



Published in final edited form as:

*Curr Drug Targets*. 2013 March 1; 14(3): 287–324.

## Matrix Metalloproteinases as Potential Targets in the Venous Dilation Associated with Varicose Veins

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### Abstract

Varicose veins (VVs) are a common venous disease of the lower extremity characterized by incompetent valves, venous reflux, and dilated and tortuous veins. If untreated, VVs could lead to venous thrombosis, thrombophlebitis and chronic venous leg ulcers. Various genetic, hormonal and environmental factors may lead to structural changes in the vein valves and make them incompetent, leading to venous reflux, increased venous pressure and vein wall dilation. Prolonged increases in venous pressure and vein wall tension are thought to increase the expression/activity of matrix metalloproteinases (MMPs). Members of the MMPs family include collagenases, gelatinases, stromelysins, matrilysins, membrane-type MMPs and others. MMPs are known to degrade various components of the extracellular matrix (ECM). MMPs may also affect the endothelium and vascular smooth muscle, causing changes in the vein relaxation and contraction mechanisms. ECs injury also triggers leukocyte infiltration, activation and inflammation, which lead to further vein wall damage. The vein wall dilation and valve dysfunction, and the MMP activation and superimposed inflammation and fibrosis would lead to progressive venous dilation and VVs formation. Surgical ablation is an effective treatment for VVs, but may be associated with high recurrence rate, and other less invasive approaches that target the cause of the disease are needed. MMP inhibitors including endogenous tissue inhibitors (TIMPs) and pharmacological inhibitors such as zinc chelators, doxycycline, batimastat and marimastat, have been used as diagnostic and therapeutic tools in cancer, autoimmune and cardiovascular disease. However, MMP inhibitors may have side effects especially on the musculoskeletal system. With the advent of new genetic and pharmacological tools, specific MMP inhibitors with fewer undesirable effects could be useful to retard the progression and prevent the recurrence of VVs.

### Keywords

MMP; endothelium; vascular smooth muscle; extracellular matrix; chronic venous insufficiency disease; TIMP

### INTRODUCTION

Chronic venous disease (CVD) is a common disorder of the lower extremity that depending on its severity could have different manifestations including varicose veins (VVs). VVs commonly affect the superficial veins of the lower extremity and manifest as abnormally dilated, twisted, and tortuous veins. If untreated VVs could lead to several complications including thrombophlebitis and chronic venous insufficiency (CVI) with venous leg ulcers.

VVs are characterized by incompetent valves, venous reflux and vein wall dilation. A primary valve dysfunction may cause significant reflux, increased venous hydrostatic pressure, chronic venous hypertension, and vein wall dilation. Also, a primary vein wall dilation to expand to neighboring valves causing valve distortion, dysfunction, and incompetence, and the resulting venous reflux leads to increased venous pressure and further vein wall dilation [1, 2].

Despite the major medical and socio-economical consequences of VVs, the pathophysiological mechanisms involved are not fully understood. Matrix metalloproteinases (MMPs) are proteolytic enzymes that have been identified in many tissues and organs including the venous system. MMPs play a major role in tissue remodeling and the continuous turnover of collagen, elastin and other proteins of the extracellular matrix (ECM), and have been implicated in cardiovascular remodeling and vascular disease. The last two decades have witnessed great advances in our understanding of the role of MMPs in the development and progression of VVs. Studies on venous tissue from experimental animals and human have shown marked changes in the expression/activity of various MMPs in association with vein wall remodeling. Also, studies on wound ulcer fluid environment have suggested possible correlation between the activity of MMPs and the development of skin lesions and venous leg ulcers. An imbalance between MMPs expression/activity and endogenous tissue inhibitors of MMPs (TIMPs) could cause pathological changes in the vein wall and valves and lead to CVD. However, the upstream mechanisms causing elevation of MMPs in VVs, and the downstream mechanism linking MMPs to vein wall dilation are not clearly understood. Several studies have shown that increased mechanical stretch or pressure in human tissues is associated with increased expression of MMPs [3–6], and increased venous hydrostatic pressure could be a primary cause of elevated MMPs levels in VVs. In addition to their proteolytic properties on ECM, MMPs may have early effects on other cellular components of the vein wall including ECs and vascular smooth muscle (VSM) [7–9]. Also, prolonged increases in venous hydrostatic pressure may cause EC injury and increase cell permeability, leading to leukocyte infiltration and vascular inflammation [5], which in turn lead to tissue fibrosis, wall resolution, valve degradation and irreversible vein damage characteristic of late stages of CVI.

In this review we will discuss reports published in the Pubmed database and experimental data from our laboratory to highlight the role of MMPs in VVs. The review will discuss VVs, the predisposing factors and the clinical and experimental evidence for a role of MMPs in VVs, thrombophlebitis and venous ulcers. We will describe how increases in venous pressure could lead to increased MMPs expression, and the potential transcription factors involved. We will also describe the effects of MMPs on ECM and the newly-discovered effects on ECs and VSM. The review will then discuss various strategies for management of VVs and their complications, and provide a perspective on new strategies and future directions to target the MMP pathway in order to retard the progression and prevent the recurrence of VVs.

## Varicose Veins

VVs are a common health and socioeconomical problem affecting over 25 million of the adult population in the United States [10]. The first national screening program in the United States identified VVs in 32% of participants, and many participants were considered at risk of developing venous thromboembolism during abdominal or orthopedic surgery [11].

VVs are superficial lower extremity veins that are abnormally twisted, dilated and often associated with incompetent valves (Fig. 1). If untreated, VVs could cause complications such as thrombophlebitis, deep venous thrombosis and venous leg ulcer. According to

CEAP (clinical-etiology-anatomy-pathophysiology) classification, CVD has seven clinical stages C0-6, with C0 indicating no visible sign of venous disease, C1 telangiectasies (spider veins), C2 VVs, C3 edema, C4a skin pigmentation or eczema, C4b lipodermatosclerosis or atrophie blanche, C5 healed ulcer, and C6 active ulcer. C4-6 are often designated as chronic venous insufficiency (CVI), reflecting the advanced stage of the disease [2, 12] (Table 1) (Fig. 2).

### Predisposing Factors for Varicose Veins

Several genetic and environmental factors have been associated with CVD including age, gender, pregnancy, estrogen therapy, obesity, family history of VVs, phlebitis and prior leg injury (Fig. 1). Environmental and behavioral factors associated with CVD include prolonged standing and sedentary lifestyle [13–15]. Family history and familial hereditary factors suggest a genetic component of VVs [16]. For instance, genetic mutations in iron metabolism genes may play a role in VVs. Prolonged venous reflux is associated with iron overload and dermal hemosiderin deposition that is directly correlated with clinical symptoms of CVI including skin changes and lipodermatosclerosis [17]. Iron deposition may induce the formation of free radical which could cause further tissue injury, and progression to advanced forms of CVI and leg ulcers [18, 19]. Also, Factor XIII is a cross-linking protein that plays a key role in ulcer healing [20]. Mutations in hemochromatosis C282Y (HFE) gene and Factor XIII V34L gene variants have been identified in patients with CVD and have been associated with increased risk of severe forms of CVI, skin changes and the size of venous ulcers [21, 22]. Interestingly, specific FXIII genotypes (L34 variant) have favorable ulcer healing rates, while HFE gene mutations may increase the risk of venous ulcer, but have no influence on healing time [23].

Some clinical conditions support a genetic component of VVs. Patients with Klippel-Trenaunay Syndrome have congenital venous anomalies in the form of atresia, agenesis of the deep venous system, valve insufficiency, venous aneurysms, and embryonic veins [24]. These patients have impaired venous muscle pump function and valve competence, and often present with VVs, limb hypertrophy, and dermal capillary hemangiomas (port wine stain) [25]. Primary lymphedema-distichiasis is a rare syndrome involving mutation in the FOXC2 gene, and one of its features is VVs at an early age, supporting a role of FOXC2 gene in the pathogenesis of VVs and a heritable element of the disease [26, 27]. A study of genealogical trees has shown that in nine families studied, VVs are linked to the candidate marker D16S520 on chromosome 16q24, which may account for the linkage to FOXC2 gene. Families of affected patients with the D16S520 marker have shown evidence of saphenofemoral junction reflux. The linkage to a candidate marker for the FOXC2 gene suggests that there is a functional variant within or in the vicinity of the gene that predisposes to VVs, and that CVI could be heritable in an autosomal dominant mode with incomplete penetrance [28]. Heterozygous mutation in the Notch3 gene has also been identified in the CASADIL (cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy) pedigree with VVs [29]. Microarray analysis of 3,063 human cDNAs from VVs and control veins have shown upregulation of 82 genes, particularly those regulating ECM, cytoskeletal proteins, and myofibroblasts production [30]. Ehlers-Danlos syndrome comprises more than ten connective tissue disorders with abnormal collagen synthesis, joint hypermobility, distensible skin, ocular disease, bone deformities and fragility, and cardiovascular disorders that render the blood vessels and visceral tissue walls fragile and susceptible to rupture. Patients with Ehlers-Danlos syndrome type IV are prone to vascular pathology and may present with VVs [31, 32]. Recently, single nucleotide polymorphisms in the promoter regions of MMP-9 genes have been identified among the Chinese population. In MMP-9, a 1562 C to T substitution has been associated with increased promoter activity and plasma levels of MMP-9, and

polymorphisms in the promoter region of MMP-9 are associated with VVs in the Chinese population [33].

In support of a genetic component of CVD, the elasticity of the lower limb vein wall is reduced not only in patients with venous insufficiency, but also in individuals with a high risk of developing VVs and in children of patients with VVs [34, 35]. Also, VVs pathology may not be confined to lower extremity veins, and could affect other tissues and cells in a generalized fashion. In patients with VVs, arm veins also show abnormal increase in distensibility, suggesting a systemic disease of the vein wall [36]. Both cultured dermal fibroblasts from dermal biopsies of patients with chronic vascular disease and cultured VSMCs from VVs show increased synthesis of type I collagen and decreased synthesis of type III collagen despite normal gene transcription, suggesting that VVs patients may have a systemic abnormality in collagen production and post-translational inhibition of type III collagen synthesis in various tissues [37, 38]. Also, in both VSMCs and fibroblasts of patients with VVs there is an increase in hydroxyproline content, indicating increased collagen, as compared to control. However the proportion of collagen type III was reduced despite normal mRNA transcription. These data may explain the loss of distensibility in VVs, and suggest a generalized defect in collagen metabolism and a genetic component of VVs pathology [39].

Age and gender are important factors in the development of VVs, and CVD may be more prevalent in females [2]. The Framingham Study has shown that the incidence of VVs is 2.6% in women and 1.9% in men [40]. The Edinburgh Vein Study screened 1566 subjects 18–64 years old for CVD and found that women were more likely to report VVs-related leg symptoms [41]. However, in a follow-up study the age-adjusted prevalence of truncal VVs was 40% in males and 32% in females, and the prevalence of VVs and CVD increased with age [42]. Also, studies measuring vein reflux using duplex ultrasound found CVD in 9.4% of men and 6.6% of women, which rose with age (21.2% in men and 12.0% in women older than 50) [43]. Interestingly,  $\alpha$ -adrenergic, AngII-, depolarization-induced, and  $[Ca^{2+}]$ -dependent contraction are reduced in female compared with male rat inferior vena cava (IVC), possibly due to increased estrogen receptor expression/activity and enhanced endothelium-dependent relaxation pathways in females. These data suggested inherent sex differences in venous tissue function, whereby an enhanced estrogen receptor-mediated venous relaxation and decreased venous contraction would lead to more distended veins in females [44].

Pregnancy is associated with physiologic changes that could contribute to venous distension and the development of VVs. During pregnancy plasma levels of estrogen and progesterone increase [45]. Early in pregnancy there is an increase in blood volume and plasma volume expansion [46]. Also, fetal growth and weight gain increase intra-abdominal pressure and central venous return [47, 48], and the increased venous pressure could lead to vein valve failure and progression of varices. Studies have also shown that overweight and obese women are more likely to develop VVs [49]. Compared with non-overweight women, moderately overweight women (BMI = 25.0–29.9 kg/m<sup>2</sup>) were more likely to report VVs, and obese women (BMI  $\geq$  30.0 kg/m<sup>2</sup>) were three times more likely to report the presence of VVs. However, no relationship between BMI and venous disease was observed among males [50]. Importantly, overweight and obese women have greater plasma levels of total and bioavailable circulating estrogens than non-overweight women particularly after menopause [51], further supporting a relationship between estrogen levels and VVs formation.

The ergonomics and physical activity of an occupation may be a factor in the epidemiology of VVs. In a community-based study of VVs conducted in Jerusalem among men and

women aged 20 to 64 years, the prevalence of VVs was higher among individuals who spent much of their work day standing. Women were more likely to report occupations requiring prolonged standing compared with men (31.4% vs. 13.6%). However, the prevalence ratio related to workplace posture (standing vs. sitting) was higher in men (1.88) than in women (1.53) [52].

### MMPs and Varicose Veins

MMPs may play a role in the vascular remodeling associated with vascular disease including VVs [53]. MMPs are endopeptidases discovered in 1962 as a collagen proteolytic activity during the ECM protein degradation associated with resorption of the tadpole tail [54]. Since then, the MMP family has grown to include at least 28 members in vertebrates, 23 in humans, and 14 in blood vessels. MMPs, also called matrixins, are multidomain zinc ( $Zn^{2+}$ ) metalloproteinases that degrade various components of ECM and belong to the larger superfamily of proteases called metzincins, which also includes adamalysins, serralysins, and astacins. Typically MMPs consist of a propeptide of about 80 amino acids, a catalytic metalloproteinase domain of about 170 amino acids, a linker peptide (hinge region) of variable lengths and a hemopexin domain of about 200 amino acids [55, 56] (Fig. 3). Members of the MMP family have 3 distinguishing features: 1) Sequence homology with collagenase-1 (MMP-1); 2) Cysteine switch motif PRCGXPD in the prodomain that maintains MMPs in the proMMP zymogen form, and chelates the active  $Zn^{2+}$  site, except MMP-23 which lacks the cysteine switch motif; and 3)  $Zn^{2+}$ -binding motif bound by 3 histidine molecules with the conserved sequence HEXGHXXGXXH located in the catalytic domain. MMPs are commonly classified based on domain organization and substrate preference into collagenases, gelatinases, stromelysins, matrilysins, membrane-type (MT)-MMPs and others [55, 57] (Table 2).

Collagenases include MMP-1, -2 (neutrophil collagenase), -13 and -18 (*Xenopus*). These MMPs cleave fibrillar collagen type I, II and III into characteristic 3/4 and 1/4 fragments. They first unwind triple helical collagen then hydrolyze the peptide bonds. The MMPs hemopexin domain is essential for cleaving native fibrillar collagen while the catalytic domain is needed for cleaving noncollagen substrates [58, 59]. MMP-13 (collagenase 3) is overexpressed in cartilage tissues of osteoarthritis patients and is very efficient in degrading type II collagen [60].

Gelatinases include gelatinase A (MMP-2) and gelatinase B (MMP-9). Gelatinases digest denatured collagens (gelatins), and have 3 type II fibronectin repeats inserted in the catalytic domain, which bind to gelatin, collagens and laminin [61]. MMP-2 cleaves collagen in two phases, the first resembling that of the interstitial collagenases, followed by gelatinolysis, which is promoted by the fibronectin-like domain [62, 63]. The collagenolytic activity of MMP-2 is much weaker than collagenases. However, because proMMP-2 is recruited to the cell surface and activated by membrane-bound MT-MMPs, it may accumulate pericellularly and induce substantial localized collagenolytic activity [64].

Stromelysins 1, 2 and 3, also known as MMP-3, -10, and -11, respectively, have the same domain arrangement as collagenases, but do not cleave interstitial collagen. MMP-3 and -10 are similar in structure and substrate specificity, while MMP-11 is distantly related. Despite their similar substrate specificity, MMP-3 has higher proteolytic efficiency than MMP-10. MMP-3 and MMP-10 digest a number of ECM components and participate in proMMP activation, but MMP-11 has very weak activity toward ECM molecules. Also, MMP-3 and -10 are secreted from the cells as inactive proMMP, but MMP-11 is activated intracellularly by furin and secreted from the cells as an active enzyme [65].



Matrilysins, include matrilysin-1 (MMP-7) and matrilysin-2 (MMP-26, endometase), which lack the linker peptide or hinge region and the hemopexin domain [66, 67]. MMP-7 acts intracellularly in the intestine to process procryptidins to bactericidal forms. MMP-7 degrades ECM components, and cleaves cell surface molecules such as Fas–ligand, pro-tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), syndecan 1 and E-cadherin to generate soluble forms [68]. MMP-26 is expressed in breast cancer cells [69].

Membrane-Type MMPs (MT-MMPs) include 4 transmembrane MMPs, MT1-, MT2-, MT3- and MT5-MMP (MMP-14, -15, -16 and -24, respectively), and the glycosylphosphatidylinositol-anchored proteins MT4- and MT6-MMP (MMP-17 and -25, respectively). MT-MMPs have a furin-like pro-protein convertase recognition sequence at the C-terminus of the propeptide. They are activated intracellularly and the active enzymes are expressed on the cell surface. All MT-MMPs except MT4-MMP (MMP-17) can activate proMMP-2 [70]. MT1-MMP digests collagen-I, -II and -III and other ECM proteins, and in the presence of TIMP-2 it activates proMMPs such as proMMP-13 on the cell surface [71, 72] (Table 2).

Other MMPs include MMP-12, -20 and -27 which have a domain arrangement and chromosome location similar to stromelysins. MMP-12 (metalloelastase) is expressed in macrophages and is essential for macrophage migration [73] and is also found in hypertrophic chondrocytes and osteoclasts [74, 75]. MMP-12 digests elastin and other ECM proteins. MMP-19 is a potent basement membrane-degrading enzyme that plays a role in tissue remodeling, wound healing and epithelial cell migration by cleaving laminin5- $\gamma$ 2 chain [76–79]. MMP-19 deficient mice develop diet-induced obesity due to adipocyte hypertrophy, but are less susceptible to skin cancers induced by chemical carcinogens [80].

