

Published in final edited form as:

*Matrix Biol.* 2012 January ; 31(1): 3–16. doi:10.1016/j.matbio.2011.10.001.

## Molecular functions of syndecan-1 in disease

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### Abstract

Syndecan-1 is a cell surface heparan sulfate proteoglycan that binds to many mediators of disease pathogenesis. Through these molecular interactions, syndecan-1 can modulate leukocyte recruitment, cancer cell proliferation and invasion, angiogenesis, microbial attachment and entry, host defense mechanisms, and matrix remodeling. The significance of syndecan-1 interactions in disease is underscored by the striking pathological phenotypes seen in the syndecan-1 null mice when they are challenged with disease-instigating agents or conditions. This review discusses the key molecular functions of syndecan-1 in modulating the onset, progression, and resolution of inflammatory diseases, cancer, and infection.

### Keywords

Syndecan; Proteoglycan; Heparan sulfate; Inflammation; Infection; Cancer

## 1. Introduction

Syndecans comprise a major family of cell surface heparan sulfate proteoglycans (HSPGs) (Bernfield et al., 1999; Park et al., 2000b; Couchman, 2010). Syndecans are expressed on the surface of all adherent cells and on many non-adherent cells. The mammalian syndecan family consists of 4 members, each encoded by distinct genes. Although cloning of the first syndecan, syndecan-1, was reported in 1989 (Saunders et al., 1989), its existence was known prior to this report as the Bernfield (Rapraeger and Bernfield, 1983) and Hook (Kjellen et al., 1981) groups have identified an HSPG intercalated in the plasma membrane of mouse mammary gland epithelial cells and rat hepatocytes, respectively. This HSPG was found to be expressed predominantly on the basolateral surface of epithelial cells (Rapraeger et al., 1986; Hayashi et al., 1987), bind to extracellular matrix (ECM) components, such as collagens I, III and V (Koda et al., 1985), fibronectin (Saunders and Bernfield, 1988) and thrombospondin (Sun et al., 1989), and associate with the actin cytoskeleton (Rapraeger et al., 1986). Because these properties indicated that this HSPG is well positioned to function as an anchor that stabilizes the morphology of epithelial sheets by connecting the ECM to the intracellular cytoskeleton, the name 'syndecan' (from the Greek *syndein*, which means to bind together) was given to this HSPG. Subsequently, syndecan-2 (fibroglycan) (Marynen et al., 1989; Pierce et al., 1992), syndecan-3 (N-syndecan) (Carey et al., 1992; Gould et al., 1992), and syndecan-4 (amphiglycan/ryudocan) (David et al., 1992; Kojima et al., 1992) were identified.

Syndecans are type I transmembrane proteins, consisting of an extracellular domain where heparan sulfate (HS) chains attach distally to the plasma membrane, followed by the signature transmembrane and cytoplasmic domains which are highly homologous among the different syndecans and across species. The transmembrane domain contains a GxxxG dimerization motif and mediates both homotypic and heterotypic dimerization of syndecans (Dews and Mackenzie, 2007). The syndecan cytoplasmic domain contains several conserved signaling and scaffolding motifs, including one invariant Ser, three invariant Tyr, and a Glu-Phe-Tyr-Ala PDZ binding domain at the C-terminus (Bernfield et al., 1999; Lambaerts et al., 2009; Couchman, 2010). Syndecans are expressed on different cells and locations at different times and levels (Bernfield et al., 1992). In adult tissues, syndecan-1 is predominantly expressed by both simple and stratified epithelial cells and plasma cells, although it is expressed at a lower level in several other cell types and its expression can be induced in these cells. Syndecan-2 is typically expressed by endothelial cells and mesenchymal cells, whereas syndecan-3 expression is mostly restricted to neural crest-derived cells. Syndecan-4 is expressed ubiquitously, but it is expressed at lower levels than other co-expressed syndecans on any given cell type.

Syndecans bind to and regulate many HS- and heparin-binding molecules (Bernfield et al., 1999; Fears and Woods, 2006). However, much debate surrounds the biological significance and specificity of syndecan interactions due to the large number of potential ligands. Available data indicate that syndecans can regulate the biological activity of ligands by affecting their stability, conformation, oligomerization, or compartmentalization (Park et al., 2000b). Further, although all syndecans contain the primary ligand-binding HS chains, they show distinct temporal and spatial expression patterns (Hayashi et al., 1987; Kim et al., 1994) and, thus, are likely to function specifically *in vivo*. In addition, syndecans can bind to ligands through their core protein (McFall and Rapraeger, 1998; Chen et al., 2004; Hayashida et al., 2006; Dews and Mackenzie, 2007; Beauvais et al., 2009; Couchman, 2010; Purushothaman et al., 2010), suggesting that extracellular, intramembrane, and intracellular interactions mediated by the core protein are also important in the biological functions of syndecans.

On the cell surface, syndecans function primarily as coreceptors that catalyze the encounter between ligands and their respective signaling receptors (Bernfield et al., 1999; Park et al., 2000b). Syndecans can also function as soluble HSPGs as their extracellular domain, replete with all their HS chains, can be proteolytically released from the cell surface by a process known as ectodomain shedding (Jalkanen et al., 1987; Bernfield et al., 1999; Hayashida et al., 2010). Syndecan shedding is thought to be an important post-translational mechanism that regulates syndecan functions as it both rapidly reduces the amount of cell surface HS and generates soluble syndecan ectodomains that can function as autocrine or paracrine effectors. Syndecan shedding is induced *in vitro* by several inflammatory factors (Fitzgerald et al., 2000; Park et al., 2004; Charnaux et al., 2005; Brule et al., 2006; Chung et al., 2006; Mahtouk et al., 2007; Yang et al., 2007) and *in vivo* under certain pathological conditions (Kainulainen et al., 1998; Kato et al., 1998; Park et al., 2001; Yang et al., 2002; Haynes et al., 2005; Xu et al., 2005; Hayashida et al., 2008a, 2009a, 2009b; Kliment et al., 2009; Hayashida et al., 2011; Haywood-Watson et al., 2011). The majority of syndecan shedding agonists induce shedding by activating outside-in signaling pathways (Fitzgerald et al., 2000; Park et al., 2000a, 2004; Hayashida et al., 2008b), and several metalloproteinases can shed syndecan ectodomains (Li et al., 2002; Endo et al., 2003; Ding et al., 2005; Brule et al., 2006; Fears et al., 2006; Purushothaman et al., 2008; Pruessmeyer et al., 2010). These data suggest that the timing, location, and extent of syndecan shedding are regulated by a hierarchical mechanism involving shedding agonists, intracellular signaling pathways, and metalloproteinases.

Surprisingly, mice lacking syndecan-1 (Alexander et al., 2000; Park et al., 2001; Stepp et al., 2002) or syndecan-4 (Echtermeyer et al., 2001; Ishiguro et al., 2001) are healthy and fertile, and do not show major pathologies. However, dramatic pathological phenotypes emerge when both syndecan-1 null (*Sdc1*<sup>-/-</sup>) (Table 1) and *Sdc4*<sup>-/-</sup> (Echtermeyer et al., 2001; Ishiguro et al., 2001; Kon et al., 2008; Echtermeyer et al., 2009; Jiang et al., 2010; Ikesue et al., 2011) mice are challenged with disease-causing agents or conditions. These observations suggest that other syndecans or HSPGs can functionally compensate for the loss of syndecan-1 or -4 during development, but not in certain post-developmental processes, such as the pathogenesis of diseases. This review seeks to provide an overview of the key molecular functions of syndecan-1 in inflammatory diseases, cancer, and infection.

## 2. Inflammatory diseases

Inflammation is a fundamental host response to endogenous or exogenous agents that have the potential to cause tissue injury. The inflammatory response removes or sequesters harmful agents, and facilitates the restoration of the normal structure and function of damaged tissues. Multiple regulatory mechanisms have evolved to actively contain and resolve the inflammatory response in a timely manner to prevent excessive inflammatory tissue injury. However, when one or several of these mechanisms fail, the inflammatory response can be exaggerated or sustained, and lead to several acute and chronic diseases, such as acute lung injury, sepsis, colitis, arthritis, asthma, and fibrosis of the lung, heart, skin, kidney, and liver. Syndecan-1 binds to many factors that mediate and regulate the inflammatory response (Bartlett et al., 2007). The significance of syndecan-1 interactions in inflammatory diseases is highlighted in several studies examining the response of animals or cells whose expression or function of syndecan-1 has been experimentally altered.

### 2.1. Syndecan-1 in leukocyte recruitment

The acute inflammatory response, which occurs in minutes to hours, involves the coordinated delivery of plasma proteins and recruitment of leukocytes to the site of infection or injury. Although exaggerated or sustained leukocyte responses lead to unwanted tissue injury, an adequate and timely recruitment of leukocytes is essential for repair as wounds tend to heal poorly and may even lead to lethal outcomes in individuals with insufficient leukocytes (Lekstrom-Himes and Gallin, 2000). Available data indicate that syndecan-1 regulates several steps of leukocyte recruitment in non-infectious inflammatory diseases (Fig. 1).

