Review

The Journal of Cardiovascular Aging

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Extracellular vesicles: the key to unlocking mechanisms of age-related vascular disease?

Meredith Whitehead, Marco Antonazzi, Catherine M. Shanahan ២

BHF Centre of Research Excellence, School of Cardiovascular and Metabolic Medicine & Sciences, King's College London, London SE5 9NU, UK.

Correspondence to: Prof. Catherine M. Shanahan, School of Cardiovascular and Metabolic Medicine & Sciences, King's College London, 125 Coldharbour Lane, London SE5 9NU, UK. E-mail: cathy.shanahan@kcl.ac.uk

How to cite this article: Whitehead M, Antonazzi M, Shanahan CM. Extracellular vesicles: the key to unlocking mechanisms of age-related vascular disease? *J Cardiovasc Aging* 2024;4:12. https://dx.doi.org/10.20517/jca.2023.49

Received: 5 Dec 2023 First Decision: 3 Jan 2024 Revised: 18 Jan 2024 Accepted: 19 Jan 2024 Published: 1 Feb 2024

Academic Editor: Ali J. Marian Copy Editor: Fangyuan Liu Production Editor: Fangyuan Liu

Abstract

Aging is associated with the development of two ubiquitous, detrimental pathologies, vascular calcification and amyloidosis. These pathologies are characterized by the accumulation of toxic aggregates in the vessel extracellular matrix (ECM) in the form of crystalline calcium and phosphate mineral or insoluble protein fibrils, respectively. These aggregates impact ECM integrity, drive vascular stiffening, and can also cause cell death and phenotypic change in the cells that interact with them. The deposition of both calcification and amyloid requires a nucleus that can mediate the mineralization of calcium and phosphate, or amyloid aggregation from precursor proteins or peptides. Emerging evidence suggests that changes in the composition of the ECM associated with cellular senescence, as well as extracellular vesicle (EV) release, cargo-loading, trapping, and aggregation within the ECM, are common and synergistic mechanisms that regulate the development of these pathologies. Importantly, vascular smooth muscle cells (VSMCs) orchestrate the formation of both pathologies that commonly co-occur in the aging vasculature. Here, we outline the commonalities and differences in what is known about the genesis of calcification and amyloid, and highlight key questions and areas that remain unknown and require further investigation. The complex relationship between senescence, EVs, and the ECM, mediated by VSMCs, which drives the accumulation of HA and amyloid, could be a target for therapeutic intervention.

Keywords: Senescence, extracellular vesicles, calcification, amyloid, vascular smooth muscle cells



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INTRODUCTION

Advancing age is the major risk factor for cardiovascular disease (CVD). With age, the occurrence and prevalence of coronary heart disease, hypertension, and atherosclerosis increase dramatically, making CVD the leading cause of death worldwide, representing 30% of all deaths^[1]. By 2050, the incidence of CVD could double in conjunction with increases in life expectancy and damaging trends, such as obesity and smoking^[2]. A significant feature of vascular aging is increased vascular stiffness, which can worsen chronic conditions, such as hypertension, contributing to CVD-related mortality rates^[3,4]. Pulse wave velocity (PWV), a clinical measure of arterial stiffness, is an independent indicator of cardiovascular risk and is used as a marker of vascular aging^[5]. Age-related diseases, such as chronic kidney disease (CKD)^[6], atherosclerosis^[7], Alzheimer's disease (AD)^[8], and dementia^[9], are all associated with increased PWV, which is a predictor of all-cause mortality as well as cardiovascular mortality^[10,11], highlighting the significance of vascular stiffening in age-related decline.

Vascular calcification and vascular amyloidosis are two age-related pathologies that are highly prevalent in patients over the age of 60 years^[12-14]. These pathologies affect both the intimal and medial layers of the vessel wall and accelerate vascular stiffening, thus promoting significant negative outcomes^[15,16]. Both pathologies are mediated by vascular smooth muscle cells (VSMCs) and emerging data suggest that similar mechanisms may contribute to their progression with age. However, despite this dramatic association with age, their co-occurrence and possible common and/or interacting mechanisms have not been systematically investigated.

Vascular calcification is characterized by the accumulation of hydroxyapatite (HA), a calcium phosphate mineral $[Ca_3(PO_4)_2]$. There is a strong association between age and medial calcification^[17], with vascular calcification also being closely correlated to biological aging, as younger patients with calcification have advanced aging of the vasculature and the survival risk of a person significantly older^[18]. Previously considered a passive, degenerative process associated with aging, it is now recognized as an active and dynamic process orchestrated by VSMCs and involving the upregulation of genes associated with both osteogenesis and chondrogenesis, with many similarities to the formation of bone^[19].

Calcification occurs at two anatomical sites in the vessel wall, the intima and the media. Intimal calcification is linked with atherosclerosis and is characterized by HA accumulation with lipids, macrophages, and VSMCs. This causes inflammation, growth, and potentially destabilization of atherosclerotic plaques [Figure 1]^[20,21]. Conversely, medial calcification has been associated with arteriosclerosis, an age-associated disease that is accelerated in association with metabolic disorders such as diabetes and renal failure^[22,23]. Medial calcification is associated with structural and ECM changes in the arterial tunica media, usually augmenting vascular stiffness^[20].

Amyloidosis is the accumulation of insoluble, fibrous amyloid, formed from the aggregation of soluble proteins or peptide monomers into β -sheets^[24]. Vascular amyloidosis occurs in the vessel intima and media and several amyloids have been described in the cardiovascular system^[25]. Cardiovascular amyloids are commonly classified as age-related and are often not associated with antecedent disease, with age being the only risk factor^[26,27]. In the vessel wall, the most common amyloids are aortic medial amyloid (AMA), amyloid transthyretin (ATTR), apolipoprotein A-1, and amyloid light chain (LC)^[28,29]. To date, there has been limited research investigating the pathological role of amyloid deposits in the vasculature. The two amyloids that have received the most attention are medin, the cause of AMA, which is present in the aorta of over 97% of Caucasian people over 50 years old^[30], and A β , the same amyloid that causes Alzheimer's Disease (AD), that accumulates in leptomeningeal vessels of the brain in cerebral amyloid angiopathy



Figure 1. The vessel wall is organized into 3 layers. The intimal layer is made up of one layer of endothelial cells and the internal elastic lamina. The medial layer of a healthy vessel consists of contractile vascular smooth muscle cells (VSMCs), the extracellular matrix (ECM), and elastic fibers. The adventitia is the outer layer of connective tissue. Vascular calcification can occur at the intimal and medial layers with different risk factors and outcomes. Intimal calcification is commonly associated with atherosclerosis and is characterized by formation of a plaque with a fibrous cap and accumulation of lipids and inflammatory cells. Medial calcification often results in arteriosclerosis and vascular stiffening. VSMCs are the main mediators of medial calcification and can undergo phenotypic differentiation into a more synthetic, osteoblast-like phenotype, characterized by extracellular matrix (ECM) remodeling and extracellular vesicle (EV) secretion. (Created with BioRender.com)

(CAA). Previous studies have suggested that medin accumulation in the vessel wall can reduce arterial elasticity through its interaction with elastic fibers^[31], while stiffening of cerebral arteries is thought to accelerate A β accumulation in the brain^[32,33]. However, more studies are required to delineate exactly the effects of amyloidosis in the vasculature.

