

REVIEW

Role of Matrix Metalloproteinases and Therapeutic Benefits of Their Inhibition in Spinal Cord Injury

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Summary: This review will focus on matrix metalloproteinases (MMPs) and their inhibitors in the context of spinal cord injury (SCI). MMPs have a specific cellular and temporal pattern of expression in the injured spinal cord. Here we consider their diverse functions in the acutely injured cord and during wound healing. Excessive activity of MMPs, and in particular gelatinase B (MMP-9), in the acutely injured cord contributes to disruption of the blood-spinal cord barrier, and the influx of leukocytes into the injured cord, as well as apoptosis. MMP-9 and MMP-2 regulate inflammation and neuropathic pain after peripheral nerve injury and may contribute to SCI-induced pain. Early pharmacologic inhibition of MMPs or the gelatinases (MMP-2 and MMP-9) results in an improvement

in long-term neurological recovery and is associated with reduced glial scarring and neuropathic pain. During wound healing, gelatinase A (MMP-2) plays a critical role in limiting the formation of an inhibitory glial scar, and mice that are genetically deficient in this protease showed impaired recovery. Together, these findings illustrate complex, temporally distinct roles of MMPs in SCIs. As early gelatinase activity is detrimental, there is an emerging interest in developing gelatinase-targeted therapeutics that would be specifically tailored to the acute injured spinal cord. Thus, we focus this review on the development of selective gelatinase inhibitors. **Key Words:** Spinal cord injury, matrix metalloproteinases, neuropathic pain, recovery, therapeutics.

INTRODUCTION

Matrix metalloproteinases (MMPs) are involved in a wide range of proteolytic events requiring matrix remodeling in normal development, wound healing, and repair throughout life [1, 2]. In brain and spinal cord injuries, MMPs have been shown to degrade components of the basal lamina, leading to disruption of the blood-brain barrier (BBB) [3–5], and to contribute to oxidative stress [6], demyelination [4], leukocyte trafficking, and a progressive

neuroinflammatory response [3, 4, 7]. In this review, we will focus on the roles of MMPs and the therapeutic implication of their inhibitors in spinal cord injury (SCI).

For the past decade, MMPs have been studied in a variety of neurodegenerative diseases, as well as in experimental models of central nervous system (CNS) trauma [3, 4, 8]. Studies of the latter have demonstrated that MMPs, expressed acutely after injury, are key mediators of pathogenesis. Their short-term blockade with either broad-spectrum MMP inhibitors or more specific gelatinase inhibitors is neuroprotective and results in neurologic recovery [5, 6]. Given these exciting findings, there is considerable interest in defining those MMPs that mediate early pathogenesis by establishing the therapeutic window for intervention and by developing more specific inhibitors for MMPs.

Here we provide a brief overview of MMPs and their regulation, we address their complex roles in both the acute and chronically injured spinal cord, and we

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consider the emergence of specific MMP inhibitors that will likely lead to a more complete understanding of the role of these proteases in the injured spinal cord and to the development of therapeutics that are specifically tailored to the patient with SCI.

OVERVIEW OF THE BIOLOGY OF MMPs

MMPs are best known for their ability to cleave constituents of the extracellular matrix (ECM). In more recent years, their targets have greatly expanded to include other proteinases, proteinase inhibitors, clotting factors, chemotactic molecules, latent growth factors, growth factor-binding proteins, cell surface receptors, cell-cell adhesion molecules, and almost all structural ECM proteins [9]. Processing of these molecules influences cell functions as diverse as cellular differentiation, migration, regulation of growth factor activity, survival or apoptosis, angiogenesis, inflammation, and signaling [1, 2].

Classification and protein structure

The MMPs constitute a family of both a zinc- and a calcium-dependent endopeptidase that includes 23 human (24 murine) MMP members [1]. Based on protein structure and substrate specificity, MMPs are divided into collagenases (MMP-1, MMP-8, MMP-13, and MMP-18), gelatinases (MMP-2 and MMP-9), stromelysins (MMP-3, MMP-10, and -11), membrane-type MMPs (MMP-14 [MT1-MMP], MMP-15 [MT2-MMP], MMP-16 [MT3-MMP], MMP-17 [MT4-MMP], MMP-24 [MT5-MMP], and MMP-25 [MT6-MMP]), and other MMPs [1, 2]. In general, MMPs contain three structural domains: 1) the *N*-terminal propeptide domain, 2) an internal catalytic domain, and 3) a *C*-terminal hemopexin domain. MMPs are initially produced as inactive zymogens, with a cysteine residue at the propeptide region that binds the zinc ion present at the catalytic site. Activation requires removal of the propeptide domain to expose the active catalytic site. The transmembrane domain is found only in the membrane-type MMPs (MT-MMPs) and is linked to the plasma membrane either by a transmembrane domain or by a glycosylphosphatidylinositol linkage attached to the hemopexin domain.

Modulation of MMPs

MMPs are regulated by various mechanisms that include gene transcription, mRNA stability, translational control, zymogen activation via proteolysis, inhibition by endogenous inhibitors, and their localization [9]. MMPs are regulated at the transcriptional level and by post-translational modification. Most MMPs are not constitutively expressed at detectable levels. Transcription of

MMP genes is thought to be mediated by intracellular signals in response to growth factors, cytokines, chemokines, and components of the ECM [9].

Post-translational modifications provide a second level of regulation. Although most MMPs are secreted as inactive zymogens, a subset of MMPs including MMP-11, MMP-27, and the MT-MMPs are activated primarily intracellularly by serine proteases of the pro-protein convertase class such as furin [9]. Some MMPs are cleaved in their propeptide domains, by serine proteinases such as the uPA-plasmin system and trypsin. Activation of other pro-MMPs is mediated by already activated MMPs. For example, pro-MMP-7 is activated by MMP-3, and MMP-7 activates pro-MMP-1 and pro-MMP-9 [10]. MT1-MMP activates pro-MMP-13 and this activated MMP-13, which may then go on to activate MMP-9 [11, 12]. MMP-2 is activated at the cell surface through a unique and complex mechanism involving MMP-14 (MT1-MMP) and tissue inhibitors of matrix metalloproteinase (TIMP)-2 [9]. Indeed, the transmembrane MT-MMPs (MT1-MMP, MT2-MMP, MT3-MMP, and MT5-MMP) are all able to activate pro-MMP-2, except for MT4-MMP and human, but not mouse). MT2-MMP are the only MT-MMPs that are unable to activate MMP2 [9].