In oxazolone-induced delayed-type hypersensitivity (DTH), a widely used model of allergic contact dermatitis, *Sdc1*<sup>-/-</sup> mice showed increased leukocyte recruitment and prolonged edema formation (Kharabi Masouleh et al., 2009). The expression of pro-inflammatory cytokines (TNF $\alpha$ , IL-6), chemokines (CCL5, CCL3), and the adhesion molecule ICAM-1 was significantly increased in *Sdc1*<sup>-/-</sup> mice compared to wild type (Wt) mice. Syndecan-1 expression was reduced in Wt mice during the DTH response. Further, blocking antibodies against CD18 inhibited the increased adhesion of *Sdc1*<sup>-/-</sup> leukocytes to ICAM-1 (Kharabi Masouleh et al., 2009). Similarly, in dextran sodium sulfate-induced colitis, an exaggerated and prolonged recruitment of leukocytes was observed in *Sdc1*<sup>-/-</sup> mice relative to Wt mice (Floer et al., 2010). This was accompanied by increased expression of TNF $\alpha$ , CCL3 and VCAM-1, impaired mucosal repair, and higher mortality, consistent with the reduced expression of syndecan-1 and impaired gut mucosal repair seen in patients with inflammatory bowel disease (Day and Forbes, 1999). Enhanced leukocyte adhesion and transendothelial migration, and increased expression of pro-inflammatory factors (*e.g.*, MCP-1) were also observed in *Sdc1*<sup>-/-</sup> mice subjected to myocardial infarction by permanent ligation of the left coronary artery (Vanhoutte et al., 2007).

These data suggest that one of the primary functions of syndecan-1 in inflammation is to negatively regulate leukocyte adhesion and migration, possibly by inhibiting the interactions between leukocyte integrins and endothelial ICAM-1 and VCAM-1. Interestingly, *Sdc1*<sup>-/-</sup> neutrophils showed increased adhesion to endothelial cells and ICAM-1 *in vitro*, which were reduced to Wt levels by the addition of low molecular weight heparin (Kharabi Masouleh et al., 2009), suggesting that syndecan-1 on neutrophils negatively regulate neutrophil adhesion to endothelial cells in an HS-dependent manner. However, as evident from the usage of syndecan-1 (CD138) as a marker for plasma cells and tumors of the B cell lineage (Sanderson and Borset, 2002), leukocytes other than plasma cells do not normally express syndecan-1. Induction of syndecan-1 in activated macrophages has been reported (Yeaman and Rapraeger, 1993), but this has not been observed in other leukocytes. Thus, it is likely that syndecan-1 on endothelial cells or syndecan-1 derived from epithelial cells functions as inhibitors of leukocyte adhesion (Fig. 1A). Alternatively, the loss of syndecan-1 may have led to the generation of functionally abnormal leukocytes, but the effects of syndecan-1 on leukocyte maturation and differentiation have not been explored.

Syndecan-1 also regulates the generation and activity of chemokine gradients in inflammatory diseases (Fig. 1B). Most chemokines bind to HSPGs through discrete domains containing a stretch of positively charged amino acids (Lortat-Jacob et al., 2002; Handel et al., 2005). It is generally thought that this interaction is important in tethering the chemokines to endothelial cell surfaces at the site of infection or injury, activating the weakly bound leukocytes and inducing their firm adhesion onto the endothelium, and generating a chemokine gradient that guides the directional migration of leukocytes (Handel et al., 2005; Bao et al., 2010). However, HSPGs can also directly potentiate (Webb et al., 1993; Middleton et al., 1997; Dias-Baruffi et al., 1998) or inhibit chemokine activities (Kuschert et al., 1999), illustrating the complexity of how HSPGs regulate chemokines.

In a mouse model of bleomycin-induced acute lung injury, syndecan-1 shedding was found to facilitate the generation of a CXC chemokine gradient that directs the transepithelial migration of neutrophils (Fig. 1C) (Li et al., 2002). Here, CXCL1 binds to cell surface syndecan-1 and the complex is shed by MMP-7 into the alveolar space, generating a chemotactic gradient across the alveolar epithelium in the basal to apical direction. Hence, in both *Sdc1*<sup>-/-</sup> and *Mmp7*<sup>-/-</sup> mice, there was a significant reduction in CXCL1 levels and neutrophils in bronchoalveolar lavage fluids (BAL), whereas CXCL1 was found complexed to syndecan-1 ectodomains in Wt BAL (Li et al., 2002). This mechanism is thought to confine neutrophilic inflammation to specific sites of tissue injury where it is needed for normal repair, and to limit excessive inflammatory tissue damage.

In a mouse model of allergic lung disease, syndecan-1 ectodomains shed by intranasal allergen challenge bound to CC chemokines (CCL7, CCL11, CCL17) and inhibited the recruitment of Th2 cells to the lung, dampening the Th2 inflammatory response (Xu et al., 2005). The syndecan-1 HS chains mediate this antiinflammatory activity, as HS attenuated parameters of allergic lung disease, but syndecan-1 core proteins lacking HS did not. Consistent with this mechanism, allergen-instilled *Sdc1*<sup>-/-</sup> mice showed overwhelming lung inflammation compared to Wt mice (Xu et al., 2005).

Increased T cell influx was also observed in *Sdc1*<sup>-/-</sup> mice subjected to animal models of Gram-positive toxic shock (Hayashida et al., 2008a) and anti-glomerular basement membrane (GBM) nephritis (Rops et al., 2007). *Sdc1*<sup>-/-</sup> mice injected intraperitoneally with staphylococcal superantigens showed increased serum levels of TNF and IL-6, tissue levels of CXCL9, multi-organ injury, and lethality as compared to Wt mice (Hayashida et al., 2008a). T cell depletion rescued *Sdc1*<sup>-/-</sup> mice from Gram-positive toxic shock, indicating that increased T cell infiltration, likely mediated by CXCL9, is the cellular cause

of enhanced susceptibility. Intraperitoneal injection of neutralizing anti-IFN $\gamma$  antibodies partially, yet significantly reduced disease parameters in *Sdc1*<sup>-/-</sup> mice. Because HS binds to and inhibits IFN (Fritchley et al., 2000; Sarrazin et al., 2005), these data suggest that syndecan-1 suppresses Gram-positive toxic shock by inhibiting the capacity of IFN to induce the expression of pro-inflammatory factors (e.g., TNF $\alpha$ , CXCL9) and subsequent T cell infiltration. The molecular cause of increased T cell influx in the autologous phase of anti-GBM nephritis is not known, but there was increased expression of Th2 cytokines and chemokines in *Sdc1*<sup>-/-</sup> kidneys (Rops et al., 2007), suggesting that syndecan-1 regulates the Th1/Th2 balance in this inflammatory disease.

## 2.2. Syndecan-1 in the resolution of inflammation

Nonresolving inflammation is a major cause of many diseases (Serhan, 2007; Nathan and Ding, 2010). The acute inflammatory response occurs at the expense of the normal functioning of tissues as leukocytes help to heal tissues as well as to destroy them. Thus, it is imperative to rapidly halt the inflammatory response once the harmful agents are removed. In a mouse model of endotoxic shock, syndecan-1 shedding was found to facilitate the resolution of neutrophil inflammation by removing sequestered CXC chemokines in an HS-dependent manner (Hayashida et al., 2009b). Endotoxemic *Sdc1*<sup>-/-</sup> mice showed significantly increased multi-organ injury and dysfunction and lethality compared to endotoxemic Wt mice. Interestingly, the increased morbidity and mortality of endotoxemic *Sdc1*<sup>-/-</sup> mice were due to an exuberant local inflammatory response and not because of an enhanced systemic inflammatory response. High levels of CXCL1 and CXCL2 were sustained in multiple *Sdc1*<sup>-/-</sup> organs at later times after LPS, and this was associated with prolonged neutrophil inflammation and enhanced organ damage. Inhibition of syndecan-1 shedding in endotoxemic Wt mice inhibited the removal of tissue-associated CXCL1 and CXCL2 and impeded the resolution of neutrophilic inflammation in multiple organs, leading to more severe disease. In contrast, administration of HS facilitated the removal of tissue-associated CXCL1 and CXCL2, halted neutrophil accumulation, and attenuated multi-organ injury and lethality in endotoxemic *Sdc1*<sup>-/-</sup> mice.

Although the precise manner in which syndecan-1 shedding facilitates the removal of tissue-associated CXCL1 and CXCL2 is unclear, the newly synthesized CXC chemokines may be tethered to syndecan-1 on the endothelial cell surface, and syndecan-1 shedding may release the sequestered chemokines, resulting in dispersion of the CXC chemokine gradient for neutrophil infiltration (Fig. 1D). Also, because syndecan-1 is expressed in excess of CXCL1 and CXCL2, syndecan-1 shedding may release large amounts of unbound syndecan-1 lectodomains that can displace CXC chemokines tethered to endothelial syndecan-1 and other HSPGs (Fig. 1D). Altogether, these findings suggest that syndecan-1 shedding is an important post-translational mechanism that confines, attenuates, or resolves inflammation by modulating HS-binding pro-inflammatory factors.