A key feature of aging is the accumulation of senescent cells^[34]. These cells show a number of specific features including exit from the cell cycle, morphology change, persistent DNA damage, and activation of the senescence-associated secretory phenotype (SASP), leading to the release of inflammatory mediators and proteases that can act remotely to accelerate cellular dysfunction and aging. In the vessel wall, senescent cells accumulate with age and premature senescence of VSMCs is a feature of young patients with calcified arteries, such as children on dialysis^[18,35]. Extensive research has been conducted on the impact of senescence on vascular calcification^[36]; however, there remains a lack of knowledge regarding the relationship between senescence, calcification, and amyloidosis, despite several forms of pathological amyloidosis being associated with aging.

Senescent VSMCs have also been shown to increase the secretion of extracellular vesicles (EVs)^[37]. The role of EVs in calcification and amyloidosis in the vasculature is interesting as both pathologies require a nucleus to initiate either calcification or amyloid formation. VSMC-derived EVs and exosomes have been implicated in the deposition and nucleation of both HA and amyloid^[38-40], but the precise mechanisms governing what initiates EVs to mineralize, aggregate protein, or both are unknown. However, it is clear that EVs are an important trigger for both pathologies and could account for their concomitant development in the vessel wall with age.

Building on these key facts, in this review, we will focus on the role of senescence and the involvement of EVs in driving these pathologies. The co-occurrence of calcification and amyloidosis, particularly in the vasculature, and their common and reciprocal mechanisms need further investigation to tease apart their roles in CVD and identify novel therapeutic targets to alleviate vascular stiffening.

MECHANISMS OF VASCULAR CALCIFICATION AND AMYLOID DEPOSITION: A KEY ROLE FOR VSMCS

Vascular calcification

VSMCs in the arterial tunica media have a contractile phenotype, providing dynamic adaptability to blood volume changes occurring in the vasculature. In healthy adults, VSMCs exhibit a contractile phenotype with low proliferation, but with aging and the occurrence of chronic inflammation and structural damage, VSMCs switch to a more proliferative/synthetic phenotype and migrate to the site of injury [Figure 2]^[41,42]. This synthetic characteristic provides cytokines and growth factors and induces overexpression of elastin, collagens, proteoglycans, and other ECM components to repair the damaged wall^[43]. Synthetic VSMCs secrete EVs containing calcification inhibitors, β 1 integrin, and ECM proteins such as fibronectin and collagen VI, which stimulate cell migration and invasion^[44].

Calcification is a dynamic process involving VSMC osteogenic differentiation, with overexpression of proosteogenic genes, enhancing the accumulation of HA [Figure 2]^[16,45]. During the initial stages of calcification, runt-related transcription factor (Runx2), which regulates osteogenic differentiation during the physiological formation of bones, is upregulated. In addition, other transcription factors (TF) and morphogens that can drive osteo/chondrogenic pathways, such as Sox9 or Msx2, which are involved in cartilage differentiation and bone maturation, respectively, and bone morphogenetic protein 2 (BMP2) are upregulated^[46,47]. These TFs also regulate the expression of a number of proteins such as matrix metalloproteinases (MMPs), osteocalcin (OCN), collagen type 1-alpha1 (COL1 α 1), bone sialoprotein (BSP), and alkaline phosphatase (ALP) that can regulate mineral growth^[48]. VSMC osteogenic differentiation is accompanied by downregulation of VSMC contractile genes, such as smooth muscle protein 22- α (SM22 α) and α -smooth muscle actin (α -SMA), which also control cell motility and shape^[49].

VSMCs can respond to pathological stimuli by expressing calcification inhibitors. For instance, matrix Gla (γ -carboxylate glutamate) protein (MGP) is a potent calcium inhibitor that inhibits BMP2. It has been documented that during vascular calcification, these protective mechanisms can become defective, and VSMCs start to release EVs, including apoptotic bodies and exosomes, which enhance calcification by providing an essential nucleation point for HA formation^[50,51]. In the healthy vasculature, these EVs are loaded with mineralization inhibitors, but during calcification, these are deficient and the EVs become mineralization-competent, acting as a nucleation nidus [Figure 2].

Vascular amyloid

Like HA, amyloid fibril formation is a nucleation-dependent mechanism. It can be initiated by soluble, cleaved peptide monomers or destabilized protein structures, which are susceptible to aggregation into oligomers, which may induce cell death, or further aggregation into amyloid. Partial unfolding of proteins can occur at high temperatures or low pH^[52]. Aggregation of denatured monomers or amyloidogenic peptides occurs slowly during the lag phase and occurs when hydrophobic regions of proteins, which are usually embedded in the protein structure, become exposed through cleavage or denaturation and begin to aggregate with other hydrophobic regions or peptides [Figure 3]. The growth of unstructured aggregates and amyloid seeds into mature fibrils occurs exponentially during the growth phase to more thermodynamically stable fibrils^[53]. Amyloid plaques generally comprise several components, the fibrils



Figure 2. Contractile vascular smooth muscle cells (VSMCs) express high levels of contractility markers, including α -smooth muscle actin (α -SMA) and smooth muscle protein 22 α (SM22 α), to regulate vascular tone. Synthetic VSMCs have low contractility, proliferation and secrete extracellular vesicles (EVs) that can promote VSMC migration and invasion to drive vascular repair after injury. Elevated calcium and phosphate levels (Ca/P) can induce an osteogenic VSMC phenotype with bone-like markers, extracellular matrix (ECM) deposition, and apoptosis. Osteogenic VSMCs secrete calcifying EVs, which form a nidus for mineralization due to the loss of calcification inhibitors and being rich in annexins and phosphatidylserine.



