Molecular targets in aortic aneurysm for establishing novel management paradigms

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Abstract: Aortic aneurysm (AA) is a lethal disease and presents a large challenge for surgeons in the clinic. Although surgical management remains the major choice of AA, operative mortality remains high. With advances in understanding of the mechanisms of AAs, molecular targets, such as matrix metalloproteinases (MMPs), D-dimer, and inflammation markers, including C-reactive protein, interleukins and phagocytes, are important in the pathology of development of AA. These markers may become important for improving the diagnostic quality and provide more therapeutic choices for treatment of AA. Although these new markers require long-term trials before they can be translated into the clinic, they can still be helpful in determining new directions. The main aim of this review is to discuss the current findings of molecular targets in progression of AA and discuss the potential application of these new targets for managing this disease.

Keywords: Aortic aneurysm (AA); molecular targets; inflammation; remodeling; nanomedicine

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Introduction

Aortic aneurysm (AA) is a potential lethal disease with an increasing incidence rate that reaches approximately 10.4/100,000 people each year (1,2). The general feature of AA is a lack of obvious evidence or clinical symptoms for determining its presence (3), and death due to ruptured AA is common (4,5). Progression of AA leads to unpredictable features of this disease and can lead to unexpected rupture. The annual incidence of aortic dissection and rupture is 3.5/100,000 patients (6). The various histological, anatomical, and clinical presentations need to be carefully considered before deciding on treatment options for AA. Surgical repair of AA is the main treatment with a mean 30-day mortality rate of 8.2% (95% confidence interval: 6.4–10.6) (7). In the clinic, physicians usually use medicine to delay the progression of AA, such as traditional betaadrenergic receptor blockers, when AA does not reach the standard for surgery. Recently, molecular targets of development of AA have been found with technological innovations. These targets include matrix metalloproteinases (MMPs), elastin-peptides (SEP), C-reactive protein, and PIIINP-collagen, which have provided a new direction for diagnosis and management of AA.

Pathophysiology of AA

The aorta is a heterogeneous vessel with different components in thoracic aortic aneurysm (TAA) and abdominal aortic aneurysm (AAA), as reported in

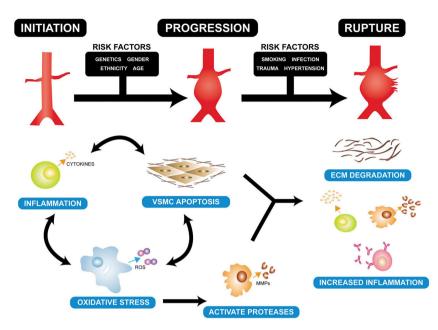


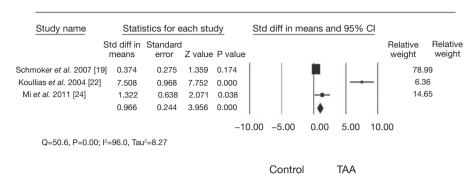
Figure 1 Schema showing the inflammation process in AA (10). AA, aortic aneurysm.

previous reviews (8,9). However, matrix remodeling and inflammation are similar between TAA and AAA. Formation of the AA is correlated with inflammatory infiltrates in apoptosis of vascular smooth muscle cells and extracellular matrix degradation (Figure 1) (10). Additionally, a change in matrix proteins and a variety of transforming growth factor (TGF)- β signals appear to be vital for development of TAA (9,11,12). The weaken of the aortic ultrastructure and increased the risk of dilatation, dissection, and rupture may be caused by this alteration (13,14). TGF- β can affect matrix degradation by regulating alternate pathways, especially in Marfan syndrome. A previous study showed that in patients with TAA, the TGF-B2 signal is decreased through cells expressing TGF-βRIIb and mutations in this receptor result in an increase in TGF- β 2 signal (15). This suggests that an increase in TGF-β signal may lead to development of a rtic pathogenesis. The TGF- β 2 signal finding has aroused a wide concern in the intracellular signaling pathway (16,17). Moreover, pro-inflammatory chemokines and cytokines expedite the inflammatory process. MMPs, which are a large family of enzymes, are derived from smooth muscle cell production and response to cyclic strain that can also progress degradation of the extracellular matrix (18). The function of increased MMP levels in the progression of AAs is important (19,20). In certain cases, an imbalance in medial expression of MMPs, specifically the gelatinases (MMP-2 and MMP-9) and tissue

