Review

IL-1β in atherosclerotic vascular calcification: From bench to bedside

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Abstract

Atherosclerotic vascular calcification contributes to increased risk of death in patients with cardiovascular diseases. Assessing the type and severity of inflammation is crucial in the treatment of numerous cardiovascular conditions. IL-1β, a potent proinflammatory cytokine, plays diverse roles in the pathogenesis of atherosclerotic vascular calcification. Several large-scale, population cohort trials have shown that the incidence of cardiovascular events is clinically reduced by the administration of anti-IL-1β therapy. Anti-IL-1β therapy might reduce the incidence of cardiovascular events by affecting atherosclerotic vascular calcification, but the mechanism underlying this effect remains unclear. In this review, we summarize current knowledge on the role of IL-1β in atherosclerotic vascular calcification, and describe the latest results reported in clinical trials evaluating anti-IL-1β therapies for the treatment of cardiovascular diseases. This review will aid in improving current understanding of the pathophysiological roles of IL-1β and mechanisms underlying its activity.

Key words: IL-1β, vascular calcification, cardiovascular events, signaling pathways

Introduction

Atherosclerosis is initiated by disturbances in blood flow and numerous systemic factors including history of smoking, hypertension, diabetes, or hyperlipidemia [1-3]. Atherosclerosis is part of the common pathophysiological basis for most cerebrovascular and cardiovascular diseases [4].

Vascular calcification is a ubiquitous pathological process in atherosclerosis [5, 6]. The annual incidence of atherosclerotic vascular calcification in the general population ranges from less than 5% in individuals under 50 years of age to greater than 12% in individuals over 80 years of age [7]. Imbalance in the calcium dynamics of atherosclerotic vessels can lead to reduced arterial compliance and impaired vascular hemodynamic responses [8, 9]. Clinically, atherosclerotic vascular calcification (Figure 1) is implicated in aortic valve stenosis, congestive heart failure, myocardial infarction, peripheral arterial occlusion, and arterial hypertension [10], which lead to high rates of morbidity and mortality [11, 12].

Calcification of the aortic arch was first reported
in relation to the risk of coronary heart disease in a cohort study examining 116,309 individuals with a median follow-up of 28-years [12]. That study was also used to assess calcification of the coronary artery in asymptomatic individuals who experienced coronary events such as myocardial infarction or coronary death [11], and calcification of the abdominal aorta in individuals with increased risk for coronary heart disease [13]. Under conditions of atherosclerosis, vascular calcification progresses, and is associated with cardiovascular events and increased mortality [7, 13-15]. A meta-analysis examining 218,080 individuals with a mean follow-up of 10.1-years indicated that vascular calcification results in 4.63-fold higher risk for all-cause mortality, a 3.94-fold higher risk for cardiovascular mortality, and a 3.74-fold higher risk for any coronary events [15].

Although vascular calcification was previously considered passive and degenerative, it is currently recognized as an active and regulated pathobiological process. Vascular calcification, which shares many features with inflammatory atherosclerosis, may be treatable and preventable. The initial feature of calcified atherosclerotic vessels is activation of inflammation [16, 17]. Cytokines secreted by inflammatory cells result in smooth muscle cell (SMC) apoptosis or SMC trans-differentiation into an osteochondrogenic cellular phenotype. Both of these events may contribute to mineral deposition in the plaque [18, 19]. Among the numerous currently known inflammatory signaling pathways, those involving IL-1β are particularly implicated in atherosclerotic vascular calcification. In this review, we summarize the specific roles of IL-1β in atherosclerotic vascular calcification. This information will help delineate the pathogenesis of atherosclerosis, and will help uncover other, currently unknown, mechanisms involving IL-1β, in order to develop anti-IL-1β therapeutics for reducing the incidence of cardiovascular events [20].