Endogenous inhibitors of MMPs

There are four known mammalian TIMPs: 1) TIMP-1, 2) TIMP-2, 3) TIMP-3, and 4) TIMP-4. The *N*-terminal domain of TIMP proteins contains the MMP inhibitory domain, whereas the *C*-terminal domain of the TIMPs mediates important protein-protein interactions, in particular with the hemopexin domains of pro-MMPs [13]. Following activation, TIMPs modulate MMP activity by binding to the catalytic sites of MMPs, forming tight 1:1 noncovalent complexes [9, 13]. The TIMPs differ in their affinity for specific MMPs, and their interaction does not always lead to inhibition. The latter is exemplified by TIMP-1, which binds pro-MMP-9, thus protecting this protease from MMP-3 cleavage.

MMPs AND THE INJURED SPINAL CORD

Here we profile those MMPs that have been studied in the injured spinal cord, focusing on their unique temporal and cellular profiles in the injured cord (Table 1).

Time course and cellular localization

One of the earliest studies of MMPs in SCI focused on the gelatinases MMP-9 and MMP-2 in the contused cords of rats [14]. Gelatin zymography revealed prominent MMP-9 activity by 12- to 24-h postinjury followed by a rise in MMP-2 by 5 days postinjury. These findings have since been confirmed in other species and models,

including hemisection and compression. Zymography and Western blots revealed a transient increase in MMP-9 at 1-day postinjury followed by a gradual rise in MMP-2, which then remained elevated for weeks thereafter [6, 15–18]. Although less studied, MMP-1 is also up regulated 1-day postinjury in a similar model of SCI; however, its time course has yet to be studied [18]. Together, these findings highlight a unique temporal profile for MMP-1, MMP-9, and MMP-2 in the injured cord, with the first and second dominating in the more acute phase, and the third associated with wound healing.

Several studies have profiled the expression of virtually all known MMPs after SCI by evaluation of mRNA transcripts. In a murine model of spinal cord compression, there is up-regulation of mRNA transcripts encoding MMP-9, MMP-3, MMP-7, MMP-10, MMP-11, MMP-13, MMP-19, and MMP-20 within 24 h of injury, whereas increased expression of MMP-2, MMP-12, and MMP-13 is delayed in onset until 5 days after injury [19]. A similar trend is reported in a spinal cord contusion model in the rat [20].

During their peak expression in the acutely injured spinal cord, both MMP-1 and MMP-9 are localized to neurons and glia [18]. In addition, MMP-9 is detected in blood vessels, neutrophils, and macrophages [5, 6, 18]. MMP-2 localizes to reactive astrocytes and neurons in the chronically injured cord [17, 20]. Finally, MMP-12 is expressed primarily in microglia/macrophages [19]. Similar to other MMPs, its temporal and cellular expression varies according to type and severity of injury.

MMPS AND THEIR INHIBITORS IN INJURY AND RECOVERY PROCESSES

SCI produces secondary tissue damage that continues to evolve days and weeks after the initial insult, accompanied by corresponding functional impairments. Reducing the extent of progressive tissue loss following SCI represents an essential step toward recovery after SCI. MMP inhibitors have been tested in several animal models of acute and chronic injury (Table 2). The overall impression is that early blockade of MMPs stabilizes the barrier, reducing apoptotic cell death, and confers both early and long-term neuroprotection.

Inflammation and barrier breakdown

Experimental models of SCI suggest that MMPs support infiltration of inflammatory cells into the injured cord and most likely contribute to early disruption of the blood-spinal cord barrier. As leukocytes transmigrate across the vascular wall, they release MMPs, which in turn degrades tight-junction related proteins and the

surrounding basal lamina. Zonulae occludens-1, VE-cadherin, and occludin are substrates for MMP-2, MMP-3, MMP-7, and MMP-9 [21–24]. Basal lamina proteins, such as fibronectin, laminin, and heparan sulfate, are also degraded by MMPs [25]. Involvement of MMPs in barrier disruption is supported by the finding that the broad-spectrum MMP inhibitor BB-3103 decreases endothelial gap formation and occludin loss [26].

Of the MMPs expressed by leukocytes, MMP-9, conveyed by infiltrating neutrophils, is a key mediator of early pathogenesis in the injured cord. Immunologically depleting neutrophils prior to SCI results in reduced MMP-9 activity in the injured cord, suggesting that these leukocytes are the principal source of this protease in the injured tissue [14]. In a murine model of spinal cord contusion injury, barrier disruption to the protein luciferase, is maximal at 24-h postinjury [27], a time point that corresponds to peak activity of MMP-9 [5]. Moreover, barrier disruption is reduced in MMP-9 null mice, as well as mice treated with the broad-spectrum MMP inhibitor GM6001 early during the maximal expression 3 h to 3 days after injury. Similar to findings with GM6001, intrathecal administration of the selective gelatinase inhibitor SB-3CT at 2 h before injury to the rat spinal cord reduces both MMP-9 activity and barrier disruption by 1-day postinjury and decreases apoptotic cell death [6].

Similar findings supporting greater stabilization of the barrier have been reported in spinal cord injured MMP-12 null mice [19]. MMP-12, also known as macrophage metalloelastase, is critical for the migration of blood-borne macrophages across the endothelial basement membranes into inflammatory sites [28]. Spinal cord injured MMP-12 null animals show attenuation of blood-spinal barrier breakdown. It is likely that MMP-12 also influences the migration of macrophages into the injured cord. Comparisons of cell density of Iba-1-positive elements reveal fewer macrophages and microglia in MMP-12 null mice compared with wild type animals [19].