Figure 3. Amyloid fibrils are formed by denaturation of a native state protein to form a denatured monomer or cleavage to form amyloidogenic peptides. These aggregate slowly to form amyloid seeds during the lag phase. The amyloid seeds rapidly aggregate into β -sheets in the exponential growth phase to form insoluble amyloid fibrils. Extracellular vesicles have been identified as mediators of amyloid formation as they contain several different amyloid precursor proteins or peptides. They can accelerate the aggregation of amyloid peptides by acting as a nucleus for aggregation and fibril growth. (Created with BioRender.com)

from denatured protein or peptide precursor, non-fibrillar serum amyloid P component (SAP), glycosaminoglycans (GAGs), heparin sulfate proteoglycans (HSPGs), and apolipoprotein E (apoE). Previous research suggests SAP and apoE stabilize amyloid fibrils and protect them from degradation by increasing fibril strength and density^[54].

AMA is a localized, aortic, age-related amyloid. It is a major cardiovascular amyloid, comprising a 50-amino acid peptide medin, produced by cleavage of milk fat globule epidermal growth factor (EGF) 8 (MFGE8)^[30,31,55]. MFGE8 is highly expressed by VSMCs and accumulates in the vessel wall with age^[56], with its accumulation also correlating with PWV and cardiovascular risk factors in diabetic patients^[57]. It contains a domain with an RGD peptide that binds integrins to mediate several processes in VSMCs, including proliferation^[58], osteogenic differentiation^[59], and inflammation^[60]. Medin is composed of a fragment from the C2-like domain of MFGE8, which can bind phospholipids. Amyloid deposits in the medial layer of the thoracic aorta are present in over 97% of Caucasian people over 50 years old^[30]. Both MFGE8 and medin have been shown to bind to elastin, accounting for its accumulation in large arteries^[61], and previous studies have suggested that medin accumulation in the vessel wall can reduce arterial elasticity through its interaction with elastic fibers^[31].

Amyloid- β (A β) peptides are recognized as the primary driver of AD. Their accumulation leads to the formation of characteristic plaques within the aging brain parenchyma. The aggregation of A β peptides has also been identified in crucial cerebral blood vessels, such as the internal carotid artery, resulting in CAA^[62]. A β accumulates predominantly in the medial and adventitial layers of leptomeningeal arteries, indicating a role of VSMCs in its deposition^[63]. Cerebral arteries play a key role in the clearance of A β through the perivascular drainage pathway, which is reduced with age and in AD, leading to A β deposition in the ECM and its uptake by VSMCs, causing disruption of the vessel wall^[64,65]. Furthermore, substantial amounts of A β can be detected in the circulation and accumulate in atherosclerotic plaques in the carotid artery and aorta^[66,67]. Interestingly, circulating A β levels have been shown to be predictive of cardiovascular mortality and vascular disease progression in patients with coronary heart disease and were significantly associated with arterial stiffness and atherosclerosis^[68,69]. However, despite its presence in the cardiovascular system and association with vascular disease, its pathological role and mechanisms of accumulation have not been investigated.

AGING AND SENESCENCE

Cellular senescence is a hallmark of vascular aging, calcification, and amyloidosis^[34]. This physiological process is characterized by telomere shortening, DNA damage, and arrest of the cell cycle, with cells unable to respond correctly to mitotic stimuli. Cellular senescence triggers several cell cycle regulators, such as p53/ p21 and p16/retinoblastoma protein tumor suppressor activation^[70,71], and plays a crucial role in vascular aging through increased secretion of inflammatory cytokines, proteases, and growth factors, comprising the SASP.

Recent studies have demonstrated that VSMCs rapidly undergo osteogenic differentiation during aging^[72,73], which could both be driven by, and contribute to, cellular senescence. VSMC senescence *in vitro* has been shown to enhance calcification and increase the expression of osteogenic genes, such as Runx2, ALP, and collagen 1^[73]. In klotho knockout mice, a mouse model of aging, significantly upregulated expression of RUNX2 was shown in the calcified lesions in the aorta^[74]. Interestingly, CKD patients have decreased klotho levels, which decline further as the disease progresses^[75]. Senescent VSMCs and apoptotic bodies have also been detected in the medial layer of calcified arteries from children with CKD, with their vascular phenotype more closely resembling that of old subjects^[45,76,77], highlighting the role of senescence in driving calcification during aging.

VSMC senescence also contributes to calcification by the release of inflammatory cytokines and growth factors. *In vitro* replicative senescence of VSMCs shows correlations with age-related morphological changes and DNA damage accumulation, which can lead to the overexpression of pro-osteogenic genes and downregulation of calcification inhibitors^[72,73]. Inflammatory cytokines, particularly interleukins (ILs), are highly activated during the SASP, with IL-6 promoting calcification through osteogenic gene upregulation (Runx2, ALP, and MMP2) via extracellular related kinase (ERK) signaling pathways^[78].

There is strong evidence showing the accumulation of senescent VSMCs in atherosclerotic plaques. VSMC senescence in atherosclerosis is thought to be induced by replication, resulting in telomere shortening, or stress-induced premature senescence (SIPS), caused by DNA damage and oxidative stress^[79,80]. While it is not clear what role senescence-induced calcification has in plaque formation and stability, it has been shown to be closely correlated with cardiovascular events, providing further evidence for the importance of VSMC senescence in both the intima and media of the vessel wall^[81,82].

Extracellular proteases are also components of the SASP. MMPs play an important role in both bone and vascular tissue remodeling^[83]. Several studies have identified MMPs as promoters of vascular calcification^[84,85]. MMPs degrade several key ECM components that are important in maintaining integrity of the vessel wall, such as elastin, fibronectin, and laminin. Previous research showed that the degradation of elastin preceded VSMC loss and vascular calcification in a model of CKD^[86]. Therefore, enhanced matrix degradation, induced by SASP, could contribute to vascular calcification. In addition to matrix degradation, MMP2 and MMP9 have been shown to enhance vascular calcification through upregulation of BMP2^[85].

To date, there is limited understanding of the contribution of senescence to amyloidosis, particularly in the cardiovascular system. Although AMA is an extremely prevalent and age-associated form of amyloidosis, the role of cellular aging in its development is under-researched. Accumulation of medin in the medial layer of human aortas has been strongly correlated with age^[36], and a recent study showed VSMC senescence increased the production and secretion of medin in sEVs^[37]. This resulted in increased accumulation of medin in the ECM, promoting its aggregation into amyloid fibrils, indicating a role of senescence in triggering AMA development.

Senescent cells have been detected in the vasculature in both AD patients and AD mouse models and are thought to contribute to increased permeability and loss of integrity of the blood-brain barrier^[87]. VSMC senescence has not yet been investigated thoroughly in CAA, which overlaps significantly with AD. Cellular senescence plays a role in AD, although its precise mechanisms are not fully understood^[88]. Various cell types, such as astrocytes, neurons, and endothelial cells, express senescence-associated proteins in AD brains, indicating a link between senescence and Aβ pathophysiology^[89,90].