## 2.3. Syndecan-1 in matrix remodeling

Remodeling of the ECM is critical in the restoration of the normal structure and function of injured tissues. Fibrogenic factors induce the synthesis of matrix components, and epithelial, mesenchymal, and inflammatory cells assemble the developing ECM using cross-linking enzymes, matrix receptors, and matrix-degrading enzymes. Syndecan-1 has been shown to modulate matrix assembly in several models of inflammatory diseases. *Sdc1*<sup>-/-</sup> mice subjected to myocardial infarction showed abnormal infarct healing, associated with the assembly of a disorganized matrix with a predominance of smaller and fragmented collagen fibers (Vanhoutte et al., 2007). Increased inflammatory cells in *Sdc1*<sup>-/-</sup> mice contributed to increased MMP-2 and MMP-9 activity and degradation of tissue transglutaminase by MMP-2 (Vanhoutte et al., 2007). Consistent with the role of tissue transglutaminase in

collagen fiber assembly (Lorand and Graham, 2003), these deficiencies led to impaired collagen cross-linking and increased cardiac dilatation and failure in *Sdc1*<sup>-/-</sup> mice. Further, overexpression of syndecan-1 reduced inflammation and protected against cardiac dilatation and failure in *Sdc1*<sup>-/-</sup> mice (Vanhoutte et al., 2007), suggesting that syndecan-1 plays a key role in the normal remodeling of injured cardiac tissues.

However, syndecan-1 can also exacerbate cardiac fibrosis, as seen in a mouse model of angiotensin II-induced cardiac fibrosis (Schellings et al., 2010). Syndecan-1 expression was increased predominantly in the fibrotic regions of the left ventricle after angiotensin II treatment. The loss of syndecan-1 protected mice against angiotensin II-induced cardiac dysfunction and fibrosis by blunting the effects of angiotensin II and TGF $\beta$ 1 on cardiac expression of connective tissue growth factor (CTGF) and collagens I and III. HS chains of syndecan-1 are important as protamine decreased CTGF and collagen I levels, while recombinant syndecan-1 without HS did not affect CTGF levels (Schellings et al., 2010). Further, HS binds to angiotensin II (Koppel et al., 2003) and the loss of syndecan-1 mitigated cardiac dysfunction and fibrosis in a Smad2-dependent manner (Schellings et al., 2010). These findings suggest that syndecan-1 may promote cardiac fibrosis by amplifying angiotensin II and TGF $\beta$ 1 signaling in an HS-dependent manner.

Once solubilized, syndecan-1 ectodomains can exhibit functions similar to or different from cell surface syndecan-1. In idiopathic pulmonary fibrosis (IPF), syndecan-1 expression is increased in fibrotic lungs and syndecan-1 ectodomains are increased in BAL and lung homogenates from IPF patients (Kliment et al., 2009). Syndecan-1 ectodomains are also increased in BAL and lung homogenates in extracellular superoxide dismutase (EC-SOD) null mice subjected to asbestos- or bleomycin-induced lung fibrosis (Kliment et al., 2009), consistent with the protective role of EC-SOD in oxidative stress-induced syndecan-1 shedding (Kliment and Oury, 2011). Interestingly, cell surface syndecan-1 was found to promote alveolar re-epithelialization, whereas syndecan-1 ectodomains shed by oxidative stress increased neutrophil chemotaxis and inhibited re-epithelialization (Kliment et al., 2009). Syndecan-1 ectodomains also increased fibroblast proliferation and the release of TGF $\beta$ 1 (Kliment et al., 2009). Thus, cell surface syndecan-1 apparently attenuates fibrosis by promoting repair, whereas syndecan-1 ectodomain promotes fibrosis by inhibiting epithelial repair and up-regulating fibrotic factors and processes. The underlying mechanism of how this is accomplished is not known. Because reactive oxygen species (ROS) can bind to and modify HS (Rees et al., 2005), syndecan-1 ectodomains shed by oxidative stress may exhibit different HS structures and function from those of membrane-bound syndecan-1. Further, because EC-SOD directly binds to HS and heparin (Sandstrom et al., 1992; Kliment et al., 2008), this anti-oxidant may not only protect syndecan-1 from shedding but also from modification by ROS. Future studies directed at defining the structure: function relationship of oxidized syndecan-1 HS should address these potentially important mechanisms in fibrotic diseases.

#### 2.4. Other functions of syndecan-1 in inflammatory diseases

Syndecan-1 has also been shown to regulate protein leakage in the intestinal epithelium (Bode et al., 2008), smooth muscle cell (SMC) proliferation in vascular injury (Fukai et al., 2009), keratinocyte proliferation in dermal injury (Stepp et al., 2002), and lipoprotein metabolism (Stanford et al., 2009). Protein-losing enteropathy (PLE) is characterized by excessive efflux of plasma proteins into the intestinal lumen, and is often associated with viral infections, increased IFN $\gamma$  and TNF $\alpha$  levels, and a pro-inflammatory state (Lenz et al., 2003). Loss of syndecan-1 or HS from the basolateral surface of intestinal cells was found to be closely associated with PLE episodes, whereas both were induced when the disease resolved (Bode et al., 2008). More importantly, *Sdc1*<sup>-/-</sup> and *Sdc1*<sup>-/-</sup> intestinal epithelium showed increased basal protein leakage and were susceptible to protein loss upon IFN $\gamma$  and

TNF $\alpha$  challenge and increased venous pressure (Bode et al., 2008). Because HS/heparin can bind to and inhibit IFN $\gamma$  (Fritchley et al., 2000; Sarrazin et al., 2005) and TNF $\alpha$  (Lantz et al., 1991), syndecan-1 may potentially attenuate protein leakage by blocking the activities of these pro-inflammatory cytokines in an HS-dependent manner. Addition of 2-*O*- and 3-*O*-desulfated heparin rescued the pathological phenotypes (Bode et al., 2008), suggesting that these sulfate modifications in syndecan-1 HS are not required. However, precisely how loss of syndecan-1 or HS leads to increased intestinal epithelial leakage is unclear as epithelial tight junctions were determined to be intact at the EM level (Bode et al., 2008). Perhaps syndecan-1 is required for the maintenance of normal epithelial morphology and of other epithelial junctions, such as adherens junctions, as syndecan-1 expression has been linked to E-cadherin expression and loss of syndecan-1 induces an epithelial to mesenchymal transition (EMT)-like process in mammary gland epithelial cells (Kato et al., 1995). Alternatively, because IFN $\gamma$  and TNF $\alpha$  synergize to induce syndecan-1 shedding (Henry-Stanley et al., 2006) and because HS has been proposed as an impermeable molecule that seals epithelial gaps (Watson et al., 2005), syndecan-1 ectodomains shed by the pro-inflammatory cytokines may serve this function in preventing PLE.

Arterial injury induces SMC proliferation and migration, and intimal accumulation of cells and ECM. Syndecan-1 was found to regulate these processes in a mouse model of ligation-induced carotid artery injury (Fukai et al., 2009). *Sdc1*<sup>-/-</sup> mice developed a larger neointimal lesion after injury compared to Wt mice that developed little to no lesion. Injured *Sdc1*<sup>-/-</sup> arteries also showed a significant increase in SMCs in both the media and intima, suggesting that the primary function of syndecan-1 in arterial injury is to inhibit abnormal SMC proliferation. Interestingly, cultured *Sdc1*<sup>-/-</sup> SMCs showed increased proliferation in response to PDGF-BB, thrombin, FGF-2, EGF and serum, and this was accompanied by increased mRNA levels of PDGF-B chain. DNA synthesis in *Sdc1*<sup>-/-</sup> SMCs treated with thrombin, PDGF-BB, or serum was inhibited by the down-regulation of PDGF-BB or PDGFR $\beta$ . Because HS can inhibit arterial SMC proliferation and migration (Bingley et al., 1998), these findings suggest that HS chains of syndecan-1 regulate SMC proliferation in arterial injury by inhibiting PDGF signaling and possibly signaling by other HS-binding growth factors, such as FGF-2 and EGF. In contrast, keratinocyte proliferation was reduced in *Sdc1*<sup>-/-</sup> mice in a model of dermabrasion (Stepp et al., 2002). This was accompanied by reduced localization of  $\alpha$ 9 integrin in the injured *Sdc1*<sup>-/-</sup> epidermis. A subsequent study showed that TGF $\beta$ 1 increases the cell surface expression of  $\alpha$ 2 $\beta$ 1,  $\alpha$ v $\beta$ 6,  $\alpha$ v $\beta$ 8 and  $\alpha$ 6 $\beta$ 4 integrins in cultured Wt keratinocytes, but not in *Sdc1*<sup>-/-</sup> keratinocytes (Stepp et al., 2007). These data suggest that syndecan-1 modulates keratinocyte proliferation by regulating integrin expression and the cross-talk between integrins and growth factor signaling.

Elevated plasma triglyceride is a key risk factor for atherosclerosis. Syndecan-1 was identified as the primary HSPG receptor that mediates hepatic clearance of triglyceride-rich lipoproteins (TRLs) in mice (Stanford et al., 2009). *Sdc1*<sup>-/-</sup> mice showed increased plasma levels of TRLs after fasting and delayed clearance of TRLs during consumption of normal dietary fat. This was specific to the loss of syndecan-1, as *Sdc3*<sup>-/-</sup> and *Sdc4*<sup>-/-</sup> mice did not show these phenotypes and adenovirus-mediated syndecan-1 overexpression rescued *Sdc1*<sup>-/-</sup> mice from the pathological phenotypes (Stanford et al., 2009). The abundant expression of syndecan-1 on the microvilli of hepatocyte basal membranes suggests that syndecan-1 is well positioned to capture circulating TRLs. Binding of TRLs to the HS chains of syndecan-1 could occur through electrostatic interactions between negatively charged sulfate and carboxyl groups with positively charged groups in apolipoproteins or lipases (Stanford et al., 2009), and TRLs may be internalized when syndecan-1 undergoes endocytosis (Fuki et al., 1997).