Oxidative stress and apoptosis

Oxidative stress contributes to pathogenesis in the injured spinal cord [4]. MMPs are regulated by reactive oxygen species, including nitric oxide and hypochlorous acid [29, 30]. Transgenic rats that over-express the antioxidant enzyme superoxide dismutase 1 (a crucial endogenous antioxidant enzyme responsible for eliminating superoxide) showed enhanced neuroprotection after SCI [31]. Active MMP-9 is increased at days 1, 3, and 7 after SCI in wild type rats, whereas there is no increase in the superoxide dismutase 1 transgenic rats. The end result is reduced disruption of the barrier and decreased apoptosis [6].

Table 1. Summary of MMPs in Experimental Models.

Type	Common Name	Species	Type of Injury	Changes in MMP	Peak Time for Change	Localization	Methods for Detecting MMP	References
MMP-1	Collagenase 1	Rat	Contusion	Increased activity	24 h	Neurons, glia	Westerns, ICC	[18]
MMP-2	Gelatinase A	Rat	Contusion	Increased transcripts	1–21 days	Not studied	qtPCR	[20]
		Mice	Contusion	Increased activity	5 days	Not studied	Gelatin zymography	[14]
			Contusion		7–14 days	Astrocytes	<i>Gelatin/in situ</i> zymography, ICC	[16, 17]
		Rat	Hemisection		1 day–4 weeks	Not studied	Gelatin zymography	[15]
		Rat	Compression		3–7 days	Not studied	Gelatin zymography	[6]
		Mice	Nerve ligation		7–21 days	Astrocytes	Gelatin zymography,	[43]
							ICC	
		Mice	Compression	Increased transcripts	5 days	Not studied	qtPCR	[19]
		Mice	Contusion		5 days	Not studied	qtPCR	[32]
		Rat	Contusion		3–21 days	Neurons	qtPCR, gelatinase zymography, westerns, ICC	[20]
MMP-9	Gelatinase B	Rat	Contusion	Increased activity	12–24 h	Not studied	Gelatin zymography	[14]
		Mice	Contusion		24 h	Astrocytes, vessels, macrophages	<i>Gelatin/in situ</i> zymography, ICC	[5, 16]
		Rat	Contusion		24 h	Neurons, glia	Westerns, ICC	[18]
		Rat	Hemisection		24 h–48 h	Not studied	Gelatin zymography	[15]
		Rat	Compression		24 h	Neutrophils, endothelial cells, neurons	<i>Gelatin/in situ</i> zymography, westerns, ICC	[6]
		Mice	Nerve ligation		6 h–1 day	Dorsal root ganglion neurons	Gelatin zymography, ICC	[43]
		Rat	Nerve crush	Increased transcripts and activity	24 h	Not studied	qtPCR, gelatin zymography	[44]
MMP-12	Metalloelastase	Mice	Compression	Increased transcripts	1 day	Not studied	qtPCR	[19]
		Mice	Compression	Increased transcripts	5–14 days	Macrophages/microglia	qtPCR, ICC	[19]
MMP-3	Stromelysin 1	Rat	Contusion	Increased transcripts	21 days	Not studied	qtPCR	[20]
MMP-7	Matrilysin	Mice	Compression		24 h	Not studied	qtPCR	[19, 20]
		Rat	Contusion		3 days			
MMP-10	Stromelysin 2	Mice	Compression		24 h			
		Mice	Compression		3 days			
MMP-11	Stromelysin 3	Rat	Contusion		24 h			
		Mice	Compression		3–21 days			
MMP-13	Collagenase 3	Rat	Contusion		24 h			
		Mice	Compression		3 days			
MMP-14	MT1-MMP	Rat	Contusion		5 days			
MMP-16	MT3-MMP	Rat	Contusion		3 days			
MMP-17	MT54-MMP	Rat	Contusion		3 days			
MMP-19		Rat	Contusion		1–21 days			
					1–21 days			
					24 h			

MMP-20	Enamelysin					24 h				
MMP-21	XMMP					21 days				
MMP-23	Femalysin	Mice	Compression	Decreased transcripts		1-2 days	Not studied	qtPCR		[19]
MMP-24	MT5-MMP					2-5 days				
MMP-15	MT2-MMP	Rat	Contusion	Changed transcripts		1-3 days ↓ 7-21 days ↑ 1 day ↓ 3-21 days ↑	Not studied	qtPCR		[20]
MMP-8	Collagenase 2									
MMP-25	MT6-MMP									
MMP-28	Epilysin									

ICC = immunocytochemistry; qtPCR = quantitative Polymerase Chain Reaction; MMP = matrix metalloproteinase; MT = membrane type; XMMP = Xenopus laevis matrix metalloproteinase.

Table 2. Injury Phenotypes of Mice Deficient in MMP.

MMP	Knock-Out Mice	Model	Mechanism	Phenotype	References
MMP-2 -/-		SCI	Elimination of MMP-2; compensatory increase in MMP-9	Reduced white matter sparing and fewer serotonergic fibers, widespread reactive astrogliosis, and impairment in locomotor recovery.	[17]
MMP-12 -/-		SCI	Elimination of MMP-12	Improved functional recovery, attenuation of barrier disruption, and reduces microglial/macrophages.	[19]
MMP-9 -/-		SCI	Elimination of MMP-9	Improved function recovery and reduced barrier disruption, neutrophil infiltration, neuropathic pain and inflammatory pain.	[5, 43]
Candidate Therapeutics					
MMP-2/MMP-9 inhibitor, subcutaneous		SCI	Inhibit MMPs, especially MMP-9 and MMP-2	Reduced apoptosis in neurons and glia.	[32]
GM6001, intraperitoneal		Spinal nerve crush		Increased cell survival and attenuation of mechanical allodynia, degradation of myelin basic protein, macrophage influx, and glial activation.	[44]
Inhibitor I, intrathecal		Spinal nerve ligation	Inhibit MMP-9	Ameliorated early neuropathic pain behavior	[43]
Inhibitor-III, intrathecal		Spinal nerve ligation	Inhibit MMP-2	Ameliorated late neuropathic pain behavior	[43]
SB-3CT, intrathecal		SCI	Inhibit MMPs, especially MMP-9 and MMP-2	Reduced MMP activity, barrier disruption, and apoptosis.	[6]

MMP = matrix metalloproteinases; SCI = spinal cord injury.