EXTRACELLULAR VESICLES

An important mechanism of vascular calcification is the secretion and deposition of EVs from VSMCs. EVs have also been linked to amyloid-related diseases in both physiological and pathological contexts, including $AD^{[91,92]}$ and $AMA^{[37,93]}$.

EVs are key mediators of intercellular communication, acting as vehicles for transporting proteins, lipids, and RNA^[94]. Importantly, EVs comprise different subtypes of vesicles, often categorized by size, biogenesis, or isolation technique. Microvesicles (200-1,000 nm) outwardly bud off directly from the plasma membrane

of cells, while small EVs (sEVs) or exosomes (10-200 nm) are formed within endosomes [Figure 4]. Apoptotic bodies (1-5 μ m) are formed from cell fragments during apoptosis.

Early endosomes are formed from inward budding of the cell membrane, forming clathrin-coated vesicles (CCVs), or from vesicles from the Golgi apparatus. Intraluminal vesicles (ILVs) are small, lipid bilayerbound vesicles, which inwardly bud from the early endosomal membrane^[95]. This becomes a multivesicular endosome (MVE) and the ILVs are released as exosomes, sometimes described as sEVs, when the MVEs fuse with the plasma membrane^[96]. Alternatively, the ILVs are degraded by the fusion of the MVEs with lysosomes^[96]. Exosomes are released from a variety of cell types, including blood cells, endothelial cells, and VSMCs and can interact with recipient cells through three different mechanisms: direct interaction between exosome transmembrane proteins and target cell receptors; fusion with the target plasma membrane, followed by the release of the contents into the cytosol; and internalization of exosomes into the cell^[97]. Internalized exosomes can merge into endosomes, be released again into neighboring cells, or mature into lysosomes, leading to degradation^[98].

Exosomes and sEVs consist of a lipid bilayer with several membrane proteins: major histocompatibility complexes (MHC), which are involved in antigen presentation; integrins and tetraspanins, for cell targeting and adhesion; annexins and flotillin, for membrane trafficking; and lipid rafts, which influence membrane fluidity and membrane protein and receptor trafficking^[96]. They are loaded with a variety of proteins both on the surface and within the lumen, such as adhesion molecules and matrix-degrading enzymes, cytoskeletal proteins and enzymes, mRNA, and microRNAs.

Microvesicles are formed from direct budding of the plasma membrane and comprise a heterogeneous population of vesicles, differing based on cell type and function. The plasma membrane may undergo remodeling, meaning the composition of microvesicles can appear distinct from the parent cell^[99]. Microvesicles can be packed with miRNAs, mRNAs, and proteins and, like exosomes or sEVs, can act as a mechanism for the transport of bioactive molecules and signaling.

Apoptotic bodies are formed during the disassembly and fragmentation of apoptotic cells. During apoptosis, cells shrink, and the intracellular contents become more densely packed. The plasma membrane then undergoes blebbing, forming apoptotic bodies, which contain components of the apoptotic cells, such as cytoplasm, organelles, chromatin, and other nuclear remnants^[100]. The cargo of apoptotic bodies is very heterogeneous as contents are randomly packed into the vesicles.

EVs in vascular calcification

The change of VSMCs to a synthetic phenotype is a characteristic of vascular repair in calcifying conditions and aging and is associated with increased EV secretion^[38]. Specifically, EV production and contents change with disease and stress, highlighting their roles in both normal, physiological, and pathological processes^[101].

A previous study on vascular calcification demonstrated how calcium phosphate deposition in the medial layer of major arteries was mediated by exosomes. During HA deposition, there was a decreased concentration of calcification inhibitors and a disruption of vascular homeostasis^[4]. Interestingly, VSMC-derived exosomes contain key calcification inhibitors, such as MGP and fetuin-A^[102-104]. In chronic kidney disease, high concentrations of calcium and phosphate induce mineral stress, which causes a reduction of calcification inhibitors secreted by exosomes^[39].



Figure 4. Blebbing of the plasma membrane causes the formation of microvesicles or medium extracellular vesicles (EVs). Exosomes or small EVs are formed in early endosomes from clathrin-coated vesicles or Golgi apparatus, where budding of the membrane to create intraluminal vesicles results in maturation into a multivesicular endosome (MVE). The MVE either fuses with lysosomes to be degraded or fuses with the plasma membrane to release the intraluminal vesicles as exosomes. Exosomes or small EVs contain a myriad of proteins and molecules, including microRNAs (miRNAs), extracellular matrix components, matrix metalloproteinases (MMPs), annexins, and several different amyloid proteins. (Created with BioRender.com)

Elevated levels of calcium induced sphingomyelin phosphodiesterase 3 (SMPD3) expression and secretion of VSMC-derived exosomes, while chemical inhibition of SMPD3 inhibited the calcification process^[39]. Exosomes contain calcium and are rich in calcium-binding annexins [Table 1], so during calcification, exosomes act as nucleation sites for the formation of HA deposits through binding of two exosomal membrane proteins: annexin A6 and phosphatidylserine (PS)^[38]. Aggregation of EVs, through annexins on their surface, and their tethering to ECM components actively enhance their mineralization capacity^[105,106]. Recently, senescence has been implicated in promoting the nucleation of EVs as a study showed EVs from senescent cells were high in calcium and contained increased levels of annexin A2, annexin A6, and BMP2^[12,107].

In addition to exosomes, other types of EVs have been shown to be involved in calcification; PS in apoptotic bodies helped the formation of calcium phosphate, providing a nucleation site for mineral growth^[51]. In addition, a study found microvesicles extracted from the plasma of a group of elderly people had higher concentrations of annexin (which may act as possible nucleation sites), promoting vascular calcification; this correlated with high concentrations of calcium and increased quantities of bone-related proteins^[107].