Enamelysin (MMP-20) is a tooth-specific MMP expressed in newly formed tooth enamel and digests amelogenin [81]. Amelogenin imperfecta, a genetic disorder with defective enamel formation involves mutation at MMP-20 cleavage sites [82]. MMP-21 is an MMP with measurable gelatinolytic activity expressed in various fetal and adult tissues, macrophages of granulomatous skin lesions, fibroblasts in dermatofibromas, and in basal and squamous cell carcinomas [83, 84]. MMP-22 was cloned first from chicken fibroblasts, and a human homologue was later identified, but its function and substrate are unclear [85].

MMP-23 is a type II membrane protein regulated by a single proteolytic cleavage for both its activation and secretion [86]. It harbors a transmembrane domain and a furin recognition motif (convertase cleavage site which activates MMP) in the propeptide, and is therefore cleaved in the Golgi and released as an active enzyme into the extracellular space [86]. MMP-23 is unique among the matrixins as it lacks the cysteine switch motif in the propeptide and the linker peptide or hinge region, and the hemopexin domain is substituted by cysteine-rich immunoglobulin-like domains immediately after the C-terminus of the catalytic domain [64, 71, 87, 88] (Fig. 3). MMP-23 is expressed predominantly in ovary, testis and prostate, suggesting a specialized role in reproduction [87]. MMP-27 is expressed in B-lymphocytes and is overexpressed in cultured human lymphocytes treated with anti-(IgG/IgM) [89]. Epilysin (MMP-28) is one of the latest MMPs to be identified [90]. Epilysin was first cloned from human keratinocyte and testis cDNA libraries, and is expressed in the lung, placenta, heart, gastrointestinal tract and testis [91, 92]. MMP-28 is increased in cartilage from patients with osteoarthritis and rheumatoid arthritis [93, 94].

Studies have shown increased MMPs in lower extremity venous blood of patients with VVs. The plasma and venous tissue levels of MMP-1, -2, -3, -9 and -13 are elevated in VVs [95–97]. In a study examining MMP-1, -3 and -13 in proximal and distal segments of VVs vs. control veins MMP-1 and -13 mRNA was not different in VVs vs. control veins or in

proximal vs. distal segments of VVs. MMP-3 mRNA was not amplified in any of the vein segments studied. However, when protein levels were measured, MMP-1 was elevated in VVs compared to control veins, and MMP-1 and -13 was increased in proximal vs. distal VVs segments. These findings suggested that MMPs protein levels are increased in VVs, and that their post-transcriptional modification may explain their differential distribution in VVs [95]. Another study has linked the changes in the thickness of VVs to the balance between MMPs and TIMPs and showed greater expression of TIMP-2 and connective tissue accumulation in tunica media of VVs compared with arm and neck veins of control subjects. TIMP-2 and -3 expression was greater in hypertrophic than atrophic segments, and in the thicker proximal than distal segments of VVs [98]. Immunohistochemical studies have shown variable distribution of MMPs in the venous tissue intima, media and adventitia. VVs show increased distribution of MMP-1 in all layers, while in normal veins MMP-1 is localized in the endothelium and adventitia. MMP-9 is expressed throughout the vein wall in both control and VVs, with increased levels in the VSM layer in VVs. TIMPs were not detected in any of the veins examined [97, 99]. These findings suggest that MMPs may affect all layers of the vein wall including the ECM leading to vein wall degradation and VVs formation [97]. Other studies have localized MMP-9 immunostaining in VSMCs of VVs but not control veins [99]. While MMPs may contribute to the pathophysiology of VVs, the presence of MMPs in VVs does not imply causation, and the diverse location of different MMPs in various layers of the vein wall including ECM, ECs and VSM suggest that MMPs may have different effects at different stages of VVs.

### **MMPs in Thrombophlebitis**

One of the complications of VVs is thrombophlebitis characterized by endothelial, vein wall and valve inflammation and leukocytes infiltration leading to disruption of vein function and venous thrombosis [2, 100, 101]. Thrombophlebitis may occur in the course of VVs or may be induced during treatment of VVs with sclerosing agents [101, 102]. Specimens of saphenous vein from patients with CVD have shown increased monocytes/macrophage infiltration in the vein wall and valves [100, 103]. Also, intercellular adhesion molecule-1 (ICAM-1) was elevated in CVD specimens, but other cytokines were not [101].

VVs with thrombophlebitis may have different MMPs expression compared with VVs. In a study examining MMP-1, -2, -3 and -9 in control veins, VVs and VVs complicated by thrombophlebitis, thrombophlebitic VVs showed an elevated content of MMPs in the vein wall, and increased MMP-1, -2 and -9 activity. VVs showed increased activity of MMP-2. These marked changes in MMPs content and activity in VVs especially those affected with thrombophlebitis could lead venous tissue remodeling and alterations in the mechanical properties of the vein wall [96]. Other studies have shown a higher count of mast cells, T cells and B cells in thrombotic compared with non-thrombotic VVs [104], further implicating inflammation in this stage of the disease.

### **MMPs in Lipodermatosclerotic Skin and venous Leg Ulcer**

Some of the features of advanced stages of CVD are skin changes, lipodermatosclerotic skin and venous leg ulcer. Lipodermatosclerotic skin shows increased MMP activity and ECM turnover. Dermal biopsies from lipodermatosclerotic skin showed increased mRNA expression and protein levels of MMP-1 and -2 and TIMP-1, and increased activity of MMP-2 as compared to healthy skin. In addition, there was an increase in proMMP-1:TIMP-1 complex indicating the overexpression of proteinases and enhanced binding to TIMP [105]. Immunohistochemistry experiments revealed prominent expression of MMP-1 and -2 in the basal and suprabasal layers of the epidermis, perivascular region and reticular dermis, and reduced expression of TIMP-2 in the basement membrane of

lipodermatosclerotic skin [105], supporting excessive and unrestrained MMP activity and ECM turnover.

Analysis of the components of venous leg ulcer microenvironment has shown dermal fibroblasts, keratinocytes, inflammatory cells, ECM, growth factors, cytokines and bacteria, and circulating ECs in the microcirculation. Also, the chronic venous ulcer wound fluid (VUWF) shows marked protease activity, with increased collagenase activity 116-fold over that in acute wound fluid, and decreased activity in venous ulcer that shows healing [106–108].

In both acute and chronic wounds, ECM provides a milieu for keratinocytes to migrate [109], and changes in protease activity could affect the ECM properties in wounds. Chronic VUWF has up to 10-fold increase in the levels of MMP-2 and -9 and increased MMP-1 and gelatinase activity as compared with acute wound fluid, suggesting increased ECM turnover [110, 111]. Inhibition studies with the MMP inhibitor doxycycline suggested that the source of collagenase and gelatinase activity in VUWF was from fibroblasts and mononuclear cells, but not neutrophils, which are involved in acute wound healing [111]. Studying the source of collagenase activity is important particularly because bacteria, a component of the venous ulcer microenvironment, produce collagenase. Human collagenase degrades collagen in a specific 3/4 and 1/4 fragments, whereas bacterial collagenase degrades collagen randomly in a non-specific manner. Collagenase from VUWF degrades collagen in the specific 3/4 and 1/4 fragments, supporting the presence of human collagenase activity [111]. MMP-1 is also found in migrating keratinocytes of the wound [112]. MMP-12 is abundant in fibroblasts in the ulcer bed, but not in the epidermis or in acute wounds. Increased expression of MMP-1, -3 and -13 with concomitant reduction in gene expression and immunoreactivity of TIMP-1 and -2 was also observed in the acute phase of skin ulcer with inflammation and dermatitis [113].

MMPs in wound ulcer could also cause abnormalities in tissue perfusion or affect angiogenesis and the microvasculature. When the effects of VUWF and acute wound fluid from donor skin graft sites were tested on ECs culture model of angiogenesis and tubule formation, VUWF caused marked reduction in the formation and length of tubules as compared with control fluid. Addition of the synthetic MMP-2/MMP-9 Inhibitor I to VUWF restored angiogenesis [114], suggesting that MMPs in VUWF have anti-angiogenic effects that disrupt the microcirculation in the perivascular regions and inhibit wound healing.

The regulation of MMP production in lipodermatosclerotic tissue and venous ulcer involves several factors. Dermal fibroblasts and leukocytes are major sources for MMPs especially MMP-2 [115]. MMP production in lipodermatosclerotic tissue and venous ulcer may involve mitogen-activated protein kinase (MAPK) which regulates MMP expression and proteolytic activity in dermal fibroblasts [116, 117]. In fibroblasts, TNF- $\alpha$  induces MMP-19 expression, which is inhibited by blocking the ERK<sub>1/2</sub> pathway with PD98059 or p38 MAPK pathway with SB203580. Adenovirus-mediated induction of ERK<sub>1/2</sub> and p38 MAPK in fibroblasts increases MMP-19 expression. Also, activation of c-JNK increases proMMP-19, highlighting the role of MAPK in regulating MMP expression and proteolytic activity in dermal fibroblasts, with potential implications in the pathogenesis of venous ulcer [116–118]. Post-translational modifications of MMPs are also essential for their activity and are regulated by TGF- $\beta$ 1 [115].

MMPs are thought to regulate not only the development of skin changes and venous ulcer but also wound healing. An important element in wound healing is Factor XIII (FXIII) which promotes collagen cross-linking, and thereby modulates the effects of MMPs. *In vitro* studies evaluated the effects of increasing concentrations of collagenase and FXIII on



fibroblast survival. At high collagenase concentrations (2 mg/mL) 95% of fibroblasts were nonviable, and FXIII did not inhibit the effects of collagenase. However, at lower collagenase concentrations (0.5–1 mg/mL) FXIII abrogated the effects of collagenase and increased fibroblasts survival. Interestingly, topical application of FXIII improved venous ulcer healing [20]. In addition to FXIII, iron overload has been found in the serum and dermis of limbs of patients with venous ulcer compared with control subjects. Increased MMP-9 activity was also observed in venous ulcer. Iron overload can cause oxidative stress and increased production of reactive oxygen species (ROS). The iron deposits in the limbs could be released in the serum causing overload and activation of MMPs and ROS and lead to impaired ulcer healing [18].

MT1-MMP and extracellular MMP inducer (EMMPRIN, CD147) are important inducers/activators of MMPs [55, 56, 119]. Immunohistochemistry studies in venous ulcer biopsies have shown increased expression of MMP-2, MT1-MMP, MT2-MMP and EMMPRIN in venous ulcer dermis compared to control dermis. MMP-2 and EMMPRIN were overexpressed in the perivascular regions, which would favor unrestrained MMP activation and ECM turnover in the perivascular region of the venous ulcer [119]. Comparison of healing and non-healing venous ulcers has shown greater levels of PDGF AA in healing ulcers, but no difference in MMPs or EMMPRIN. Also, VUWF of healing ulcers had elevated levels of PDGF AA and TIMP-2 and low levels of MMP-2 as compared with non-healing ulcer, supporting that increased protease activity favors a non-healing environment [120]. Decreased MMPs levels favor enhanced ulcer wound healing, and compression treatment for 4 weeks was associated with reduced levels of MMP-1, -2 and -3 and greater rate of venous ulcer healing [121].

Wound healing involves an orderly process of inflammation, re-epithelisation, matrix deposition and tissue remodeling and the expression of MMPs and TIMPs varies in different CEAP stages of CVI [115]. Regulation of MMPs profile in CVI occurs at gene expression, transcriptional and post-transcriptional levels [115]. Biopsies from patients with stasis dermatitis (C4a) show upregulation of MMP-1, 2 and 13, but downregulation of TIMP-1 and TIMP-2 as compared with healthy controls [113]. The distribution of MMP/TIMP in acute wounds and chronic venous ulcer also varies. MMP-9 and MMP-13 are localized in the chronic venous ulcer bed [112], MMP-1 is localized in migrating cells in chronic venous ulcer [112], and TIMP-1 and TIMP-3 are localized in proliferating cells in the edges of acute wounds [122], suggesting that MMP-1, TIMP-1 and TIMP-3 are vital for re-epithelialization, while MMP-9 and MMP-13 may be involved in the remodeling of collagenous matrix in chronic wounds.

Although MMPs may be increased in VVs, the upstream mechanisms causing upregulation of MMPs and the downstream mechanisms linking the increased MMPs to the structural and functional changes in the various components of the vein wall are not clearly established and will be discussed in detail below.

### **Increased Venous Hydrostatic Pressure, MMPs Expression, and Vein Wall Dilation**

Veins are capacitance vessels that play a major role in determining the venous return and preload. Although the vein wall is relatively thin, it has three layers; the intima with lining ECs, media with several layers of VSM and adventitia which contains fibroblasts embedded in ECM of various proteins including collagen and elastin. The lower extremity has an intricate network of superficial and deep veins connected by perforator veins that pass through fascial spaces [123] (Fig. 2). Blood flows from the superficial to deep veins, except in the foot where the flow is reversed. The superficial veins include small saphenous vein, which starts from the ankle, runs in the posterior of the leg and joins the popliteal vein at the saphenopopliteal junction. The great saphenous vein starts from the ankle in the medial side

of the lower limb and joins the common femoral vein at the saphenofemoral junction. Deep leg veins include common femoral, deep femoral, femoral, popliteal and tibial veins. In the upright posture, the blood in the lower extremity veins travels against gravity and the fluctuating thoracoabdominal pressure towards the central circulation. The saphenous veins and their tributaries are subjected to high hydrostatic pressure. In all individuals in the standing position the column of blood in the lower extremity venous system reflects a venous pressure at the ankle of 90–100 mmHg [124, 125], causing marked increases in vein wall tension. To cope with the high hydrostatic pressure, the leg superficial and deep veins are equipped with bicuspid valves that maintain blood flow in the cephalic direction and prevent its return toward the feet (Fig. 2). Vein valves function in concert with muscles mainly in the calf but also in the foot and thigh, to pump blood against gravity towards the heart. Leg veins also cope with the high hydrostatic pressure by compensatory changes in the vein wall function/structure. Persistent increases in venous pressure could lead to alterations in the vein wall structure and venous tissue remodeling.

Studies have shown that increased mechanical stretch or pressure in human tissues is associated with increased expression of MMPs not only in fibroblasts but also in ECs and VSMCs [3–6]. These observations have suggested that increases in venous pressure or wall tension may increase MMPs expression/activity, and that MMPs may then affect different wall components and cell types including ECM, fibroblasts, VSMCs and ECs. In later stages of the disease, severe increases in venous hydrostatic pressure cause EC damage, leukocyte infiltration, and superimposed venous inflammation characteristic of advanced stages of CVI.

MMPs appear to be induced by postural changes in patients with VVs. Comparison of plasma from the brachial vein and lower extremity VVs in patients in the standing position following 30 minutes of stasis have shown an increase in proMMPs such as proMMP-9 in plasma from VVs compared to arm vein. The increased proteolytic activity was associated with increased plasma levels of endothelial and leukocyte activation markers including ICAM-1, vascular cell adhesion molecule-1 (VCAM-1), angiotensin converting enzyme, and L-selectin levels of suggesting EC and polymorphonuclear cell activation and enzymatic granule release in VVs during periods of postural blood stasis [126, 127]. Importantly, upregulation of these markers in VVs was inversely proportional to the PO<sub>2</sub> of the blood sampled [127]. These observations support a role of MMPs as important proteolytic enzymes and in the interaction between leukocytes and ECs in VVs, and a potential link to blood PO<sub>2</sub> level.

Studies in animal models have further examined the relationship between increased venous pressure and MMP expression in the vein wall [128]. In a rat model of acute venous hypertension produced by creating a femoral arterio-venous fistula and examined at three weeks, most rats had venous reflux and increased venous pressure in the ipsilateral compared to the contralateral control femoral vein. Also, the pressurized veins were dilated and the valve leaflets length and width were reduced. Long term, the pressurized veins demonstrated marked inflammatory changes and leukocyte infiltration with increased expression of P-selectin and ICAM-1. There were no differences in MMP-2 or MMP-9 at three weeks, and the number of apoptotic cells in the vein wall and valves was increased [129]. In a follow-up study evaluating the effects of chronic venous hypertension there was an increased pressure in the femoral vein and progressive reflux at 42 days post arterio-venous fistula induction. The valves distal to the fistula demonstrated increased diameter, decreased height, and valve fibrosis in the media and adventitia. Valve obliteration and elevated MMP-2 and -9 levels were observed after 21 and 42 days of venous hypertension [130]. The presence of MMP-2 and -9 in the vein wall with valve destruction supports a link between MMPs and VVs formation. However, in this model only proximal segments of

veins were analyzed, and whether the venous changes are caused by venous hypertension, venous arterialization, or a combination needs to be examined in order to evaluate if these venous abnormalities are transmitted to distal vein segments as observed in human CVD. Also, an increase in apoptosis in this model is not a feature of human VVs, highlighting the differences between the pathological changes in VVs as compared to those resulting from adaptation to an AV fistula.

Increased venous hydrostatic pressure may induce ECs permeability/injury, leukocyte infiltration and attachment, and initiate inflammation [5] (Fig. 1). Leukocyte wall infiltration and inflammation in turn activate MMPs and lead to ECM degradation, vein wall weakening and wall/valve fibrosis. Persistent venous wall dilation and valve dysfunction lead to further increases in hydrostatic pressure and CVD. The early stages of CVD are contained within the vasculature leading to clinical sign of VVs, while more advanced CVD causes progression to CVI affecting surrounding tissues and leading to skin changes and venous leg ulcer.

Experiments on isolated rat inferior vena cava (IVC) have shown that prolonged increases in vein wall tension are associated with decreased vein contraction and increased amount of MMP-2 and -9. The decreases in vein contraction were prevented in veins pretreated with MMP inhibitors supporting a role of MMPs as a link between increased venous pressure/wall tension, decreased vein contraction and increased venous dilation [8].