### 3. Cancer

Syndecan-1 expression is dysregulated in many cancers, including carcinomas of the prostate (Kiviniemi et al., 2004), breast (Lendorf et al., 2011), pancreas (Juuti et al., 2005), ovary (Kusumoto et al., 2010), and colon (Hashimoto et al., 2008). Low syndecan-1 expression in tissue is associated with a worse prognosis in head and neck (Anttonen et al., 1999), lung (Anttonen et al., 2001) and colorectal (Lundin et al., 2005) cancers, whereas high levels of shed syndecan-1 in serum correlate with a poor prognosis in lung cancer (Joensuu et al., 2002) and myeloma (Seidel et al., 2000b). These findings suggest that syndecan-1, like many other proteoglycans, has important functions in tumor biology (Iozzo and Sanderson, 2011). Indeed, accumulating evidence suggests that syndecan-1 modulates several key processes of tumorigenesis, such as cancer cell proliferation and apoptosis, angiogenesis, and metastasis (Fig. 2).

#### 3.1. Syndecan-1 in cancer cell proliferation and apoptosis

Syndecan-1 can affect tumorigenesis by regulating mediators of tumor cell survival and proliferation (e.g., oncogenes, growth factors). For example, *Sdc1*<sup>-/-</sup> mice were protected against Wnt-1 induced mammary tumorigenesis (Alexander et al., 2000). Further, syndecan-1 ectodomains, but not syndecan-1 core protein devoid of HS, potentiated the activity of Wingless, a *Drosophila* Wnt-1 homologue, in transgenic *Drosophila* S2 cells expressing the Wnt receptor DFz2 in a dose-dependent manner (Alexander et al., 2000). These findings suggest that syndecan-1 functions as a coreceptor for Wnt signaling in an HS-dependent manner. A subsequent study suggested that syndecan-1 affects the capacity of Wnt to induce the accumulation of mammary progenitor cells (Liu et al., 2004), but it is not known if syndecan-1 similarly regulates Wnt signaling in other target tissues of Wnt.

Hepatocyte growth factor (HGF) binds to the HS chains of syndecan-1 on myeloma cells (Seidel et al., 2000a; Derksen et al., 2002). The signaling receptor for HGF is Met, and Met signaling has been implicated in the growth, survival, and spread of various cancers (Birchmeier et al., 2003; Christensen et al., 2005). Importantly, the interaction of HGF with syndecan-1 potentiated Met signaling, which involves the activation of PI3 kinase-protein kinase B and Ras-MAP kinase pathways (Derksen et al., 2002). Syndecan-1 shedding adds another dimension to the regulation of HGF-Met signaling in myeloma biology (Fig. 2A). Heparanase induced syndecan-1 shedding in myeloma cells through the upregulation of MMP-9 and urokinase-type plasminogen activator (uPA), both of which can act as a syndecan-1 sheddase (Purushothaman et al., 2008). Heparanase also induced HGF expression in myeloma cells, and newly synthesized HGF associated with syndecan-1 ectodomains and the HGF-syndecan-1 ecto-domain complex enhanced Met signaling in osteoblasts and stromal cells (Ramani et al., 2011). This paracrine Met signaling activated IL-11 and upregulated the receptor activator of NF- $\kappa$ B ligand (RANKL) in osteoblast-like Saos2 cells. IL-11 inhibits bone formation (Hughes and Howells, 1993) and supports bone resorption (Shaughnessy et al., 2002). Further, heparin enhances IL-11 signaling through activation of ERK MAP kinase (Rajgopal et al., 2006), suggesting that syndecan-1 functions in an HS-dependent manner. Together, these data suggest that syndecan-1 synergizes with heparanase to promote myeloma bone disease by activating the HGF-Met-IL-11-RANKL signaling axis.

The tumor microenvironment provides a conducive *milieu* for the growth and progression of tumor cells. Syndecan-1 expression is elevated in the reactive stroma of breast carcinoma tissue (Stanley et al., 1999). Invasive breast cancer cells induced the expression of syndecan-1 in mouse embryonic fibroblasts (MEFs) *in vitro*, but this was not observed with normal or less invasive cell lines. Moreover, syndecan-1 expressing MEFs enhanced the growth of breast cancer cell lines in co-culture and promoted breast carcinoma progression



*in vivo* (Maeda et al., 2004), both in an HS-dependent manner. Syndecan-1 expression in reactive stromal fibroblasts could store and present heparin-binding growth factors such as FGFs, HGF, and EGFs to cancer cells. Indeed, shedding of syndecan-1 ectodomains from stromal fibroblasts stimulated tumor cell growth through the activation of FGF2 and SDF1 signaling (Su et al., 2007). A subsequent study identified membrane type 1 matrix metalloproteinase (MT1-MMP) as the stromal syndecan-1 sheddase in breast carcinoma (Su et al., 2008). These studies reveal an interesting mechanism where tumor cells utilize syndecan-1 shedding to derive growth factors from the neighboring stroma to maintain its growth conducive environment.

Syndecan-1 was also found to regulate tumor cell apoptosis. Syndecan-1 knock-down in myeloma cells induced growth arrest and apoptosis (Khotskaya et al., 2009). This could be due to diminished pro-survival signals as a result of reduced levels of cell surface HSPG coreceptors for growth factor signaling. However, several studies suggest that syndecan-1 can also directly promote tumor cell apoptosis. For example, recombinant human syndecan-1 ectodomains induced apoptosis in MCF-7 breast cancer cells (Sun et al., 2008). Syndecan-1 expression was determined to be upregulated through n-3 polyunsaturated fatty acids (n-3 PUFA)-activated peroxisome proliferator-activated receptor  $\gamma$  (PPAR $\gamma$ ) signaling, which promoted apoptosis in breast cancer cells. Docosahexaenoic acid (DHA, an n-3 PUFA)-mediated upregulation of syndecan-1 expression and direct addition of syndecan-1 ectodomains also led to apoptosis of cultured human prostate cancer cells (Hu et al., 2010).

It is not clear how both cell surface and shed syndecan-1 can stimulate growth and survival in myeloma cells and yet also elicit apoptosis in other cancer cells. Perhaps this apparent paradox could be explained by tumor cell-specific requirements for growth factor signaling. In addition, the bell-shaped activity curve of HS in regulating HS-binding ligands may explain the opposing functions of syndecan-1 in tumor cell apoptosis. HS generally serves as a scaffold that brings both ligand and its binding partner to proximity. In other words, HS potentiates the interaction by increasing the local concentration of the ligand and the bell-shaped curve may be concave or convex depending on whether HS-binding is inhibitory or stimulatory for the ligand. Thus, at either ends of the optimal HS concentration, HS activity is low or absent and will not potentiate growth factor signaling. Further, the tumor microenvironment is likely unique to the distinct tumors, and HS chains of syndecan-1 may be differentially modified to either promote or attenuate tumor cell growth. Indeed, both growth promoting and growth inhibiting sequences are present in HS (Liu et al., 2002a). Whether one or several of these mechanisms control the growth-modulating activity of syndecan-1 in cancer remains to be explored.

### 3.2. Syndecan-1 in cancer cell invasion and metastasis

Metastasis of cancer cells is a multi-step process that involves cell adhesion and migration and ECM degradation (Hanahan and Weinberg, 2011). Cell surface syndecan-1 is thought to promote cell adhesion to the ECM and retard cancer cell migration. For example, head and neck squamous cell carcinoma (HNSCC) cells expressing high levels of syndecan-1 migrated poorly and were less invasive in collagen I matrices as compared to HNSCC cells expressing lower levels of syndecan-1 (Ishikawa and Kramer, 2010). Loss of syndecan-1 diminished cell adhesion to collagen substrates but enhanced cell motility and invasion through 3-D collagen gels. Inhibition of the  $\alpha 2\beta 1$  integrin by anti- $\alpha 2$  monoclonal antibodies blocked attachment of syndecan-1 silenced HNSCC cells to collagen I (Ishikawa and Kramer, 2010), suggesting that syndecan-1 is a coreceptor for  $\alpha 2\beta 1$  binding to collagen I.

Integrins are key regulators of cell adhesion, proliferation, and migration (Hynes et al., 2002; Desgrosellier and Cheresh, 2010; Rathinam and Alahari, 2010). There is ample

evidence suggesting that syndecan-1 cooperates with integrins in mediating these cellular processes (Fig. 2B) (Iba et al., 2000; Morgan et al., 2007; Ogawa et al., 2007; Gama-de-Souza et al., 2008; Vuoriluoto et al., 2008; Beauvais et al., 2009; Ishikawa and Kramer, 2010). Syndecan-1 can potentiate the activation and signaling of  $\alpha v \beta 3$  and  $\alpha v \beta 5$  integrins through the engagement of  $\alpha v$  ligands such as vitronectin (Beauvais et al., 2009). Integrins and syndecan-1 can also form a dynamic linkage between the ECM and cytoskeleton, allowing for cell migration that involves formation of cell adhesion at the leading edge and disruption of cell adhesion at the rear edge, together with active polymerization and depolymerization of the actin cytoskeleton (Ridley, 2011). Syndecan-1 gene silencing reduced  $\alpha 2 \beta 1$  integrin-dependent activation of FAK that led to reduced HNSCC cell adhesion to collagen I (Ishikawa and Kramer, 2010). In addition, gene silencing of syndecan-1 in HNSCC cells reduced levels of active RhoA but enhanced Rac1 and triggered filopodia and lamellopodia formation at the cell's leading edge (Ishikawa and Kramer, 2010). These data suggest that integrin and syndecan-1 function cooperatively to stabilize focal adhesions and actin cytoskeleton, resulting in restricted cell locomotion. These findings also suggest that the loss of cell surface syndecan-1, seen in many carcinomas, favors acquisition of the meta-static phenotype by cancer cells.