inhibitors of metalloproteinases (TIMP-2 and TIMP-1), result in accelerated proteolytic degradation of elastin and collagen fibers (21,22). They have found that MMP-9 levels were increased in TAA (Figure 2) compared with controls and the levels of MMP-2 was no obviously change in TAA compared with controls (Figure 3). Additionally, TIMP-1 and TIMP-2 levels were significantly lower in TAA compared with controls (Figure 4). All studies showed that the ratio of MMP-9 to TIMP-1 was 3.7 times in TAA compared with controls. The ratio of MMP-9 to TIMP-2 in TAA was 26.5 times that of controls. Notably, MMP-2 and -9 have elastolytic and collagenolytic properties (23). Elastolytic and collagenolytic properties can increase stiffness of the thoracic aortic wall and decrease elastic ability, which lead to the appearance of AA. Therefore, inflammation, matrix degradation and remodeling weaken aortic tensile strength and lead to formation of AA.

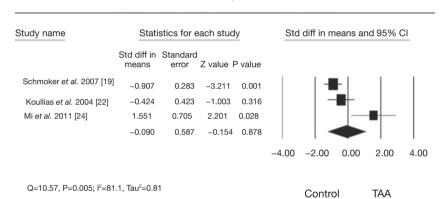
Current parameters in the diagnosis of AA

Although AA usually develops asymptomatically, once a certain size is reached, the risk of dissection, rupture, and death could be sharply increases. Due to the asymptomatic nature, many patients with AA would not be diagnosed until complications occur. Therefore, there are many parameters that are used to help diagnosis and management of complications.



MMP-9 in TAA compared to control

Figure 2 MMP-9 levels are increased in TAA compared with controls (21). TAA, thoracic aortic aneurysm.



MMP-2 in TAA compared to control

Figure 3 MMP-2 levels are not significantly different between TAA and controls (21). MMP, matrix metalloproteinase; TAA, thoracic aortic aneurysm.

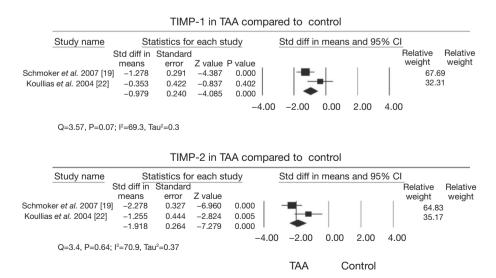


Figure 4 TIMP-1 and TIMP-2 are significantly decreased in TAA compared with controls (21). TIMP, tissue inhibitors of metalloproteinase; TAA, thoracic aortic aneurysm.

Size

Size is one of the important characteristics in AA. However, AA grows in a generally indolent manner, increasing by approximately 1 mm each year (24). Interestingly, aneurysms with larger diameters incline to expand more rapidly. The annual growth rate with a 4.0 cm of ascending TAA is 0.10 cm, while the annual growth rate with an 8.0 cm of ascending TAA is 0.19 cm (25). In AAA, ruptures appear in 25–41% of AAAs with a diameter >50 mm over 5 years (26). In TAA, large aneurysms, especially the size greater than 5 to 6 cm, expand more rapidly compared with small aneurysms (27-32). Therefore, the size of the AA is a good predictor for aortic rupture.

Genes

The AA is divided into atherosclerotic, syphilitic, bacterial, traumatic and congenital aneurysms, and dissecting aneurysms in the etiology. In the congenital aneurysms, the family factor is a major risk factor in the AA appearance and growth, the rate of diagnosed AA was lower in patients without family history than ones with first-degree relatives have diagnosed AA (33,34). The congenital AA forms are associated with Marfan syndrome, Loeys-Dietz syndrome and Ehlers-Danlos syndrome and relevant genes contains FBN1, ACTA2, PRKG1, TGFBR1, TGFBR2 genes. The mutation of this genes can benefit to the AA appearance and progression. For example, the Marfan syndrome has associated with the mutation of FBN1 and the TGF-β activation signal moreover the different position mutation of FBN1 has the different mechanism in the pathogenesis (35). The family genetic examination may have an application prospect to screen AA in individuals who have the family history using PCR amplification. But detection rate for genes mutations in familial AA is <20% and many individuals with family history of AA have normal diameter of aortic. Thus, the effectively way to screen family AA is ultrasounds. In current, there is article suggested the standard of screen family AA is the age of screen first-degree relatives is 50 for male and 55 for female rather than the less than 60 (36-38).