After SCI, MMP-2 contributes to apoptotic cell death, and is up-regulated along with neuronal and glial apoptosis. This temporal relationship raises a question as to whether or not MMP2 is a determinant of apoptosis after SCI [32]. Supporting this possibility, spinal cord injured mice, treated with an MMP-2/MMP-9 inhibitor delivered subcutaneously daily for 5 days, showed reduced levels of apoptosis in both neurons and glial [32]. Such a reduction in total cell death may contribute to long-term recovery.

Collectively, these studies provide strong evidence for pathological MMP-directed disruption of the blood-brain barrier, leukocyte infiltration, and cell apoptosis after SCI. Because early blockade of MMPs stabilizes the barrier, reduces leukocyte infiltration, and confers both early and long-term neuroprotection, it suggests that early inhibition of MMPs may be an efficacious strategy for SCI.

Glial scar formation

Prolonged inhibition of MMPs, especially with broad-spectrum inhibitors during wound healing can be detrimental to neurological recovery [33]. One reasonable explanation is that some MMPs, expressed during wound healing, are beneficial. There is growing literature to support this hypothesis. In the more chronically injured cord, MMPs are key regulators of the local microenvironment, degrading the ECM, and modulating the formation of an inhibitory glial scar [34].

After SCI, a variety of growth-inhibitory molecules are up-regulated [35]. A glial scar, an interface that is inhibitory to axonal regeneration, forms at the site of injury and is composed of reactive astrocytes, microglia/macrophages, and ECM molecules, especially chondroitin sulfate proteoglycans (CSPGs) [35]. In response to injury, astrocytes, oligodendrocyte progenitors, and macrophages increase the expression of CSPGs, which in turn inhibit neurite outgrowth *in vitro* and regeneration *in vivo* [36, 37]. MMPs degrade the core protein of some CSPGs, as well as other growth-inhibitory molecules, such as Nogo and tenascin-C [34]. CSPGs, such as neurocan and versican, are degraded by MMP-2, whereas tenascin-C, brevican, neurocan, NG2, phosphacan, and versican are degraded by MMP-3 [34]. By degrading CSPG and other inhibitory molecules, MMPs support axonal regenerative potential in the injured CNS [2, 34].

In a rat hemisection model, *in situ* zymography reveals MMP-related gelatinase activity in the injured site, which is spatially and temporally correlated with scar formation [15]. *In vitro* and *in vivo* data support the possibility that MMPs facilitate migration of astrocytes. *In vitro* scratch wound assays show attenuated migration of cultured MMP-9 null astrocytes or astrocytes treated with an MMP-9 inhibitor [38]. Moreover, in spinal cord injured

MMP-9 null mice, glial scar formation is abrogated along with reduced CSPG immunoreactivity at the lesioned epicenter [38].

The importance of MMP-2 in promoting functional recovery in the chronically injured spinal cord has been evaluated in MMP-2 deficient mice. MMP-2 null mice show greater CSPG immunoreactivity, fewer serotonergic fibers caudal to the injury site, and significantly reduced motor recovery compared with wild-type mice after a contusive SCI [17]. Such a finding may result from reduced sprouting across the lesioned site.

Finally, the complexity of MMPs in SCI is further realized in the context of axonal dieback. After SCI, infiltrated macrophages mediate long-distance axonal retraction from the initial site of injury. The involvement of MMPs in dieback is supported by *in vitro* studies showing that a broad-spectrum inhibitor or a specific MMP-9 inhibitor prevents macrophage-induced axonal retraction [39].

In summary, MMPs limit the formation of an inhibitory glial scar and degrade the inhibitory proteins, as well as cleave extracellular proteins that sequester growth factors [34, 40], thus supporting recovery processes. They also mediate adverse responses including axonal dieback.

MMPS AND NEUROPATHIC PAIN

Neuropathic pain manifests as an unpleasant somatosensory experience evoked by lesion or dysfunction in the nervous system and occurs in as many as 85% of SCI patients [41]. Types of neuropathic pain include: allodynia, which is a painful response to an innocuous stimulus; and hyperalgesia, which is an exaggerated pain response to a noxious stimulus. After SCI, neuropathic pain can be localized in dermatomes above, at, or below the level of injury. The complexities of neuropathic pain create a barrier for therapeutic intervention.

Peripheral nerve injury (PNI) models are perhaps the most common method of studying neuropathic pain, as mechanisms can be examined without direct damage to central pathways. Recent evidence shows that similar inflammatory mediators of pain exist for SCI and PNI [42], suggesting that PNI may help elucidate pain mechanisms after SCI. In PNI, MMPs appear to induce and maintain neuropathic pain, but a similar role after SCI is unknown [43–45]. Prominent similarities in cytokine profiles and microglial activation between SCI and peripheral injury make an MMP mechanism with at or below-level pain plausible [42].

Neuropathic pain modulation in PNI

The gelatinases (MMP-9 and MMP-2) appear to regulate neuropathic pain after PNI through degenerative

and proinflammatory mechanisms. Nerve injury induces a rapid increase of active MMP-9 in the axon and dorsal root ganglia (DRG) by 24 h. MMP-9 gene expression precedes and outlasts the active form [44, 45]. Active MMP-2 in the DRG is delayed until day 7 and persists through day 21 [43]. Although a distinct time course has not been described for gelatinase activity in the dorsal horn after PNI, an early MMP-9 and delayed MMP-2 pattern occurs [43]. A differential time course suggests distinct roles of MMP-9 and MMP-2 in the development and maintenance of neuropathic pain.