The spatiotemporal expression pattern of syndecan-1 also allows it to differentially modulate tumor cell migration. Cell surface syndecan-1 inhibited, whereas syndecan-1 ectodomains promoted the invasion of MCF-7 breast cancer cells into Matrigel (Nikolova et al., 2009). Similarly, cell surface syndecan-1 inhibited migration of HT1080 fibrosarcoma cells on collagen while syndecan-1 ectodomains stimulated HT1080 cell migration (Endo et al., 2003). Treatment with MMP inhibitors increased syndecan-1 expression on the surface of HT1080 cells, concurrent with the formation of actin stress fibers that led to inhibition of cell migration (Endo et al., 2003). Similar to carcinoma cells, the loss of cell surface syndecan-1 likely leads to reduced cell adhesion and spreading, thereby allowing the sarcoma cells to acquire a more motile and invasive phenotype. These findings suggest that membrane-bound syndecan-1 and soluble syndecan-1 may play unique roles at different stages of cancer progression, and provide insights into targeting cell surface vs. shed syndecan-1 in cancer therapy.

Epithelial–stromal interactions are crucial in the acceleration of carcinoma growth and progression. During neoplastic transformation, stromal fibroblasts and the ECM are altered to allow a more supportive microenvironment for cancer progression. Syndecan-1 is aberrantly expressed by fibroblasts in the surrounding stroma of infiltrating breast carcinoma (Stanley et al., 1999; Maeda et al., 2004). In addition to supporting breast carcinoma growth and angiogenesis, increased stromal syndecan-1 expression altered fibronectin production and ECM organization (Yang et al., 2011). The ECM produced by syndecan-1-negative fibroblasts consisted of an intersecting meshwork, which was in stark contrast to the parallel pattern of fibronectin and collagen I fibers produced by syndecan-1-positive fibroblasts (Yang et al., 2011). Importantly, the structurally abnormal ECM produced by the syndecan-1-expressing fibroblasts promoted breast carcinoma cell motility. How stromal syndecan-1 modulates ECM assembly is not fully understood, but syndecan-1 may modulate fibronectin matrix assembly directly or together with integrins (Stepp et al., 2010), as was observed for syndecan-2 (Klass et al., 2000) and syndecan-4 (Midwood et al., 2004).

### 3.3. Syndecan-1 in angiogenesis

Tumor angiogenesis generates new vascular beds that provide nutrients and oxygen for the highly metabolic tumor mass. Syndecan-1 can bind to pro-angiogenic factors like FGF-2 and VEGF, and subsequently present these factors to their respective receptors on endothelial cells to initiate endothelial invasion and budding. The broader functional implications of syndecan-1 in angiogenesis may also be to allow soluble syndecan-1 ectodomains with

bound pro-angiogenic factors to foster angiogenesis at premetastatic niches. For example, in myeloma, shedding of syndecan-1 ectodomains by heparanase facilitated endothelial invasion and subsequent angiogenesis (Fig. 2C) (Purushothaman et al., 2010). Heparanase also upregulated HGF and VEGF in myeloma cells, and syndecan-1 ectodomains bound to VEGF and presented VEGF to endothelial cells (Purushothaman et al., 2008, 2010). Binding of syndecan-1 ectodomains to  $\alpha v \beta 3$  and  $\alpha v \beta 5$  integrins is apparently necessary for its pro-angiogenic function, as a short inhibitory peptide that mimics the syndecan-1 ectodomain core protein (synstatin) abrogated syndecan-1 interactions with both integrins and inhibited endothelial cell invasion as well as tumor growth *in vivo* (Beauvais et al., 2009; Purushothaman et al., 2010).

HS fragments generated by heparanase can stimulate melanoma angiogenesis *in vivo* (Roy and Marchetti, 2009). However, there is a clear need for an intact syndecan-1 ectodomain in heparanase-induced angiogenesis in myelomas. Rat aortas were grown in Matrigel in the presence of medium conditioned either by heparanase-low or heparanase-high myeloma cells. Addition of syndecan-1 ectodomains and VEGF to medium conditioned by low heparanase-expressing myeloma cells (low syndecan-1 ectodomain levels) enhanced endothelial sprouting. Conversely, immunodepletion of syndecan-1 from the conditioned medium of high heparanase-expressing myeloma cells (high syndecan-1 ectodomain) reduced endothelial sprouting (Purushothaman et al., 2010), indicating that an intact syndecan-1 ectodomain promotes angiogenesis.

Increased syndecan-1 expression in stromal fibroblasts is observed in several carcinomas, such as those of the breast (Stanley et al., 1999; Maeda et al., 2004), stomach (Wiksten et al., 2001), and thyroid (Barbareschi et al., 2003). In a xenograft model of human breast carcinoma cells and syndecan-1-transfected fibroblasts implantation into mice, stromal syndecan-1 expression was associated with significantly elevated microvessel density and larger vessel area (Maeda et al., 2006). Expression of syndecan-1 in stromal fibroblasts of human breast carcinomas also correlated significantly with high microvessel density and larger vessel area (Maeda et al., 2006). These findings raise the possibility that syndecan-1 in the reactive stroma may sequester pro-angiogenic factors and increase the local concentration of these factors to promote angiogenesis.

## 4. Infectious diseases

Syndecan-1, as the major cell surface HSPG of epithelial cells, is thought to be targeted by microbial pathogens especially during the early phase of infection. Several studies suggest that syndecan-1 is subverted in several steps of infection, including the initial attachment and subsequent entry of pathogens into host cells, and inhibition of host defense mechanisms (Fig. 3). Evidence that these syndecan-1 functions are indeed important in the pathogenesis of infectious diseases is provided by studies using animal and cell culture-based models of infection, where it was observed that loss or reduction of syndecan-1 enables mice or cells to significantly resist infection by several viral and bacterial pathogens (Table 1). The underlying molecular mechanisms have yet to be precisely defined, but several studies have revealed interesting details of how both cell surface and shed syndecan-1 can promote pathogenesis through distinct molecular mechanisms.

### 4.1. Cell surface syndecan-1 in pathogen attachment and entry

A wide variety of pathogens, including viruses, bacteria and parasites, use cell surface HSPGs as attachment receptors (Rostand and Esko, 1997; Spillmann, 2001; Rusnati and Urbinati, 2009; Bartlett and Park, 2010). For example, hepatitis E virus (HEV) appears to target syndecan-1 for its attachment to host cells (Kalia et al., 2009). HEV is a positive-sense, non-enveloped single-stranded RNA virus that causes hepatitis E, a recently described

inflammatory liver disease of humans. HEV uses its major capsid protein pORF2 to bind to hepatoma cells, and this is significantly inhibited by heparinase III treatment and syndecan-1 knock-down (Kalia et al., 2009). Syndecan-4 knockdown or treatment with phosphatidylinositol-specific phospholipase C (PI-PLC) that releases GPI-anchored glypicans from the cell surface did not inhibit pORF2 binding to hepatoma cells (Kalia et al., 2009). These results suggest that HEV binds specifically to the HS chains of syndecan-1 for its attachment to host cells.

Further evidence supporting the role of syndecan-1 as a viral attachment receptor comes from a study examining its role in a cell culture-based model of human papillomavirus (HPV) infection. HPV is a non-enveloped, double-stranded DNA virus that infects skin and mucosal epithelial cells, and hijacks the host cell cycle to cause both benign and malignant epithelial tumors. Induced expression of syndecan-1 in erythroleukemia cells significantly increased HPV attachment and internalization compared to the induced expression of syndecan-4 or glypican-1. (Shafti-Keramat et al., 2003). Further, HPV attachment and internalization were inhibited by heparinase III treatment of host cells and addition of antibodies against the HPV capsid protein L1 (Shafti-Keramat et al., 2003). L1 binding to cell surface HSPGs induces a conformational change in the minor capsid protein L2, exposing residues in L2 that bind to a secondary entry receptor, which mediates HPV internalization (Richards et al., 2006; Day et al., 2008). These observations suggest a mechanism where HPV attaches to host cells via the L1-syndecan-1 interaction, which leads to the activation of L2 and L2-mediated HPV entry.