The observations that increased mechanical stretch or pressure is associated with increases MMPs expression in fibroblasts, ECs, and smooth muscle cells [3–6], have suggested that increased venous hydrostatic pressure could be a primary cause of elevated MMPs levels in VVs. In search for the upstream mechanisms linking the increases in vein wall tension to the changes in MMP expression and reduction in venous contraction, studies have pointed to a potential role of the hypoxia inducible factors HIF-1 $\alpha$  and HIF-2 $\alpha$ . HIFs are nuclear transcriptional factors which regulate genes involved in oxygen homeostasis [131]. The role of HIF as a potential factor linking increased venous pressure and MMP expression in VVs has been supported by reports demonstrating upregulation of HIF-1 $\alpha$  and HIF-2 $\alpha$  transcription factors, and HIF target genes in VVs compared with non-VVs. Also, exposure of VVs and non-VVs to hypoxic conditions was associated with increased amount of HIF-1 $\alpha$  and HIF-2 $\alpha$  protein and HIF target genes. These findings suggest that the HIF pathway may be associated with several pathological changes in the VVs wall, and that hypoxia may contribute to the pathogenesis of VVs [132]. In addition to the regulation of HIF by oxygen tension, other factors such as hormones, cytokines and metallic ions as well as mechanical stretch may induce HIF expression [131, 133]. Increased expression of HIF in response to mechanical stretch has been demonstrated in several tissues and cell types including the myocardium [134], fibroblasts [135], VSMCs [136], and skeletal muscle fibres [4, 137]. In support of transcriptional regulation of HIF-1 $\alpha$  by mechanical stress, studies have shown upregulation of HIF-1 $\alpha$  mRNA by ~ two-fold in VSMCs subjected to cyclic stretch for 4 hr [136]. Other studies have shown increased HIF-1 $\alpha$  and HIF-2 $\alpha$  mRNA and proteins in rat capillary ECs of skeletal muscle fibers exposed to prolonged mechanical stretch [137]. The protein amount and activity of HIF-1 $\alpha$  and HIF-2 $\alpha$  may also be regulated by mechanical stretch [136–138]. Studies have shown an increase of HIF-1 $\alpha$  protein in response to increased mechanical stress of the left ventricular wall by induction of aortocaval shunt or intraventricular balloon expansion [134]. Similar increases in HIF-1 $\alpha$  protein have been shown in fibroblasts that are cyclically stretched for 24 hr [135]. Other studies have shown that the expression and activity of MMP-2 and MMP-9 can be regulated by HIF [139, 140]. Also, prolonged increases in rat IVC wall tension are associated with reduced vein contraction, and the reduction in vein contraction was reversed by the HIF inhibitors U0126 and echinomycin, and enhanced in the presence of the HIF stabilizer

dimethylallyl glycine (DMOG) [127]. Prolonged vein wall stretch was also associated with increased expression of HIF-1 $\alpha$  and -2 $\alpha$  and MMP-2 and -9, and the increased expression of MMP-2 and -9 was reversed by HIF inhibitors. These findings suggest that prolonged increases in vein wall tension are associated with overexpression of HIF-1 $\alpha$  and -2 $\alpha$ , increased MMP-2 and -9 expression, and reduced contraction in rat IVC. These data are consistent with the view that increased vein wall tension secondary to venous hypertension may induce HIF overexpression and cause an increase in MMP expression and reduction of venous contraction, leading to progressive venous dilation and VVs formation [127].

The mechanism of HIF regulation by mechanical stretch is unclear, but may involve PI<sub>3</sub>K and MAPK [133, 136, 137]. Cell membrane ion channels, integrins, and receptor tyrosine kinases are mechano-sensitive to stretch [141]. Mechanical stretch may stimulate PI<sub>3</sub>K by activating Ca<sup>2+</sup> influx through transient receptor potential ion channels such as TRPV4 [142]. Also, integrins may transduce mechanical stretch to initiate signaling cascades and MAPK activation [143]. Receptor tyrosine kinases and G protein-coupled receptors are also stimulated by biomechanical stress with subsequent activation of MAPK [144]. Mechanical stretch may also increase the generation of ROS which in turn activate MAPK [145, 146]. Interestingly, the increased HIF-1 $\alpha$  and HIF-2 $\alpha$  mRNA expression and the reduction in venous contraction associated with prolonged vein wall stretch were reversed in veins treated with MAPK inhibitors, supporting a role of MAPK in the regulation of HIF by mechanical stretch. As MMPs are upregulated in response to increases in venous pressure and HIFs and in VVs, an important question is how MMPs are activated and what are their potential substrates.

### MMP Cleavage, Activation and Potential Substrates

MMPs are highly homologous Zn<sup>2+</sup>-dependent endopeptidases that cleave most of the components of ECM. The basic MMP structure consists of a prodomain, catalytic domain, hinge region, and hemopexin domain [55, 56] (Fig. 3). The catalytic domain contains the Zn<sup>2+</sup> binding motif HEXXHXXGXXH and a conserved methionine, forming a 'Met-turn' 8-residues downstream, which supports the active site cleft structure around the catalytic Zn<sup>2+</sup> [147].

Matrixins are synthesized as pre-proenzymes and the signal peptide is removed during translation to generate proMMPs. ProMMPs have a 'cysteine switch' motif PRCGXPD in which the cysteine residue coordinates with the catalytic Zn<sup>2+</sup> in the catalytic domain, keeping the proMMP in an inactive form [148]. The cysteine switch interacts with the Zn<sup>2+</sup> active binding site, and thereby prevents MMP activation and substrate degradation. MMPs may function through one of three catalytic mechanisms. The first mechanism called the base-catalysis mechanism is carried out by the conserved glutamate residue and Zn<sup>2+</sup> [149]. In the second mechanism, the catalytic action involves an interaction between a water molecule and Zn<sup>2+</sup> during the acid-base catalysis [150]. In the third mechanism, a histidine from the HEXXHXXGXXH-motif participates in catalysis by dissociation of Zn<sup>2+</sup> from it, thus allowing the Zn<sup>2+</sup> ion to assume a quasi-penta coordinated state. In this state, the Zn<sup>2+</sup> ion is coordinated with the two oxygen atoms from the catalytic glutamic acid, the substrate's carbonyl oxygen atom, and the two histidine residues, and can polarize the glutamic acid's oxygen atom, proximate the scissile bond, and induce it to act as reversible electron donor. This forms an oxy-anion transition state. At this stage, a water molecule acts on the dissociated scissile bond and completes the hydrolysis of the substrate [151]. Collectively, upon binding of the substrate, the Zn<sup>2+</sup>-bound water molecule attacks the substrate carbonyl carbon, and the transfer of protons through a conserved glutamine residue to the amide nitrogen of the scissile bond results in peptide cleavage [152, 153]. The hemopexin domain confers much of the substrate specificity for MMPs. For instance, the hemopexin domain of MMP-1 collagenase is essential for the specificity of the catalytic

cleavage of collagen. Also, MMP-2 is localized to specific extracellular collagenous sites by its fibronectin domains and MT1-MMP (MMP-14) requires the hemopexin domain for cell surface clustering as part of its collagenolytic capacity and ability to activate proMMP-2. The hemopexin domain also determines MT1-MMP binding and shedding of CD44 [154]. Activation of MMPs involves cleavage of the cysteine switch and detachment of the hemopexin domain, and may require other MMPs or other proteinases. For example, MMP-3 activates proMMP-1 into fully active MMP-1 [155]. ProMMP-2 activation takes place on the cell surface by most MT-MMPs, but not MT4-MMP [156], a process that may require TIMP-2 [157, 158]. ProMMP-2 forms a complex with TIMP-2 via their C-terminal domains, thus permitting the N-terminal inhibitory domain of TIMP-2 to bind to MT1-MMP on the cell surface. The cell surface-bound proMMP-2 is then activated by an MT1-MMP that is free of TIMP-2. MT1-MMP inhibited by TIMP-2 can also act as a “receptor” for proMMP-2. The MT1-MMP-TIMP-2-proMMP-2 complex is then presented to an adjacent free MT1-MMP for activation [159]. Thus, TIMP-2 may determine the MT1-MMP choice between direct cleavage of its own substrates and activation of MMP-2 [160]. However, for a number of MMPs including membrane-bound MMP-11, -23, -28, activation occurs intracellularly via the endopeptidase furin, which selectively cleaves paired base residues [65, 87, 90, 161].

Oxidants generated by leukocytes or other cells can both activate (via oxidation of the prodomain thiol followed by autolytic cleavage) and inactivate MMPs (via modification of amino acids critical for catalytic activity), providing a mechanism to control bursts of proteolytic activity. Several proMMPs are activated by ROS *in vitro* [162–165]. Foam cell derived ROS can activate proMMP-2. Nitric oxide (NO) may also activate proMMP-9 during cerebral ischemia by reacting with the thiol group of the cysteine switch and forming an S-nitrosylated derivative [165]. Also, hypoxia may increase MMP-2 and -9 mRNA expression [166]. Other MMPs such as MMP-9 depend predominantly on plasmin for activation [167]. MMP-7 is activated both by MMP-3 and by hypochlorous acid, a product of myeloperoxidase found in plaque macrophages. MMP-7 can activate MMP-1 [162, 168]. Also, serine proteinases such as neutrophil elastase may favor matrix breakdown by inactivating TIMPs [169, 170].

MMPs can be activated by heat, low pH, thiol-modifying agents such as 4-aminophenylmercuric acetate, mercury chloride, and N-ethylmaleimide, oxidized glutathione, sodium dodecyl sulfate, and chaotropic agents by causing disturbance of the cysteine-Zn<sup>2+</sup> interaction at the cysteine switch [171].

Collagen type I, II, III, IV, V, VI, VII, VIII, IX, X, and XIV are known substrates of MMPs, with different efficacies (Table 2). MMP substrates also include other ECM proteins such as fibronectin, vitronectin, laminin, entactin, tenascin, aggrecan, myelin basic protein. Certain forms of ECM proteins may require the cooperative effects of several MMPs to accomplish their complete degradation. MMP-1 and -8 degrade fibrillar helices into fragments and unfold their triple helix conformation. The so-formed single  $\alpha$ -chain gelatins are then further degraded by the gelatinases MMP-2 and -9 into oligopeptides [63].

Casein and gelatin are the most common substrates used to study MMP activity. Gelatin is a valid substrate particularly for MMP-2 and -9. While casein is not a physiologically relevant MMP substrate, it is a generic proteinase substrate digested by a wide range of proteinases.

### Biological Effects of MMPs

MMPs affect different receptors and pathways such that the overall effects of MMPs vary depending on the predominant receptors or pathways in the tissue examined. Also, individual MMPs vary in their proteolytic activity and tissue substrates, further contributing



to the discrepancy in the effects of MMPs [172]. MMPs play a role in many biological processes including tissue remodeling and growth, wound healing and angiogenesis, cell proliferation, migration, differentiation and apoptosis, as well as tissue defense mechanisms and immune responses [173, 174]. Defensins, a family of polar antimicrobial peptides that contribute to the innate immune system of some animals, are synthesized in an inactive proform that are activated by the proteolytic removal of the pro-domain by MMP-7, thus allowing them to insert into the bacterial membrane and disrupt its integrity [175, 176]. MMP-3 and -7 can also cleave all IgG proteins, an important process that prevents the initiation of the complement cascade and helps in the removal of IgG from damaged or inflamed tissue [177]. Also, the receptor of the complement component C1q (C1qR) exists in both a membrane-bound form and a soluble form that inhibits the hemolytic activity of C1q. MT1-MMP releases the membrane bound C1qR, thus allows tumor cells to avoid targeted destruction by the complement system and thereby facilitates tumor-cell survival [178–180]. MMPs also modulate many bioactive molecules at the cell surface [181], and may regulate cell signaling by interaction with G-protein-coupled receptors [182]. Changes in MMP expression/activity may also be involved in vascular remodeling and placentation during pregnancy and in the vascular changes associated with preeclampsia [57]. Increased expression of MMPs has been observed during different stages of mammalian development, from embryonic implantation [183] to the morphogenesis of different tissues including lung, bone and mammary gland [184, 185].

### MMPs and ECM Degradation

ECM is the extracellular component of mammalian tissue that provides support and anchorage for cells, segregating tissues from one another, regulating cell movement and intercellular communication, and providing a local depot for cellular growth factors. Formation of ECM is essential for biological processes involved in maintaining tissue integrity and regeneration including wound healing and fibrosis. ECM consists mainly of fibers, proteoglycans and polysaccharides. Fibers are mostly glycoproteins and include collagen and elastin. Collagen is the main extracellular protein while elastin, which is exceptionally unglycosylated, provides flexibility for the skin, arteries and lungs. Proteoglycans are glycoproteins containing more carbohydrate than protein. Proteoglycans attract water to keep the ECM environment hydrated and also bind and store growth factors. Proteoglycans include chondroitin sulfate which provides tensile strength to cartilage, ligaments and aortic wall, heparan sulfate which regulates biological activities such as angiogenesis and blood coagulation, and keratan sulfate in cartilage and bone. Syndecan-1 is a proteoglycan and integral transmembrane protein that binds chemotactic cytokines and plays a role in the inflammatory process. Other components of ECM include laminin in the basal lamina of epithelia, and fibronectin which binds cells to ECM, modulates the cell cytoskeleton and facilitates cell movement. ECM also contains polysaccharides such as hyaluronic acid, and proteolytic enzymes which cause continuous turnover of ECM proteins [186–188].

ECM and its protein components provide the structural scaffolding necessary for blood vessel support, cell differentiation, signaling and function, as well as cell migration, epithelialization and wound repair. A key feature of the vein wall structure and integrity is its content of collagen and elastin, which are important MMP substrates. ECM contains other proteins and glycoproteins including fibronectin, vitronectin, aggrecan, entactin, proteoglycans, tenascin, fibrin, and laminin [189, 190]. The identification of MMPs in venous tissues undergoing remodeling supports their role in ECM turnover. MMPs are either secreted from the cell or anchored to the plasma membrane with heparin sulfate glycosaminoglycans. MMPs play an important role in the control of tissue architecture and in degradation of ECM and tissue remodeling [55]. MMPs degrade different components of

ECM including collagen, casein and laminin. The collagenases MMP-1, -8, -13 and -14 efficiently degrade fibrillar collagens type I, II and III in their triple-helical domains [191]. Cleavage by these MMPs renders the collagen molecules thermally unstable so that they unwind to form gelatin, which is then degraded by other members of the MMP family including the major gelatinases MMP-2 and -9. MMP-2 localize at the cell surface by binding via its carboxyl terminus to integrin  $\alpha_v\beta_3$  or the MMP-14-TIMP-2 complex [67]. When bound, the catalytic site of MMP-2 is exposed and can then be cleaved and activated. The  $\alpha_2$  chains of collagen IV bind MMP-9 with a high affinity even when MMP-9 is inactive [192]. This juxtaposition of enzyme and substrate makes a pool of the enzyme that is rapidly available upon activation for any remodeling events.

In human saphenous vein grafts intended for bypass surgery, MMP production and ECM turnover have been implicated in intimal hyperplasia, a leading cause of graft stenosis and failure [193]. In a study evaluating intact versus endothelium-denuded human saphenous vein in culture for 14 days, veins with intact endothelium developed thicker neointima and increased production of MMP-9, that was mainly localized in the vicinity of internal elastic lamina. Inhibition of MMPs with doxycycline abrogated neointima formation and MMP-9 expression [193]. Other studies examined the effects of perfusing saphenous vein segments *ex vivo* and demonstrated increased expression of MMP-2 and -9 when exposed to venous pressure, but a 50% decrease in the expression of gelatinases when veins were exposed to arterial pressure for up to 3 days. These data suggested an association between hemodynamic changes in venous pressure and shear stress, saphenous vein remodeling, and MMP expression [194]. Studies on saphenous vein grafts intended for coronary artery bypass surgery have also shown that the expression and activity of MMP-2 are higher in vein segments exhibiting increased pathologic venous remodeling and lower patency [195]. In addition to the role of MMPs in saphenous vein remodeling, MMPs have been implicated in the invasion of vena cava by metastatic pancreatic cancer. Specifically, liver metastasis is associated with vein invasion and overexpression of MMP-2 and -9 in vena cava [196].

Type III collagen is important for blood vessels elasticity and distensibility, and alterations in collagen synthesis and collagen type I to type III ratio may affect the vein wall architecture and lead to structural weakness, venous dilation and VVs formation. The levels of collagen I and III are co-regulated in fibroblasts, and addition of collagen III to cultured VSMCs from VVs decreases collagen I synthesis [37, 38]. Interestingly, among the various MMPs evaluated, proMMP-2 is increased in dermal fibroblasts cultured from patients with venous disease [197].

Both VVs and saphenous vein in the vicinity of varices show imbalance in ECM [198]. Decreased elastin has been implicated in the pathogenesis of VVs [199]. In contrast, studies have shown increased [198], decreased [200], or unchanged [201] collagen content in VVs wall. The net collagen content represents a balance between its biosynthesis and its degradation by MMPs [198]. In a study examining ECM proteins in 372 vein specimens from 17 patients and 36 control vein specimens from 6 patients, VVs showed marked increase in the ECM proteins collagen, laminin and tenascin, and borderline increase in fibronectin [202]. A study in 8 patients with VVs and 8 control subjects showed an increase in the elastic network, accumulation of collagen type I and fibrillin-1 and overproduction of MMP-1, -2 and -3 in the veins and skin of patients with VVs [203]. Importantly, normal appearing vein segments just inferior to VVs had the same ECM profile as the adjacent VVs [202].

The activity of MMPs in various tissues is tightly regulated by endogenous tissue inhibitors of MMPs (TIMPs) [55]. Alteration of the fine balance between MMPs and TIMPs contribute to the pathological changes in vascular diseases such as atherosclerosis,

aneurysms and VVs [57]. MMPs have been proposed as biomarkers of certain pathological conditions including vascular disease, and are often assessed by measuring their plasma levels, tissue gene expression and protein amount, as well as proteolytic activity using gel zymography.