1. Vascular calcification in atherosclerosis

Vascular calcification was previously believed to result from passive deposition of calcium and
phosphorus on blood-vessel wall. Recent studies have shown, however, that vascular calcification is an active, reversible, highly regulated, and preventable process that is similar to physiological bone development [21]. The two pathobiological mechanisms currently known to underlie vascular calcification are induction of osteogenesis and loss of inhibition of mineralization [22]. Abnormalities in Ca\(^{2+}\) and phosphate metabolism [23], combined with increased oxidative stress [24], stimulation of inflammatory factors [25], dysregulation of certain miRNAs such as miR-34a [26], and disorders of lipid metabolism [27] lead to decreased expression of α-SMA, SM22α, and smooth muscle-myosin heavy chain, which are necessary for the maintenance of vascular function. Meanwhile, the expression of Runx2, SOX9, SP7, MSx2, and OPN, which are factors related to bone formation, is upregulated. This cascade promotes the activity of ALP and expression of BMP-2, resulting in osteogenic or chondral differentiation of VSMCs [8, 28]. Inhibition of autophagy [29], matrix remodeling [30], cellular apoptosis, and matrix vesicles [31] may also accelerate vascular calcification. Recent studies in mouse models and humans have shown that blood vessels can produce and secrete factors, including PPI, OPN, OPG, MGP, FET-A, and Smad 6, which inhibit mineralization or vascular calcification [32-35]. Inhibition of expression in these factors can lead to the initiation of vascular calcification.

Atherosclerosis usually leads to vascular calcification [36]. In early stages of atherosclerosis, bone-related proteins can be detected histologically, and both the occurrence and development of vascular calcification are associated with the process of atherosclerosis [37]. In the past decade, studies on coronary atherosclerotic calcification were mainly focused on lumen stenosis and plaque vulnerability. While calcification of atherosclerotic plaque core does not increase the vulnerability of the plaque [38], microcalcification of the fibrous caps on atherosclerotic plaques increases circumferential stress, which can, indeed, increase the vulnerability of the plaque. Overall, the size, shape, and location of the microcalcifications can directly define the vulnerability of the plaques [39]. Electron beam computed tomography (CT) and intravascular ultrasound are currently used to detect calcifications in the arteries of 90% of patients with coronary atherosclerotic heart disease. Thus, the degree of vascular calcification may be directly related to the degree of vascular stenosis and risk for cardiovascular events in patients with atherosclerotic diseases [40].

Hence, an accurate, safe, and reproducible clinical detection technique is extremely important for the diagnosis, prevention, and treatment of vascular-calcification-related diseases. In the early stages of vascular calcification, molecular imaging technology, such as optical near-infrared fluorescence imaging [41], can be used to detect osteogenesis early at the (sub)cellular levels [42]. The detection and quantification of advanced vascular calcification can be performed by CT, intravascular ultrasound (IVUS), transthoracic echocardiography, pulse wave velocity measurement, planar radiographs, and magnetic resonance imaging (MRI) [43]. However, there are currently no satisfactory therapeutic approaches to vascular calcification in clinical practice. Even statins, which have been shown to decrease osteogenesis in vivo and vitro [44, 45], have failed to prove beneficial in clinical trials [46]. Preventive measures are critical to decreasing the occurrence of vascular calcification and delaying the progression of vascular calcification. Although the modification of risk conditions (hyperglycemia, uremia, hypertension, hyperlipidemia, secondary hyperparathyroidism, and metabolic syndrome) and other factors (dietary phosphorous, oral activated charcoal, vitamins K and D, magnesium oxide, warfarin, bisphosphonates, antioxidants, estrogen, fetuin, osteopontin, anti-inflammatory agents, mineralocorticoids) might be possible [47-52], the details of any underlying mechanisms are not fully understood, and large-scale clinical trials have been limited.