Other viral pathogens that target epithelial cells have also been shown to bind to cell surface HSPGs for their initial attachment to host cells (Spillmann, 2001; Bartlett and Park, 2010), suggesting that syndecan-1 functions prominently in many viral infections. The molecular details of syndecan-1-virus interactions have yet to be clearly defined, but because viruses bind to the HS moiety of HSPGs, structural features of syndecan-1 HS may be important. For example, initial studies revealed that the alpha herpesviruses Herpes simplex virus (HSV)-1 and HSV-2 bind to HSPGs using glycoproteins gB and gC and that these interactions mediate viral attachment but not entry (WuDunn and Spear, 1989; Laquerre et al., 1998). Gene silencing studies showed that both syndecan-1 and -2 are important in HSV-1 attachment (Bacsa et al., 2011). However, a rare form of HS containing 3-*O*-sulfated glucosamine residues was found to induce gD-mediated HSV-1 internalization (Shukla et al., 1999; Liu et al., 2002b). Soluble 3-*O*-sulfated HS was also found to trigger HSV-1 entry into CHO cells (Tiwari et al., 2007), suggesting that syndecan-1 may directly mediate HSV internalization if its HS chains are modified to contain sufficient 3-*O*-sulfated HS domains to potentiate gD signaling.

Human immunodeficiency virus (HIV), the etiologic agent of AIDS, binds to HSPGs on macrophages, dendritic cells, spermatozoa, endothelial cells, and epithelial cells primarily through gp120 (Saphire et al., 2001; Argyris et al., 2003; Bobardt et al., 2003; de Witte et al., 2007; Ceballos et al., 2009; Smith et al., 2010). HIV binds to K562 human leukemic cells transfected with syndecan-1, but not to non-transfected K562 cells, which are deficient in syndecans (Saphire et al., 2001), indicating that syndecan-1 can mediate the attachment of HIV to host cells. Further, syndecan-1 expression was increased in the gut of HIV-1-infected individuals and this was associated with increased viral translocation across the gut epithelium (Smith et al., 2010). However, several syndecans can apparently mediate HIV attachment to host cells. HIV binding to Burkitt lymphoma-derived Namalwa B cells was significantly increased when these cells were transfected with syndecan-1, -2, -3, or -4 (Bobardt et al., 2003), and syndecan-3 was shown to be the major HIV-1 attachment receptor on dendritic cells (de Witte et al., 2007). These findings suggest that syndecans expressed on target host cells of HIV may contain unique HS structures that HIV

preferentially binds to or that syndecans are expressed in cellular compartments where HIV preferentially infects. Alternatively, syndecans may be physically and/or functionally linked (e.g., via syndecan cytoplasmic domain signaling) to HIV entry receptors, such as CD4, DC-SIGN, and mannose receptors. These mechanisms have yet to be explored.

The role of the syndecan cytoplasmic domain in microbial pathogenesis was directly examined in a cell culture model of *Neisseria gonorrhoeae* infection (Freissler et al., 2000). Along with *Chlamydia trachomatis*, *N. gonorrhoeae* is one of the most common causative agents of bacterial sexually transmitted diseases in the US. *N. gonorrhoeae* binds to syndecan-1 and -4 through its outer membrane protein Opa<sub>HSPG</sub> as overexpression of syndecan-1 or -4 in HeLa cells increased *N. gonorrhoeae* infection (Freissler et al., 2000). Interestingly, *N. gonorrhoeae* attached but did not invade HeLa cells expressing syndecan-1 or -4 lacking the cytoplasmic domain. Furthermore, a syndecan-4 mutant lacking the syndecan-4-specific dimerization motif in the cytoplasmic domain that binds to PKC and phosphatidylinositol 4,5-bisphosphate, and a syndecan-4 mutant lacking the C-terminal Glu-Phe-Tyr-Ala PDZ binding domain did not support *Neisseria* invasion (Freissler et al., 2000). Mutagenesis studies with the syndecan-1 cytoplasmic domain was not performed, but syndecan-1 also contains the Glu-Phe-Tyr-Ala PDZ binding motif, suggesting that PDZ proteins that bind to the cytoplasmic domain of syndecans, such as syntenin (Grootjans et al., 1997), may be important in facilitating gonococcal invasion. In addition, *Neisseria* binding to cell surface HSPGs induces a signaling pathway that involves phosphatidylcholine-specific phospholipase C, diacylglycerol, acidic sphingomyelinase, and ceramide (Grassmé et al., 1997). Whether the cytoplasmic domain of syndecans also interacts with these signaling molecules remains to be determined. Regardless, these observations indicate that syndecan-1 can serve as an entry receptor for *N. gonorrhoeae* invasion (Fig. 3B).

#### 4.2. Syndecan-1 shedding in microbial pathogenesis and host defense

*Pseudomonas aeruginosa* (Park et al., 2000a), *Staphylococcus aureus* (Park et al., 2004), *Streptococcus pneumoniae* (Chen et al., 2007), and *Bacillus anthracis* (Popova et al., 2006) have been shown to secrete factors that induce syndecan-1 shedding. Pathogen-induced ectodomain shedding is not restricted to syndecan-1, as several pathogens induce the ectodomain shedding of a variety of inflammatory mediators from the host cell surface (Vollmer et al., 1996; Walev et al., 1996, 2000; Schmidtchen et al., 2001; Lemjabbar and Basbaum, 2002), presumably to modulate the host environment to favor pathogenesis. *P. aeruginosa* is a Gram-negative bacterium associated with infections of the skin, urinary tract, and lung. *P. aeruginosa* induces syndecan-1 shedding through LasA (Park et al., 2000a), which is a known virulence factor of *P. aeruginosa* in animal models of lung infection (Woods et al., 1982; Blackwood et al., 1983). LasA activates a PTK-regulated, metalloproteinase-mediated shedding mechanism of host cells to enhance syndecan-1 shedding (Park et al., 2000a). *P. aeruginosa*-induced syndecan-1 shedding is a virulence mechanism as newborn *Sdc1*<sup>-/-</sup> mice significantly resisted intranasal lung infection with *P. aeruginosa* compared to newborn Wt mice (Park et al., 2001). Intranasal administration of syndecan-1 ectodomain or HS enhanced bacterial virulence in *Sdc1*<sup>-/-</sup> mice, whereas inhibition of syndecan-1 shedding attenuated virulence in Wt mice. Further, *P. aeruginosa* did not bind to syndecan-1 and excess heparin had no effect on the adhesion of *P. aeruginosa* to lung epithelial cells in vitro. Syndecan-1 ectodomains were also detected in BAL after intranasal infection with *P. aeruginosa* or instillation of LasA, and syndecan-1 ectodomains inhibited the antibacterial activity of several cationic antimicrobial peptides in vitro (Park et al., 2001). Adult *Sdc1*<sup>-/-</sup> mice also showed significantly attenuated *P. aeruginosa* sepsis following thermal injury (Haynes et al., 2005). Together, these findings suggest that the

subversion of syndecan-1 shedding to inhibit host defense is an important virulence mechanism of *P. aeruginosa*.

*S. aureus* is an important Gram-positive bacterial pathogen of humans that causes a wide variety of diseases including keratitis, impetigo, pneumonia, osteomyelitis, endocarditis, toxic shock syndrome, and sepsis. *S. aureus* induces syndecan-1 shedding through its cytotoxic virulence factors  $\alpha$ -toxin and  $\beta$ -toxin (Park et al., 2004). Similar to *P. aeruginosa* LasA, both  $\alpha$ -toxin and  $\beta$ -toxin induce syndecan-1 shedding by activating the PTK-dependent, metalloproteinase-mediated shedding mechanism of host cells (Park et al., 2004). The physiological significance of *S. aureus*-induced syndecan-1 shedding was examined in a mouse model of scarified corneal infection (Hayashida et al., 2011). Scarified *Sdc1*<sup>-/-</sup> corneas significantly resisted *S. aureus* infection compared to Wt corneas that express abundant syndecan-1 in their epithelium. *S. aureus* did not bind to syndecan-1 and did not require syndecan-1 to colonize cultured corneal epithelial cells, but it induced syndecan-1 shedding in corneal epithelial cells both *in vitro* and *in vivo*. Topical administration of purified syndecan-1 ectodomains or HS significantly increased, whereas inhibition of syndecan-1 shedding significantly decreased the bacterial burden in corneal tissues. Further, depletion of neutrophils in the resistant *Sdc1*<sup>-/-</sup> mice increased the corneal bacterial burden to that of the susceptible Wt mice, suggesting that syndecan-1 moderates neutrophils to promote infection. Interestingly, neutrophils infiltrated similarly into infected Wt and *Sdc1*<sup>-/-</sup> corneas. However, syndecan-1 ectodomains and HS significantly inhibited the killing of *S. aureus* by isolated neutrophils. These observations suggest that *S. aureus* exploits the capacity of syndecan-1 ectodomains to inhibit neutrophil-mediated bacterial killing mechanisms in an HS-dependent manner to promote its pathogenesis in the cornea. Whether  $\alpha$ -toxin and/or  $\beta$ -toxin induce syndecan-1 shedding during *S. aureus* corneal infection remains to be determined, but both toxins are established virulence factors in animal models of *S. aureus* keratitis (O'Callaghan et al., 1997; Girgis et al., 2005), suggesting that syndecan-1 shedding may mediate the capacity of these exotoxins to enhance *S. aureus* virulence in the cornea.

However, because many pathogens bind to cell surface HSPGs for their attachment and entry, it is not clear why syndecan-1 shedding is not a host defense mechanism that rapidly down-regulates microbial attachment sites. In fact, *P. aeruginosa* binds to HSPGs on the basolateral surface of polarized epithelial cells (Bucior et al., 2010), which is where syndecan-1 is expressed, and *S. aureus* shows increased adherence to ARH-77 cells overexpressing syndecan-1 (Henry-Stanley et al., 2005). Perhaps only highly effective pathogens that possess redundant mechanisms for attachment, such as *S. aureus*, can use syndecan-1 shedding to promote their pathogenesis. Alternatively, some pathogens may exploit both cell surface and shed syndecan-1 to attach to host cells and to inhibit host defense mechanisms. Whether other bacterial pathogens that possess the capacity to induce syndecan-1 shedding, such as *S. pneumoniae* and *B. anthracis*, subvert this host mechanism to promote their pathogenesis remains to be determined.