The venous dilation and tortuosity associated with VVs may be related to the effects of MMPs and their inhibitors TIMPs on ECM components and subsequent vein wall remodeling. The vein wall architecture markedly changes in VVs. In further defining the morphological ECM components and the role of a systemic effect of VVs in patients with CVD, the expression of matrix proteins collagen type I, fibrillin-1, and laminin, as well as MMPs and TIMPs were examined. The control group consisted of dermal biopsies and saphenous vein specimens from patients undergoing coronary revascularization. Both dermal biopsies and VVs specimens showed elevated matrix proteins, and MMP-1, -2 and -3 but not MMP-7 or -9 nor TIMP-1, -2 and -3, suggesting similar matrix protein production and degradation in VVs and adjacent skin [203]. These observations raise the possibility that a skin biopsy may be useful to identify patients at risk for the development of VVs before overt clinical manifestations. Other studies have shown an increase in collagen, a decrease in elastin and increased collagen:elastin ratio in both VVs and competent saphenous vein segments in proximity to varicosities compared to control saphenous vein, although gelatin zymography and elastase activity did not show any differences among the tissues examined. These findings suggested that in VVs and saphenous vein segments near the varices there is an imbalance in connective tissue matrix that may occur prior to valve insufficiency, and that increases in proteolytic activity and matrix degradation are not essential for VVs formation [198]. Also, evaluation of segments at the saphenofemoral junction from patients with VVs and bypass patients demonstrated that MMPs activity was unchanged in VVs compared with control veins, most MMPs located in the adventitia, and the amount of MMP-2 decreased while the amount of TIMP-1 increased. It was concluded that structural vein wall changes and VVs formation may occur despite a decrease in proteolytic activity [204]. Other studies have shown a three-fold increase in TIMP-1/MMP-2 ratio in VVs compared with control veins, suggesting that proteolytic inhibition and ECM accumulation may account for the pathogenesis of VVs [205]. These studies highlight several important points in VVs formation, namely imbalance in connective tissue matrix (collagen:elastin), imbalance of proteolytic tissue degradation, changes in vein wall connective tissue structure prior to valve insufficiency, and proteolytic activity and matrix degradation may not be necessary for VVs formation. These observations raise the question of what role do MMPs have in VVs formation if not degradation of ECM. In addition to their proteolytic effects on ECM in late stages of CVD, MMPs may affect other cell types including ECs and VSM particularly in early stages of VVs formation.

### **Varicose Veins, MMPs, and VSM Dysfunction**

Studies have examined if the ECM changes observed in VVs are related to abnormalities in VSMCs. VSMCs cultured from VVs have decreased number of cells staining for collagen type III and fibronectin compared to VSMCs from control veins, although the transcriptional products of these two proteins were similar in VVs and control veins. The synthesis and deposition of collagen type III but not type I is markedly lower in VVs VSMC culture. Analysis of the supernatant of confluent cells showed no differences in the amount of MMP-1, -2 and -9 and TIMP-1 and -2 in cells from VVs vs. control veins. These data suggested that the regulation of collagen type III and fibronectin in VSMCs was altered at the post-transcriptional level [206]. Other studies have shown that treatment with non-selective MMP inhibitor marimastat (BB-2516) caused partial restoration of the production of collagen type III in VSMCs from VVs. Also, both gene transcription and protein amount of MMP-3, which degrades fibronectin, was elevated in cells from VVs. It was concluded

that the degradation of collagen type III and fibronectin in VSMCs cultured from VVs likely involves the expression of MMP-3 [207]. Thus, cultured VSMCs from VVs show imbalance in collagen production with increased type I collagen but suppressed type III collagen production. Because of normal mRNA expression of type III collagen, the reduction in its production may be related to post-transcriptional events such as degradation/inhibition by MMP-3 and could be manifested as changes in the mechanical properties, elasticity and distensibility of the VVs wall. Recent studies suggested that desmuslin gene expression is required for the maintenance of VSMC phenotype, and decreased desmuslin expression may affect differentiation of VSMCs and contribute to the development of VVs [208].

### Effects of MMPs on VSM Contraction

MMPs, via PI<sub>3</sub>K and ATP synthesis, may transactivate EGFR and mediate  $\alpha$ -adrenergic receptor-induced maintenance of vascular tone. Inhibition of the expression of MMP-2 or -7 blunted the phosphorylation of Akt by PI<sub>3</sub>K and thus inhibited the response to phenylephrine (Phe) in rat mesenteric artery [209]. Other studies have shown that MMP-2 and -9 inhibit Phe-induced contraction of rat aorta [210]. The MMP-induced inhibition of aortic contraction is concentration- and time-dependent and reversible, suggesting that the actions of MMPs are not solely due to irreversible degradation of ECM. Also, the inhibitory effects of MMPs on VSM contraction are not likely due to degradation of Phe or  $\alpha$ -adrenergic receptors because MMPs also inhibit prostaglandin F<sub>2</sub> $\alpha$ -induced contraction, suggesting that the effects of MMPs are not specific to a particular agonist/receptor, but likely involve direct effects on common VSM contraction pathway(s) downstream from receptor activation.

Although some studies suggest an increase in MMPs immunoreactivity in the adventitia, endothelium and VSM of the VVs wall [97], the question remains as of how MMPs cause venous dilation and VVs formation. The role of MMPs in VVs has largely been attributed to their proteolytic effects on ECM, degradation of the valve leaflets and weakening of vein wall structure [128, 211]. The localization of MMPs in the VVs wall adventitia and fibroblasts is consistent with a role in ECM degradation [97]. The proteolytic and degenerative effects of MMPs on ECM could play a role in late stages of VVs formation. On the other hand, the localization of MMPs in the vicinity of the venous endothelium and VSM raises the possibility of an additional effect of MMPs on these cell types [97, 212].

Recent evidence suggests that MMP-2 and -9 may affect aortic VSM function [210]. Also, in rat IVC MMP-2 caused acute relaxation of Phe-induced contraction [7]. VSM contraction is triggered by increases in Ca<sup>2+</sup> release from the intracellular stores and Ca<sup>2+</sup> entry from the extracellular space. MMPs do not inhibit Phe-induced contraction in Ca<sup>2+</sup>-free solution, suggesting that they do not inhibit the Ca<sup>2+</sup> release mechanism. On the other hand, MMPs inhibit Phe-induced Ca<sup>2+</sup> influx [210]. The mechanism by which MMPs inhibit Ca<sup>2+</sup> entry could involve direct effects on Ca<sup>2+</sup> channels. MMPs may also affect K<sup>+</sup> channels [7]. MMP-2 induced IVC relaxation was abolished in 96 mmol/L KCl depolarizing solution, which prevents outward movement of K<sup>+</sup> from the cell through K<sup>+</sup> channels. To further define the K<sup>+</sup> channels involved, it was found that MMP-2 caused further relaxation of IVC segments in the presence of cromakalim, activator of ATP-sensitive K<sup>+</sup> channel (K<sub>ATP</sub>), suggesting that MMP-2 was not working through K<sub>ATP</sub> channel. In contrast, blockade of large conductance Ca<sup>2+</sup>-activated K<sup>+</sup> channels (BK<sub>Ca</sub>) using iberiotoxin inhibited the MMP-2 induced venous relaxation, suggesting that MMP-2 actions involve hyperpolarization and activation of BK<sub>Ca</sub>. MMP-2 induced activation of K<sup>+</sup> channels likely causes VSM hyperpolarization and leads to decreased Ca<sup>2+</sup> influx through voltage-gated channels [7]. This is supported by reports that MMP-2 and -9 cause aortic relaxation by inhibiting Ca<sup>2+</sup> influx into aortic VSM [210] and MMP-2 inhibits Ca<sup>2+</sup>-dependent contraction in rat IVC [9]. MMPs may also induce collagen degradation and produce Arg-

Gly-Asp (RGD)-containing peptides, which could bind to  $\alpha_v\beta_3$  integrin receptors and inhibit  $\text{Ca}^{2+}$  entry into VSM [213]. MMPs may also stimulate protease-activated receptors (PARs) and activate signaling pathways that could lead to blockade of VSM  $\text{Ca}^{2+}$  channels [214]. This is supported by reports that proteases such as thrombin activate PARs and promote endothelium-dependent VSM relaxation by inhibiting  $\text{Ca}^{2+}$  influx [215]. These data demonstrate novel effects of MMPs on venous function and suggest that protracted MMP-2 induced vein relaxation could lead to progressive wall dilatation and superimposed valve dysfunction, leading to VVs formation and CVD [57]. While MMPs may affect VSM contraction and ion channels, further studies are needed to define the role of integrins and PARs as possible molecular mechanisms via which MMPs could inhibit VSM contraction.

### MMPs and VSMC Migration

MMPs play a role in VSMC migration. In rat aortic smooth muscle cells (RASMCs) cultured on collagen I gel to mimic ECM, exposure to flow enhanced cell motility. Upregulation of MMP-1 enhanced flow-induced motility, and the MMPI GM-6001 attenuated flow-induced migration. ERK<sub>1/2</sub> phosphorylation and increased expression of AP-1 transcription factors c-Jun and c-Fos appear to be involved in MMP-mediated enhancement of flow-induced cell motility [216]. Young human aortic smooth muscle cells (HASMCs) produce active MMP-2 and have higher migratory capability than aged cells, likely due to increased MT1-MMP content and activation of proMMP-2 in young cells. In contrast, aged cells produce only proMMP-2, and its activation is prevented by upregulation of TIMPs, and treatment of young cells with TIMP-1 and -2 promotes aged HASMCs migratory behavior [217]. MMP-2 activity could also influence chemokine-induced chemotaxis of human VSMC monolayers [218]. Also, MMP-2 knockout decreases VSMC migration and intima formation in mouse model of carotid artery ligation [219, 220]. MMP-9 may also play a role in VSMC migration. Tanshinone IIA, a major constituent of *Salvia miltiorrhiza bunge*, inhibits TNF- $\alpha$ -induced HASMC migration, partly through inhibition of MMP-9 activity. Tanshinone IIA also inhibits TNF- $\alpha$ -induced ERK and c-jun phosphorylation, and NF- $\kappa$ B and AP-1 DNA-binding [221]. Suppression of MMP-9 expression by downregulation of NF- $\kappa$ B mediates the inhibitory effect of curcumin on migration of HASMCs [222]. Also, MMP-9 knockout is associated with reduced VSMC migration and intima formation in mouse model of filament loop injury [223] or carotid artery occlusion [224].

Disruption of the basement membrane is essential for VSMC migration [225]. MMPs, by degrading the basement membrane, can facilitate ECM integrin interactions leading to activation of focal adhesion kinases (FAK) and increased cell migration. MMPs also cause fragmentation of membrane components such as type I collagen, thus creating new integrin-binding sites. Growth factor receptors, cadherins and integrins mediate signaling pathways that play a role in reorganizing the cytoskeleton in preparation for migration [226, 227]. MMPs cleave E-cadherin in epithelial cells, VE-cadherin in ECs and N-cadherin in VSMCs [228, 229], thus dissolving adherence junctions and allowing the cells to migrate.

MMPs not only facilitate migration by promoting proteolysis of ECM, but may also directly enhance cell migration. MMP-1 promotes growth and invasion of cells by binding to and cleavage of PAR-1 which reveals a tethered ligand that initiates signaling via a G protein-coupled receptor and activates migration [230]. This mechanism allows the cells to sense a proteolytic environment and actively move towards an area of degraded matrix. Rearrangement and migration of SMCs into the intima may be seen in VVs [231–234]. SMCs in VVs appear disorganized and dedifferentiated from contractile to synthetic and proliferative phenotype, and demonstrate features such as vacuolization and phagocytosis [231, 232].



MMPs have been used to demonstrate the effect of MMPs on VSMC migration. Gene transfer of TIMPs reduces VSMC migration *in vitro* and inhibits or delays intima thickening *in vivo*. TIMPs 1–4 delivered directly or by gene transfer inhibit migration of SMCs *in vitro* [235, 236] and reduce neointima formation in organ culture of human saphenous vein [237]. TIMP gene transfer also preserves medial basement membrane and inhibits VSMC migration to the intima. Synthetic MMPs inhibit migration of VSMC from baboon arterial explant in culture [238], and early VSMC migration in the rat carotid balloon injury model [239]. Thus MMPs enhance VSMC migration, and MMPIs could reverse or prevent cell migration.

### MMPs and VSMC Proliferation

In addition to facilitating VSMC migration, MMPs may regulate VSMC proliferation. Pretreatment of HASMCs with ethanol extract of *Buddleja officinalis* attenuates high-glucose-induced cell proliferation by suppressing MMP-9 activity [240]. Also, MMP-9 knockout is associated with inhibition of VSMC proliferation after filament loop carotid artery injury [223], but not after tying off the mouse carotid artery [224]. These inconsistent results suggest compensatory activation of other proteases [241]. When compared with SMCs derived from normal veins, SMCs derived from VVs demonstrate increased proliferation (2-fold), migration (3-fold), MMP-2 production (3-fold), and collagen synthesis (>2-fold), with decreased expression of phenotype-dependent markers [242].

MMPs can regulate VSMC proliferation by promoting permissive interactions between VSMC and components of ECM. Integrin-mediated pathways may be essential for stimulation of VSMC proliferation [243, 244]. Also, MMPs may free growth factors from attachment to ECM components or cell surface so that they can act on their receptors. ECM binds growth factors either directly or via growth-factor-binding proteins. MMPs stimulate the release of growth factors such as transforming growth factor  $\beta$  (TGF- $\beta$ ), fibroblast growth factor 1 (FGF-1), insulin-like growth factor 1 (IGF-1), TNF $\alpha$  and heparin-binding epidermal growth factor (HB-EGF) by cleaving either the growth-factor binding protein or the matrix molecule to which these proteins attach. HB-growth factors, in particular FGF-1 and FGF-2, are potent mitogens for VSMCs and are released by the action of MMPs on proteoglycan core proteins [55]. Although ADAMs are often implicated, MMPs could also be responsible for releasing cell surface HB-EGF and stimulation of VSMC proliferation [245, 246]. MMPs also activate TGF-h by cleaving off the latency-associated peptide [247]. MMPs can also liberate active IGF-1 by degrading its binding proteins. Together with signals from FAK, these processes upregulate and/or stabilize key regulators of the cell cycle. Also, MMP-induced cadherin shedding promotes dissolution of adherens junctions and translocation of h-catenin to the nucleus where it acts as a transcription factor to further promote cell proliferation [227, 229]. MMP-3 and -7 can cleave the adherens-junction protein E-cadherin [248–252]. MMP-3 can release a soluble form of the adhesion molecule L-selectin from leukocytes, and shed membrane-bound HB-EGF to exert signaling functions [248]. MMP-7 may release soluble Fas ligand which induces apoptosis [253]. Some growth factors are proteolytically inactivated by MMPs, including the chemokine connective tissue activating peptide III (CTAP-III), monocyte chemoattractant protein (MCP) and stromal cell-derived factor 1 [254, 255]. Growth factors and cytokines are also negatively regulated when MMPs cause shedding of their receptors from the cell membrane, as in the case of surface FGF receptor 1 [256].

MMPs have been used to study the role of MMPs in VSMC proliferation. Some studies have shown excess neointima proliferation in rat carotid arteries subjected to balloon injury after treatment with the MMPI GM-6001 [257, 258]. Other studies have shown that synthetic MMPIs inhibit VSMC proliferation *in vitro* [229, 259]. Also, inhibition of MMPs is associated with decreased N-cadherin shedding, increased cell membrane N-cadherin

levels, decreased h-catenin nuclear translocation and decreased proliferation of cultured human VSMCs. Dismantling of cadherin:catenin complex also occurs in balloon-injured rat carotid arteries *in vivo* leading to increased expression of the cell cycle gene cyclin D1 and VSMC proliferation [260]. Tetracycline-based MMPiS reduce VSMC migration and neointima formation after balloon injury of rat carotid artery [239, 261]. Thus, most of the experimental evidence points to a stimulatory effect of MMPs on VSMC proliferation, and inhibition of this effect by MMPiS.

Increases in wall tension may elicit a remodeling process in the vein wall similar to that during varicosis, and this biomechanical force may be sufficient to elevate the proteolytic and proliferative activity of VSMCs. Pressure-induced increase in MMP-2 expression, gelatinase, and proliferative activity is dependent on AP-1. An increase in venous filling pressure and circumferential wall tension is sufficient to activate AP-1, which in turn triggers varicose remodeling through fuelling MMP-2 activity and VSMC hyperplasia in the vein wall. Also, in both native and cultured VSMCs exposed to increased stretch inhibition of AP-1 decreased cell proliferation, MMP-2 expression and gelatinase activity by up to 80%. AP-1 also regulates the expression of many stress-response genes, including those associated with a proinflammatory phenotype of ECs and VSMCs [262, 263]. To test if AP-1 was a suitable target for therapeutic interference with varicosis, its activity was blocked *in vivo* and *in vitro* by a specific decoy ODN. These nucleic acid-based drugs have been introduced as potent inhibitors of maladaptive remodeling processes [264, 265]. Without the need for potentially harmful transfection reagents, the AP-1 decoy ODN-supplemented ointment readily penetrated the skin of the mouse auricle and locally inhibited the stretch-induced MCP-1 expression [266]. Also, application of the AP-1 decoy ODN virtually abolished outward venous remodeling and tortuosity *in vivo*, likely due to its inhibitory effects on EC and SMC proliferation, MMP-2 expression and gelatinase activity. In cardiac fibroblasts heterodimerization of the AP-1 subunits JunB-Fra1 and JunB-FosB may be crucial to control transcription of the MMP-2 gene [267]. This finding is in line with the observation that isolated perfused JunB-deficient mouse veins do not respond to increased hydrostatic pressure with an increase in MMP-2 expression. While taking into account its additional anti-inflammatory effects, therapeutic inhibition of stretch-induced MCP-1 expression attenuates the initiation and progression of VVs remodeling and may be effective in preventing recurrence. HMG-CoA reductase inhibitors — cholesterol synthesis inhibitors that inhibit AP-1 as part of their pleiotropic activity — may be potential drugs to treat VVs [268].

### MMPs and VSMC Apoptosis

Apoptosis is a form of cell death that involves activation of the intracellular cysteine proteases, caspases. Apoptosis of VSMCs plays a role in attenuating intimal thickening and destabilizing atherosclerotic plaques [269, 270]. Several factors promote apoptosis including death signals originating from outside the cell and processes within the cell such as DNA damage, cell cycle status and levels of the tumor suppressor p53 [270]. MMP-7 is involved in the cleavage of N-cadherin and thus modulates VSMC apoptosis. In contrast, survival signals maintain VSMC viability even in the face of a pro-apoptotic environment. Survival pathways are closely linked to those involved in proliferation and therefore could be influenced by MMPs. Survival factors such as PDGF, HB-EGF and IGF-1 act via tyrosine kinase receptors to stimulate the PI<sub>3</sub>-K/Akt pathway. MMP-2, -7 and -9 cleave cell surface pro-HB-EGF and liberate the soluble active growth factor which binds to EGFR and promotes growth [245, 271]. Activation of PDGFR-β and ERK1/2 is involved in the production of MMP-1 in oxLDL- and 4-hydroxynonenal-stimulated human coronary VSMCs [272]. MMP-1, -2, -8 and -9 degrade members of the IGF binding protein family and thereby increase the bioavailability of IGF-1 and its anti-apoptotic effects [55].