2. Roles of the pro-inflammatory cytokine IL-1β

IL-1β is implicated in numerous inflammation-related diseases including rheumatoid arthritis, inflammatory bowel disease, osteoarthritis, type 2 diabetes, gout, multiple sclerosis, and Alzheimer’s disease [53-55]. As a potent, pro-inflammatory cytokine with a wide range of biological effects, IL-1β is synthesized and secreted by various cells including macrophages, fibroblasts, B lymphocytes, natural killer cells, and smooth muscle cells [56]. As canonical negative feedback regulation to control IL-1β expression and secretion, IL-10 suppresses IL-1β and IL-1β-induced IL-1Ra [57, 58], the natural antagonist of IL-1β. There is also another mechanism underlying IL-1β-TGF-β1-related feedback to decrease the production of IL-1β [59, 60]. One mechanism by which downstream and upstream negative feedback modulate IL-1β via paracrine secretion of interferons has been reported [61]. Of which, IFN-II/IFNγ mediated the downstream regulation via its inhibition of the nitrosylation of NLRP3 inflammasome [62]. While IFN-I/IFNβ mediated the upstream regulation via an IL-10 and STAT-3 dependent manner. The IFN-I/IFNβ secreted predominantly by fibroblasts
responded to IL-1β elevation is a strong switch to attenuate the IL-1β induced inflammation at later stages [63]. Recently, the secretion of IL-1β was revealed to be modulated by a negative feedback loop including IL-1β/NF-κB/TIR8/IL-1β during IL-1β-induced epithelial-myofibroblast trans-differentiation [64].

IL-1β also plays an important role in the development of cardiovascular diseases [65]. Various factors can activate IL-1β production and pyrolysis, enabling the participation of IL-1β in the pathophysiological process of cardiovascular diseases [66]. Avolio et al. [67] and Qi et al. [68] discovered excessive activation of IL-1β in the hypothalamic paraventricular nucleus under conditions of hypertension, and found that inhibition of IL-1β can alleviate hypertension by reducing the activity of the sympathetic nervous system. Under conditions of hypertension, IL-1β also participates in the remodeling of aortic blood vessels by activating the renin-angiotensin-alderosterone system [69]. Coronary artery thrombosis and blockage of coronary blood flow resulting from ruptured coronary artery plaques are the main causes of acute myocardial infarction. The ischemic or necrotic myocardium can generate increased levels of ATP and oxidative stress products, which then stimulate the expression of IL-1β [70, 71].

Atrial fibrillation is a common type of arrhythmia possessing a complex mechanism [72]. Increased levels of IL-1β result in atrial myoelectrical and structural remodeling, and induction of atrial fibrillation [73]. Activated Macrophage-inducible C-type lectin (Mincle) causes increased expression of IL-1β in the microglial cells residing in the paraventricular nucleus. At 24 hours after myocardial infarction, increased levels of IL-1β are increased further, and are accompanied by excessive activation of the sympathetic nerves. This cascade can result in the occurrence of a malignant ventricular arrhythmia [74].

Recent studies have shown that therapeutic targeting of inflammatory factors can improve cardiovascular outcomes in patients with a history of myocardial infarction [75]. In the Canakinumab Anti-Inflammatory Thrombosis Outcomes Study (CANTOS), administration of the monoclonal IL-1β-neutralizing antibody canakinumab successfully reduced the rate of recurrent cardiovascular events by 17% [20]. However, not all anti-inflammatory therapies benefit in the protection from cardiovascular events [75, 76], the detailed mechanism of anti-IL-1β-mediated activity remains unclear.

3. Signaling pathways involved in IL-1β-mediated regulation of atherosclerotic vascular calcification

3.1. IL-1β induces endothelial-to-mesenchymal transition and promotes atherosclerotic vascular calcification

Endothelial-to-mesenchymal transition (EndMT), a specific form of epithelial-to-mesenchymal transition (EMT), is characterized by the loss of endothelial features and acquisition of specific mesenchymal markers in endothelial cells [77, 78]. EndMT is known to participate in the pathogenesis of atherosclerosis [79-82] and also occurs in atherosclerotic vascular calcification [83, 84].