## 5. Concluding remarks

Studies using animal models of diseases have provided clear indications that syndecan-1 plays an important role in the development of inflammatory diseases, cancer, and infection. Syndecan-1 is a critical cofactor in the pathogenesis of these diseases as *Sdc1*<sup>-/-</sup> mice only show dramatic pathological phenotypes when challenged with disease-causing agents or conditions, and do not show spontaneous pathologies. In general, syndecan-1 attenuates non-infectious inflammatory diseases by inhibiting leukocyte adhesion onto the activated endothelium, reducing the expression and inhibiting the activity of pro-inflammatory factors, confining leukocyte infiltration to specific sites of tissue injury, or by removing

sequestered chemokines and facilitating the resolution of inflammation. In contrast, in cancer and infectious diseases, syndecan-1 functions are mostly pro-pathogenic. Syndecan-1 enhances oncogene and growth factor signaling, inhibits cancer cell apoptosis, and promotes angiogenesis. However, syndecan-1 can also attenuate these processes depending on the type of tumor, site of action, or the mechanisms that it regulates. In infectious diseases, syndecan-1 clearly promotes pathogenesis as it mediates the attachment and entry of pathogens into host cells and inhibits host defense mechanisms. Moreover, loss of syndecan-1 is a gain of function mutation that increases the resistance of mice to several bacterial infections. These data suggest that one of the crucial functions of mammalian syndecan-1 *in vivo* is to assure the adequate and correct functioning of inflammation. However, this beneficial function of syndecan-1 comes at a price as certain cancer cells and microbial pathogens have either adapted or evolved to take advantage of syndecan-1 for their pathogenesis. The underlying mechanisms that govern the physiological vs. pathological functions of syndecan-1 are not known, but the unique spatiotemporal expression pattern of syndecan-1 likely mediates its prominent functions in many diseases. Future studies aimed at defining the molecular details of how syndecan-1 expression and shedding are turned on or off, and how syndecan-1 is targeted to specific cellular and tissue compartments during pathogenesis are anticipated to reveal new therapeutic targets for inflammatory diseases, cancer, and infection.

## Acknowledgments

We thank our colleagues in the Park laboratory for helpful discussions. We apologize for not citing studies that may have been relevant but was omitted due to space limitations. Our research described in this review is supported by NIH Grants HL094613 and HL107472.

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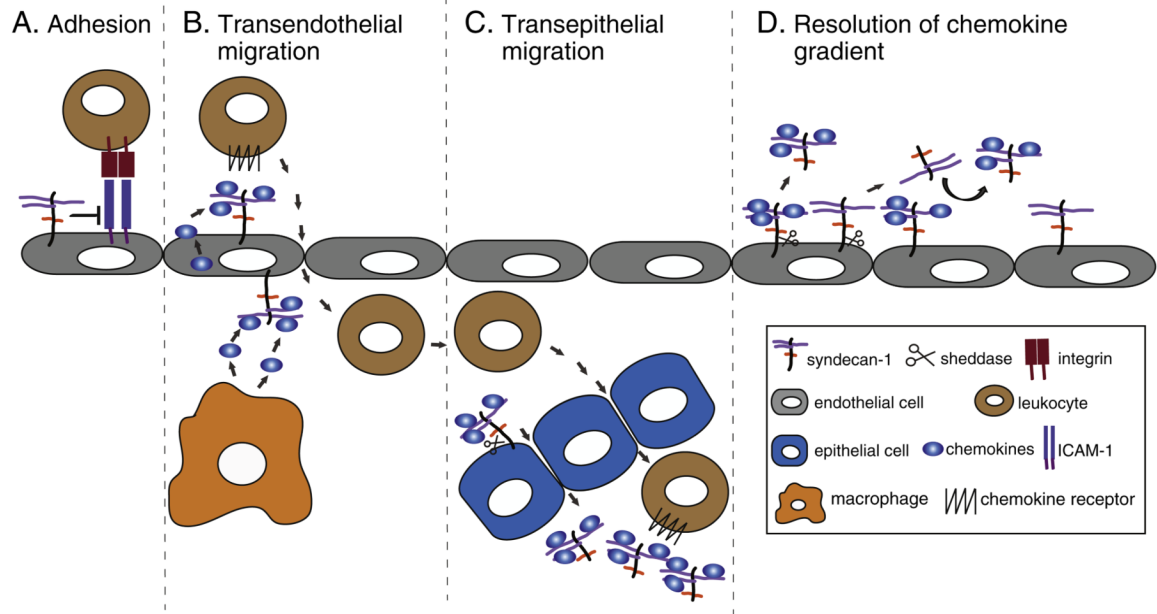
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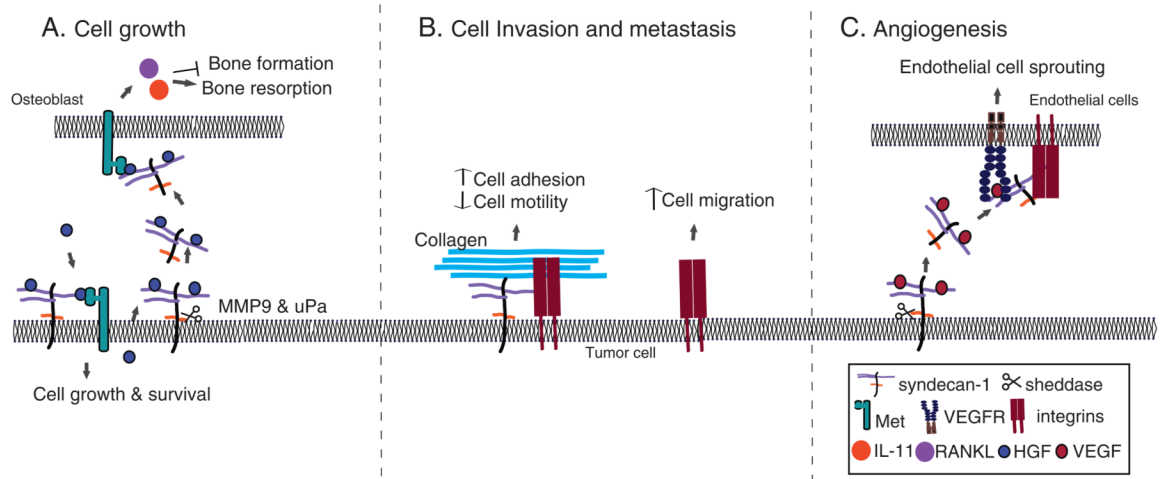


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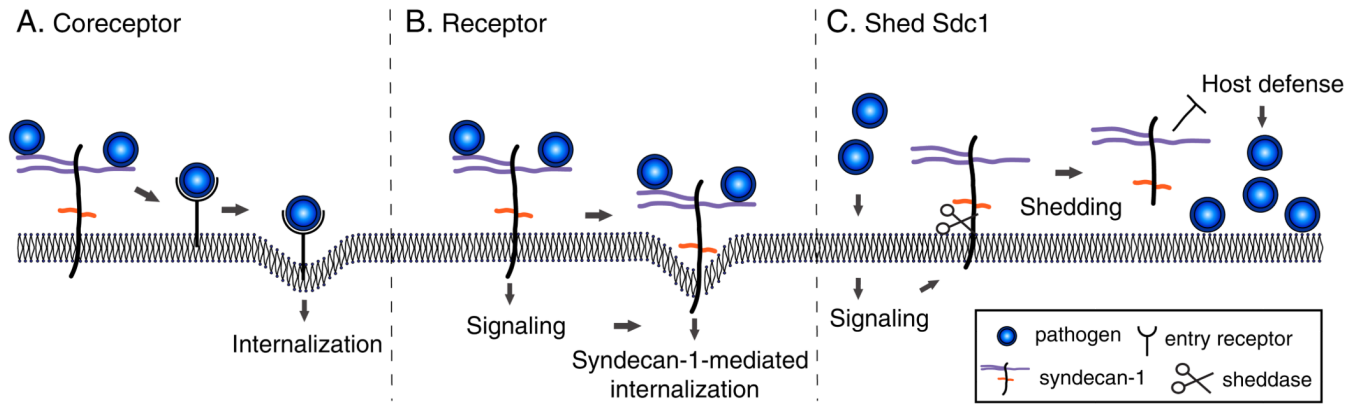
**Fig. 1. Mechanisms of syndecan-1 in inflammatory diseases**

A) Syndecan-1 inhibits leukocyte adhesion to activated endothelial cells by inhibiting the  $\beta 2$  integrin-ICAM-1 interaction. B) Endothelial syndecan-1 binds to chemokines and generates a chemokine gradient that facilitates the transendothelial migration of leukocytes. Cell surface syndecan-1 is shown in both the basal and apical surfaces as the normal polarized expression of syndecan-1 on the basolateral surface is disrupted by injury. C) MMP-7-mediated syndecan-1 shedding in alveolar epithelial cells generates a chemokine gradient for the transepithelial migration of neutrophils. D) Syndecan-1 shedding resolves chemokine gradients by removing chemokines tethered to cell surface syndecan-1 or by displacing chemokines tethered to other HSPGs through the release of unbound syndecan-1 ectodomains.