Mechanical sections

Calculations of the mechanical sections of AA can be performed using six independent variables: aortic pressure, aortic diameter, and thickness of the aorta in systole and diastole. When the size of the aneurysm attached a key point, the aorta cannot continue to stretch in systole, which could increase the stress to the aortic wall (39). Magnetic resonance imaging (MRI) has been reported to be used in measuring the mechanical sections of AA. In one study of AAA, wall stress in the control group was lower than that the AA group by using a three-dimensionally reconstructed model (40). Therefore, these measurements can be predictive factors for indicating a higher risk of rupture of regions of AAs (41,42).

Biomarkers for AA

Most patients with AA are asymptomatic until rupture of the aneurysm. Therefore, biomarkers, especially plasma proteins, might be useful for diagnosing and monitoring of AA in an early phase. Some circulating biomarkers have been established for AAA (43-45). Recent studies have shown that SEP can be used as a biomarker for predicting expansion of AAA (46-48). Additionally, matrix protease, interleukin-6, C-reactive protein, TNF- α , D-dimer, and IFN- γ have also been studied by many research groups (49-54). Although there have been positive results regarding to these biomarkers in many studies, the numbers of the patients in trials are limited. Therefore, many difficulties need to be resolved before their translation into the clinic.

Molecular targets in AA

In progression of AA, the quantity of production, such as MMP and matrix proteins, related to AA appears to change. These factors may become molecular targets for diagnosis and management of AA. Molecular target therapy has become a potential useful plan to treat AA.

Inflammation targets

Inflammation is a major component in progression of AA. Endothelial activation, recruitment of leukocytes and the up-regulation of adhesion molecules are important events in the early pathology of AA. Moreover, pro-inflammatory chemokines and cytokines can accelerate the inflammatory process of AA (55,56). Therefore, inflammatory activities can become targets for the diagnosis and management of AA.

Metabolism activities

There is a general agreement that AAA formation has a close relationship with destruction of elastin and collagen

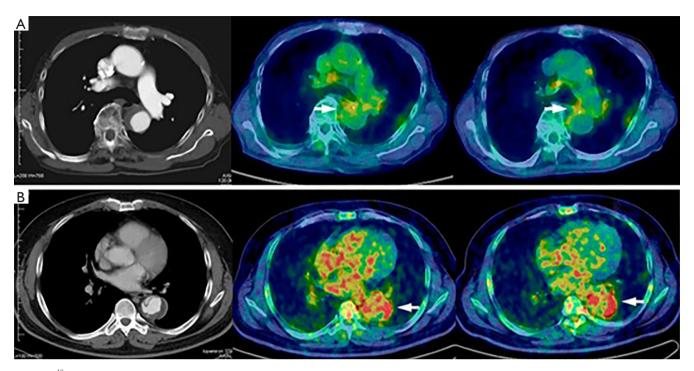


Figure 5 ¹⁸F-FDG uptake to predict short and mid-term prognosis in medically controlled patients with AA dissection by comparing uptake of ¹⁸F-FDG in controls and in patients with AAD. (A) ¹⁸F-FDG SUV_{max} and SUV_{mean} were on 50-minute images (middle) and 100-minute images (right) on patients with favorable outcome; (B) ¹⁸F-FDG SUV_{max} and SUV_{mean} were on 50-minute images (middle) and on 100-minute images (right) on patients with unfavorable outcome (68). F-FDG, F-fluorodeoxy glucose; SUV, standardized uptake value; AA, aortic aneurysm; AAD, acute aortic dissection.