	Role of EVs	EV cargo	Ref.
Vascular calcification	Nucleus for mineral growth	Annexin A1, annexin 2, annexin A5, annexin A6, BMP2, TNAP, phosphatidylserine, hydroxyapatite, Grp78, collagen I	[38,39,105 -108]
Amyloidosis	Cleavage of amyloid proteins	APP, PSEN1, BACE1	[91,109]
	Secretion of amyloidogenic proteins and peptides	Medin, MFGE8, amyloid- β , tau, prion protein, $\alpha\text{-synuclein},$ PMEL, SOD1	[37,110-113]
	Promotion and acceleration of amyloid aggregation	Apolipoprotein E, HSPG2, cellular prion protein, cholesterol, glycosphingolipids	[37,114-116]
	Facilitating uptake of amyloid seeds by neighboring cells	Phosphatidylserine	[116]
ECM aging	Degradation of ECM components	MMP2, MMP14, ADAM9, ADAM10, ADAMTS5	[117-119]
	Collagen crosslinking	LOXL2, PLOD1, PLOD2, PLOD3, P3H1, TGM2	[119,120]

Table 1. Role of EVs and their cargo in calcification, amyloidosis and ECM aging

EVs: Extracellular vesicles; BMP2: bone morphogenetic protein 2; TNAP: tissue non-specific alkaline phosphatase; APP: amyloid precursor protein; PSEN1: presenilin 1; BACE1: beta-secretase 1; MFGE8: milk fat globule EGF-factor 8; PMEL: premelanosome protein; SOD1: superoxide dismutase 1; HSPG2: heparan sulfate proteoglycan 2; MMP: matrix metalloproteinase; ADAM: a disintegrin and metalloproteinase; ADAMTS5: a disintigrin and metalloproteinase with thrombospondin motifs 5; LOXL2: lysyl oxidase homolog 2; PLOD: procollagen-lysine,2-oxoglutarate 5-dioxygenase; P3H1: prolyl 3-hydroxylase 1; TGM2: transglutaminase 2.

EVs may also contain miRNAs, small endogenous molecules that regulate mRNA transcription repression through binding to the UTR^[121,122]. Altered levels of pro-calcific miRNAs contained in VSMC-derived calcified exosomes in the extracellular space have been shown to contribute to the pro-osteogenic phenotype of VSMCs^[123]. Furthermore, many miRNAs identified in exosomes can regulate Runx2 and Smad1 gene expression^[124]. Together, these studies demonstrate a key role for EVs in vascular calcification through several distinct mechanisms.

EVs in amyloidosis

Recently, there has been growing interest in the cargo of EVs, particularly in the context of amyloid-related pathologies. Exosomes, sEVs, and MVEs have also been identified as mediators in physiological and age-related pathological amyloid metabolism^[40,125-129]. Several amyloid proteins have been identified in EVs, including A β , medin, tau, and α -synuclein [Table 1]. To date, research has focussed mainly on the role of EVs in neurodegenerative diseases. For example, prion proteins (PrP), which cause transmissible spongiform encephalopathies, have been associated with exosomes from neurons, epithelial cells, and neuroglial cells, and increased exosome secretion was shown to increase the infectiousness of abnormal prion proteins (PrP^{Sc})^[110,114,130]. Exosomes from neurons and cerebral spinal fluid can also contain α -synuclein, a component of Lewy bodies, which is a characteristic of Parkinson's disease and dementia with Lewy bodies^[111,131].

Recent research has shown that VSMC-derived sEVs drive aggregation and accumulation of medin amyloid^[37]. sEVs were shown to deposit small medin aggregates into the ECM and accelerate the aggregation of medin into amyloid fibrils by acting as a nucleus for fibril growth. Senescent VSMCs secreted more sEVs and increased deposition of medin in the ECM and inhibition of sEV secretion with 3-OMS treatment attenuated this. Interestingly, while 3-OMS decreased the overall amount of medin in the senescent ECM, it could not prevent the accumulation of medin in a fibril-like form. Furthermore, super-resolution microscopy showed sEVs from senescent VSMCs were coated with large fibril-like medin seeds, suggesting senescent sEVs may be a sufficient trigger for fibril formation.

Not all amyloids are pathological, as there have been some amyloids identified that are thought to have physiological roles^[132]. Exosomes have been shown to play a pivotal role in the formation of amyloid fibrils from fragments of the premelanosomal protein (PMEL) in melanocytes^[133]. These fibrils act as a scaffold for the polymerization of melanin, which protects cells from UV and oxidative stress. Cleavage of PMEL into amyloidogenic fragments occurs in MVEs by β -secretase 2 (BACE2) and their aggregation into fibrils is nucleated by ILVs, which are formed in MVEs prior to secretion as exosomes^[129,133].

A similar mechanism to PMEL processing is observed during the accumulation of $A\beta$ in the brain in AD. Numerous in vivo studies have shown that circulating exosomes in AD mouse models and AD patient body fluids, including blood and CSF, contain the C-terminal fragments of amyloid precursor protein (APP)^[109] and soluble Aβ42^[134,135]. Processing of APP resulting in amyloid production occurs predominantly in the endosomal pathway, while non-amyloidogenic APP processing occurs at the cell surface membrane^[125]. A β peptides are formed from cleavage of the membrane APP by β -secretase 1 (BACE1) in MVEs, followed by cleavage of the membrane-anchored C-terminal fragment by γ -secretase. The optimum activity of BACE1 is at acidic pH^[136], explaining its dominant localization at the endosomal membrane, as endosomal compartments have acidic pH $(6.1-6.8)^{[137]}$. A β peptides are sorted in MVEs and are loaded into the ILVs, which are then released as exosomes^[92]. Following secretion, exosomes could function as a nucleation site for amyloid plaque formation^[92,138]. The release of A β into the extracellular space could also infect neighboring cells and act as amyloid seeds to accelerate the progression of disease^[138,139]. Interestingly, increasing exosome secretion from neuronal cells has also been described as beneficial, with a study showing A β -laden exosomes from neurons were taken up by microglia for degradation. Despite exosomes significantly promoting amyloidosis, there may also be a role for them in $A\beta$ clearance which is a method for AD treatment that needs further investigation^[116].

EVs in senescence

Cellular senescence triggers changes in EV secretion and cargo loading, which can propagate aging processes through paracrine and endocrine signaling, driving inflammation, gene expression, and metabolic changes in recipient cells^[140]. Senescent cell-derived EVs can carry nuclear and mitochondrial DNA which activate immune signaling, such as the cyclin GMP-AMP synthase (cGAS)-stimulator of interferon genes (STING) pathway, or toll-like receptor 9 (TLR9) mediated inflammatory response^[141]. EV secretion was increased from senescent VSMCs and senescent VSMC-derived sEVs accelerated medin aggregation into fibrils through changes in sEV cargo and altered binding to HSPG2 in the ECM^[37,93]. Senescent endothelial cell-derived EVs are enriched in annexins A2 and A6, BMP2, and calcium and can promote calcification of VSMCs^[107]. Just as "senescent EVs" induce an aging phenotype and enhance both calcification and amyloidosis, EVs from young cells can improve an aging phenotype. EVs from young cardiac progenitor cells reversed cell senescence *in vitro* and *in vivo* through telomere elongation, inhibition of the DNA damage response, and reduced inflammation in cultured human fibroblasts and cartilage-derived stem/ progenitor cells in a naturally aged rat model^[142]. Previous studies have also demonstrated the role of several miRNAs, including miR-34a and miR-126b, in mediating the effects of young EVs and highlighted the potential of EVs as a novel treatment for aging pathologies^[143,144].

