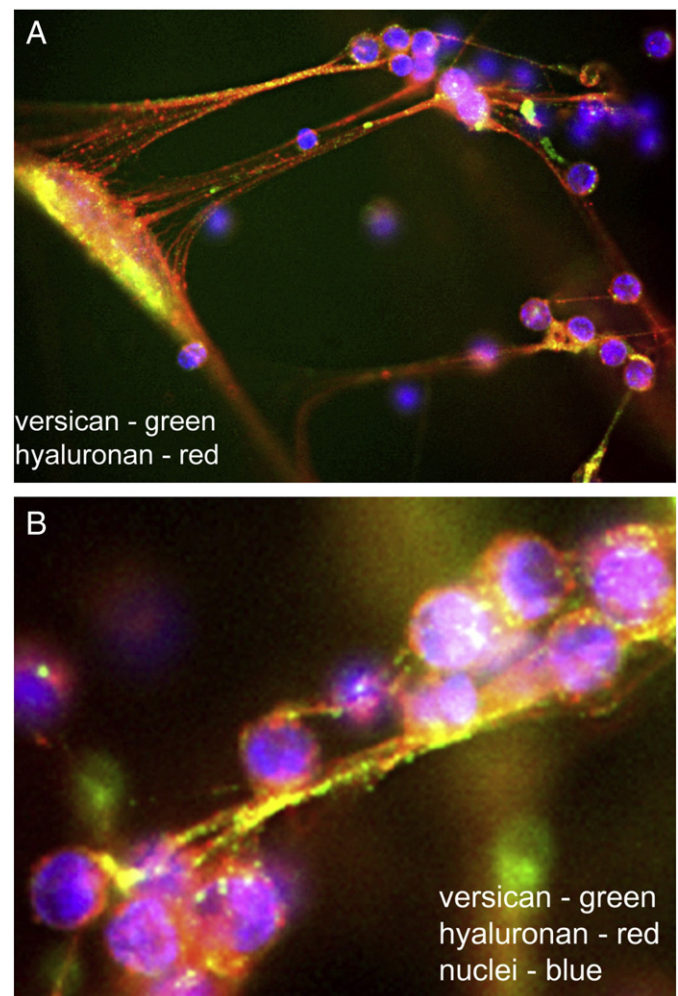


Fig. 7. A. Lung fibroblasts stimulated with poly I:C produce a copious amount of ECM enriched in HA (green) and versican (red) in the form of long cables that bind added CD4⁺ activated T cells (blue). B. Higher magnification showing the co-localization of versican and hyaluronan cables and the HA-rich surface of the T cells. Reprinted from Evanko et al., Matrix Biol, 2012 [117] with permission from Elsevier.

variant [116]. Such differential regulation of the versican isoforms may explain, in part, the migratory defect in the Splotch mouse. It is of interest that the V3 isoform lacks CS chains, which should reduce the exclusionary properties of the ECM. Clearly, expression of V3 in ASMCs decreases their migratory activity [32]. Versican can also influence the migration of lymphoid cells. Recent experiments indicate that T cells do not migrate when they are added to an ECM enriched in versican and HA, but this inhibition of migration can be reversed if this ECM is first pretreated with blocking antibodies to versican [117].

Versican influences the migration of a variety of other cell types, and this activity appears to be mostly associated with the anti-adhesive activities involving the G1 domain of the molecule. In the nervous system and in axonal growth, the V2 splice variant inhibits axonal outgrowth and migration [60,118–122]. This inhibiting activity of versican can be reduced, but not eliminated, by removing the CS chains, indicating that multiple domains of versican are involved in controlling axon regeneration. Although the V2 isoform is widely present in the CNS [123], it is predominately localized to the myelinated fiber tracts. Oligodendrocytes are the likely source of V2 [124,125]. The finding that both the GAGs and core protein domains of the molecule are involved in the inhibitory activity suggests a direct interaction with the cells, or modification of the surrounding matrix to form exclusionary boundaries. The fact that versican plays a fundamental role in axonal migration is highlighted by studies that show upregulation of versican along with other hyalactins following CNS injury [124]. These changes have been associated with the failure of nerves to regenerate. The importance of the hyalactins in preventing nerve regeneration is further highlighted by studies that show that degradation of CS chains by chondroitinase ABC lyase treatment following spinal cord injury in experimental animals promotes regeneration of both ascending and descending corticospinal-tract axons [126]. Such results suggest that manipulating versican synthesis in spinal cord injury may be a useful intervention for therapeutic treatment of this condition. No doubt versican does not act alone in creating these exclusionary boundaries and it will be important to identify other key players with which versican interacts. Failure of axons to regenerate is also characteristic of multiple sclerosis, and versican appears to increase in plaques present in the white matter of the brain from patients with multiple sclerosis [127].