at the medial level. Additionally, metabolism of elastin and collagen appear to be related to dispensability of AA (57). Several European studies have shown that the increased metabolic activity is associated with formation of AA, which can be used to evaluate inflammation of the aorta. Metabolic activity can be measured by an increase in ¹⁸F-fluorodeoxy glucose (FDG) uptake as measured by positron emission tomography (PET) or computed tomography (CT). FDG, as a glucose analog that accumulates in high metabolic activity in cells is often used for PET imaging of inflammation (58). Growing evidence show that an increase in ¹⁸F-FDG uptake is a latent signal in the aorta with active atherosclerotic inflammation (59,60). Many studies have attempted to use ¹⁸F-FDG to evaluate aortic diseases (61-67). One study used ¹⁸F-FDG uptake to predict short and midterm prognosis in medically controlled patients with AA dissection by comparing uptake of ¹⁸F-FDG in controls and in patients with AAD (Figure 5) (68). This study showed that the ¹⁸F-FDG standardized uptake value (SUV) was greater in unfavorable AA dissection groups than in favorable AA dissection groups on 50-minute images (68).

Through the observation of ¹⁸F-FDG uptake on 100-minute images, SUV_{max} and SUV_{mean} of $^{18}\mathrm{F}\text{-}\mathrm{FDG}$ in the between unfavorable and favorable AA dissection groups were no significant difference and both higher than those in controls at the proximal, distal, and maximum sites (all P<0.05). Therefore, use of ¹⁸F-FDG uptake on 50-minute to stratification patients with AA dissection may be useful for predicting short-term and mid-term prognosis and achieving better management (68). Additionally, many previous studies have shown that FDG in the AA group has higher uptake than that the normal aortic group (63,65,69). Moreover, in AA patients, uptake of FDG is higher in the symptomatic group compared with the asymptomatic group (67,70). Interestingly, sites with a positive ¹⁸F-FDG uptake have been shown to accumulate a higher amount of adventitial inflammatory cells with a reduction in smooth muscle cells in the media compared with negative ¹⁸F-FDG samples (71). Therefore, ¹⁸F-FDG might be a new pathway to study the mechanical of AA and to predict the risk of rupture. Because of FDG belongs to the glucose, diabetes whether influence FDG uptake due to the state of impaired

glucose utilization has appeared. Previous studies have suggested that diabetes cannot influence the evaluation of uptake of ¹⁸F-FDG in animal trial and human trial. At the moment, the relation of diabetes with AA has also become an attention point. A review suggested that the diabetes is an important risk factor in the coronary and peripheral artery disease, but has a negatively effect in AA due to the hyperinsulinemia can lead to up-regulation fibrinogen, collagen synthesis, plasminogen activator inhibitor-1 and down regulation fibrinolysis, inflammation and MMP (72-74).

Phagocytosis

Inflammation is a vital factor in progression of AA. There were histological studies found that lymphocytes and macrophages infiltrate into the aorta occurred during the angiotensin II infusion in apoE^{-/-}mice (75). Macrophages are associated with aneurysm growth and rupture in animal models and patients (76,77). A previous study showed that MRI is a noninvasive method that can be used to assess the morphology of AA (78). Additionally, using MRI to detect macrophages in AA can be used to reflect the degree of inflammation in AA (79). Recently, some studies have also shown that ultrasmall super-paramagnetic iron oxide (USPIO) contrast agents can be used to label atherosclerotic plaques and can indicate the macrophage load when it is used as an imaging agent. Due to the small particle size (10-30 nm), USPIO escapes recognition by the reticuloendothelial system, exists in the blood, and accumulates into the vascular inflammation sites. In these inflammatory sites, USPIO undergoes phagocytosis by tissue-resident macrophages within which it accumulates and is detectable on T2- and T2*-weighted MRI sequences (80-82). In recent research, USPIO contrast agents were used for detecting macrophage infiltration in the pre-clinical formation state of AAA (83,84). The previously study found that compared with mice without a USPIO agent, signal intensity was decreased in mice with a USPIO agent in which many macrophages were observed in remodeled adventitia (Figure 6) (83). These results indicated that with formation of AA, the process of macrophages moving to the aneurysm can be detected by using a USPIO MRI contrast agent. Because of this important discovery, many study groups have attempted to use this particle to study progression of AA (77,79). One other study attempted to show whether uptake of USPIO in the aortic wall was associated with the rate of aneurysm expansion. They divided 27 patients into three groups according to whether uptake of USPIO was in the aortic wall or in thrombus. Finally, they found that the aneurysm

expansion rate of focal uptake areas of the aortic wall was three-fold higher than in patients without uptake of USPIO or without specific uptake of USPIO (77). Therefore, USPIO may be able to be applied in the clinic to predict development of AA. However, more human trials are required before use of USPIO in the clinic.

