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Vascular calcification: from pathophysiology to biomarkers

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ABSTRACT

The link between vascular calcification (VC) and increased mortality is now well established. Over time, as clinical importance of this phenomenon has begun to be fully considered, scientists have highlighted more and more physiopathological mechanisms and signaling pathways that underlie VC. Several conditions such as diabetes, dyslipidemia and renal diseases are undoubtedly identified as predisposing factors. But even if the process is better understood, many questions still remain unanswered. This review briefly develops the various theories that attempt to explain mineralization genesis. Nonetheless, the main purpose of the article is to provide a profile of the various existing biomarkers of VC. Indeed, in the past years, a lot of inhibitors and promoters, which form a dense and interconnected network, were identified. Given importance to assess and control mineralization process, a focusing on accumulated knowledge of each marker seemed to be necessary. Therefore, we tried to define their respective role in the physiopathology and how they can contribute to calcification risk assessment. Among these, Klotho/ fibroblast growth factor-23, fetuin-A, Matrix Gla protein, Bone morphogenetic protein-2, osteoprotegerin, osteopontin, osteonectin, osteocalcin, pyrophosphate and sclerostin are specifically discussed.

Keywords: Vascular calcification Fibroblast growth factor-23 Fetuin-A Matrix Gla protein Bone morphogenetic protein-2 Osteoprotegerin

Abbreviations: 1,25 (OH)₂D, 1,25-dihydroxyvitamin D; ALP, alkaline phosphatase; AVK, vitamin K antagonists; BAX, BCL2-associated X protein; BMDM, bone marrow-derived macrophage; BMP-2, bone morphogenetic protein-2; CAC, coronary artery calcifications; CAD, coronary artery disease; CaSR, calcium-sensing receptor; Cfba1/Runx2, core-binding factor subunit 1 α / runt-related transcription factor 2; CKD, chronic kidney disease; cMGP, carboxylated MGP; CPP, calciprotein particles; dp-uc MGP, dephosphorylated uncarboxylated MGP; EPCs, endothelial progenitor cells; ePPi, extracellular PPi; Fet-A, fetuin-A; FGF-23, fibroblast growth factor-23; HD, hemodialysis; HIF-1α, hypoxia-inductible factor 1α; MGP, matrix gla protein; MVs, matrix vesicles; MVSC, multipotent vascular stem cells; ND, non dialysis; OC, osteocalcin; ON, osteonectin; OPG, osteoprotegerin; OPN, osteopontin.

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Refe	rences					
1. Inti			polymorphisms located at 9p21 locus near the cyclin genes has been suggested in the genesis of pathology [9]. These genes encode			
00	CHERA	nce of vascular calcification (VC) is not new. Arterial	cyclins that may be broadly linked to cellular senescence and inflam-			
		n has been discovered in the "Iceman" who lived	mation but the accurate causative DNA sequences remains debated			
		ago [1] and scientists had already paid attention to this				
			[3]. Cones ENDD1 and NITEE are respectively implicated in infancy and			
		on – and to its relation with renal disease – in the 19th	Genes ENPP1 and NT5E are respectively implicated in infancy and			
		VC increases with age and is notably dysregulated in	idiopathic VC. The first one encodes a protein which transforms ATP to			
		rslipidemia, renal disease and hypertension [3]. Bone for-	adenosine and pyrophosphate (PPi, inhibitor of calcification) whereas the second one converts AMP into adenosine and inorganic phosphate (Pi, accelerator of mineralization). The VC phenotype caused by muta- tions in these genes underlines the role of PPi and Pi in pathogenesis			
		olves hydroxyapatite [Ca10(PO4)6(OH)2] crystals, whose				
	-	nt begins in matrix vesicles (MVs) that bud from osteo-				
		ular smooth muscle cells (VSMCs) that have undergone				
osteo	blast c	differentiation are also able to release similar vesicles	[3]. Mutations in ABCC6, a gene encoding a nucleoside-sensitive trans-			
with s	shared	d protein content [4]. Such differentiation is restrained	porter, have also been linked to hereditary calcification. Alternative action of ABCC6 may include deficient hepatic production of inhibitory			
or inh	ibited	under normal conditions and there is a balance with os-				
		erentiation experienced by monocytes and macrophages	factor of matrix Gla protein (MGP), an important inhibitor of calcifica-			
		vascular wall. Moreover, the reaction which allows crystal	tion [3,11].			
		nermodynamically unfavorable and is inhibited by pyro-				
_		[3]. In some situations, physiological balance is broken	2.2. Osteoblastic differentiation			
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2. Overview of vascular calcification : from genetics to osteoblastic differentiation

the fact we cannot completely deny the link with ageing, VC is no longer

regarded as a passive process. Today, it is considered as an actively regulated and complex process that remains not completely understood.