After mechanical damage to the axon, Schwann cells release MMP-9, initiating macrophage infiltration and degradation of myelin basic protein [44, 46]. Exposure of the bare axon leads to increased sodium channel expression and ectopic hyperexcitability of afferents [47]. As a result, action potentials outlast the stimulus creating central sensitization, a common mechanism of neuropathic pain.

Many proinflammatory cytokines and growth factors mediate gelatinase expression in the nervous system, potentially inducing neuropathic pain. Indeed, exposure of the uninjured peripheral nerve to nerve growth factor, tumor necrosis factor alpha or interleukin-1 beta (IL-1 β) induces robust MMP-9 expression by Schwann cells [45]. Although neuronal sensitization occurs with elevated cytokines alone [48], MMP-9 expression also appears to mediate pain behavior. Gene deletion of MMP-9 reduces nociceptive pain behavior after PNI [43, 45]. In a proof-of-principle design, Kawasaki et al. [43] intrathecally injected MMP-9 and found marked allodynia accompanied by increased IL-1 β cleavage and microglial activation in the dorsal horn co-localized with phosphorylated p38 MAP kinase. Blocking IL-1 β signaling with a neutralizing antibody prevented allodynia, establishing IL-1 β as a downstream regulator of neuropathic pain by reducing microglial activation and p38 levels. Normal expression and intrathecal injection of MMP-2 also cleaves IL-1 β and activates spinal astrocytes at later time points, potentially maintaining neuropathic pain [43]. The distinct temporal activations of MMP-2 and MMP-9 provide novel opportunities for therapeutic intervention during different stages of allodynia.

MMP inhibitors and neuropathic pain

Outside of gene deletion, use of siRNAs and endogenous peptide inhibitors for general and specific inhibition of MMP-9 and MMP-2 effectively reduce allodynia after PNI. Daily systemic administration of a broad-spectrum gelatinase inhibitor (GM6001) 1 h to 12 days after PNI resulted in immediate and sustained attenuation of mechanical allodynia [44]. Cellular effects included preservation of myelin basic protein,

little macrophage influx, and low glial activation in the dorsal horn [44]. Thus, relatively short-term inhibition of MMPs limited the behavioral and cellular sequella of central pain syndromes. However, an overabundance of cells occurred in the nerve, DRG, and dorsal horn after treatment with broad-spectrum MMP inhibitors due to reduced apoptosis [44]. Less apoptosis and increased cell density is not without risk, especially after SCI. Such side effects must be managed if broad-spectrum inhibitors are used to treat neuropathic pain.

Temporal profiles of MMP-9 and MMP-2 provide specific opportunities for treating neuropathic pain at different stages. Pretreatment with MMP-9 siRNA prevented the onset of allodynia for a maximum of 5 days. When MMP-2 siRNA was delivered late after PNI during pronounced MMP-2 activity, allodynia was attenuated for at least 1 day. Moreover, behavioral benefits were accompanied by reduced cleavage of IL-1 β with both types of siRNA and reduced microglial activation in the dorsal horn with MMP-9 siRNA [43]. Even greater reversal of allodynia occurred when endogenous inhibitors of MMP-9 (TIMP-1) and MMP-2 (TIMP-2) were administered during early or late phases of MMP activation after PNI, but the effect was transient, lasting only 3 to 24 h [43]. The largest and longest reduction in allodynia after PNI occurred with intrathecal injection of specific pharmacologic inhibitors timed to periods of high MMP activity. Inhibition of MMP-9 using Inhibitor-I (Calbiochem, Gibbstown, NJ), starting 2 days before and through 5 days after the injury, delayed the onset of allodynia for at least 6 days (Table 2). Using the synthetic MMP-2 inhibitor (Inhibitor-III, Calbiochem), allodynia was attenuated for a maximum of 10 days with daily intrathecal injections (Table 2) [43]. These results suggest that intrathecal delivery effectively reduces allodynia. Timing of the delivery and selection of the type of inhibitor will be critical to clinical success in treating neuropathic pain.

Translation to SCI

Marked similarity of behavioral and cellular mechanisms of neuropathic pain exists between PNI and SCI. Given the role of gelatinases in pain development and maintenance after PNI, examination of MMP-9 and MMP-2 with at and below-level pain after SCI is warranted. While a differential time course exists with MMP-9 and MMP-2 after both types of injury, the relevance of delivering broad-spectrum MMP inhibitors early is unclear, as SCI-induced neuropathic pain onset occurs months or years after injury. Perhaps broad-spectrum inhibitors delivered at specific time points would be more effective for SCI, although serious musculoskeletal side effects may result from long-term

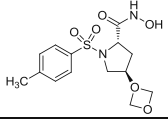
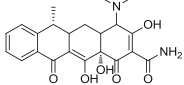
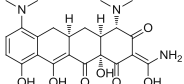
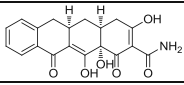
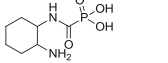
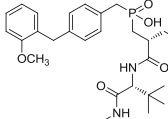
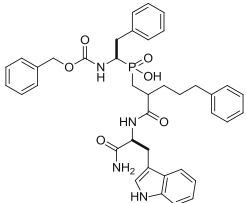
Table 3. Competitive MMP Inhibitors.