EVs have been shown to regulate vascular calcification, amyloidosis, and cellular senescence, indicating a possible link between these aging pathologies and their contribution to vascular stiffening and CVD. The role of EV cargo, their interaction with the extracellular environment, and mechanisms regulating their secretion and accumulation need clarification.

EXTRACELLULAR MATRIX AGING AND EVS

Both calcification and amyloidosis accumulate in the extracellular space in the vessel wall, regulated in part by the secretion of EVs. The ECM is a key component of the vascular wall and is essential for the maintenance of cellular homeostasis, acting as a scaffold and supporting cellular organization in many tissues and organs. It is comprised of several families of proteins, such as collagens, proteoglycans (PGs), and MMPs, which can influence its adherent, morphological, and phenotypical characteristics. The ratio of collagens, PGs, and elastin is essential to preserve its mechanical properties. The ECM is considered actively dynamic since it frequently undergoes remodeling by proteases, in both physiological and pathological events. The remodeling process is predominantly controlled by MMPs and their inhibitors, which regulate the balance between matrix degradation and production^[145]. Dysregulation of this balance is typical of aging pathologies, such as fibrosis^[146], and therefore, might be implicated in cerebrovascular amyloidosis and calcification. EVs directly impact the ECM and contribute to deposition and remodeling in both physiological and pathological environments. Through their surface-bound enzymes, which include several MMPs, elastases, and collagen crosslinking enzymes, EVs can actively degrade the ECM, resulting in the turnover of matrix components [Table 1]^[117,119].

Vascular aging is strongly associated with ECM aging, as matrix stiffness is the result of a series of processes occurring with age, with major changes to the mechanical forces affecting cellular function. For instance, when elastin starts to lose its elasticity, the mechanical force stress is transferred to collagen and fibronectin which is less elastic, hindering the function of the ECM, reducing its stability, and affecting cell interactions. Hence, this process appears deleterious, because when matrix begins to stiffen, it also activates downstream molecular pathways, which enhance apoptosis^[147]. A recent study showed VSMC senescence to be a key regulator of increased ECM stiffness, through the TF Sox9^[119]. Changes in the composition and stiffness of the senescent ECM were able to induce a senescent phenotype in young VSMCs, including cell cycle arrest and DNA damage, causing a positive feedback loop. An interesting aspect of this study was the role of EVs, as there was an accumulation of EVs in the ECM from senescent or Sox9 overexpressing VSMCs, compared to the young or control VSMCs, respectively. The EVs were loaded with procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (PLOD3), an enzyme that caused increased ECM stiffness through collagen crosslinking. Previous studies into lung diseases, such as chronic obstructive pulmonary disease (COPD), identified subsets of EVs that drive ECM-specific proteolysis by MMPs and elastases, severe enough to cause loss of alveolar units^[148,149]. EVs from subjects with COPD were also a sufficient stimulus to cause a COPD phenotype in mice primarily through surface-bound neutrophil elastase^[149], further demonstrating an important relationship between the ECM and EVs with aging and disease.

Several pathological mechanisms occur during ECM aging. Collagen crosslinking, the formation of advanced glycation end products (AGEs), and the deposition of lipids and calcium are the most studied. Interestingly, these modifications have also been detected during both arterial calcification^[150-153] and amyloidosis^[154,155].

ECM is known to have a role in bone formation and remodeling^[83] beyond its crucial signaling function^[156]. Interestingly, elastin degradation by matrix proteolytic enzymes during aging and CKD leads to vascular calcification and VSMCs switch to a synthetic phenotype^[86]. MMPs, such as MMP2 and MMP9, are considered calcification enhancers during senescence. Indeed, those enzymes upregulate BMP2, promoting VSMC calcification and degradation of other key ECM components of the vascular tunica media such as elastin, fibronectin, and collagens^[84,85].

Changes in VSMC-derived ECM and basement membrane components caused by senescence have been described in AMA, where senescence increased HSPG2 expression and extracellular deposition, increasing medin accumulation and aggregation^[37]. Knock-down of HSPG2 in the senescent ECM attenuated medin fibril formation and deposition. Interestingly, in this study, decreasing sEV secretion decreased the accumulation of medin in the senescent ECM, but did not decrease the deposition of medin in a fibrillar form. Additionally, sEVs from senescent VSMCs contained larger medin fibril seeds and preferentially bound to HSPG2 in the ECM, indicating a complex relationship between cellular senescence, sEV secretion, and changes in ECM composition.

Changes in ECM composition have also been described in many neurodegenerative diseases, such as AD and CAA. Thickening and delamination of ECM and basement membrane have been shown to be present with greater severity in the AD brain^[157]. Data suggest that a change in ECM composition of aged brains creates an amyloidogenic environment as ECM proteins, such as collagen IV and fibronectin, are increased in AD brains, with fibronectin shown to bind A β and augment amyloid deposition *in vivo*^[158].

TOXICITY OF CALCIFICATION AND AMYLOID

Both calcification and amyloid have been shown to have toxic effects on the vasculature. Previous studies have described the toxicity of HA nanoparticles on VSMCs, which reduced cell viability through disruption of cell membranes, increased reactive oxygen species production, and mitochondrial dysfunction, resulting in cell necrosis^[159]. Uptake of HA particles by VSMCs was also shown to induce expression of osteogenic genes, such as BMP2, Runx2, and ALP, and induced cell death, causing accumulation of apoptotic bodies, indicating a positive feedback cycle where calcification exacerbates calcification. HA crystals in the vessel can also be taken up by macrophages, causing the secretion of several proinflammatory cytokines, including TNF α , IL-1 β , and IL-8, mediated by protein kinase C alpha (PKC α)^[160,161]. Inflammation is a hallmark of atherosclerosis, intimal and medial calcification and can promote VSMC apoptosis or osteogenic differentiation, again demonstrating how HA accumulation can exacerbate calcification.