Versican also appears to be involved in the motility of cancer cells. A number of studies suggest that tumor stromal-derived versican can influence tumor cell motility and invasiveness to promote metastasis in several different cancers [12,107,128–133].

6. Versican and cell adhesion

Early studies showed that versican is anti-adhesive [9,134–136], and this activity appears to reside in the G1 domain of versican [96,134]. However, the carboxyl-terminal domain of versican interacts with the $\beta 1$ integrin of glioma cells, activating focal adhesion kinase (FAK), promoting cell adhesion and preventing apoptosis in this cell type [96,134,137]. The pro-adhesive property of the G3 domain of versican raises the possibility that different breakdown products of versican might affect cell adhesion in different ways. Interestingly, overexpression of V3 which lacks the CS chains, but contains the G1 and G3 domains, leads to extensive cell spreading and increased adhesion in ASMCs [32].

We and others have recently shown that versican can influence the adhesion of myeloid and lymphoid cells as part of the inflammatory response see reviews [23,39]. As myeloid and lymphoid cells enter tissues, they come into contact with specific components of the ECM, such as HA and versican, together with a number of other proteins including TSG-6 and inter-alpha-trypsin inhibitor (I α I) which promote their adhesion [39,138–142]. These components can interact to form filamentous cable-like structures that serve as scaffolds for the infiltrating cells. Using *in vitro* models and stimulating stromal cells such as fibroblasts and ASMCs with agonists that promote endoplasmic reticulum (ER) stress, such as poly I:C or tunicamycin [139,141,143,144] (Fig. 6), we and others have shown that versican is a critical player in leukocyte adhesion to the ECM (Fig. 6) [39,142,145]. In addition to monocyte adhesion, human-activated T lymphocytes adhere to this ECM as well (Fig. 7) [117]. Not only do T cells adhere, but versican also significantly reduces their ability to migrate and produce IL-10 when in contact with the versican-enriched ECM. Such studies indicate that specific components of the ECM such as versican can have dramatic effects on the phenotype of immune cells. Thus, versican should be considered a potential therapeutic target in the control of the immune response associated with inflammation in disease [27–29]. The fact that the production of this leukocyte adherent ECM is formed by cells stimulated by ER stress agonists is highly significant and more attention needs to be given to the key pathways involved in the generation of this ECM.

7. Versican and ECM assembly

ECM remodeling takes place throughout different phases of disease progression as part of an injury and/or inflammatory response. These

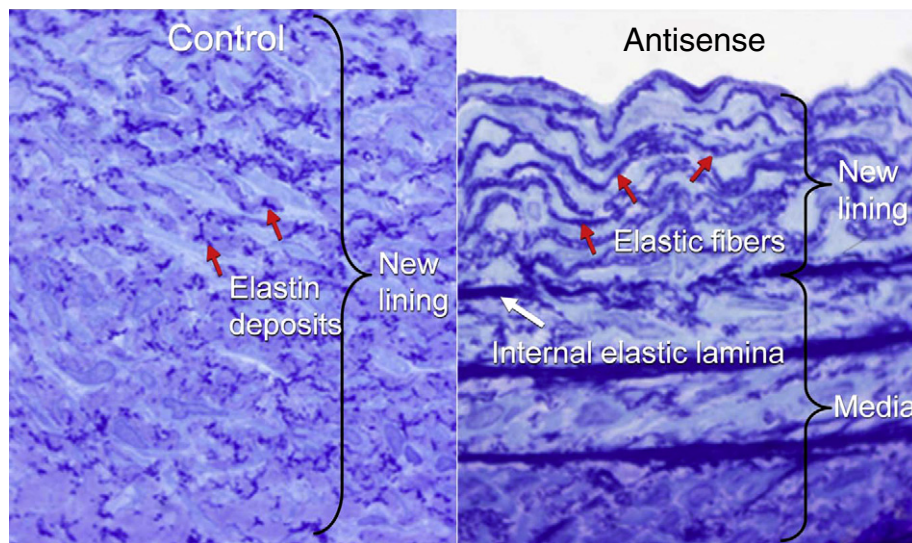


Fig. 8. Light micrographs of a 28-day neointima from a balloon-injured rabbit carotid artery seeded with ASMCs containing the LXS empty retroviral vector (left panel) or seeded with ASMCs transduced with a retroviral vector containing antisense to versican (right panel). The neointima seeded with the antisense-expressing cells contains multiple lamellae of elastic fibers.