Cell-matrix contacts promote VSMC survival, and their disruption leads to apoptosis and anoikis [273]. FAK activation triggered by ECM-integrin interactions induces p53, a survival signaling pathway [274, 275]. Regulated MMP production appears to favor FAK activation and hence survival signaling. Conversely, excessive MMP production could degrade ECM proteins or integrins and promote anoikis [276]. MMPs may also modulate apoptosis by cleaving death ligands such as TNF- $\alpha$  and Fas ligand and their receptors. MMP-1, -2, -9, -8 and -13 and the MT-MMPs MMPs 14–17 can cleave pro-TNF- $\alpha$  [55, 277]. Similarly, MMP-7 sheds Fas-L from the surface of several cell types [278, 279]. Caspase-mediated cleavage of the DNA repair enzyme poly-ADP ribosepolymerase is an important step in apoptosis, and in isolated cardiac myocytes, nuclear-localized MMP-2 can carry out this cleavage [280].

TIMP-3, but not TIMP-1 or -2, is an effective stimulator of apoptosis in many cells including VSMCs [236, 279], suggesting that an ADAM rather than an MMP is the target. TIMP-4 also stimulates VSMC apoptosis [281]. Thus, MMPs may regulate VSMC apoptosis via several pathways, and MMPIs could oppose the effects of MMPs on apoptosis. However, the role of VSMC apoptosis has not been clearly defined in VVs

### MMPs and Endothelium-Dependent Vascular Relaxation

ECs control vascular tone by releasing relaxing factors including nitric oxide (NO), prostacyclin (PGI<sub>2</sub>), and also through hyperpolarization of the underlying VSMCs by endothelial-derived hyperpolarizing factor (EDHF) [282]. EDHF-mediated responses involve an increase in the intracellular Ca<sup>2+</sup> concentration, the opening of Ca<sup>2+</sup>-activated K<sup>+</sup> channels of small and intermediate conductance and hyperpolarization of ECs. EC hyperpolarization spreads via myoendothelial gap junctions and result in EDHF-mediated hyperpolarization and relaxation of VSMCs [283–285]. MMPs may have endothelium-dependent vasorelaxation effects. MMP-2 may mediate the bacterial LPS-induced vascular hyporeactivity to vasoconstrictors in rat aorta via an endothelium-dependent mechanism [286]. MMPs could increase PGI<sub>2</sub> synthesis and activate PGI<sub>2</sub>-cAMP pathway, or increase EDHF release and enhance K<sup>+</sup> efflux via K<sup>+</sup> channels leading to venous tissue hyperpolarization and relaxation [287]. MMP-2 induced IVC relaxation is abolished in high KCl depolarizing solution, which prevents outward movement of K<sup>+</sup> from the cell through K<sup>+</sup> channels, and inhibited by the BK<sub>Ca</sub> blocker iberiotoxin, suggesting a role of a hyperpolarization pathway [7]. Long-term, these novel effects of MMPs on venous relaxation pathways could lead to progressive venous dilatation and VVs formation.

Upregulation of MMPs may also be associated with impaired vasorelaxation. Upregulation of MMP-3 and downregulation of TIMP-1 mediate the impaired endothelium-dependent vasodilation, EC apoptosis and endothelial disruption exerted by FOXO3 in HUVECs [288]. Also, upregulation of MMP-2 and -9 may be responsible for nicotine-induced endothelial disruption and unresponsiveness of blood vessels to the vasodilator acetylcholine, and the MMPI doxycycline partially reversed this effect [289]. In renovascular rat model of hypertension, antioxidant treatment inhibited the decrease of endothelium-dependent vasorelaxation and attenuated the vascular dysfunction and remodeling by inhibiting oxidative stress-induced upregulation of MMP-2 [290].

### Effects of MMPs on EC Integrity and Vascular Permeability

MMPs may regulate EC integrity and vascular permeability. MMP-1 mediates the activation of HUVECs into prothrombotic, proinflammatory, and cell-adhesive state by supernatants from cultured melanoma and colon cancer cells [291]. In mouse aorta, MMP-13 mediates the endothelial protective effect of NO by cleaving the pro-inflammatory ICAM-1 [292]. Infiltration of inflammatory cells and increased expression of surface markers including

VCAM-1 and ICAM-1 is often observed in ECs of VVs [95, 231, 293]. MMPs may also increase vascular permeability and cause vascular disruption. Upregulation of MMP-2 and -9 mediate the increase in membrane permeability and vascular disruption induced by human immunodeficiency virus-1 envelope gp120 in rat brain [294]. Also, MMP-2 and -9 decrease the barrier integrity of primary porcine brain capillary ECs [295]. In support of these findings, the MMPI GM6001 prevents degradation of the tight junction protein occludin and reduces the intercellular gap formation and permeability in porcine cerebral microcapillary ECs [296]. Thus the expression of MMPs by SMCs and ECs may influence the relaxation/constriction properties of the vein wall and contribute to the development of VVs. Also, the loss of endothelial integrity may expose the components of the vein wall to degradation by MMPs.

### Management of Varicose Veins

Management of VVs comprises compression therapy, pharmacological treatment, sclerotherapy and surgical intervention (Table 3). Graduated compression stockings relieves the edema and pain, controls progression to more severe forms of CVI with skin changes and ulceration, and prevents venous thromboembolism following surgery [297–301]. The benefits of compression garments in patients with CVI may not be related to improvement of the deep venous hemodynamics [302]. Instead, graduated compression stockings with an ankle pressure of 21 mmHg affect the residual venous volume fraction, which is linearly correlated to the amount of reflux and a measure of venous hypertension in the limb [303].

Pharmacological treatment of VVs and CVD comprises naturally occurring plant extracts and glycosides including  $\alpha$ -benzopyrones (coumarins),  $\gamma$ -benzopyrones (flavonoids), saponosides (escin, horse chestnut seed extract), and plant extracts (blueberry and grape seed, Ginkgo biloba) (Table 3). Although the precise mechanism of action of these venoactive drugs is not known, they appear to improve venous tone and capillary permeability, and induce venotonic stimulation and leukocyte modulation. Flavonoids may affect leukocytes and the endothelium and modify the degree of inflammation and reduce edema [304]. Saponosides may limit the distensibility and morphologic changes in the vein wall [305]. In rat aortic rings, escin (horse chestnut seed extract) protects the endothelium, causes  $\text{Ca}^{2+}$ -dependent venous contraction [306, 307], and decreases contraction to venoconstrictors such as  $\alpha$ -adrenergic and AngII ( $\text{AT}_1\text{R}$ ) receptor stimulation and membrane depolarization [307]. Diosmin may improve venous tone, microvascular permeability, lymphatic activity, and microcirculatory nutritive flow [308]. Also, rutosides enhance endothelial function in patients with CVI [309].

In patients with CVI there is elevated circulating ECs, a marker of venous injury and ischemia [310]. In a randomized clinical trial Ginkgo biloba extract, troxerutine, and heptaminol (Ginkor Fort) were tested for their venoprotective effects in patients with CVI. The treated patient had decreased circulating ECs, supporting venoprotective effects of these compounds against EC injury associated with CVD [311]. Daflon 500, a combination of the active venotonic diosmin and the inactive compound hesperidin, is formulated as a micronized purified flavonoid fraction for increased absorption. In 3 randomized clinical trials, involving 183 patients and equal number of controls, hemodynamic and clinical improvement of symptoms were observed in patients receiving Daflon vs. placebo [312]. Also, a trial of 231 patients with CVI showed that a combination of coumarin and the flavonoid troxerutin, with compression garments for 12 weeks resulted in less edema and pain compared with placebo [313]. Another trial evaluating the use of venoruton (oxerutin) vs. Daflon in patients with CVI for 8 weeks has shown that venoruton improved symptoms as assessed by a validated venous quality of life tool [309]. Also, a controlled trial in patients with venous ulcer demonstrated that using Daflon 500 in addition to compression increased the number and rate of healed venous ulcers and improved venous symptoms

[314]. Horse chestnut seed extract is as effective as compression stockings in reducing the CVD symptoms of heaviness, itching, leg edema and pain in the short-term, but its long-term safety and efficacy have not been established [315, 316].

Other compounds have been tested in advanced CVI. In clinical trials pentoxifylline increased healed ulcers and healing rate over placebo, but its mechanism of action was unclear [317, 318]. Also, red vine leaves (AS 195) may increase microcirculatory blood flow and transcutaneous oxygen tensions and alleviate edema in CVD [319]. Prostaglandins that inhibit platelet aggregation, leukocyte activation and capillary permeability and promote venodilation and fibrinolysis can be useful. In a randomized trial, infusion of prostaglandin E1 vs. placebo for 20 days showed healed venous ulcers at 4 months, and a decrease in ulcer area at 80 days, suggesting beneficial effects in venous ulcer healing, possibly by improving the microcirculation and reducing the hypoxemia associated with venous ulcer [320].

Sclerotherapy is widely used for treatment of small telangiectasias (spider veins), reticular veins, and large VVs. For smaller diameter veins diluted forms of sclerosing agents are used to avoid tissue inflammation and necrosis. Concentrated sclerosing agents such as hypertonic saline, sodium morrhuate, and ethanolamine oleate are less commonly used because of their side effects. Sodium tetradecyl sulfate (STS) is a detergent that can be used as liquid or foam. Sodium morrhuate and STS are approved for treating varicosities in the United States. Polidocanol is superior to normal saline in obliterating incompetent VVs and improving venous hemodynamics [321]. Sclerotherapy with polidocanol foam under duplex ultrasound guidance is a standard treatment of intracutaneous telangiectasias, subcutaneous VVs, transfacial perforating veins and venous malformations in Europe [322]. Polidocanol and STS are used in Europe and South America as a mixture of 1:4 liquid:air, which increases the surface area and contact with the vein wall. Because of some concerns such as air emboli, visual disturbances, cough and migraines, this form of therapy is not approved in the United States [323, 324].

Surgical treatment of CVI includes endovenous ablation, surgical stripping and phlebectomy of superficial veins. Endovenous ablation is performed with either radiofrequency or infrared laser at wavelengths 810 to 1320 nm, but could go up 1470 and 1550 nm. The resulting high intensity endoluminal thermal heat causes endothelial protein denaturation and vein occlusion [325]. Endovenous ablation has relatively good occlusion rates and long term benefits. Laser therapy has only 3–7% recurrence rates at 2–3 years follow up [326, 327], while radiofrequency therapy has 2% recanalization rate at 4 years follow up [328]. Surgical stripping of the saphenous vein with high ligation of the saphenofemoral junction is considered durable and the standard for many patients with CVD [329]. Large varicose clusters that communicate with the incompetent saphenous vein can be avulsed by “stab phlebectomy”. An alternative to open phlebectomy is the use of transilluminated power phlebectomy to remove clusters of varicosities using fewer incisions and thereby decrease the operation time [330]. In a clinical trial, 500 patients with incompetent superficial and deep venous systems and demonstrable CVI and venous ulcer were randomized to surgery only to the superficial veins plus compression vs. compression alone. Patients having surgery of the superficial veins demonstrated marked reduction in ulcer recurrence (12%) compared with compression alone (28%) at 12 months [331].

### **MMP Inhibitors in Varicose Veins**

The current strategies for management of VVs have variable success rates, and recurrence of varicosities is observed in many cases. Also, most of the current therapies are used in overt and advanced cases of VVs, and few therapies are available for prevention or limiting the progression of early manifestations of CVD. The failure/recurrence of some of the current CVD treatment strategies is largely related to the fact that most of these techniques are



designed to treat the manifestations rather than the cause of the disease. MMP inhibitors (MMPIs) have been used to investigate the role of MMPs in different physiologic processes and in the pathogenesis of specific diseases. Loss of certain cellular function following treatment with MMPIs has supported a role of MMP in the regulation of specific cellular functions. MMPIs have also been investigated as potential therapeutic tools for arthritis, cancer and vascular disease. The observations that MMPs expression/activity is modulated during the development of VVs and its progression to thrombophlebitis and wound ulcers have raised interest in the potential use of MMPIs in the management of CVD.

MMPIs can be macromolecules such as endogenous tissue inhibitors of MMPs (TIMPs) and monoclonal antibodies or small molecules (natural and synthetic products) (Table 4). Synthetic MMPIs include Zn<sup>2+</sup> binding globulins (ZBG), non-ZBG, and mechanism-based inhibitors. Most MMPIs contain a Zn<sup>2+</sup> binding group (e.g. hydroxamic acid, carboxylic acid, sulfhydryl group) that is either substituted with a peptide-like structure that mimics the MMP substrate that they cleave or appended to a smaller side chain substituent that interact with specific subsite or pocket (e.g., P1', P2', P3') within the MMP active site [332]. The potency and specificity of an MMPI can be determined by the substituent groups. Substituent groups include  $\alpha$ -substituents (increase the inhibitory activity against specific MMPs), P1' substituents (major determinant of activity and selectivity), P2' substituents (variable range of substituent groups; steric bulk proximity to amide provides oral bioavailability), and P3' substituents (variable range of substituent groups; charged or polar groups affect biliary excretion) [333]. While a large number of MMPIs have been developed, many of them lack specificity. Novel pharmacological tools and genetic engineering have improved the specificity of MMPIs to target specific MMPs in vascular disease [53].

### Endogenous MMP Inhibitors

TIMPs and  $\alpha$ 2-Macroglobulin are two major endogenous inhibitors of MMPs. TIMPs are endogenous naturally occurring MMPIs that bind MMPs in a 1:1 stoichiometry. To date, 4 homologous endogenous TIMPs, TIMP-1, -2, -3 and 4, have been identified. TIMP-1 and -3 are glycoproteins, while TIMP-2 and -4 do not contain carbohydrates. TIMPs have an N-terminal domain (125 aa) and C-terminal domain (65 aa), each containing 3 disulfide bonds. The N-terminal domain folds as a separate unit and is capable of inhibiting MMPs [334, 335]. The TIMP molecule wedges into the active-site cleft of MMP in a manner similar to that of the substrate. Cys1 is instrumental in chelating the active site Zn<sup>2+</sup> with its N-terminal  $\alpha$ -amino group and carbonyl group, thereby expelling the water molecule bound to the catalytic Zn<sup>2+</sup>. TIMPs have poor specificity for a given MMP, and each TIMP can inhibit multiple MMPs. An exception is the ability of TIMP-2 and -3 to inhibit MT1-MMP and MT2-MMP, while TIMP-1 has poor inhibitory effect on MT1-MMP, MT3-MMP, MT5-MMP and MMP-19 [336, 337]. TIMPs also inhibit a broader spectrum of metalloproteinases. ADAMs are members of the disintegrin and metalloproteinase family. In contrast with ADAMs, ADAMTS (ADAMs with a thrombospondin motif) are secreted and lack a membrane anchor. Both ADAM and ADAMTS have similar active and binding sites as MMPs, and hence are inhibited by broad spectrum MMPIs [333]. TIMP-1 inhibits ADAM-10 while TIMP-2 inhibits ADAM-12 [338] and TIMP-3 has a much broader inhibition profile including ADAM-10, -12 [339] and -17 [338] and ADAMTS-1, -2, -4 and -5 [340, 341]. Because of its broad inhibitory spectrum, TIMP-3 ablation in mice causes lung emphysema-like alveolar damage [342] and faster apoptosis of mammary epithelial cells after weaning [343], whereas TIMP-1-null mice and TIMP-2-null mice do not exhibit obvious abnormalities. Binding studies have shown that native TIMPs can distinguish between stromelysins. TIMP-1 and -2 binding to human MMP-10 (stromelysin-2) catalytic domain is 10-fold weaker than their binding to the MMP-3 (stromelysin-1) catalytic domain.

The x-ray crystal structure of TIMP-1 bound to MMP-10 catalytic domain demonstrated substantial differences with the structure of the TIMP-1.MMP-3 catalytic domain at the binding interface. These structural differences may help in the design of more selective TIMP-based inhibitors toward members of the stromelysin family and increase their therapeutic applications [344]. TIMP-1 has a threonine 2 residue that interacts with the specific P<sub>1</sub>' substituent of MMP-3. Substitutions at threonine 2 of TIMP-1 affect the stability of complexes with MMPs and influence the specificity for different MMPs. For example, a substitution of alanine for threonine 2 causes a 17-fold decrease in binding of TIMP-1 to MMP-1 relative to MMP-3 [345, 346]. A functional and intact catalytic domain and the ability for the histidine residues to bind zinc are necessary for the interaction and complex formation TIMP with MMP [347].

MMP activity is partly regulated by  $\alpha$ 2-macroglobulin and related proteins. Human  $\alpha$ 2-macroglobulin, a glycoprotein consisting of four identical subunits and found in blood and tissue fluids, acts as a general proteinase inhibitor.  $\alpha$ 2-Macroglobulin inhibits most endopeptidases by entrapping the peptidase within the macroglobulin. The complex is then rapidly cleared by endocytosis via a low density lipoprotein receptor-related protein-1 [348].

Other proteins inhibit selected members of MMPs: a secreted form of  $\beta$ -amyloid precursor protein inhibits MMP-2 [349]; a C-terminal fragment of procollagen C-proteinase enhancer protein inhibits MMP-2 [350]; tissue factor pathway inhibitor-2, a serine proteinase inhibitor, inhibits MMP-1 and -2 [351], and RECK, a GPI-anchored glycoprotein, inhibits MMP-2, -9 and -14. However, the mechanism of action of these inhibitors is not well-described [352].

### Exogenous (Synthetic) MMP Inhibitors

Because the mechanism by which MMPs cleave their substrates requires catalytic Zn<sup>2+</sup> ion, the design of MMPI has traditionally utilized Zn<sup>2+</sup> binding globulin (ZBG). ZBGs displace the Zn<sup>2+</sup>-bound water molecule and inactivate the enzyme [353]. ZBG also acts as an anchor to lock the MMPI in the active site and direct the backbone of the inhibitor into the target substrate-binding pockets [354] (Table 4).