The formation of amyloid fibrils from oligomeric aggregates has previously been discussed as a protective mechanism^[24,162,163]. Small, pre-fibrillar aggregates are toxic to cells, whereas mature amyloid fibrils, the precursor protein or monomeric form may not^[164]. A possible mechanism for this is the exposure of residues normally embedded inside the protein being exposed to and interacting with cell membranes, intracellular and extracellular components^[24,165,166]. In Aβ pathology, the accumulation of intracellular or extracellular oligomeric aggregates precedes the formation of extracellular plaques and contributes to the main pathological effects of AD^[167]. Similarly, previous research into the effects of medin showed that medin aggregates induced several pathological effects, such as endothelial dysfunction, vascular inflammation, and VSMC death^[168,169]. Acceleration of medin fibril formation by heparin decreased the toxic effects of medin, suggesting the formation of mature, benign amyloid fibrils in AMA could be a protective mechanism against the toxic nature of smaller medin aggregates^[170].

Both calcification and amyloidosis can contribute to ECM stiffening. HA accumulation and ECM remodeling during calcification both directly impact arterial stiffening, through increased turnover of collagen and degradation of elastin, causing weakening of the ECM^[171,172]. In addition, medial calcification is associated with increased PWV and increased pulse pressure, which are linked to the accelerated development of CVD^[22]. The pathological nature of amyloid is more complex as research has shown that fibrous amyloid deposits can contribute to reduced elasticity of vessels^[65]. However, a previous study showing AMA was associated with weakening of the aortic wall in the development of thoracic aortic aneurysm, demonstrated the main pathological effects of medin can be attributed to toxic pre-fibrillar

aggregates^[169]. This suggests that toxic aggregates could contribute to vascular pathologies, which may then be exacerbated by diminished elasticity and vessel wall weakening by amyloid fibrils.

CO-OCCURRENCE OF CALCIFICATION AND AMYLOID

Even though vascular calcification and amyloidosis are age-related pathologies occurring in patients over the age of 60, there have been few systematic studies investigating their co-occurrence^[12-14]. In other disease contexts, localized amyloid deposits have been associated with calcification in skin, larynx, and lungs, forming calcified tumoral amyloids, known as calcified amyloidomas^[173-175].

Amyloid deposits in the cardiovascular system have been observed in calcified regions of the coronary artery and aortic valves and have been associated with aortic stenosis and calcific aortic valve disease (CAVD)^[176]. Recent studies have suggested that amyloid deposition might play a direct role in the pathogenesis of CAVD. Analysis of the morphology of calcified valves described circular cavities within mineralized structures, filled with fibrous protein, which exhibited birefringence, indicating the presence of amyloid at the core^[177]. A study showed 88% of aortic valves from CAVD patients had amyloid accumulation, with all the patients over 50 years having amyloid deposition^[178]. Interestingly, only one-third of subjects under 30 years had amyloid deposits, indicating a potential link with aging, CAVD, and amyloid. Immunohistochemical staining was used to try to identify the amyloid protein involved, with apoA-I being detected in the mineralized areas of the valves. In vitro treatment of valve interstitial cells with amyloid extracts from calcified valves enhanced mineralization and apoptosis^[177]. These studies highlighted the potential effect of apoA-I amyloid as an "interface" between the organic and mineral phases in CAVD. A previous study identified sEVs, described as matrix vesicles (MVs), as involved in CAVD, with annexin-rich MVs detected in calcified human aortic valves^[179]. An interesting aspect of CAVD to investigate further would be the mechanisms regulating the accumulation of both mineral and amyloid in the valve to identify common features, such as EVs, and identify novel therapeutic targets.

In the early 1960s, *Ratinov* first used *ex vivo* radiography and microscopic analysis to reveal the presence of intracranial arterial calcification (IAC) in the extradural region of human internal carotid arteries^[180]. IAC has been documented in many Caucasian and Asian populations and was shown to be a major element of cerebral atherosclerosis^[181-183]. Recently, IAC has gained scientific interest as it is thought to be a significant cause of ischemic stroke^[184]. IAC shares some common risk factors with amyloidosis and other cardiovascular diseases. Aging appears to be a major risk factor for IAC; an early study on the pathology did not observe any sign of IAC in patients who were 20 to 30 years old. Moreover, further research has demonstrated that IAC clinical outcomes worsen in concomitance with increasing age^[181,182,185]. In addition, the imbalance of calcium and phosphate levels can be considered a risk factor as a high percentage of IAC cases have been recorded in patients with CKD^[186-188].

Recent investigations have brought to light a significant association between a hereditary variant of CAA, known as the Dutch type, and IAC^[189]. Studies have revealed a connection between osteopontin (OPN), and the TGF β 1/phospho-SMAD2/3 signaling pathway in calcified arteries affected by hereditary CAA^[190,191]. Interestingly, the researchers observed a correlation between the pro-calcific accumulation of collagen 1, OPN, and the activation of the pSMAD2/3 pathway with the severity of CAA. This observation underscored the vital role of the ECM in calcified vessels impacted by hereditary CAA^[189]. Based on their findings and the spatial relationship between vascular calcification and the deposition of A β in the brain vessel medial layer, the researchers proposed that calcification is more likely a characteristic of advanced-stage CAA rather than a cause of amyloid deposition.

Investigations into hereditary CAA cases evidenced a consistent involvement of basal ganglia vessels in IAC^[192]. The calcification patterns observed in vessels affected by hereditary CAA are categorized as a subtype of microvasculopathies associated with CAA, presenting as a secondary manifestation of the disease^[189]. Nonetheless, unraveling the underlying mechanisms that underpin the calcification of certain vessels in the context of CAA remains a mystery that warrants further exploration.

COMMON AND RECIPROCAL MECHANISMS OF CALCIFICATION AND AMYLOIDOSIS

The role of EVs in calcification and amyloidosis in the vasculature is interesting, as VSMC-derived sEVs and exosomes have been implicated in the deposition and nucleation of both HA and amyloid. The precise mechanisms governing what initiates EVs to mineralize, aggregate protein, or both are unknown, but it is clear that EVs are an important trigger for both pathologies and could account for their concomitant development in the vessel wall with age.

As for the reciprocal effects of calcification and amyloid, evidence suggests amyloid could enhance vascular calcification through different mechanisms. Oligomers can disrupt cell membranes and create pores, which can cause mineral ion imbalance, and oligomeric forms of medin have been shown to be toxic to cells and able to upregulate the activity of MMP2^[168,169]. Apoptosis can induce vascular calcification by the formation of apoptotic bodies, which can act as nucleation sites for mineral formation^[51]. Previous research has shown that MMPs can also enhance vascular calcification through upregulation of BMP2 and ECM degradation^[85,86,193]. Therefore, extracellular deposition of medin aggregates, which are increased with aging, could exacerbate vascular calcification.