No.	Structure	Other Name	IC ₅₀ (nM) or *K _i (nM)						Indication/Comments	Ref.
			MMP-1	MMP-2	MMP-3	MMP-7	MMP-9	MMP-14		
1		Batimastat, BB-94	3	4	20		10	19	TACE: IC ₅₀ = >20,000 nM. Cancer. Administered ip and intrapleurally in clinical trials. Produced musculoskeletal (MSS) syndrome. Discontinued after phase I (poor oral bioavailability).	(80)
2		Marimastat, BB-2516	0.78	0.41	14	4.1	0.79		MMP-13: IC ₅₀ = 1.2 nM. Cancer. Orally bioavailable. Produced MSS. Discontinued after phase III (lack of efficacy).	(54)
3		Ilomastat, GM6001, Galardin®	0.4*	0.5*	27*	3.7	0.2*	13.4*	Cancer, macular degeneration, COPD. Discontinued after phase I (poor oral bioavailability).	(81)
4		Rebimastat, BMS-275291, D-2163	9	39	157	23	27	40	Cancer. Musculoskeletal toxicity. Discontinued after phase III (poor response).	(82)
5		Tanomastat, BAY 12-9566	>5,000*	11*	134*		301*		MMP-13: K _i = 1,470 nM. Arthritis, cancer, kidney transplant. Produced MSS, hematological toxicity, thrombocytopenia. Discontinued Phase III (poor survival rate).	(83)
6		Prinomastat, AG-3340	5.7	0.048	3.5	72	0.048		MMP-13: IC ₅₀ = 0.2 nM. Cancer, macular degeneration. Musculoskeletal toxicity. Discontinued after phase III (poor survival rate).	(54)
7		Cipemastat, Trocade®, Ro-32-3555	1.77	3,420	694		150	7.84	MMP-13: IC ₅₀ = 7.28 nM. Rheumatoid arthritis, osteoarthritis. Produced MSS. Discontinued after phase II (lack of efficacy).	(55)
8		CGS-27023, MMI-270	1* 15		16*		3* 9		MMP-13: K _i = 5 nM TACE: K _i = 54 nM Cancer. Oral bioavailability: 44% dog. Produced some MSS. Discontinued after phase II.	(84)
9		MMI-166	>1,000	0.4	>1,000	>1,000	90	100	MMP-8: IC ₅₀ = 400 nM Orally active in animal models of tumor growth and metastasis.	(85)
10		Tolyksam		5	5,558		49		MMP-8: IC ₅₀ = 397 nM MMP-12: IC ₅₀ = 34 nM MMP-13: IC ₅₀ = 201 nM	(86)
11		FR255031	77.8	91.1	>10,000		3.94	1.80	MMP-13: IC ₅₀ = 8 nM TACE: IC ₅₀ = 231 nM	(55)
12		RS-130830	590* 233	0.22*	9.3*		0.58* 3		MMP-13: K _i = 0.52 nM TACE: K _i = 176 nM Osteoarthritis. Discontinued after phase I.	(84)
13		SC-78080/SD-2590	>10,000	<0.1	28.7	7,000	0.18	13	MMP-13: IC ₅₀ = 0.1 nM Oral bioavailability 52%-82%.	(87)

Table 3. (Continued).

14			>10,000	400	370		1,230	>10,000	MMP-8: IC ₅₀ = >10,000 nM MMP-13: IC ₅₀ = 9 nM Oral bioavailability: rat 4.2%	(88)
15			835				228		MMP-13: IC ₅₀ = 77 nM TACE: IC ₅₀ = 16 nM	(89)
16			>10,000	20			5.5		Protein binding 83%, high clearance, low systemic exposure	(90)
17			>10,000	5.2	137		1.9		MMP-13: IC ₅₀ = 3.5 nM Protein binding 58%, t _{1/2} = 6 hr	(90)
18				9.3			2.9		Protein binding 74%. Reduced ischemia-induced brain edema in rat MCAO model.	(90)
19		ABT-770	4,600	4	42	>10,000	120		Oral bioavailability: rat 93%, dog 83%, monkey 95%. Efficacious in several cancer animal models. Produced phospholipidosis.	(91)
20		ABT-518	8,900	0.78	12	11,000	0.5		MMP-13: IC ₅₀ = 3.3 nM Oral bioavailability >70% in rat, dog, and monkey. Solid tumors. Orally bioavailable. No further development reported after phase I.	(54)
21			>25,000	1			152	>10,000	MMP-13: IC ₅₀ = 1,130 nM	(92)
22		Ro-28-2653	16,000	10	1,800		12	10	MMP-8: IC ₅₀ = 15 nM MMP-16: IC ₅₀ = 23 nM TACE: IC ₅₀ = >20,000 nM	(93)
23			>5,000*	0.23*	3,200*		0.72*		MMP-13: K _i = 2.7 nM TACE: K _i = >1000 nM	(94)
24			3,245				7		MMP-13: IC ₅₀ = 4 nM TACE: 32% inhibition at 1 μM	(95)
25			155				1		MMP-13: IC ₅₀ = 0.8 nM TACE: IC ₅₀ = 122 nM	(96)
26				5.2					APN: IC ₅₀ = 75.2 nM	(97)

Table 3. (Continued).

27				3					APN: IC ₅₀ = 48.5 nM	(98)
28		Doxycycline Adoxa®, Doryx®, Vibra-Tabs®, Periostat®	>100000	>100000	>100000	>100000	>100000		Approved for acne, bacterial infections, gum disease, and rosacea. Available as iv infusion and as oral tablets and capsules.	(53)
29		Minocycline Minocin®				125,000	180,000		Approved for acne, bacterial infections, and rosacea.	(57)
30		Incyclinide, Metastat®, COL-3		5,500					Acne, AIDS-related Kaposi's sarcoma, cancer. Phase II.	(99)
37			>100000	4,000	>100000		20,000		MMP-13: IC ₅₀ >100000 nM TACE: IC ₅₀ >100000 nM	(64)
38			1,200	6.6	1,600				MMP-8: IC ₅₀ = 2.4 nM MMP-12: IC ₅₀ = 5 nM MMP-13: IC ₅₀ = 4.5 nM	(66)
39		RXP-03	>2,000*	20*		8% @ 2 μM	10*	105*	MMP-8: K _i = 2.5 nM MMP-11: K _i = 5 nM MMP-13: K _i = 16 nM	(65)

Values with asterisk (*) means K_i

use. Alternatively, early specific gelatinase inhibitors may prevent the development of pain. The fact remains that there have been no studies of the role of gelatinases in neuropathic pain after SCI. The debilitating nature of neuropathic pain after SCI and the potential of MMP-9 and MMP-2 to alter the course of its development and maintenance places greater emphasis on explorative studies of selective gelatinase inhibitors.