Early MMPIs included hydroxamic acids (ZBG1), carboxylates (ZBG2), thiols, and phosphonic acids (phosphorus-based ZBGs) [152]. Of these MMPIs, hydroxamic acids are preferred due to their relative ease of synthesis and potent binding [355]. Hydroxamates have a collagen mimicking structure which facilitates MMP inhibition by bidentate chelation of the active site Zn<sup>2+</sup> ion [356]. A contributing factor to the effectiveness of hydroxamates is the hydrogen bonding that results between the heteroatoms of the ZBG and neighboring amino acid residues that are conserved in all MMP active sites. Although hydroxamates are potent MMPIs and some of them may show selectivity among different MMPs [357–361], many of them have musculoskeletal side effects and poor oral bioavailability [362, 363].

Succinyl hydroxamates such as batimastat (BB-94) and marimastat (BB-2516) are broad spectrum compounds that inhibit a wide range of MMPs. Batimastat has low water solubility and limited use in treating human inflammatory and malignant conditions. Marimastat is soluble in aqueous conditions and is orally bioavailable. An *in vitro* study showed a decrease in type III collagen and upregulation of MMP-3 in SMCs isolated from human VVs, with partial reversal of these changes with marimastat [207]. A limiting factor of this class of MMPIs is the non-specific cross-reactivity with other MMPs besides MMP-2 and -9. Because gelatinases have been implicated in tumor invasion and rheumatoid arthritis, attempts have been made to increase the selectivity of hydroxamates by designing compounds with greater interaction with the active site and the fibronectin type II domain of MMP-2 [364, 365].

Sulfonamide hydroxamates inhibit a variety of MMPs [366], but inhibit other enzymes with an active site  $Zn^{2+}$  moiety such as carbonic anhydrases and tumor necrosis factor converting enzyme (TACE), a member of ADAMs [333, 367]. Sulfam hydroxamates are a group sulfonamide hydroxamate MMPIs with increased specificity toward MMP-2, -9 and -13 and an  $IC_{50}$  for MMP-2 (1 nM) 1000 fold more selective over MMP-1 [357]. Sulfam hydroxamates can also be chemically modified to become potent TACE inhibitors ( $IC_{50}=3.7$  nM) without any cross reactivity with MMP-1, -2, -9 and -13 [368]. Sulfonamide phosphonates are structural analogs of hydroxamates in which incorporation of a favorable structure to attain affinity at the  $Zn^{2+}$  binding group and an arylsulfonamino backbone leads to nM potency and selective inhibition of MMP-8 [369]. Addition of other substituent groups to such as a tetrahydroisoquinoline moiety to sulfonamide hydroxamates led to a compound that inhibits a large number of MMPs, except MMP-7 [370]. Phosphinamide hydroxamates are potent MMPIs with  $IC_{50}$  of 20.5 nM for MMP-1 and 24.4 nM for MMP-3, but the hydrolysis of the phosphinamide bond in acidic conditions leads to inadequate oral bioavailability and limit their clinical use [333, 371].

Other small molecule MMPIs include carboxylates and thiols. In contrast with hydroxamates which have great affinity for MMPs and their binding is pH independent, carboxylates binding and MMP inhibition is dependent on the pH [333]. The  $IC_{50}$  for MMP-1 inhibition by carboxylates is in the  $\mu$ M range, i.e. 1000 fold less potent than hydroxamates [333, 372]. Several carboxylate inhibitors have been developed from the parent compound N( $\alpha$ )substituted 2,3-diaminopropionic acid to improve their MMP specificity. The 2-arylsulfonyl-1,2,3,4-tetrahydro-isoquinoline-3-carboxylates form MMP-8 inhibitor complex directed at the  $S1'$  hydrophobic specificity region near the catalytic  $Zn^{2+}$  ion. The molecular scaffold for  $Zn^{2+}$  binding and the hydrophobic region lie in the 1,2,3,4-tetrahydro-isoquinoline and the aryl aromatic rings, respectively, and enhance MMP-8 hydrophobic specific binding [373]. Several carboxylates such as PGE-2909492, PGE-6292544 have specificity for MMP-13 [374]. Other carboxylates prepared by the reaction of arylsulfonyl isocyanates with a 5-dibenzo-suberenyl/suberyl moieties, followed by the conversion of the carboxy-o-methyl to the carboxylate moieties are powerful MMPIs with activities in the low nM range, and selectivity for the deep pocket MMP-2, -8 and -9 over the shallow pocket MMP-1 [375].

Thiols are another class of small molecule MMPIs which are highly dependent on the mercapto group and the substitution for the  $P1'$  and  $P2'$  amino groups on the MMP molecule. Thiols are potent MMPIs with  $IC_{50}$  in the nM  $IC_{50}$  range, and stereotactic substitutions at the thiol group can change their potency by 100 fold [376]. Thiol MMPIs such as 4-(4-phenoxphenylsulfonyl)butane-1,2-dithiol and 5-(4-phenoxphenylsulfonyl)pentane-1,2-dithiol show selectivity for MMP-2 and -9. Both inhibitors chelate the catalytic  $Zn^{2+}$  ion of MMP-2 by 2 sulfur atoms and lead to conformational changes at the catalytic  $Zn^{2+}$  binding group [377].

By improving the potency and selectivity of ZBGs, some of these MMPIs have been considered as potential therapies for certain degenerative and vascular disease. Site specific delivery is another approach that allowed the use of MMPIs with low potency. Using these approaches, a series of biphenyl sulfonamide carboxylate MMPIs with high selectivity for MMP-13 were designed for treatment of osteoarthritis [378]. The carboxylic acid scaffold of these MMPIs was also used to develop selective MMP-12 inhibitors for treatment of chronic obstructive pulmonary disease [379]. Also, a series of MMPIs with improved selectivity towards MMP-12 over MMP-13 were generated by using a fused ring system. Selective hydroxamate inhibitors of MMP-2 have also been developed as potent anti-angiogenic agents, and compound 7 is the most selective inhibitor of this series. Compound 7 (UK-370, 106) is also a selective inhibitor of MMP-3 and has been developed for topical treatment of

CVU and chronic non-healing wounds [361], but has not been tested in clinical trials. Systemic administration of compound 7 may also modify the MMP profile in CVU because when it is administered intravenous in rats, it clears rapidly from plasma but slowly from dermal tissue [380]. Another class of arylsulfonamido-based type A hydroxamic acid compounds substituted on their sulfonamido nitrogen with an oxyalkyl side chain, instead of the hydrogen atom of type B compounds, or an alkyl side chain of type C compounds produced potent and selective MMPIs of MMP-2 and -9 that block *in vitro* tumour cell invasion. The prototype showed a very good MMP inhibitory profile, with high potency towards MMP-2 (IC<sub>50</sub> = 0.41 nM), MMP-9 (IC<sub>50</sub> = 16 nM) and MMP-14 (IC<sub>50</sub> = 7.7 nM), and blocked angiogenesis in the chemoinvasion model of HUVECs at submicromolar concentrations [359].

### Derivatives of Early ZBGs

Hydrazide (ZBG3) and sulfonylhydrazide (ZBG4) analogs of the hydroxamate MMPI illomastat (GM-6001) have been developed [381] (Table 4). Sulfonylhydrazide 9 is a potent inhibitor of MMP-1, -2 and -9 [382]. Mercaptosulfide (ZBG8) inhibitors target MMP-14. MMPIs with phosphorus-based ZBGs have demonstrated improved selectivity, and compound 18 is a potent phosphonate inhibitor that exhibits selectivity for MMP-8 [383]. Other phosphorus-based ZBGs include the carbamoyl phosphonate ZBG9 [354].

The net negative charge on ZBGs prevents cell penetration and restricts these MMPIs to the extracellular space, and therefore lowers their toxicity [384]. Several MMPIs based on these ZBGs are selective for MMP-2 in both *in vitro* and *in vivo* models of tumor invasion and angiogenesis. Compound 20 (N-methylcarbamoylphosphonic3Acids) shows marked specificity towards MMP-2 with little inhibition of MMP-1, -3, -8 and -9. Chronic intraperitoneal administration of compound 20 in a murine model of melanoma markedly inhibits lung metastasis [385]. Compound 21 (*cis*-2-Aminocyclohexylcarbamoylphosphonic acid, *cis*-ACCP) was introduced as a carbamoyl phosphonate MMPI that targets MMP-2 and -9, but spares MMP-1, -3, -8, -12 and -13. Compound 21 dose-dependently inhibits cell invasion in a Matrigel assay and prevents tumor colonization in the murine melanoma model, and reduces tumor growth and metastasis in the more aggressive murine model developed by implantation of human prostate tumor cells in immunodeficient mice. These ZBGs have the advantage of being water soluble at physiological pH, show efficacy both orally and intraperitoneally, and are not acutely toxic at the concentrations used in the murine models [384].

Other ZBGs have been developed to improve selectivity, bioavailability, and pharmacokinetics, and include oxygen, nitrogen, and sulfur donor–atom ligands and monodentate, bidentate, and tridentate chelators.

### Nitrogen-Based ZBGs

Nitrogen-based ZBGs (ZBG10–16) have binding preference to late transition metals and improved selectivity towards Zn<sup>2+</sup>-dependent enzymes [386, 387]. An example of these ZBGs is compound 22, a modest inhibitor of MMP-9 that does not inhibit MMP-1, -2 or -12. The most extensively studied nitrogen-based ZBGs are the pyrimidine-2,4,6-trione and dionethione compounds. Pyrimidinetrione-based inhibitors have 100-fold selectivity for MMP-13 over MMP-2, -8 and -12 [388]. The pyrimidine-2,4,6-trione group is a known constituent of several FDA-approved drugs such as barbiturates, and therefore its metabolic disposition and bioavailability have been well-studied [389]. The pyrimidine-2,4,6-trione MMPIs have been optimized for gelatinase specificity and as anticancer drugs [390] and also to inhibit MMP-13 as part of the development of anti-osteoarthritis drugs [391–393].

Other MMPIs with high selectivity for specific MMPs include compound 23 (Ro 28-2653) with high selectivity for MMP-2, MMP-9, and MT1-MMP [394]. Ro 28-2653 has been evaluated for its anti-invasive, anti-tumorogenic, and anti-angiogenic activity and has shown inhibition of chemoinvasion by 85% at concentrations as low as 10 nM and anti-cancer efficacy in several *in vitro* and *in vivo* models [395]. In a rat model of tumorigenesis, Ro 28-2653 showed marked reduction in tumor growth and spread, and angiogenesis, with very few side effects [395]. Another compound Ro 32-3555 (trocade) has broad specificity for the collagenases MMP-1, -8 and -13 with affinity and IC<sub>50</sub> in the nM range and good oral bioavailability, making it a potential therapeutic drug for rheumatoid and osteoarthritis [396]. Ro 32-3555 also inhibits MMP-3, and the gelatinases MMP-2 and -9.

### Heterocyclic Bidentate ZBGs

A series of heterocyclic bidentate chelators ZBG20–30 was developed as alternative ZBGs and MMPIs [397]. These ZBGs have some features in common with hydroxamic acids but with better biostability and tighter Zn<sup>2+</sup> binding due to ligand rigidity and, in some cases, the incorporation of sulfur donor atoms [398, 399]. These ZBGs inhibited MMP-1, -2 and -3 with greater potency than acetohydroxamic acid [397], and low cell toxicity [400]. Compound 25 (AM-6) is a pyrone-based inhibitor with greater selectivity toward MMP-3 over MMP-1 and -2 [401]. Compound 26 (1,2 -HOPO-2) inhibits MMP-12 at low concentrations, and is a potent inhibitor of MMP-2, -3 and -8, but less effective against MMP-1, -7, -9 and -13 [402]. In an *ex vivo* rat heart model of ischemia and reperfusion, hearts treated with compound 26 recovered more than 80% of their contractile function compared with 50% of untreated hearts [402]. Compound 30 (AM-2) inhibits MMP-3 with an IC<sub>50</sub> of 240 nM [401, 402], while the structural isomer compound 32 (hydroxypyrrone 9b) shows weaker inhibition (~30%) at concentrations as high as 100 μM [403].

### Non-Zinc-Binding MMPIs

Because the Zn<sup>2+</sup> binding site is the most conserved feature in all MMPs, eliminating or minimizing the interaction of an MMPI with the catalytic Zn<sup>2+</sup> could improve MMP selectivity [354, 404-409] (Table 4). Several non-Zn<sup>2+</sup>-binding MMPIs show a noncompetitive inhibition mechanism [408]. Nearly all non-Zn<sup>2+</sup>-binding MMPIs show high MMP-13 selectivity and effectiveness in the treatment of animal models of osteoarthritis [404, 410]. Compound 37 inhibits MMP-13 but does not appreciably inhibit MMP-1, -2, -3, -7, -8, -9, -12, -13, -14 or -17. Binding of these inhibitors may rigidify the MMP active site into a specific conformation that is less conducive for substrate binding. The flexibility of MMP-13, relative to other MMPs, may allow for a favorable conformation that is not accessible in other MMPs [406, 410].

While these non-Zn<sup>2+</sup>-binding MMPIs are potent and selective, their hydrophobicity is critical in maintaining their protein–inhibitor interaction and high potency, but also results in poor water solubility. To improve the solubility without affecting potency, derivatives were developed to modify the solvent-exposed portions of the molecule and improve the compound solubility while maintaining the hydrophobic core structures and drug properties [404, 407].

Preclinical studies with compound 37 have shown encouraging results in models of osteoarthritis. Compound 37 has an efficacy at doses as low as 0.1 mg/kg in MMP-13-induced rat model of cartilage knee joint damage. Also, oral administration of compound 37 twice daily at 30mg/kg resulted in a 68% reduction in the cartilage lesion area of tibial plateaus in rats with surgically induced cartilage knee damage. When the rat joints were subsequently examined for evidence of fibroplasias and expanded inner synovial lining, which are indicative of musculoskeletal syndrome, fibroplasias were absent in rats treated



with compound 37, but present in rats treated with broad-spectrum MMPi. These highly selective MMPi may minimize the side effects associated with broad-spectrum MMPi, although it is not clear if their selectivity is due to their non-ZBG properties or other factors [406, 410].

### Mechanism-Based MMPi

Compound 40 (SB-3CT) was introduced as the first mechanism-based inhibitor of MMPs. SB-3CT binds in the active site and forms a covalent bond with the substrate protein upon activation by  $Zn^{2+}$  coordination [354] (Table 4). The formation of the covalent bond impedes inhibitor dissociation as compared to the traditional chelating competitive inhibitors. This reduces the rate of catalytic turnover and decreases the amount of MMPi needed to saturate the enzyme active sites. SB-3CT is a selective inhibitor of MMP-2 and -9. The structure of SB-3CT is relatively simple, as reflected by its low molecular weight. The mechanism of inhibition of SB-3CT is similar to that of a “suicide substrate” in which a functional group is activated, leading to covalent modification of the enzyme active site [411]. SB-3CT exhibits slow-binding kinetics with MMP-2, -3 and -9, with a time scale for establishing equilibrium between the enzyme and inhibitor and the enzyme–inhibitor complex in the order of seconds to minutes. Although binding rate can vary in speed, slow-binding inhibition is characterized by slow dissociation rates [412]. Progress curves, which display the enzyme activity of MMP-2, -9 and -3 with SB-3CT over time, are non-linear [411]. The curves show that the initial enzyme rate is not maintained and is instead reduced to a new “steady-state rate” of MMP activity. This indicates a slow-binding mechanism of inhibition, characteristic of an interaction between an enzyme and an inhibitor that resists dissociation. The selectivity of SB-3CT towards MMP-2 and -9 may be related to its slow-binding mechanism nearly irreversible inhibition. Following SB-3CT induced 95% inhibition, MMP-2 regains 50% activity only after 3 days with dialysis, but this slight reversibility may distinguish it from a true suicide inhibitor, which operates strictly by an irreversible mechanism [411, 412].

In preclinical studies, SB-3CT has shown anti-cancer effects in T-cell lymphoma and prostate cancer models [413–415] and reduced the invasion ability of human prostate cancer cells in *in vitro* Matrigel tests [414]. In a mouse model of T-cell lymphoma, SB-3CT administered at 5–50 mg/kg/d causes dose-dependent reduction in the number of liver metastases [416]. At 50 mg/kg/d, SB-3CT inhibited liver metastases by 73% and reduced the colony size of the metastases, while the broad-spectrum inhibitor batimastat increased metastasis. SB-3CT also showed promising results in a bone metastasis model of prostate cancer demonstrating reduced tumor growth and angiogenesis [414]. Also, SB-3CT provides neuronal protection in a murine stroke model [417]. In mice treated with SB-3CT either prior to or 2 h following cerebral ischemia induced by right middle cerebral artery occlusion, the infarct volume is decreased to 30% of the control. Administration of SB-3CT is protective up to 6 h after the ischemic event in mice. Also, neurological behavioral scores evaluated 24 h after reperfusion show that SB-3CT-treated mice exhibit marked improvement as compared to control mice, and the improvement is correlated with the infarct volume.

Although SB-3CT has marked *in vivo* activity, it undergoes rapid metabolism in mice [418], suggesting that a metabolite of the parent compound may be responsible for the *in vivo* activity [419]. Analysis of the different MMPi metabolites has led to the design of derivatives that have better *in vivo* stability and longer systemic effects [420]. Compound 43 is a more potent inhibitor of MMP-2, -3, -7, -9 and -14 than SB-3CT, and demonstrates slow-binding kinetics with MMP-2, -9 and -14 [420]. Compound 45 is a slow-binding and more potent inhibitor of MMP-9 than MMP-2, but a competitive inhibitor of other MMPs.

Metabolites of compound 45 are 75% more stable than those of SB-3CT, resulting in prolonged systemic effects [420].

Mechanism-based slow-binding inhibitors such as SB-3CT and its successors have shown clinical promise. Other types of covalent modification in the active site may help in the design of more selective MMPi [354]. However, even with the marked improvements in the design of MMPi, therapeutic inhibition of MMPs is still a challenge and the antibiotic doxycycline remains the only FDA-approved MMPi [362, 363, 421, 422]. One limitation of many MMPi is that they may cause musculoskeletal side effects including joint stiffness, pain, inflammation and tendinitis [354, 363, 423, 424].