IL-1β-mediated EndMT contributes to the pathogenesis of various diseases. Lee et al. have shown that injury-induced IL-1β expression induces EndMT in corneal fibrosis by upregulating the expression of FGF-2 [85]. Recombinant IL-1β induces EndMT in human esophageal microvascular endothelial cells, highlighting the important role of IL-1β in early-stage esophageal adenocarcinoma [86]. IL-1β-induced EndMT also impairs the angiogenic potential of human umbilical vein endothelial cells (HUVECs) [87].

The relationship between IL-1β and EndMT in atherosclerotic vascular calcification is still poorly understood. Sanchez-Duffhues et al. reported that IL-1β-sensitized bone morphogenetic protein-9 (BMP-9)-induces osteogenic differentiation via induction of EndMT. This process is mediated by the downregulation of bone morphogenetic protein receptor type II (BMPR2) expression and subsequent inactivation of the c-Jun N-terminal kinase (JNK) signaling pathway in human primary aortic endothelial cells [88]. This hypothesis was further corroborated in vivo and in patient-derived atherosclerotic tissues [88]. These findings suggest that IL-1β may induce EndMT and promote atherosclerotic vascular calcification (Figure 2A).

3.2. IL-1β inhibits the mobilization and infiltration of bipotent mesodermal progenitor cells (MPCs), thereby accelerating atherosclerotic vascular calcification

Stem or progenitor cells, and their dynamics, play an important role in cardiovascular diseases [89-92]. Using identification of cell surface markers, such as platelet-derived growth factor receptor alpha (PDGFRα) and stem cell antigen-1 (Sca-1), Cho et al. [93] discovered a cluster of vascular calcifying progenitor cells residing in the arterial adventitia. These cells are derived from the bone marrow and
mobilize to the inflamed atherosclerotic lesions. Mesodermal progenitor cells (MPCs), such as Lin-CD29+/Sca-1+/PDGFRα- cells, possess bidirectional differentiation, which enables the development of MPCs into osteoblasts (OBs) or osteoclasts (OCs). MPCs isolated from the adult bone marrow are also progenitors of Sca-1+/PDGFRα+ cells, which can potentially differentiate into OBs [94]. IL-1β, which is elevated in the sera and arteries of hypercholesterolemic ApoE-/− mice, enhances the mobilization and infiltration of Sca-1+/PDGFRα+ cells. Conversely, IL-1β inhibits bipotent MPCs and accelerates atherosclerotic vascular calcification [94]. Thus, current studies suggest that IL-1β is likely a key regulator of bipotent MPCs in vascular calcification.

Activation of peroxisome proliferator-activated receptor γ (PPARγ) promotes the differentiation of bipotent MPCs into OCs; this process alleviates vascular calcification in vitro and in vivo [93]. Collectively, the homeostasis between MPCs and Sca-1+/PDGFRα+ cells may play an important role in vascular calcification under conditions of atherosclerosis (Figure 2B). Induced pluripotent stem (iPS) cells originating from MPC-related cells [95], and monoclonal antibodies specific for IL-1β [96-98], are currently available as sources for the development of new therapeutics for the treatment of patients with atherosclerotic vascular calcification [93].

3.3. IL-1β activates tissue-nonspecific alkaline phosphatase to exacerbate atherosclerotic vascular calcification

Tissue-nonspecific alkaline phosphatase (TNAP) is an enzyme that degrades extracellular pyrophosphate (PPI) and promotes vascular calcification by downregulating the expression of PPI. PPI is a potent endogenous inhibitor of hydroxyapatite [Ca_{10}(PO_{4})_{6}(OH)] [99-101], which is the main component of the calcified aorta [102]. This decreased level of plasma PPI is associated with genetic or metabolic conditions, such as Hutchinson-Gilford progeria syndrome (HGPS) [99], generalized arterial calcification of infancy (GACI) [103], and advanced chronic kidney disease [104], which predispose vulnerable individuals to vascular calcification [105-107]. Transgenic overexpression of TNAP in VSMCs or in endothelial cells results in pathological calcification in vitro and in vivo [108-110]. Moreover, the upregulation of TNAP expression has also been demonstrated in dialysis-related calcification of human vessels [111].