Matrix remodeling targets

Extracellular matrix degradation is a major factor in formation, dilation and rupture of AA. Additionally, elastin fibers and collagen especially types I and III are vital for retaining the integrity of structure and stability of arteries. Development of AA is associated with degradation of collagen and elastin fibers (85). Cysteine proteases, serine proteases and MMPs produced by inflammatory cells show higher expression in AA than in the normal aorta.

MMP

MMPs are endopeptidases that can degrade many extracellular proteins (86). Comprehension of regulation of MMP activity is vital for understanding various pathogeneses of AA, and for production of new MMPrelated medicines. All the MMPs have a catalytic center that includes three complexes of the zinc ion in the active place. However, when only MMPs are activated, the zinc ion can be exposed, which is not emergence in inactivated MMPs and MMP proenzymes. Therefore, the zinc ion site may be used to target activation of MMPs (87). Recent data have demonstrated that MMP-1, -2, -3, -9, -12, and -13 play roles in progression of AA (85). Their activation can lead to progression of AA and subsequent rupture or dissection. However, in different types of AA, MMPs significantly increase at various rates. Compared with tricuspid aortic valves (TAVs), MMP-2 levels are increased by 34% in patients with bicuspid aortic valves (BAVs). However, in TAVs samples, MMP-13 levels are increased by 140% compared with BAVs (88). Before using MMPs as the target for imaging, the different designs of probes for assessing MMP need to be understood.

Antibodies were used in the earliest attempt to design probes for assessing MMP, and various antibodies have been developed. These antibodies only bind to other epitopes rather than the zinc ion. Therefore, antibodies are not an ideal approach for imaging MMPs (87). Substrates bind to active locations of MMPs and can be cleaned by using enzymes. Moreover, different MMPs have special substrates, which can help to select target MMPs. When Hu et al. Molecular targets in AA for establishing novel diagnostic and management paradigms

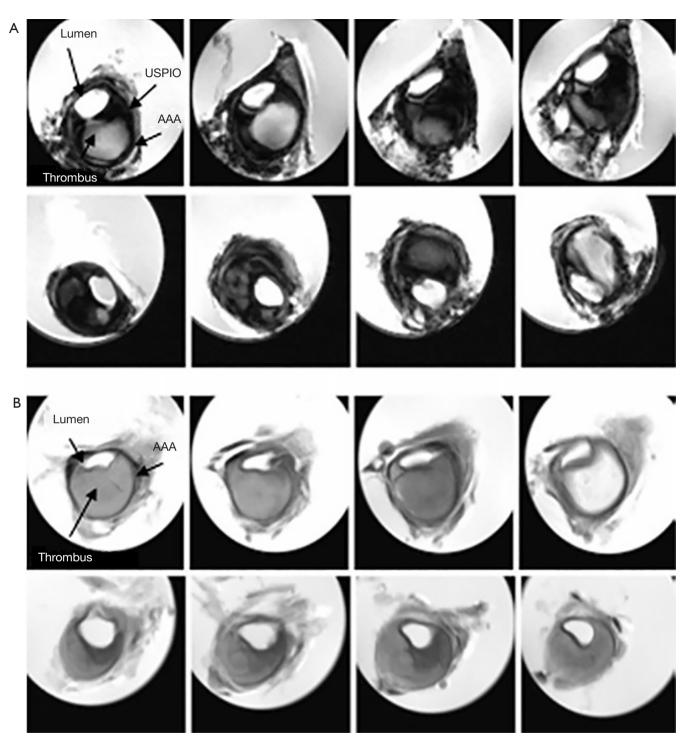


Figure 6 *Ex vivo* images of AAAs of mice with (A) USPIO administration and (B) without USPIO administration. A macrophage-rich area has decreased signal intensity in USPIO administered mice (83). AAA, abdominal aortic aneurysm; USPIO, ultrasmall super-paramagnetic iron oxide.