SYNTHETIC GELATINASE INHIBITORS

Gelatinase has been implicated in a variety of pathological processes and diseases, including inflammation, cardiovascular disease, infection, neurodegenerative disease, immune response, cancer cell invasion, and metastasis [2, 8, 49]. Thus, selective gelatinase inhibitors are highly sought. However, due to the structural similarities of

MMPs, selective inhibition of gelatinase has been challenging. As recent reviews of MMP inhibitors are available [50–52], this section will focus on gelatinase inhibitors.

Competitive inhibitors

The first-generation MMP inhibitors were broad-spectrum peptidomimetics, containing a hydroxamate moiety that chelated the catalytic zinc and inactivated the protein. Examples of early peptidomimetics are batimastat (**1**), marimastat (**2**), and ilomastat (**3**) (Table 3). Although these small molecules inhibited MMPs at nanomolar concentrations, they were not selective, and in many cases also inhibited the a disintegrin and metalloproteinase ADAMs (tumor necrosis factor alpha converting enzyme TACE) proteases. Moreover, the hydroxamate moiety is readily metabolized [51], contributing to the poor oral bioavailability and limited *in vivo*

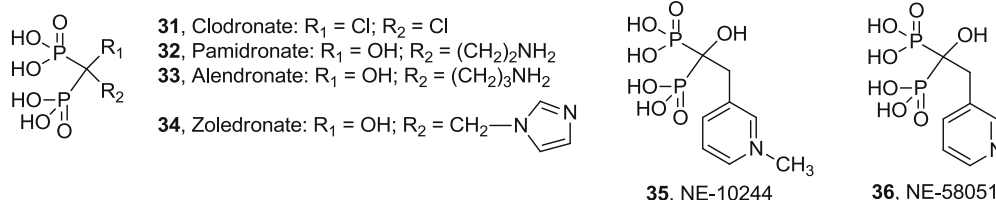
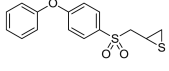
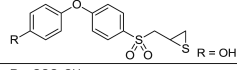
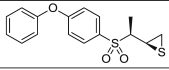
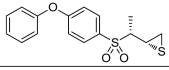
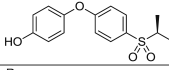
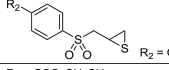

FIG. 1. Phosphorus-based inhibitors.

Table 4. Mechanism-based MMP Inhibitors.

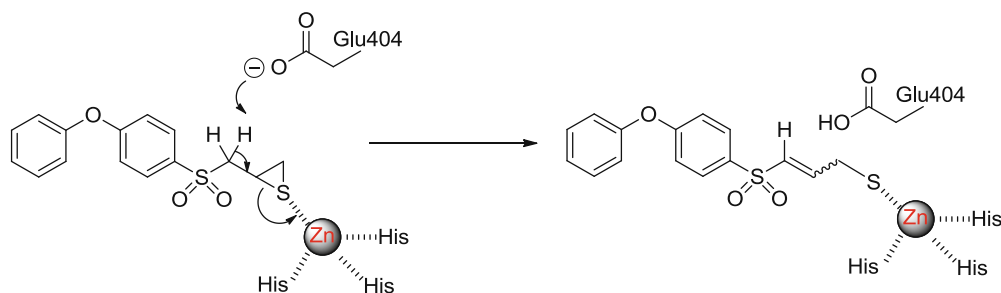
No.	Structure	Other Name	K_i (nM)						Comments	Ref.
			MMP-1	MMP-2	MMP-3	MMP-7	MMP-9	MMP-14		
40		SB-3CT	73,000	28	4,000	67,000	400	110	Slow-binding inhibitor of MMP-2 and -9. Inhibits MMP-14 as a competitive (reversible) inhibitor. Does not inhibit ADAM-10 -17.	(68, 100)
41			128,000	6	2,200	31,000	160	90	Primary metabolite of SB-3CT. Slow-binding inhibitor of MMP-14.	(75)
42	R = OSO ₂ CH ₃		140,000	23	600	18,200	5	145	Inhibits MMP-3, MMP-7, and MMP-14 as a competitive inhibitor.	(76)
43	R = OSO ₂ CH ₂ CH ₃		NI	70	38% at 20 μM	8% at 40 μM	330	360		(77)
44	R = OSO ₂ (CH ₂) ₂ CH ₃		NI	34	22% at 20 μM	5% at 20 μM	520	240		(77)
45	R = OSO ₂ CH(CH ₃) ₂		NI	61	5% at 20 μM	NI	380	790		(77)
46	R = NHSO ₂ CH ₃		NI	16	3,600	295,000	180	900	Slow-binding inhibitor of MMP-2, -9, and -14.	(78)
47	R = NHCOCH ₃		5,400	110	12,200	39,000	130	680	Slow-binding inhibitor of MMP-2 only.	(78)
48	R = CH ₂ COOCH ₃		11,000	50	8,700	13,000	40	590	Slow-binding inhibitor of MMP-2 only.	(78)
49	R = CH ₂ COOH		4,500	460	540,000	250,000	4,100	53,000	Slow-binding inhibitor of MMP-2 only.	(78)
50			NI	180	8% at 60 μM	NI	3,500	740	Slow-binding inhibitor of MMP-2, -9, and -14.	(79)
51			NI	390	9% at 60 μM	NI	3,300	2,100	Slow-binding inhibitor of MMP-2, -9, and -14.	(79)
52			NI	220	NI	NI	1,900	2,100	Slow-binding inhibitor of MMP-2, -9, and -14. Minor metabolite of 54.	(79)
53			41,000	390	29% at 200 μM	11,000	3,900	480	Slow-binding inhibitor of MMP-2, -9, and -14.	(77)
54	R ₂ = OSO ₂ CH ₂ CH ₃		7% at 200 μM	90	62% at 200 μM	26,000	12,000	11,000		(77)
55	R ₂ = OSO ₂ (CH ₂) ₂ CH ₃		30% at 200 μM	280	58% at 200 μM	35,000	6,200	3,200		(77)
56	R ₂ = OSO ₂ CH(CH ₃) ₂		NI	240	25% at 200 μM	120,000	3,500	20,000		(77)

NI=non inhibitory

efficacy. Another drawback is that these broad-spectrum inhibitors produced musculoskeletal syndrome [53].