Figure 2. Signaling pathways involved in IL-1β—mediated regulation of atherosclerotic vascular calcification. (A) IL-1β activates JNK pathway by downregulating BMPR2 expression and subsequent BMPR2-dependent inhibition of JNK, which promotes endothelial to mesenchymal transition (EndMT), leading to BMP-9-induced osteogenic differentiation. (B) IL-1β inhibits the mobilization and infiltration of mesodermal progenitor cells (MPCs), which can bi-directionally differentiate into osteoblasts (OBs) or osteoclasts (OCs). While in hypercholesterolemia IL-1β enhances the mobilization and infiltration of Sca-1+/PDGFRα+ cells, which are differentiated from MPCs and the progenitor cells of OBs. PPARγ promotes the differentiation of MPCs into OCs. (C) IL-1β stimulates tissue-nonspecific alkaline phosphatase (TNAP) expression and activity in both vascular smooth muscle cells (VSMCs) and mesenchymal stem cells followed by pyrophosphate (PPI) degradation. PPI is an effective endogenous inhibitor of Ca_{10}(PO_{4})_{6}(OH), which is a major component of the calcified aorta. This chain of events accelerates atherosclerotic vascular calcification.
Ding et al. [112] reported that IL-1β can upregulate TNAP activity and calcification in human mesenchymal stem cells via RUNX2-independent signaling. Lencel et al. [113] utilized VSMCs, stimulated by TNF-α and IL-1β, to demonstrate the cell-specific effects of TNAP, and suggested that PPARγ may mediate these differences in TNAP activity. For these reasons, IL-1β is considered a stimulator of vascular calcification in the context of atherosclerosis. Few studies have examined the roles of IL-1β and TNAP in vascular calcification. However, whether IL-1β can promote vascular calcification via a TNAP-mediated signaling pathway needs further study (Figure 2C).

4. Stimulators that regulate IL-1β expression accelerate atherosclerotic vascular calcification

4.1. Rac2 mediates atherosclerotic vascular calcification via regulation of IL-1β production in macrophages

Rac proteins are a subfamily of the Rho family, which consists of small guanosine triphosphate-binding proteins Rac 1, 2, 3, and RhoG [114, 115]. Similar to other small GTPases, Rac switches between a GTP-bound active and GDP-bound inactive state. Stimulus-induced activation of Rac is mediated by guanine nucleotide exchange factors (GEFs) [116-118]; this subfamily of proteins is critical in numerous inflammation-mediating pathological processes. Rac1 and Rac2, which are important signal transducers in inflammatory cells, affect the expression of several cytokines and growth factors [119, 120]. Recently, Ceneri et al. [121] reported a significant decrease in Rac2, and increase in IL-1β, expression, in the aortas of ApoE−/− mice and in calcified plaques in human coronary segments. They also showed that Rac2 deletion can aggravate vascular calcification by increasing macrophage expression of IL-1β, which is associated with NF-κB activation. Rac2 was also found to enhance the production of reactive oxygen species (ROS) elicited by elevation of activated Rac1 expression in Rac2−/−ApoE−/− mice. Thus, Ceneri et al. identified a novel inflammatory signaling pathway that depends on Rac2-mediated regulation of Rac1-dependent IL-1β expression in macrophages (Figure 3A) [121]. This study outlines a pathophysiological signaling mechanism to understand the rationale of the action of IL-1β in the development of atherosclerotic vascular calcification.