phases involve breakdown and disassembly of various ECM components and reassembly of particular components as part of the pathogenesis of these diseases. The sequence of changes is not unlike what is seen during wound repair in which the early ECM changes are characterized by ECM deposits which create a loose, open, and watery matrix (referred to as a “provisional ECM”) [146–148] which allows for cellular invasion and repair. This provisional matrix is then replaced by a more fibrous ECM enriched in collagens and assorted glycoproteins. Versican interacts with several different ECM molecules and, in part, plays a central role in ECM assembly. The domain structure of versican lends itself to multiple types of interactions through either protein–protein or protein–carbohydrate interactions. Perhaps the best known of these interactions involves a specific interaction between the amino terminal domain of versican (G1) and HA [149]. This interaction is stabilized by another protein – link protein – which exhibits selective binding specificity for both HA and versican [150].

In addition to HA, versican interacts with other ECM molecules and controls their organization. Versican interacts with tenascin-R through the lectin-binding domain of versican and involves protein–carbohydrate interactions [17,151,152]. The lectin-binding domain participates in other ligand interactions as well. For example, versican interacts with fibulin-1 and fibulin-2 [153,154], a growing family of ECM proteins that are expressed in particularly high levels in the developing heart valve. In adults, however, fibulin-1 and -2 are found associated with microfibrils that are part of elastic fibers. Versican also can interact with proteins associated with elastin in elastic fibers. For example, versican interacts with the elastic fiber-associated protein fibrillin [19,153–155], and versican has been shown to co-localize with elastic fibers in the skin [19]. Furthermore, fibrillins bind fibulin-2, and fibulin is preferentially localized to the

elastin/microfibril interface in some tissues, but not in others [156]. It may be that fibulin serves as a bridge between versican and fibrillin, forming high-ordered multi-molecular structures important in the assembly of elastic fibers. The relationship of versican to elastic fiber assembly is interesting and unusual. What is conspicuously absent in newly remodeled ECM is elastic fibers. In fact, elastic fibers are conspicuously absent from atherosclerotic and restenotic lesions. The importance of elastic fibers in regulating vascular disease is highlighted by studies of the elastin KO mouse. Disruption of elastin synthesis by targeting the promoter and first exon of the tropoelastin gene [157] leads to subendothelial proliferation of ASMCs and obstructive vascular closure. In fact, cells from the elastin KO animals have been shown to proliferate more rapidly than their normal littermate cells *in vitro*, but normal proliferation is restored upon addition of elastin to the KO cells [158,159]. Elastin peptides have been shown also to regulate ASMC proliferation and migration [160] and elastin has been used to dampen the restenotic response in experimental animals [159]. Thus, factors regulating elastic fiber formation may be critical to controlling vascular lesion formation. One factor that appears to inhibit elastic fiber assembly is CS, which is part of the versican molecule [161]. We have found that blocking versican expression by antisense (Fig. 8) or using forced expression of the versican variant that lacks CS, V3, in ASMCs leads to changes in tropoelastin expression and accumulation of elastic fibers in long-term ASMC cultures [34]. When these V3-transduced ASMCs are seeded into balloon-injured rat carotid arteries, a compact and highly structured neointima enriched in elastic lamellae develops [34]. Furthermore, in a recent study, Merrilees and colleagues demonstrated that injecting rabbit V3-transduced ASMCs into injured rabbit carotid arteries in animals placed on a lipid rich diet prevented lipid build up and monocyte ingress over an 8-week period [33] (Fig. 9).