Other zinc-binding groups have been used. Rebimastat (BMS-275291, **4**) contains a mercaptoacyl as a zinc-binding group and tanomastat (BAY 12-9566, **5**) has a zinc-binding carboxylate group. More selective MMP inhibitors have been developed, such as prinomastat (**6**) [54] and cipemastat (**7**) [55]. Examples of additional peptidomimetic and nonpeptidomimetic competitive inhibitors are listed in Table 3. However, as zinc chelators these inhibitors target the gelatinases, as well as other zinc-dependent enzymes, including other MMPs.

Since their discovery in the 1950s, tetracyclines have been widely used for their antimicrobial properties. However, tetracyclines also have nonantimicrobial properties, and among them is the ability to inhibit MMPs [56]. Doxycycline (Periostat®, **28**) is the only drug approved as an MMP inhibitor for the treatment of periodontitis. Minocycline (**29**) is a second-generation tetracycline analog with weak broad-spectrum MMP inhibitory activity [57] approved for acne, bacterial infections, and rosacea. Minocycline has been reported to have neuroprotective effects in various animal models of neurological diseases [58, 59]. Incyclinide (Metastat®,

**FIG. 2.** Mechanism of gelatinase inhibition by SB-3CT.

COL3 (**30**)) is a second-generation chemically modified tetracycline that lacks antimicrobial properties, inhibiting collagenases (MMP-1, MMP-8, and MMP-13), gelatinases (MMP-2 and MMP-9), and MT1-MMP, as well as other MMPs [60].

Bisphosphonates are potent inhibitors of osteoclastic activity and are widely used clinically for arresting or preventing bone loss associated with osteoporosis, metastatic bone disease, and other bone fragility conditions. Bisphosphonates also inhibit various MMPs [61, 62]. Clodronate (**31**), pamidronate (**32**), alendronate (**33**), and zoledronate (**34**), inhibit MMP-1, MMP-2, MMP-3, MMP-7, MMP-8, MMP-9, MMP-12, MMP-13, and MMP-14, with IC_{50} s ranging from 50 to 150 μ M. Analogs of risedronate, NE-10244 (**35**), and NE-58051 (**36**) (refer to FIG. 1 for their structures), inhibit MMP-2, MMP-9, and MMP-12, with IC_{50} s of 40, 160, and 80 μ M, respectively [63]. Carbamoylphosphonates (**37**) [64] and phosphinates (**38, 39**) have been reported [65, 66] as MMP inhibitors.

Mechanism-based inhibitors

In 2000, Brown et al. [67] reported on the design and synthesis of SB-3CT (**40**) (Table 4), the first mechanism-based inhibitor for gelatinases. In mechanism-based or slow-binding inhibition, the inhibitor-enzyme complex undergoes a requisite conformational change that does not readily reverse. SB-3CT is selective for MMP-2 and MMP-9, does not inhibit other MMPs, and inhibits MMP-14 (MT1-MMP) as a simple competitive (reversible) inhibitor [68]. Although the biphenyl ether motif is reminiscent of other MMP inhibitors, the basis for MMP inhibition by SB-3CT is mechanistically unique and different from all other reported MMP inhibitors. This is due to the ability of gelatinases to facilitate the requisite rate-limiting deprotonation event leading to thirane-ring opening with the active-site zinc ion (FIG. 2) [69]. Because the reactive thiolate is only generated within the active site of gelatinases, indiscriminate inhibition of other MMPs does not exist. SB-3CT has shown efficacy in several animal models of disease, including stroke [70], vascular permeability [71], vascular remodeling [72], subarachnoid hemorrhage [73], cardiopulmonary resuscitation [74], and SCI [6].

Although SB-3CT has demonstrated efficacy, it is rapidly and extensively metabolized by hydroxylation at the terminal phenyl ring (**41**) [75] to a more potent gelatinase inhibitor than the parent SB-3CT. The sulfonate (**42**) was designed to block the primary site of metabolism of SB-3CT [76]. Additional sulfonates (**43–45**) are slow-binding selective gelatinase inhibitors [77]. Other groups have been introduced into the terminal phenyl ring of SB-3CT to block metabolism, including *N*-methanesulfonate (**46**), *N*-acetate (**47**), methyl acetate (**48**), and methyl acetic acid (**49**) [78]. In efforts to reduce metabolism at the α -position to the sulfonyl group in SB-

3CT, 4 methyl diastereomers were prepared [79]. Of these, **50** and **51** were active slow-binding inhibitors of MMP-2, MMP-9, and MMP-14. The *p*-hydroxy methyl derivative **52** had a similar inhibition profile as **50**. To increase water solubility and metabolic stability, sulfonates **53–56** were prepared in which the terminal phenyl ring was eliminated [77]; these compounds showed potent inhibition of MMP-2 only.

CONCLUSIONS AND FUTURE STUDIES

It has become clear that MMPs have differing roles in both pathogenesis and recovery after SCI. How they influence injury and recovery processes is dependent on a number of factors that include when and where they are expressed and the profile of available substrates. As such, specific gelatinase inhibitors, targeting specific cellular and temporal profiles, may be more efficacious than their broad-spectrum counterparts. The potential to regulate neuropathic pain after SCI through gelatinase inhibition warrants detailed examination. Although studies of SCI support the view that early gelatinase activity is detrimental, in part, by promoting barrier dysfunction and early inflammation, involvement of MMPs in the more chronically injured cord is more complex. During wound healing, gelatinases modulate the formation of an inhibitory glial scar, support axonal regeneration, promote cell survival, and attenuate mechanical allodynia. Thus, the application of broad-spectrum MMP inhibitors in the more chronically injured cord should be approached with caution because any benefit may be outweighed by untoward effects.

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