Amyloid fibrils display distinct stereochemical properties, for example, the presence of charged residues along the fibril surface, which could enhance its capacity for binding calcium and phosphate and nucleating mineral growth^[176]. This is demonstrated in cases such as mineralization of tooth enamel, where amyloid-like structures of amelogenin form a nidus for mineralization of calcium-phosphate mineral^[194]. HSPGs, including HSPG2, are common components of amyloid plaques and have been shown to directly enhance the aggregation of amyloid proteins, such as medin^[37]. HSPGs are also known to bind calcium, providing another scaffold for mineralization with amyloid deposits.

Conversely, while evidence suggests conditions that promote calcification, such as mineral stress, could also promote amyloid formation through ECM changes and EV secretion, no studies have described a direct effect of mineralization on amyloidosis. Both amyloid and HA formation are nucleation-dependent mechanisms, with evidence showing amyloid could nucleate mineral growth, yet any effect of HA or mineral deposits on enhancing amyloid growth has not been discussed.

FUTURE DIRECTIONS FOR VASCULAR AGING RESEARCH

There are many questions remaining regarding the accumulation of HA and amyloid with age and if there are connections between the processes. While there is strong evidence for an association between pathologies, direct causal links, particularly in the vasculature, are not understood. From studies into CAVD, IAC, and tooth enamel formation, it could be suggested that amyloid acts as a scaffold for mineral growth. In the aorta, medin amyloid and HA accumulate extracellularly with age, both regulated by VSMC-derived sEVs and exosomes; however, to date, there has been no research into a causal relationship. Therefore, investigating whether medin and other vascular amyloid proteins, such as $A\beta$, can nucleate minerals to drive calcification or vice versa would be valuable. Along with being a physical template for mineralization, there is evidence to indicate that amyloid could enhance vascular calcification through cellular pathways, such as oxidative stress, membrane pore formation, and ECM remodeling through



Figure 5. Young, healthy vascular smooth muscle cells (VSMCs) have a contractile phenotype and a healthy extracellular matrix (ECM), which undergoes turnover and remodeling to maintain its compliance and regulate vascular tone and integrity. VSMCs undergoing senescence alter the composition and stiffness of the ECM, increasing collagen and fibronectin deposition and decreasing elastin. Senescent VSMCs secrete more small extracellular vesicles (sEVs), which promote both calcification and amyloid aggregation and deposition. There is increased endosomal activity and formation of multivesicular endosomes (MVE) which contain sEVs that are degraded by lysosomes or released by binding to the membrane. sEVs from senescent VSMCs contain more medin in a fibrillar form and can accelerate its accumulation as fibrils in the ECM. Senescent sEVs also contain more calcium and annexins, which enhance mineralization. What triggers sEVs to either mineralize or aggregate amyloid is unknown. (Created with BioRender.com)

increased MMP secretion. However, there is currently a lack of consensus as to what form of amyloid contributes to vascular dysfunction, as studies have discussed pathological outcomes of both oligomeric and fibrillar amyloid species. Previous studies have discussed whether aggregation of toxic oligomers into "benign" fibrils is a protective mechanism, in which case therapies to prevent amyloid formation may be detrimental to vascular function. Further investigation is therefore needed to understand the contribution of different amyloid species during vascular aging and in vascular calcification.

Identification of therapeutic targets of amyloidosis and calcification could attenuate vascular stiffening with age, which is known to contribute to CVD morbidity and mortality. In recent years, there has been a huge push towards high throughput screening (HTS) as a platform to test thousands of drugs in an *in vitro* setting. To date, there has been limited use of HTS to identify novel drugs and targets for calcification. However, HTS has been used to identify compounds that affect amyloid formation, particularly $A\beta$, α -synuclein, and $TTR^{[195]}$. In the past, small compounds that could block $A\beta$ aggregation could be counteracted by the presence of a lipid membrane, but a recent study identified compounds that were effective with vesicles present to confirm inhibition of amyloid formation at the membrane interface^[196]. Additionally, as previously discussed, the effects of blocking amyloid fibril formation may also result in cellular dysfunction caused by small aggregates. These are two important aspects of HTS that should be considered in the future for identification of potential therapies for amyloidosis.

VSMC senescence has been implicated in calcification and vascular amyloidosis and drugs that induce apoptosis of senescent cells, senolytics, have been shown to be effective in AD mouse models, with A β or tau pathology^[85,197]. There is less known about the efficacy of senolytics in calcification, although one study showed that treatment with Dasatinib + Quercetin caused clearance of senescent cells in the medial layer and reduced aortic calcification in both aged and hypercholesterolaemic mice^[198]. This was associated with a reduction in osteogenic gene expression and significantly improved vasomotor function. Therefore, senolytics may be a potential therapeutic intervention for both vascular calcification and amyloidosis.

In conjunction with senescence, sEVs are mediators and potential initial triggers for both HA and amyloid accumulation in the vasculature. There has been some research into targeting EV secretion to alleviate both pathologies in both *in vitro* and *in vivo* studies. In a mouse model of CKD, inhibition of sEV secretion with a neutral sphingomyelinase inhibitor, GW4869, resulted in reduced aortic calcification and osteogenic differentiation of VSMCs^[123]. Blocking sEV secretion also attenuated the calcification of human and mouse VSMCs *in vitro*, under calcifying or diabetic conditions^[199]. In AD mouse models, GW4869 significantly reduced A β plaque load in the brain^[135,200]. These studies highlight not only the therapeutic potential of targeting sEV secretion but also the importance of sEVs in the initiation and development of calcification and amyloidosis.

CONCLUSION

This review has highlighted that amyloid and HA deposition in the vasculature share mechanistic processes and display synergies, which may account for their co-localization in the vessel wall [Figure 5]. Mineral stress and cellular senescence can increase EV production and promote osteogenic changes in VSMCs, which could also promote amyloid deposition, while amyloid in both oligomeric and fibrillar forms could nucleate minerals and enhance calcification. Importantly, there is evidence showing EV secretion, senescence, and ECM modifications can enhance both pathologies, suggesting therapies that target these common mechanisms may alleviate age-associated vascular stiffening.

DECLARATIONS

Authors' contributions

Conceptualization, Writing and editing : Whitehead M, Antonazzi M, Shanahan CM

Availability of data and materials Not applicable.

Financial support and sponsorship

This work was supported by a Grant from the British Heart Foundation to C.S. BHF: RG/F/21/110064, https://researchgrants.bhf.org.uk. The funding body had no role in the experiment design, collection, analysis, and interpretation of data, and writing.

Conflicts of interest

All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate Not applicable.

Consent for publication Not applicable.

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