![Figure 3. Stimulators regulating IL-1β expression accelerate atherosclerotic vascular calcification. The NLRP3 inflammasome participates in the regulation of the progression of atherosclerotic vascular calcification via two signaling pathways described as follows.](http://www.ijbs.com)
4.2. NLRP3 inflammasome initiates the release of IL-1β by upregulating the expression of caspase-1, leading to atherosclerotic vascular calcification

The NLR family pyrin domain-containing 3 (NLRP3) inflammasome, which is an ROS-sensitive multiprotein complex, accelerates IL-1β maturation via activation of caspase-1 [122, 123]. NLRP3 plays a pivotal role in the pathogenesis of atherosclerosis [124-126]. Numerous studies have reported that NLRP3 is essential for atherogenesis, and that silencing the expression of NLRP3 prevents the rupture of atherosclerotic plaques [127, 128].

Wen et al. [129] found that the expression of NLRP3 inflammasome and IL-1β are upregulated in VSMCs cultured in calcification medium supplemented with β-glycerophosphate (β-GP) and in human calcified popliteal arteries having elevated expression of caspase-1. Downregulation of NLRP3 expression using treatment with NLRP3 siRNA reduces IL-1β secretion, which subsequently reduces vascular calcification, in vitro. However, whether the knockdown of IL-1β can counteract the effect of NLRP3 inflammasome in vascular calcification remains unclear and needs to be studied in vivo. Tangi et al. [130] demonstrated that TNF-α-mediated induction of IL-1β release in aortic smooth muscle cells is also NLRP3-dependent. Additionally, the NLRP3 inflammasome-mediated promotion of IL-1β secretion via upregulation of caspase-1 expression may play an important role in atherosclerotic vascular calcification (Figure 3B). Further investigation is required to uncover the details of this mechanism.

4.3. Hypercholesterolemia promotes the release of IL-1β to induce atherosclerotic vascular calcification

The effect of hypercholesterolemia on vascular calcification was confirmed when Awan et al. [131] reported that patients with familial hypercholesterolemia (FH), who possess mutations in the low-density lipoprotein receptor (LDLR) gene, show severe and extensive vascular calcification in their thoracoabdominal aortas; this process usually commences at 20 years of age in homozygous FH patients. In contrast, vascular calcification in thoracoabdominal aorta is delayed by two decades in individuals affected by heterozygous FH [132]. Although a comparably high level of plasma cholesterol is observed in Ldlr−/−C57BL/6 mice fed a standard-chow diet and wild-type C57BL/6 mice fed a high-cholesterol, high-fat diet, Ldlr−/− mice still develop considerably more extensive aortic vascular calcification compared with that of wild-type mice [133]. None of the currently available therapeutics, including statins, can stop or regress vascular calcification in a clinical setting [134]. Collectively, these findings suggest that vascular calcification in early hypercholesterolemia may involve as of yet undiscovered mechanisms, and that LDLR deficiency may be a key step in vascular calcification.

Duewell et al. [125] and Rajamaki et al. [135] found that crystals of cholesterol are absorbed by macrophages, which then express the activated inflammasome complex. These events result in cleavage and activation of pro-IL-1β and secretion of IL-1β into plasma. Consequently, the binding of plasma IL-1β to IL-1β receptors on vascular endothelial cells promotes the release of various cytokines, SMC proliferation, and macrophage activation. The steps in this cascade contribute to the development of atherosclerosis [136], but the mechanism underlying hypercholesterolemia and IL-1β activity in the context of vascular calcification remains to be uncovered.

The levels of total cholesterol are increased in LDLR-deficient (Ldlr−/−) mice and LDLR-attenuated proprotein convertase subtilisin/kexin type 9 (Pcsk9) transgenic (Tg) mice [137]. However, Awan et al. [138] reported that IL-1β plasma levels in Ldlr−/− mice are twice as high as those in Pcsk9(Tg) mice. They also found that while anti-IL-1β monoclonal antibodies considerably inhibit atherosclerotic vascular calcification in Ldlr−/− mice with hypercholesterolemia, they induce only an insignificant change in the atherosclerotic vascular calcification of Pcsk9(Tg) mice with indiscriminate hypercholesterolemia. This finding suggests a potential mechanism accounting for why Pcsk9(Tg) mice show significantly less calcification compared with that in Ldlr−/− mice. This finding indicates that IL-1β plays a key role in atherosclerotic calcification associated with LDLR deficiency. Although a hypothesis linking IL-1β, LDL-R, and the Wnt/β-catenin signaling pathways to vascular calcification in the setting of hypercholesterolemia has been proposed [133], the detailed mechanism is unknown and further investigation is still needed (Figure 3C).