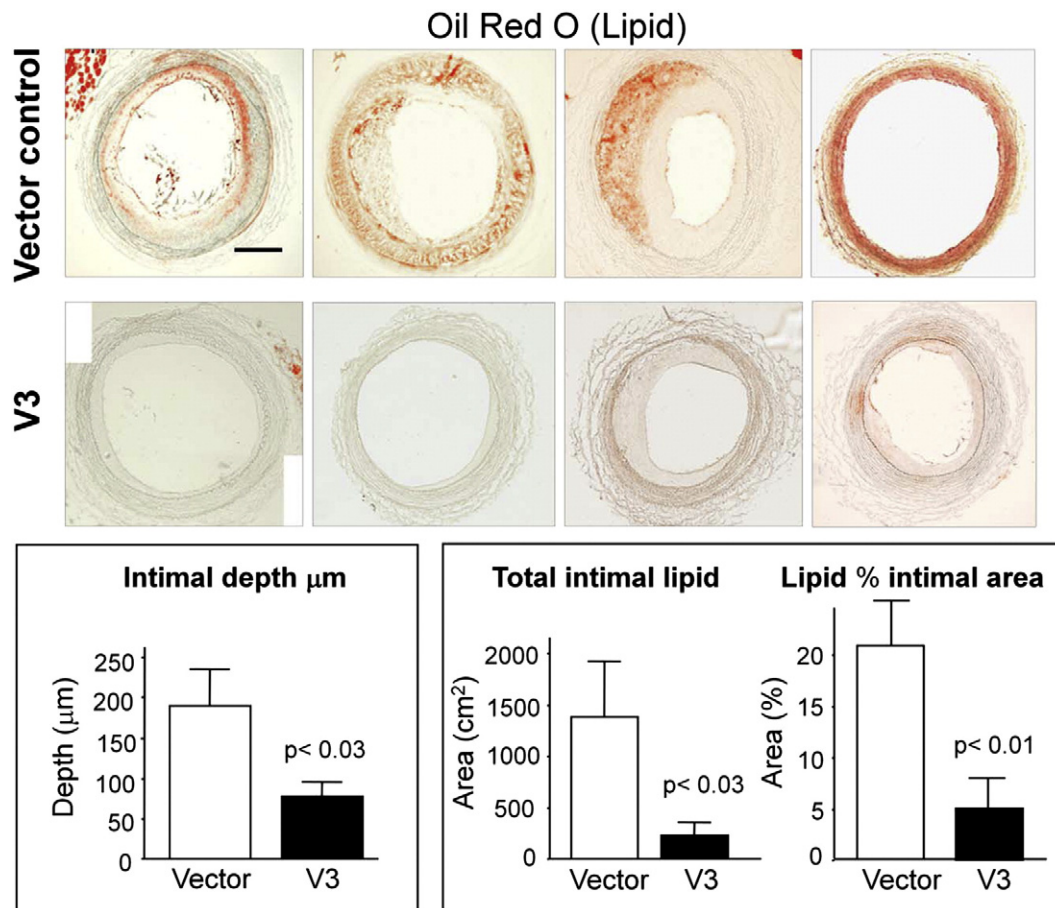


Fig. 9. Lipid accumulation in balloon-injured rabbit carotid arteries seeded with ASMCs transduced with an empty retroviral vector (top row) or with ASMCs transduced with a retroviral vector containing the V3 gene (middle row) in animals fed a high fat diet for 8 weeks. Lipid is excluded from the arteries expressing the V3 gene. Bottom row is the quantitation of lipid using the oil red O stain and morphometric analyses. [From Merrilees et al., 2011.]

Previous studies had shown that monocytes do not adhere well to elastin, but adhere avidly to collagen [162], suggesting that elastin is a poor adhesive substrate for monocytes. Thus, V3 may promote the formation of an ECM that resists monocyte adhesion and accumulation and may be an effective anti-inflammatory treatment in the prevention of cardiovascular disease. The impact of V3 on elastogenesis is further highlighted by studies that show that V3 transduction of skin fibroblasts taken from patients exhibiting defective elastogenesis, such as in Costello Syndrome and Hurler's Disease, reverses their phenotype and corrects impaired elastogenesis that characterizes these cells [30].

8. Conclusion

Versican is an ECM macromolecule that is critical to maintaining tissue integrity and homeostasis in the living organism. As reviewed in the studies cited above, not only is versican a structural component of the ECM, but also a molecule that interacts with cells to control their behavior in part by influencing cell shape and thus their phenotype in the pathogenesis of a number of diseases.

Versican can impact cell phenotype from the outside of the cell in different ways and it will be important in the future to decipher the molecular mechanisms(s) either directly and/or indirectly involved in versican determining cell fate. What is emerging is that not only does versican possess unique biological activity as an intact molecule or as part of complexes, but also it is likely that versican fragments generated as part of disease pathogenesis will affect cell phenotype in very unique ways. The fact that versican plays a central role in controlling cell behavior in disease suggests that it could become a potential selective target promising wide therapeutic benefits.

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