### Other MMP Inhibitors

Tetracyclines such as doxycycline and monocyclusins are known antibiotics that also inhibit MMPs. Tetracyclines bind the divalent  $Zn^{2+}$  and  $Ca^{2+}$  ions and have an effect on MMP gene transcription [333]. Clinical trials have evaluated the tetracycline analog COL-3, a potent inhibitor of MMP-2 and -9, in patients with advanced metastatic soft tissue sarcoma, but minimal changes in tumor progression or survival have been observed [425].

In VVs organ cultures, statins such as simvastatin and pravastatin suppress the production and activity of MMP-9, but not MMP-2, in a dose-dependent manner. However, the doses of simvastatin and pravastatin required to influence MMP-9 are much higher than the peak plasma concentrations achieved after a typical oral dose [426].

Studies have evaluated the effects of glycosaminoglycan sulodexide, an antithrombotic/profibrinolytic drug, on the activity and release of MMPs in human blood. In a prospective non-randomized study, zymography and ELISA were used to analyze the *in vitro* effects of sulodexide on pro-enzyme, complexed, and active MMP forms in plasma and serum from 60 healthy donors, and in U-937 leukemia cell line [427]. Pro- and complexed forms of MMP-9 were markedly affected by sulodexide treatment, with marked decrease in MMP-9 secretion from white blood cells in a dose-dependent manner, without any displacement of MMP prodomains. The mechanism of action of sulodexide likely involves direct inhibition of the active  $Zn^{2+}$  binding site of the proMMP-9 molecule [428]. The levels and zymographic profile of MMP-2 did not show marked differences among samples and with sulodexide treatment. The reduced release of MMP-9 forms from leukocytes and inhibition of proteolytic activity by sulodexide treatment support that inhibitors of MMP-9 activity may provide a therapeutic tool for the underlying pathological changes in ECM, and offer novel pharmacologic approaches for chronic inflammatory vascular diseases associated with enhanced MMP activation in blood and limbs, including VVs and CVD [427].

Topical products have been developed to modify the MMP/TIMP balance and improve wound healing. Hydroactive wound dressings containing polyacrylate superabsorber particles inhibit MMPs, including MMP-2 and -9, by competing for  $Zn^{2+}$  and  $Ca^{2+}$  [429]. Also, in a randomized controlled trial, oxidized regenerated cellulose/collagen matrix (ORC/collagen matrix) accelerate the healing rate in venous leg ulcer [430]. ORC/collagen matrix modifies wound microenvironments by binding and inactivating excess proteases. Exudates obtained from CVU treated with ORC/collagen matrix showed marked reduction in elastase, plasmin and gelatinase activity as compared with those treated with hydrocolloid dressings [431]. Although few clinical trials have been conducted, a 12-week multi-centred randomized trial of 117 patients comparing the efficiency of nanooligosaccharide factor (NOSF) and ORC in the local management of CVU demonstrated that NOSF was superior wound healing [432]. A case report has shown that combining topical tacrolimus and oral doxycycline successfully treated 3 resistant venous ulcers [433].

Monoclonal antibodies target specific MMPs with high affinity. The monoclonal antibodies REGA-3G12 and REGA-2D9 are specific for MMP-9 and do not cross-react with MMP-2. MMP inhibition by REGA-3G12 does not involve the Zn<sup>2+</sup> binding domain or the fibronectin region, but instead binds to the catalytic domain [434]. REGA-1G8 is not as specific, and cross-reacts with serum albumin. In addition to inhibiting MMPs, monoclonal antibodies are useful to detect MMPs in the body fluids and tissues of patients with inflammation or cancer [434–436].

Small interfering RNA (siRNA) inhibits the transcriptional product of specific MMPs with efficient inhibition of the target molecule. Research in cancer therapy using siRNA is evolving. Adenovirus-mediated siRNA 21 base-pair for MMP-9 caused a dose-dependent cell cycle arrest of medulloblastoma cells and inhibited tumor growth [437]. Other tumor cell lines such as human chondrosarcoma have shown decreased tumor invasion of type I collagen matrix *in vitro* by siRNA directed inhibition of MMP-1 [438]. Gene silencing using siRNA targeted for MT1-MMP to reduce activation of proMMP-2 in bone marrow derived stromal cells attenuated the induction of cell necrosis [439]. The use of siRNA allows for precise and specific gene knockdown and minimizes the potential for non-specific inhibition by longer double stranded antisense RNAs. Because siRNA require transfection into the cell, the efficiency of transfection and duration of siRNA inhibition may vary, and may result in only transient gene knockdown. Also, siRNA may cause nonspecific activation of innate immune responses or off-targeting of MMP genes and inadvertent downregulation of other genes [440].

### Conclusions and Perspectives

CVD including VVs is not a minor disease with only cosmetic and social implications. If untreated, VVs could lead to thrombophlebitis and CVI with skin changes and chronic leg ulcer. The pathogenesis of VVs is a complex multifactorial process involving both valve insufficiency and vein wall changes. VVs can form in any vein segment of the lower extremity, and tributaries that connect to axial veins may show varicosities without any reflux in the communicating vein. Also, normal vein segments adjacent to VVs have the same biochemical properties as the VVs wall, supporting the role of vein wall changes in the pathogenesis of VVs. VVs pathology may affect not only the lower extremity veins, but also other tissues, suggesting that VVs may be a localized manifestation of a systemic venous disease.

Most of the current therapies aim at enhancing the cosmetic appearance and comfort of the patient's symptoms, and interventional and surgical approaches in advanced cases of CVD. Because current treatment strategies are directed towards the manifestations or complications rather than the cause of CVD, they often show variable success and high recurrence rates. MMPs play an important role in ECM metabolism and other physiological processes, and increased MMP expression/activity has been associated with connective tissue remodeling in autoimmune disease. MMPs and their naturally occurring TIMPs have also been implicated in several degenerative and vascular disease including VVs. The abundance of MMPs in venous tissue supports a role in VVs, and MMPs may have distinct roles at different stages of CVD. In the early stages of CVD, MMPs may cause venous dilation by inducing hyperpolarization of the vein wall, and inhibiting balanced production of collagen subtypes. In later stages of CVD, MMPs may alter the vein wall matrix composition leading to further venous dilation and tortuosity.

The role of MMPs is often examined by measuring their plasma levels, and tissue expression and activity. Molecular imaging of MMP function *in vivo* is an emerging area of research. In murine tumors a near infrared fluorescent MMP substrate detects the activity of MMPs *in vivo* [441]. Similar approaches have identified active cathepsin B in atherosclerotic mice

and increased MMP-2 and -9 activities after murine myocardial infarction [442]. Although this technology presents challenges such as limited tissue penetration and autofluorescence of elastin in arteries, initial results have provided proof of principle for proteinase imaging *in vivo*.

Several MMPIs have been used as investigative tools of the role of MMPs in physiological and pathological processes. MMPIs have also been examined as potential therapeutic tools in cancer, arthritis, autoimmune and vascular disease. Although several MMPIs have been developed, their limited specificity and oral bioavailability as well as potential cytotoxicity prevented their further development for clinical use. In animal models, MMPIs prevent the development and progression of early disease, but show little effect in advanced disease. Also, clinical trials using synthetic MMPIs in cancer patients have shown little success, mainly due to the lack of efficacy and untoward side effects including musculoskeletal pain and tendonitis [443]. In many studies MMPIs was used as single agent therapy in patients with advanced disease, which may explain the poor performance of MMPIs in clinical trials [423].

Several approaches have been suggested to overcome the drawbacks of classic MMPIs. Specific antibody fragments could target the MMP active site in a more specific way than chemical inhibitors and could identify sites on the MMP molecule that determine their substrate specificity and extracellular location. Understanding of the nature of these interactions will allow the development of specific inhibitors of MMP-substrate binding or fragment antibodies that target these interactions. The availability of specific MMPIs may provide a new tool to limit the progression and complications of CVD and reduce the recurrence of VVs

Several other strategies may potentially downregulate MMPs. Both the intracellular signaling pathways and the downstream transcription factors which induce gene expression are being studied. Blockade of MAPK pathways, NF- $\kappa$ B or AP-1 have shown some efficacy *in vitro* or in animal models of arthritis [444]. The use of biological reagents to block inflammatory cytokines also reduces MMP expression in many tissues. Tetracyclines are weak inhibitors of MMP, but they could influence MMP synthesis and have been tested successfully in rheumatoid arthritis [445]. Gene therapy has been successful in animal models, and overexpression of TIMPs may have future application after overcoming the problems of safe and efficient delivery of genes into target tissues [446].

Certain gene mutations and variants may play a role in CVD. Identifying the genetic basis of VVs will not only help elucidate the molecular abnormalities involved, but also identify individuals at risk and the most appropriate prevention and treatment strategies. Also, while significant progress has been made in identifying the role of MMPs in VVs, their upstream regulation and downstream signaling at the different stages of CVD need to be examined. Understanding of the molecular basis of VVs formation and MMPs-induced changes in ECs and VSM function and vein wall remodeling will provide valuable information on the mechanisms involved in CVD development and progression. Also, the interaction of MMPs with other proteolytic enzymes such as ADAM and ADAMTS in the pathogenesis of VVs and stretch induced increase in AP-1, which triggers varicose remodeling through MMPs and VSMC hyperplasia in the vein wall need to be examined. Further research of the molecular mechanisms of VVs and CVD would help identify possible targets for genetic manipulation and more efficient pharmacologic interventions of the disease.

## Acknowledgments

This work was partly supported by grants from National Heart, Lung, and Blood Institute (HL-65998 and HL-98724).

## List of abbreviations

|                                |  |
|--------------------------------|--|
| <b>AP-1</b>                    | activator protein-1                          |
| <b>CVD</b>                     | chronic venous disease                       |
| <b>CVI</b>                     | chronic venous insufficiency                 |
| <b>EC</b>                      | endothelial cell                             |
| <b>ECM</b>                     | extracellular matrix                         |
| <b>EMMPRIN</b>                 | extracellular MMP inducer                    |
| <b>HUVECs</b>                  | human umbilical vein endothelial cells       |
| <b>IVC</b>                     | inferior vena cava                           |
| <b>MAPK</b>                    | mitogen-activated protein kinase             |
| <b>MMP</b>                     | matrix metalloproteinase                     |
| <b>MMPI</b>                    | matrix metalloproteinase inhibitor           |
| <b>Phe</b>                     | phenylephrine                                |
| <b>ROS</b>                     | reactive oxygen species                      |
| <b>TIMP</b>                    | tissue inhibitor of matrix metalloproteinase |
| <b>TNF-<math>\alpha</math></b> | tumor necrosis factor- $\alpha$              |
| <b>VSMC</b>                    | vascular smooth muscle cell                  |
| <b>VUWF</b>                    | venous ulcer wound fluid                     |
| <b>VVs</b>                     | varicose veins                               |

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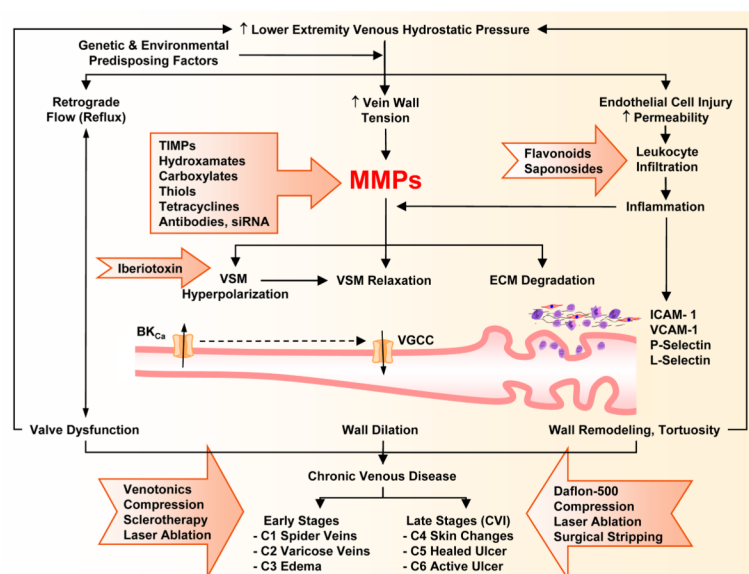


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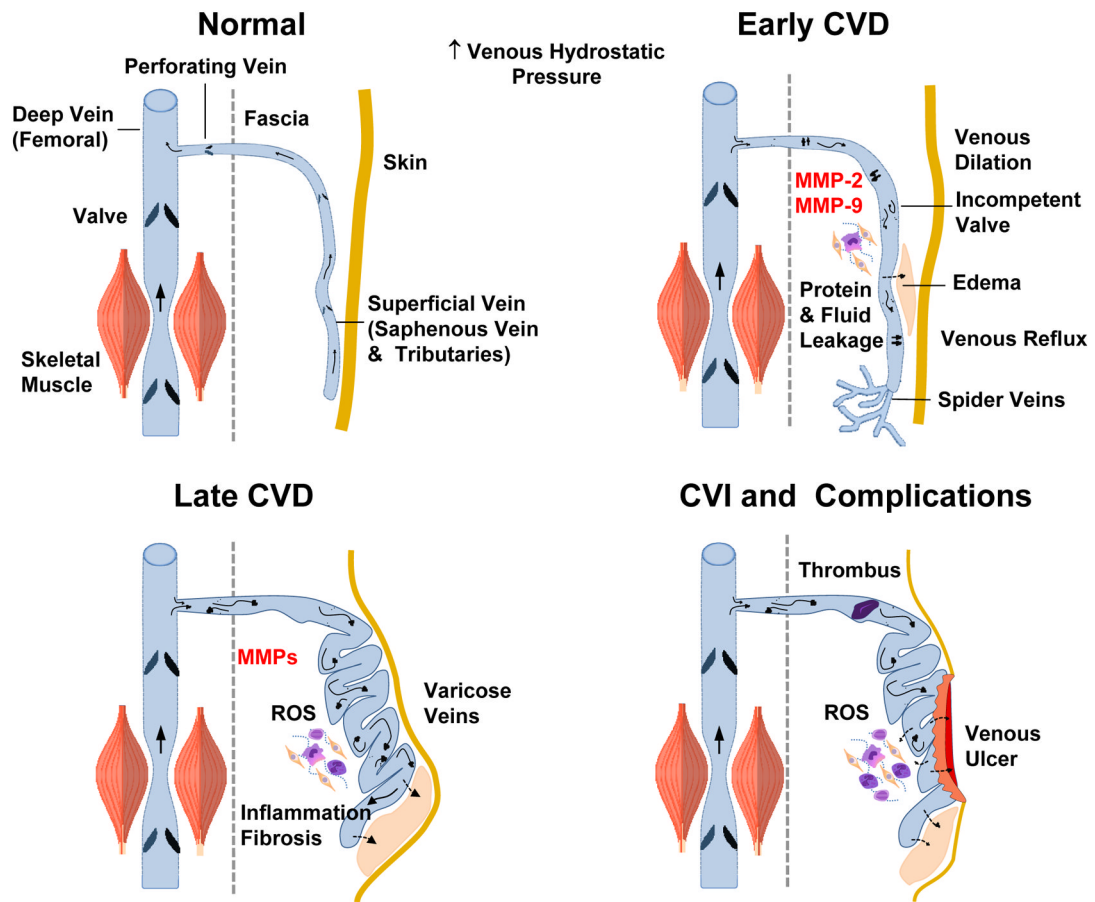
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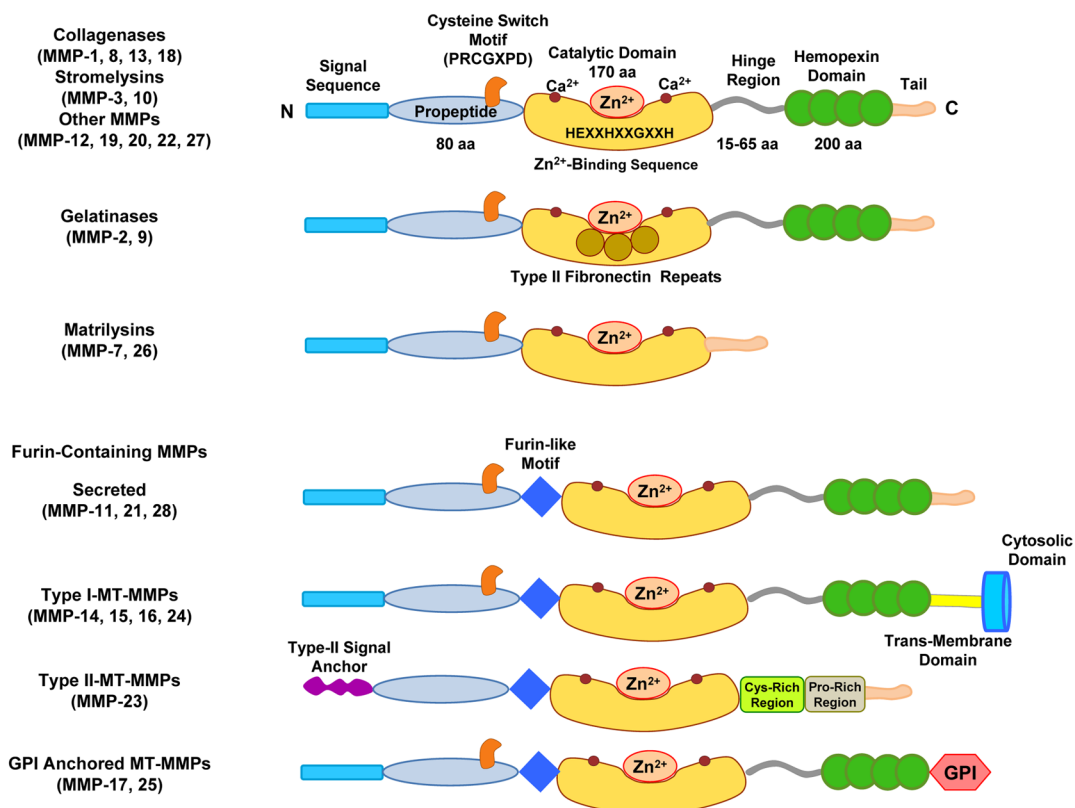


**Fig. 1.** Pathophysiology and management of CVD. In the presence of genetic and environmental risk factors, increases in lower extremity venous hydrostatic pressure could cause valve dysfunction and venous reflux as well as increased wall tension, and increased expression/activity of MMPs. In addition to degradation of extracellular matrix (ECM) proteins, MMPs may cause venous smooth muscle (VSM) hyperpolarization and activation of  $K^+$  Channels, which in turn could cause inhibition of  $Ca^{2+}$  entry through voltage-gated  $Ca^{2+}$  channels (VGCC), VSM relaxation, vein wall dilation, and early manifestations of chronic venous disease (CVD). Persistent increases in venous pressure could cause ECs injury, leukocyte infiltration and inflammation of the vein wall, which could further increase MMPs expression/activity and lead to vein wall remodeling and tortuosity and late manifestations of chronic venous insufficiency (CVI). Several pharmacological and surgical approaches (presented in block arrows) are used for treatment of early and late manifestations of CVD. Different classes of MMP inhibitors may provide a new approach for management of CVD.