**Fig. 2. Mechanisms of syndecan-1 in cancer**

A) In myeloma, cell surface syndecan-1 binds to HGF via HS chains and promotes Met activation, leading to myeloma cell growth and survival. Shed syndecan-1 also binds to and presents HGF to osteoblasts. This results in the activation of IL-11 and upregulation of RANKL, leading to inhibition of bone formation and promotion of bone resorption. B) Cell surface syndecan-1 mediates cancer cell adhesion to the ECM and retards cancer cell migration. C) Syndecan-1 shedding releases the VEGF-syndecan-1 ectodomain complex, which binds to and activates VEGFRs on endothelial cells. Syndecan-1 can also activate  $\alpha\beta3$  integrin, which mediates the activation of VEGF signaling. This transactivation mechanism stimulates endothelial cell sprouting and subsequent angiogenesis.



**Fig. 3. Mechanisms of syndecan-1 in infectious diseases**

A) Syndecan-1 is subverted by many microbial pathogens as a coreceptor that localizes pathogens to the host cell surface and promotes their interactions with specific entry receptors. B) Select pathogens, such as *N. gonorrhoeae*, can use cell surface syndecan-1 as a direct entry receptor by stimulating intracellular signaling mediated by the syndecan-1 cytoplasmic domain. C) Several bacterial pathogens induce syndecan-1 shedding and exploit the capacity of syndecan-1 ectodomains to inhibit the antibacterial mechanisms of neutrophils and antimicrobial peptides.

Table 1

Partial list of syndecan-1 functions in animal and cellular models of diseases.

| Disease                     | <i>Sdc1</i> <sup>-/-</sup> mouse and <i>Sdc1</i> KD cell phenotype   | Target molecule              | Suggested function  | References                                 |
|-----------------------------|--|------------------------------|---|--|
| <i>Inflammation</i>         |  |                              |   |  |
| Acute lung injury           | Decreased CXCL1 in alveolar space, increased PMNs in perivascular space  | CXCL1                        | Generates CXCL1 chemokine gradient for transepithelial PMN migration into the alveolar space    | (Li et al., 2002)                          |
| Allergic lung inflammation  | Exaggerated Th2 response, AHR, eosinophilia, glycoprotein hypersecretion                                       | CCL7, CCL11, CCL17           | Inhibits CC chemokine-mediated Th2 cell homing to the lung                                      | (Xu et al., 2005)                          |
| Endotoxic shock             | Sustained high levels of CXCL1 and CXCL2 in tissues, increased PMN accumulation, multi-organ injury, lethality | CXCL1, CXCL2                 | Facilitates removal of sequestered CXCL1 and CXCL2, resolution of PMN inflammation              | (Hayashida et al., 2009b)                  |
| Gram-positive toxic shock   | Increased serum TNF $\alpha$ and IL-6, T cell infiltration, multi-organ injury, lethality                      | IFN $\gamma$ , MIG           | Inhibits amplification of cytokine storm by IFN $\gamma$ , MIG-mediated T cell recruitment      | (Hayashida et al., 2008a)                  |
| Protein-losing enteropathy  | Increased intestinal protein leakage   | IFN $\gamma$ , TNF $\alpha$  | Inhibits IFN $\gamma$ and/or TNF $\alpha$   | (Bode et al., 2008)                        |
| Colitis                     | Impaired mucosal repair, increased expression of inflammatory factors, leukocyte recruitment, lethality        | FGF-2, ICAM-1                | Promotes intestinal epithelial wound repair, inhibits leukocyte adhesion onto endothelial cells | (Day and Forbes, 1999; Floer et al., 2010) |
| Vascular injury             | Increased neointimal injury, vascular SMC proliferation  | PDGF-B, PDGFR $\beta$        | Inhibits SMC PDGF-B expression and PDGFR $\beta$ activation                                     | (Fukai et al., 2009)                       |
| Skin injury                 | Delayed wound repair, reduced keratinocyte proliferation   | $\alpha$ 9 integrins         | Regulates expression and localization of $\alpha$ 9 integrins                                   | (Stepp et al., 2002)                       |
| Anti-GBM nephritis          | Increased T cell influx, albuminuria, serum creatinine   | Th2 cytokines and chemokines | Regulates Th1/Th2 balance   | (Rops et al., 2007)                        |
| Allergic contact dermatitis | Increased expression of pro-inflammatory factors, leukocyte recruitment, edema formation                       | $\beta$ 2 integrins, ICAM-1  | Inhibits $\beta$ 2 integrin-ICAM-1 interaction  | (Kharabi Masouleh et al., 2009)            |
| Myocardial infarction       | Increased leukocyte influx, MMP activity, cardiac dysfunction, collagen fragmentation                          | MMPs, adhesion molecules     | Inhibits leukocyte recruitment, collagen degradation  | (Vanhouette et al., 2007)                  |
| Cardiac fibrosis            | Reduced CTGF, collagens, cardiac dysfunction   | Angiotensin II, TGF $\beta$  | Amplifies angiotensin II and TGF $\beta$ signaling  | (Schellings et al., 2010)                  |

| Disease                       | <i>Sdc1</i> <sup>-/-</sup> mouse and Sdc1 KD cell phenotype   | Target molecule                | Suggested function   | References   |
|-------------------------------|---|--------------------------------|--|--|
| Idiopathic pulmonary fibrosis | Impaired repair in epithelial cells treated with Sdc1 siRNA, reduced migration, increased proliferation in cells treated with Sdc1 ectodomain | EC-SOD, TGFβ                   | Promotes epithelial repair (cell surface Sdc1), inhibits epithelial repair, promotes PMN migration, TGFβ release, fibroblast proliferation (shed Sdc1) | (Kliment et al., 2009)                             |
| Lipoprotein metabolism        | Increased TRLs in plasma  | Plasma lipoproteins            | Mediates uptake of hepatic and intestinal TRLs   | (Stanford et al., 2009)                            |
| <i>Cancer</i>                 |   |                                |  |  |
| Tumorigenesis                 | Reduced Wnt-induced mammary gland hyperplasia   | Wnt                            | Promotes Wnt signaling   | (Alexander et al., 2000)                           |
| Cell growth and survival      | Altered HGF-Met signaling when HS or Sdc1 expression is manipulated in cancer cells   | HGF                            | Enhances HGF-Met signaling   | (Derksen et al., 2002; Ramani et al., 2011)        |
| Apoptosis                     | Increased apoptosis in cancer cells treated with Sdc1 shRNA or Sdc1 ectodomains   | Growth factors                 | Sequesters growth factors, inhibits signaling  | (Sun et al., 2008; Khotskaya et al., 2009)         |
| Angiogenesis                  | Diminished heparanase-induced endothelial invasion when Sdc1 immunodepleted   | VEGF                           | Activates integrins and VEGF receptors on adjacent endothelial cells   | (Purushothaman et al., 2008)                       |
| Invasion and metastasis       | Increased motility and invasion, decreased adhesion to collagen when cancer cells treated with Sdc1 siRNA                                     | α2β1, αvβ3, αvβ5 integrins     | Potentiates integrin activities  | (Beauvais et al., 2009; Ishikawa and Kramer, 2010) |
| <i>Infection (pathogen)</i>   |   |                                |  |  |
| HEV                           | Reduced viral attachment and entry in cells with reduced Sdc1   | pORF2                          | Promotes viral attachment  | (Kalia et al., 2009)                               |
| HSV                           | Reduced viral attachment and entry in cells with reduced Sdc1   | gB, gC, and gD                 | Promotes viral attachment and entry  | (Bacsa et al., 2011)                               |
| Neisseria gonorrhoeae         | Decreased invasion in cells expressing Sdc1 lacking the cytoplasmic domain  | Op <sub>HSPG</sub>             | Promotes signaling required for bacterial invasion   | (Freissler et al., 2000)                           |
| <i>Pseudomonas aeruginosa</i> | Reduced bacterial burden and lethality in lung and skin infection   | Cationic antimicrobial factors | Inhibits antimicrobial factors   | (Park et al., 2001; Haynes et al., 2005)           |
| <i>Staphylococcus aureus</i>  | Reduced corneal bacterial burden  | PMNs                           | Inhibits extracellular killing mechanisms of PMNs  | (Hayashida et al., 2011)                           |
| <i>Trichinella spiralis</i>   | Similar extent of infection between Wt and <i>Sdc1</i> <sup>-/-</sup> mice, reduced Th2 responses   | Th2 cytokines                  | Moderates Th2 responses  | (Beiting et al., 2006)                             |

| <b>Disease</b>         | <b><i>Sdc1</i><sup>-/-</sup> mouse and <i>Sdc1</i> KD cell phenotype</b>                                 | <b>Target molecule</b> | <b>Suggested function</b>                       | <b>References</b>              |
|------------------------|--|------------------------|---|--------------------------------|
| <i>Plasmodium</i> spp. | Similar extent of infection between Wt and <i>Sdc1</i> <sup>-/-</sup> mice in <i>P. yoelii</i> infection | PfEMP1                 | Promotes parasitic attachment and dissemination | (Bhanot and Nussenzweig, 2002) |