a probe is cleaved, the fluorescent signal is amplified (87). CGS 27023A, a non-specific inhibitor that binds to the active catalytic domain of MMPs, has dominant uptake in the apolipoprotein E-deficient mouse by using micro-PECT/CT. Although this technique can be used to detect activation of MMPs in aortic disease, it cannot differentiate the type of effect of MMPs in inflammation of the aorta (89). Compared with substrates, inhibitors interact with MMPs in a 1:1 manner and have no signal magnification. Research groups established two nonpeptidic MMP inhibitor-based probes (hydroxamates and barbiturates) to evaluate MMP activity and obtain an ideal outcome (90-92). By using this approach, MMP inhibitors can be used for imaging targets.

Therefore, fluorescence substrates and MMP inhibitors can be used to image inflammatory activity (93-98). RP782 and RP805, special tracers for activated MMPs, are usually used to detect activation of MMPs. Previous studies on cardiovascular disease showed that ^{99m}Tc-RP805 had significant uptake in the inflammatory area compared with controls by using micro-SPECT. Additionally, ^{99m}Tc-RP805 has a potential value to diagnose changes in MMPs in progression of cardiovascular disease (96-98). Recently, a study reported that a novel MMP inhibitor, RYM, has a faster blood clearance and higher water solubility compared with ^{99m}Tc-RP805 (93). This finding indicates that ^{99m}Tc-RYM1 can image in the early time, and can improve vessel wall-to-blood contrast. In mice models, micro-PECT/CT imaging showed higher RYM in AAA compared with no dilated aortas (93). Therefore, using MMPs as a target in imaging to predict inflammatory activity in AA has obvious superiority, and this provides a basis for clinical trials.

Matrix proteins

The aorta has three layers, including the endothelial cells layer, the elastic media layer, and the adventitia. Collagen and elastin form the strength of the aorta and elastic properties of the aorta, respectively (99). Therefore, disruption of these components may change the mechanics of the vessel wall and have a large effect on the pathology of AA. Elastic fibers are highly extensible networks of cross-linked elastin that provide elastic energy storage in tissues and redistribute it during diastole and maintain normal pressure (100). Therefore, elastic degradation can decrease extensibility and increase stiffness. An increased pressure in systole can result in damage of the intima of the aortic wall. Collagen has a large stiffness modulus of approximately 1,200 MPa (approximately 1,000 times greater than elastin) and a low extensibility of approximately 13% (100). Therefore, the role of collagen is to reinforce the strength of the wall to avoid rupture of a weak elastic wall. Consequently, destruction of elastin and collagen can lead to apoptosis of vascular smooth muscle cells, which can impair the aortic wall. CAN-35 is a collagen-binding bacterial protein, which can bind to the disorder collagen. It was used as a marker to detect the quantity of collagen in AAA in an animal model (101). A study reported that, using MRI in mice, injection of CAN-35 resulted in higher MR signal compared with mice with no injection mice (102). Therefore, matrix proteins are important targets for evaluating aortic inflammation.

Potential molecular therapeutic targets

When the diameter of AA exceeds 5.5 cm in man and 5.0 cm in women, clinicians always choice artificial vessel replacement or endovascular AA repair (103). However, when the small aneurysm (diameter >3 cm) was detected, current clinicians always choice follow-up CT or ultrasonography in annual 6 to 12 months due to the risk of rupture for aneurysms is low when the diameter of AA smaller than 4 cm (104,105) and the operative to early stage of AA has no survival advantage (106,107). Thus, the early medicine intervention may become a useful way. The traditional medicine treatment for preventing AA growth is beta-adrenergic receptor blockers (108), and these are useful in treating Marfan syndrome (109). In addition, calcium channel blockers, antiplatelet agents, lipid-lowering drugs, and other antihypertensive drugs are also used to prevent the growth of AA, but there is no obvious impact on aneurysm expansion (110,111). Despite advances in understanding the mechanisms of development of AA, current pharmacological treatment of AA is limited. Research of molecular targets and the area of molecular therapy of AA has become prevalent and there are articles have reported molecular therapy used in the Marfan syndrome and atherosclerotic disease.