5. Discussion

Inflammation is proved to be an independent risk factor for cardiovascular disease, even after the lipid-lowering therapies. The SPIRE-1 and SPIRE-2 trials [139] using the statins and PCSK9 antibody bococizumab to reduce atherogenic lipids showed residual inflammation and no improvement for cardiovascular events. Since 2018, several anti-inflammation clinical trials revealed promising and challenging results for the treatment of
cardiovascular disease, such as CANTOS [20] and CIRT [76]. The CANTOS trial has shown that treatment with the monoclonal IL-1β-neutralizing antibody canakinumab can reduce the risk of recurrent cardiovascular events in patients with prior heart attack. While the CIRT trial in 5,000 patients with previous coronary disease showed no benefit for the reduction of cardiovascular events with the treatment of methotrexate – an promising anti-inflammatory approach in which once an association with fewer cardiovascular events was observed in patients with rheumatoid. These results provide us with informative implications that more unknown mechanisms exist in the anti-inflammation therapeutics for the prevention of cardiovascular events. A hypothesis of ‘innate immune training’ of IL-1β in epigenetic reprogramming of myeloid progenitor cells [140] was once proposed to understand the mechanism of the action of IL-1β. However, disputes of this issue exist [75].

Accumulating clinical evidence indicates that vascular calcification is an independent predictor of the occurrence of atherosclerosis-related myocardial infarction and stroke [7, 11-13, 15, 141-143]. Recent studies, including those summarized in this review, have shown that IL-1β plays prominent roles in the development of atherosclerotic vascular calcification. Although the incidence of atherosclerosis-related cardiovascular events has been clinically reduced by the use of anti-IL-1β therapy, as shown in several large-scale population cohort trials (Table 1), the clinical effect of IL-1β inhibition on atherosclerotic vascular calcification remains unclear. It is also unclear whether the beneficial effect induced by anti-IL-1β therapeutics in the setting of atherosclerosis is related to the effect of these therapeutics on atherosclerotic vascular calcification.

Further prospective randomized controlled trials (RCTs) are needed to evaluate the effects of IL-1β-related antibodies and drugs (Table 2) on atherosclerotic vascular calcification and incidence rate of cardiovascular events. Even in the well-known CANTOS trial [20, 144], there is no subgroups treated with anti-IL-1β therapy showing variance in the levels of vascular calcification in the setting of cardiovascular events. Therefore, further clinical trials, especially RCT studies, are needed to explore the roles of IL-1β and related therapeutics in atherosclerotic vascular calcification.

IL1B−/−, IL1R1−/−, IL-1 receptor antagonist-deficient (IL1Ra−/−), and transgenic mice overexpressing either secreted IL-1Ra or intracellular IL-1Ra1, are valuable animal models employed in basic in-vivo and in-vitro research [145-150]. These models are useful to mimic cardiac infarction and ischemic stroke in order to uncover the mechanism linking IL-1β activity, atherosclerotic vascular calcification, and cardiovascular events. New technologic advances, such as using artificial intelligence to conduct big-data analysis, will help us delineate the detailed mechanisms of cardiovascular calcification and design potential therapeutics [151]. These future studies will provide us with a broad understanding of IL-1β roles in atherosclerotic vascular calcification, and will define the value of IL-1β as a potential therapeutic target in the treatment of patients with cardiovascular diseases.