**Fig. 2.** Early and late manifestations of CVD. In normal veins with intact valves, blood flows from superficial to deep femoral vein. Increases in lower extremity venous hydrostatic pressure and vein wall tension are associated with increased MMP expression/activity, venous dilation and venous reflux and early manifestations of CVD such as edema and spider veins. Late stages of CVD are associated with further increases in MMPs, varicose veins, edema, vein tissue remodeling, inflammation and fibrosis. CVI is complicated by thrombophlebitis, further increases in ROS, and venous wound leg ulcer.

**Fig. 3.**

Major classes and structure of matrix metalloproteinase. Typically, MMPs consist of a propeptide of about 80 amino acids, a catalytic metalloproteinase domain of about 170 amino acids, a linker peptide (hinge region) of variable lengths and a hemopexin domain of about 200 amino acids. The catalytic domain contains the  $\text{Zn}^{2+}$  binding motif HEXXHXXGXXH. Matrilysins are exceptions as they lack the linker peptide and the hemopexin domain. Membrane-bound MMPs (MT-MMPs) have a furin-like proprotein convertase recognition sequence at the C-terminus of the propeptide and some of them have a glycosylphosphatidylinositol (GPI) anchor.

**Table 1**

CEAP classification of clinical stages of CVD

| Stage | Clinical Signs  |
|-------|---|
| C0    | No visible or palpable signs of venous disease                                  |
| C1    | Telangiectasia, reticular veins, or spider veins                                |
| C2    | Varicose veins. Distinguished from reticular veins by a diameter of $\geq 3$ mm |
| C3    | Edema   |
| C4    | Changes in skin and subcutaneous tissue secondary to CVD                        |
| C4a   | Skin pigmentation/eczema  |
| C4b   | Lipodermatosclerosis or atrophie blanche  |
| C5    | Healed venous ulcer   |
| C6    | Active venous ulcer   |

Table 2

Members of the MMP family and their substrates

| MMP (Other Name) Chromosome Location                       | MW KDa Proform | MW KDa Active Form | Tissue Distribution   | Collagen Substrates                   | Non-Collagen ECM Substrates  | Non-Structural ECM Component Substrates  |
|--|----------------|--------------------|---|---------------------------------------|--|--|
| <b>Collagenases</b><br>MMP-1 (Collagenase-1) 11q22.3       | 55             | 45                 | Fibroblasts, interstitial, tissue collagenase                                     | I, II, III, VII, VIII, X, and gelatin | Aggrecan, casein, nidogen, serpins, versican, perlecan, proteoglycan link protein, tenascin-C                                  | $\alpha$ 1-antichymotrypsin, $\alpha$ 1-antitrypsin, $\alpha$ 1-proteinase inhibitor, IGFBP-3, IGFBP-5, IL-1 $\beta$ , L-selectin, ovostatin, recombinant TNF- $\alpha$ peptide, SDF-1   |
| MMP-8 (Collagenase-2) 11q22.3                              | 75             | 58                 | Neutrophil, or PMNL collagenase   | I, II, III, V, VII, VIII, X           | Aggrecan, laminin, nidogen   | $\alpha$ 2-antiplasmin, pro-MMP-8  |
| MMP-13 (Collagenase-3) 11q22.3                             | 60             | 48                 | VV's, SMC, preeclampsia, breast cancer  | I, II, III, IV                        | Aggrecan, fibronectin, laminin, perlecan, tenascin   | Plasminogen activator 2, pro-MMP-9, pro-MMP-13, SDF-1  |
| <b>Gelatinases</b><br>MMP-2 (Gelatinase-A) 16q13-q21       | 72             | 66                 | Aortic aneurysm, VV's   | I, II, III, IV, V, VII, X, XI         | Aggrecan, elastin, fibronectin, laminin, nidogen, proteoglycan link protein, versican  | Active MMP-9, active MMP-13, FGF R1, IGF-BP3, IGF-BP5, IL-1 $\beta$ , recombinant TNF- $\alpha$ peptide, TGF- $\beta$  |
| MMP-9 (Gelatinase-B) 20q11.2-q13.1                         | 92             | 86                 | Aortic aneurysm, VV's   | IV, V, VII, X, XIV                    | Fibronectin, laminin, nidogen, proteoglycan link protein, versican   | CXCL5, IL-1 $\beta$ , IL2-R, plasminogen, pro-TNF $\alpha$ , SDF-1, TGF- $\beta$   |
| <b>Stromelysins</b><br>MMP-3 (Stromelysin-1) 11q22.3       | 57             | 45                 | VSMC, coronary artery disease, hypertension, tumor invasion, synovial fibroblasts | II, III, IV, IX, X, XI                | Aggrecan, casein, decorin, elastin, fibronectin, laminin, nidogen, perlecan, proteoglycan, proteoglycan link protein, versican | $\alpha$ 1-antichymotrypsin, $\alpha$ 1-proteinase inhibitor, antithrombin III, E-cadherin, fibronogen, IGF-BP3, L-selectin, ovostatin, pro-HB-EGF, pro-IL-1 $\beta$ , pro-MMP-1, pro-MMP-8, pro-MMP-9, pro-TNF $\alpha$ , SDF-1 |
| MMP-10 (Stromelysin-2) 11q22.3                             | 57             | 44                 | Atherosclerosis, uterine, preeclampsia, arthritis, carcinoma cells                | III, IV, V                            | Fibronectin, laminin, nidogen  | Pro-MMP-1, pro-MMP-8, pro-MMP-10   |
| <b>Matrilysins</b><br>MMP-7 (Matrilysin-1) 11q21-q22       | 28             | 19                 | Uterine   | IV, X                                 | Aggrecan, casein, elastin, enactin, laminin, proteoglycan link protein   | $\beta$ 4 integrin, decorin, defensin, E-cadherin, Fas-L, plasminogen, pro-MMP-2, pro-MMP-7, pro-MMP-8, pro-TNF $\alpha$ , transferrin, and syndecan $\alpha$ 2-antiplasmin  |
| MMP-26 (Matrilysin-2, endometase) 11p15                    | 28             | 19                 | Breast cancer cells, human endometrial tumor                                      | IV, gelatin                           | Casein, fibrinogen, fibronectin  | Fibrin, fibronectin Pro-MMP-2 $\beta$ 1-proteinase inhibitor   |
| <b>Membrane-Type</b><br>MMP-14 (MT1-MMP) Chr: 14-14q11-q12 | 66             | 56                 | Human fibroblasts, SMC, VSMC, uterine, angiogenesis                               | I, II, III                            | Aggrecan, dermatan proteoglycan, fibrin, fibronectin, laminin, nidogen, perlecan, tenascin, vitronectin                        | $\alpha$ 3 integrin, CD44, gC1qR, pro-MMP-2, pro-MMP-13, pro-TNF $\alpha$ , SDF-1, tissue transglutaminase   |
| MMP-15 (MT2-MMP) 16q13                                     | 72             | 50                 | Human fibroblasts, leukocytes, preeclampsia                                       | I                                     | Aggrecan, fibronectin, laminin, nidogen, perlecan, tenascin, vitronectin   | Pro-MMP-2, pro-MMP-13, tissue transglutaminase   |

| MMP (Other Name) Chromosome Location                 | MW KDa Proform | MW KDa Active Form | Tissue Distribution                                | Collagen Substrates | Non-Collagen ECM Substrates                                   | Non-Structural ECM Component Substrates                          |
|--|----------------|--------------------|--|---------------------|---|--|
| MMP-16 (MT3-MMP) 8q21.3                              | 64             | 52                 | Human leukocytes, angiogenesis                     | I                   | Aggrecan, casein, fibronectin, laminin, perlecan, vitronectin | Pro-MMP-2, pro-MMP-13  |
| MMP-24 (MT5-MMP) 20q11.2                             | 57             | 53                 | Leukocytes   | None identified     | Chondroitin sulfate, dermatin sulfate, fibronectin            | Pro-MMP2, pro-MMP-13   |
| <b>Other MMPs</b><br>MMP-11 (Stromelysin-3) 22q11.23 | 51             | 44                 | Angiogenesis, uterine                              | Does not cleave     | Laminin   | $\alpha$ 1-antitrypsin, $\alpha$ 1-proteinase inhibitor, IGFBP-1 |
| MMP-12 (matrilysin) 11q22.3                          | 54             | 45-22              | Macrophages  | IV                  | Elastin   | Plasminogen  |
| MMP-17 (MT4-MMP) 12q24.3                             | 57             | 53                 | Brain specific cerebellum, breast cancer           | Gelatin             | Fibrin  |  |
| MMP-18 (Xenopus Collagenase-4) 12q14                 | 70             | 53                 | Xenopus (amphibian)                                | I, gelatin          |   |  |
| MMP-19 (RASI-1) Chr: 12-12q14                        | 54             | 45                 | Liver  | I, IV, gelatin      | Aggrecan, casein, fibronectin, laminin, nidogen, tenascin     |  |
| MMP-20 (Enamelysin) Chr: 11-11q22.3                  | 54             | 22                 | Tooth enamel                                       | V                   | Aggrecan, amelogenin, cartilage oligomeric protein            |  |
| MMP-21 (X-MMP) 10q26.13                              | 62             | 49                 | Macrophages, fibroblasts, human placenta           |                     |   | $\alpha$ 1-antitrypsin   |
| MMP-22 (C-MMP) 1p36.3                                | 51             |                    | Chicken fibroblasts                                | Gelatin             |   |  |
| MMP-23 (CA-MMP)                                      | 28             | 19                 | Ovary, testis, prostate                            | Gelatin             |   |  |
| MMP-25 (Leukolysin, MT6-MMP) 16p13.3                 | 34             | 28                 | Leukocytes, anaplastic astrocytomas, glioblastomas | IV, gelatin         |   | Fibrin, fibronectin, pro-MMP-2                                   |
| MMP-28 (Epilysin) 17q21.1                            | 56             | 45                 | Skin keratinocytes                                 |                     | Casein  |  |

MW, Molecular mass; PMNL, polymorphonuclear leukocytes; SDF-1, stromal cell-derived factor-1; VSMC, vascular smooth muscle cell; VV's, varicose veins. References: [55, 57]



**Table 3**

## Management of varicose veins

| Treatment Strategy                               | Specific Treatment   | Reference   |
|--|--|---|
| Compression Therapy<br>Pharmacological Treatment | Graduated compression stockings  | [297]   |
| Pharmacological Treatment                        | <ul style="list-style-type: none"> <li>- <math>\alpha</math>-benzopyrones (Coumarins)</li> <li>- <math>\gamma</math>-benzopyrones (Favonoids)</li> <li>- Saponosides (Escin, horse chestnut seed extract)</li> <li>- Plant extracts (Blueberry and grape seed, Ginkgo biloba)</li> <li>- Daflon-500</li> <li>- Venoruton (Oxerutin)</li> <li>- Others: Pentoxifylline, red vine leaves (AS-195), prostaglandin E1</li> </ul> | <ul style="list-style-type: none"> <li>- [447]</li> <li>- [304]</li> <li>- [306]</li> <li>- [311]</li> <li>- [312]</li> <li>- [309]</li> <li>- [317, 319, 320]</li> </ul> |
| Sclerotherapy                                    | Sodium tetradecyl sulfate (STS), sodium morrhuate with polidocanol   | [321]   |
| Surgical Intervention                            | <ul style="list-style-type: none"> <li>- Endovenous Ablation (Radiofrequency or infrared laser)</li> <li>- Surgical stripping</li> <li>- Phlebectomy (Stab phlebectomy, transilluminated power phlebectomy)</li> </ul>   | <ul style="list-style-type: none"> <li>- [325]</li> <li>- [329]</li> <li>- [330]</li> </ul>   |

Table 4

MMP inhibitors, their selectivity to MMPs ( $IC_{50}$  or  $K_i < 1$  nM to  $10 \mu\text{M}$ ), and their use in experimental trials.

| MMPi                                  | MMP Specificity, $IC_{50}$ or $K_i$ |                |                  |                     |                    | Experimental Trials  | Ref.       |
|---------------------------------------|-------------------------------------|----------------|------------------|---------------------|--------------------|--|------------|
|                                       | <1 nM                               | 1–10 nM        | 11–100 nM        | 0.1–1 $\mu\text{M}$ | 1–10 $\mu\text{M}$ |  |            |
| ZBG 1 Hydroxamates Batimastat, BB-94) |                                     | MMP-1, 2, 8, 9 | MMP-3            |                     |                    |  | [152]      |
| 2 Carboxylates, Prinomastat, AG3340   | MMP-2, 3, 9, 13, 14                 | MMP-1          | MMP-7            |                     |                    | O <sub>2</sub> -induced retinal neovascularization, neuronal hypoxic injury, ventilator-induced lung injury, lung cancer, uveal melanoma, gliomas, prostate cancer | [448–453]  |
| 3 Hydrazides                          |                                     | MMP-1          |                  |                     |                    |  | [454]      |
| 4 Sulfonyl-hydrazides RS-104966       | MMP-13                              |                | MMP-1            |                     |                    |  | [191]      |
| 5                                     |                                     | MMP-13         | MMP-3, 8         | MMP-2               | MMP-7, 9, 14       | Osteoarthritis   | [455]      |
| 6                                     |                                     | MMP-11         |                  | MMP-3, 12           | MMP-1, 9, 14       | Chronic obstructive pulmonary disease  | [456]      |
| 7                                     | MMP-2                               | MMP-8, 9, 14   |                  | MMP-1, 3            | MMP-7              | Stop tumor invasion  | [359]      |
| 8 Mercaptosulfides                    |                                     | MMP-3          |                  | MMP-2               | MMP-9              | Chronic non-healing wounds   | [361]      |
| 9 Carbamoyl phosphonates              |                                     | MMP-2, 9       | MMP-1            | MMP-7               | MMP-3              |  | [382]      |
| 10                                    |                                     |                | MMP-2, 3         |                     | MMP-1              |  | [457]      |
| 12                                    |                                     |                |                  |                     | MMP-3              |  | [458]      |
| 16                                    | MMP-9                               | MMP-2          | MMP-1, 7, 14     |                     | MMP-3              |  | [383]      |
| 17                                    | MMP-8                               | MMP-2          | MMP-3            |                     |                    |  | [369]      |
| 18                                    |                                     | MMP-8          | MMP-2, 9, 13, 14 | MMP-1, 3            | MMP-7              | Acute liver disease, multiple sclerosis, breast cancer   | [459–461]  |
| 19                                    |                                     |                |                  | MMP-11              | MMP-2, 9, 13       |  | [462]      |
| 20                                    |                                     |                | MMP-2            |                     |                    | Melanoma   | [385]      |
| 21                                    |                                     |                |                  |                     | MMP-2              | Melanoma, prostate cancer  | [384]      |
| 22                                    |                                     |                |                  |                     | MMP-11             |  | [387, 397] |
| 23 Ro-28-2653                         |                                     | MMP-2, 14      | MMP-8, 9         |                     | MMP-3              | Anti-angiogenic and anti-invasive in tumor models  | [389, 395] |
| 24                                    | MMP-13                              |                |                  | MMP-2, 9, 12        |                    | Osteoarthritis   | [388]      |
| 25 Pyrone-Based AM-6                  |                                     |                | MMP-3            |                     |                    |  | [401]      |
| 26 1,2-HOPO-2                         |                                     |                | MMP-8, 12        | MMP-2, 3            | MMP-13             | Heart ischemia and reperfusion   | [402]      |

| MMPI                      | MMP Specificity, IC <sub>50</sub> or K <sub>i</sub> |              |              |                  | Experimental Trials  | Ref.           |
|---------------------------|---|--------------|--------------|------------------|--|----------------|
|                           | <1 nM   | 1–10 nM      | 11–100 nM    | 0.1–1 μM         |  |                |
| 27                        |   | MMP-2, 9, 13 | MMP-3        | MMP-1            | Brain edema following ischemia–reperfusion                                       | [402]          |
| 28                        |   |              | MMP-3, 9, 12 | MMP-2, 13        |  | [463]          |
| 29                        |   |              |              | MMP-8, 12        |  | [402]          |
| 30 AM-2                   |   |              | MMP-8, 12    | MMP-2            |  | [402]          |
| 31                        |   |              |              | MMP-2, 8, 12, 13 |  | [402]          |
| Non-ZBG 34                |   |              | MMP-13       |                  | Osteoarthritis   | [410]          |
| 35                        |   |              | MMP-12       | MMP-2, 8, 13     |  | [405]          |
| 36                        |   |              |              | MMP-2, 8, 13     |  | [405]          |
| 37                        | MMP-13  |              |              |                  | Osteoarthritis   | [410]          |
| 38                        |   |              |              | MMP-13           |  | [406]          |
| 39                        |   |              | MMP-13       |                  |  | [406]          |
| Mechanism-Based 40 SB-3CT |   |              | MMP-2        | MMP-9, 14        | Inhibits bone metastasis in prostate cancer, liver metastasis in T-cell lymphoma | [413–415, 418] |
| 42                        |   |              | MMP-2, 9     | MMP-14           |  | [411]          |
| 43                        |   | MMP-2        | MMP-14       | MMP-9            |  | [418]          |
| 45                        |   | MMP-9        | MMP-2        | MMP-3, 14        |  | [420]          |