Marfan syndrome

In Marfan syndrome, FBN1 mutations and TGF- β results in an imbalance between MMPs and TIMPs, and these can increase proteolysis in the aortic wall and finally cause AA formation (35,112). Much effort has been made to test therapeutic agents aiming the molecular changes, and an angiotensin receptor blocker has been found to inhibit the effect of TGF- β in the vascular wall (113-116). Additionally, there is study suggested that the long-term doxycycline, the inhibitor of MMP-2 and -9, was more effective than the β -adrenergic receptor blockers (117).

Atherosclerotic disease

Atherosclerosis is a chronic progressive disease caused by the inflammatory cellular and molecular changes, such as the macrophage and immune cells accumulation, the inflammation cytokines release. The current treatment point has transferred from tradition medicine (statins) to target inflammation medicine. An article has introduced that Toll-like receptors antagonists, T cells activation inhibitors, TNF and IL-1 receptor inhibitors, kinase inhibitors can be a prospect treatment in atherosclerotic disease (118).

AA

MMPs are major factors in progression of AA by causing degradation of the extracellular matrix. MMP activity can be suppressed by the tissue inhibitors TIMPs. MMP-2 and MMP-9 are protease that degrade matrix in TAA and AAA (119). Therefore, their role in the extracellular matrix suggests that blocking of MMPs slow progression of aortic disease. Many synthetic MMP inhibitors are known to decrease MMP activity (120), such as doxycycline-based and hydroxamate-based activity. The first synthetic MMP inhibitor that was used in the clinic was BB-94 to reduce MMP activity in cancer (121). However, the feature of poor water solubility has limited the effectiveness of BB-94 when provided orally. Doxycycline is associated with positive results in clinical trials involving small AAA. Doxycycline reduces plasma MMP-9 levels and significantly lowers aneurysmal growth rates in patients (122-124). Many studies have reported that in animal models, systemic MMP inhibitors can reduce the onset of aneurysms (125,126). However, systemic delivery may limit normal MMP activity, which is not the original aim of using MMP inhibitors.

Recently, many studies have investigated how to reduce the side effects of MMP inhibitors and increase accumulated concentrations. These studies showed that MMP inhibitors that were loaded with nanoparticles were localized in inflammation or pathological regions (127-131). However, drug-loaded nanoparticles are usually toxic and can cause adverse effects because of exposed in the circular to deliver. Additionally, rapid clearance of nanoparticles from the plasma limits use of nanoparticles in humans (132,133). Therefore, many alternative drugs for delivery to cells are being studied worldwide, especially for tumors using intrinsic functional cells. Intrinsic functional cells include macrophages, mesenchymal stem cells, natural killer cells, and mature erythrocytes (134-137). The pathological features of AA are degradation of the extracellular matrix and a decrease in vascular smooth muscle cells, associated with inflammatory cell infiltration. Additionally, macrophages have a major function in inflammatory progression. Therefore, macrophages may have the chance to become drug carriers (138). A previous study attempted to use the macrophage membrane to deliver nanoparticles (emtansine liposome). This study showed that emtansine liposomes coated with macrophage membranes had long-term stability in plasma and resulted in a higher accumulation of concentrations compared with emtansine liposomes alone. Many research groups have also attempted to use membranes to treat AA (131). We hypothesize that using macrophages to coat nanoparticles with MMP inhibitors is useful for treating AA. However, long-term in vivo trials are required before using this technique in the clinic.

Conclusions

The incidence of AA has increased in the last 50 years. Techniques of diagnosis of AA need to progress to reduce the rate of morbidity and mortality. Understanding the mechanisms of development of AA could lead to discovering newer potential methods of substituting the traditional diagnosis and therapeutic approaches. In the future, molecular targets for AA may become an attractive aspect and be used in the clinic. However, target imaging for diagnosing AA requires more pre-clinical trials. Additionally, using target drugs to the diseased site is difficult because of complex blood flow and limited biofabrication technology. Nanoparticles may be an effective vehicle for delivering therapeutic agents to the target site. These particles can lead to target drug concentrations reaching a high level in specified sites and a decrease in distribution in other sites. Therefore, molecular targets may be good candidates for a future study direction in the management of AA. The new prospect treatment may available prevent the small AA dilation and reduce the rate of AA growth for more and elderly patients. Besides that, the intervention could be suggested in the early stage based on the molecular target.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

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