### Table 1. Clinical trials examining anti-IL-1β therapy for treatment of patients with cardiovascular diseases

| Trials                                      | Drug type                      | Dosage and administration | Treatment duration | Outcomes                                                                 |
|---------------------------------------------|--------------------------------|----------------------------|--------------------|--------------------------------------------------------------------------|
| Virginia Commonwealth University-Anakinra Remodeling Trial (VCU-ART) [152] | Recombinant IL-1 receptor antagonist (IL-1Ra) | Anakinra, 100 mg/day, subcutaneous injection | 14 days            | Double-blinded, randomized, placebo-controlled; 10 patients with ST-segment elevation of acute myocardial infarction (AMI). Anakinra improved the end-diastolic volume index and left ventricular end-systolic volume index (LVESVi). |
In summary, atherosclerotic vascular calcification is a vascular lesion related to morbidity and mortality worldwide. Currently, no specific and effective treatments are available for this condition because of our insufficient understanding of the detailed molecular mechanisms driving the processes of atherosclerotic vascular calcification. Future research should aim to delineate these molecular mechanisms, including those involving IL-1β-mediated pathways, as related to the development of atherosclerotic vascular calcification.

**Abbreviations**

ALP: Alkaline phosphatase; AMI: Acute myocardial infarction; ASO: Arteriosclerosis obliterans; BMP-2: Bone morphogenetic protein-2; BMP-9: Bone morphogenetic protein-9; BMPR2: Bone morphogenetic protein receptor type II; CACS: Coronary artery calcium scoring; CANTOS: Canakinumab Anti-Inflammatory Thrombosis Outcomes Study; CTA: Computed tomography angiography; EndMT: Endothelial-to-mesenchymal transition; EMT: Epithelial-to-mesenchymal transition; Fet-A: Fetuin A; FH: Familial hypercholesterolemia; GACI: Generalized arterial calcification of infancy; GEFs: Guanine nucleotide exchange factors; HGPS: Hutchinson-Gilford progeria syndrome; HUVECs: Human umbilical vein endothelial cells; IL-1β: Interleukin-1β; IL-1Ra: IL-1 receptor antagonist; iPS: Induced pluripotent stem; JNK: c-Jun N-terminal kinase; LAD: Left anterior descending; LCA: Left coronary artery; LCX: Left circumflex artery; LDLR: Low-density lipoprotein receptor; Ldlr−/−: LDLR-deficient; LVESVi: Left ventricular end-systolic volume index; MGP: Matrix Gla protein; Mincle: Macrophage-inducible C-type lectin; MPCs: Mesodermal progenitor cells; MSx2: Muscle segment homeobox homolog of 2; NLRP3: Nlpl family, pyrin domain containing 3; NSTE-ACS: Non-ST elevation acute coronary syndrome; OB: Osteoblastic; OC: Osteoclastic; OPN: Osteopontin; Pcsk9: Proprotein convertase subtilisin/kinin type 9; PDGFRα: Platelet-derived growth factor receptor alpha; PPARγ: Peroxisome proliferator-activated receptor γ; Ppi: Pyrophosphate; RCTs: Randomized controlled trials; Runx 2: Runx-related transcription factor 2; Sca-1: Surface markers stem cell antigen-1; Smad 6: Drosophila mothers against decapentaplegic protein 6; SMC: Smooth muscle cell; Sox 9: SRY-Box transcription factor 9; SP 7: Osterix; T2DM: Type 2 diabetes mellitus; Tg: Transgenic; TNAP: Tissue-nonspecific alkaline phosphatase; TNF-α: Tumor necrosis factor-α; VSMCs: Vascular smooth muscle cells.

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**Author contributions**

XLS conceptualized and designed the study, collected the clinical diagrams used in Figure 1C-E, and revised and edited the manuscript. JLS wrote the first draft of the manuscript. MZ critically revised the manuscript. CXZ conceptualized and reviewed the manuscript. All authors read and approved the final version of the manuscript.

**Competing Interests**

The authors have declared that no competing interest exists.

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