We know that many dynamic changes can trigger or promote VC and

a lot of theories emerged during the last two decades. We will describe

hereafter some of the physiopathological mechanisms. It is important to

underline that these different mechanisms are not mutually exclusive

2.1. Genetic predisposition

As shown in Fig. 1, genetic predisposition certainly plays an important role in the genesis of this phenomenon. According to Rutsch et al., [6], 40–50% of cases of coronary calcification can be attributed to genetics. Genome-wide association studies have identified several loci linked to coronary arterial calcification, like 6p21.3, 6p24, 10q21.3 and 9p21 [3,7–9]. Interestingly, some of these regions also appear to be related to coronary atherosclerosis [7] and the 6p24 locus is also linked to myocardial infarction [10]. An implication of several single

is actively mediated by Pit-1, a sodium dependent phosphate cotransporter [14,15]. This increasing phosphate concentration in the VSMC induces a phenotypic switch of VSMCs into osteoblast-like cells [5,14,16]. The protein Cfba1/Runx2 (core-binding factor subunit 1α /runt-related transcription factor 2) is a specific and indispensable transcriptional regulator for this osteoblastic differentiation. Its expression is also enhanced with high extracellular phosphate [14,16,17]. These "new" cells will express alkaline phosphatase (ALP), secrete, under the control of Cfba-1, bone-associated proteins (such as osteopontin [18], collagen type 1, osteoprotegerin, bone morphogenic protein-2

and osteocalcin [14,19]) and release mineralization-competent MVs in

the extracellular matrix [13,14,20]. VSMCs release MVs under normal

physiological conditions and these MVs are protected from mineraliza-

tion by the presence of calcification inhibitors. Under pathological

conditions, a combination of factors makes the MVs "mineralization

competent" [21]. Moreover, an increase of intracellular phosphate level

mediated by Na/Pi transporter is thought to induce VSMC apoptosis

bone formation. First, vascular smooth muscle cells (VSMCs) undergo osteogenic differentiation into phenotypically distinct osteoblast-like

cells [5,12]. In the case of renal failure, phosphate plays a key role in

this mechanism [13,14]. In vitro, high extracellular phosphate concen-

trations induce a rise in intracellular phosphate concentration which

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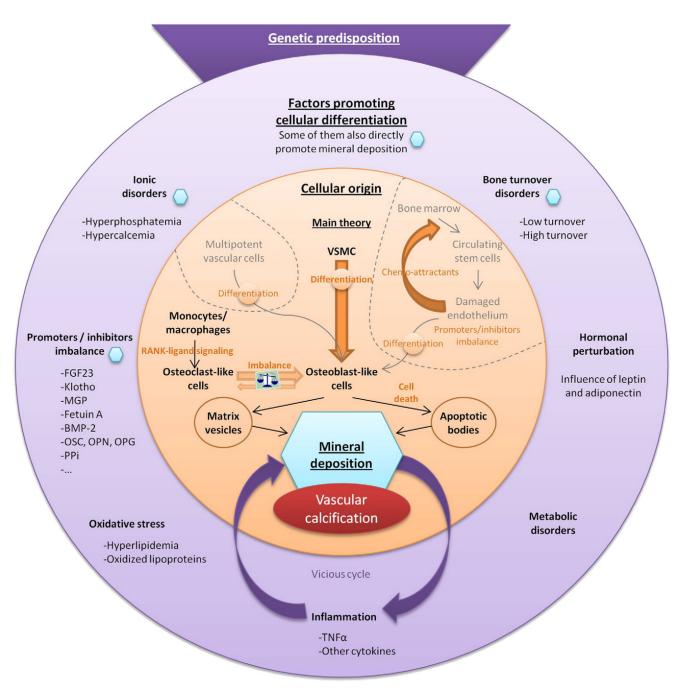


Fig. 1. Physiopathological mechanisms which promote cellular differentiation and mineral deposition.

through an unclear process that possibly involves a disruption in mitochondrial metabolism [22]. Some studies suggest that apoptosis leads to calcification [23–26]. The MVs, in which proapoptotic factor BAX (BCL2-associated X protein) have been identified [26], may be remnants of apoptotic cells. As MVs have the capacity to concentrate and crystallize calcium, apoptosis could be a key regulator of VC [22].

More recently, a different point of view has emerged according to which phenotypically distinct osteoblast-like cells might originate from stem cells rather than VSMCs. A new mechanism called "Circulating cell theory", suggesting an active role for circulating cells arising from sources such as bone marrow, has been postulated to contribute towards VC. Under the influence of chemo-attractants (released by damaged endothelium for instance), these bone marker-positive cells may home to diseased arteries. Under pathologic conditions such as an imbalance between promoters and inhibitors of VC, this population may further undergo osteogenic differentiation in the lesions, which could promote

vessel mineralization [27]. Another recent study has also claimed that multipotent vascular stem cells (MVSC) present in blood vessel wall might differentiate into osteoblast-like cells [28]. Nevertheless, this point of view is still very controversial.

Although the role of phosphate is well established in osteoblastic differentiation process, many other factors can influence this phenotype conversion and accurate causal mechanisms remained not completely understood. Under normal conditions, VSMCs produce endogenous inhibitors of calcification such as matrix gla protein (MGP), osteopontin, osteoprotegerin and pyrophosphate [29]. A long-term exposure of VSMCs to a variety of stresses can overwhelm the action of these inhibitors and induce differentiation [21]. Among these chronic stresses, ionic disorders (especially hyperphosphatemia and hypercalcemia) are incriminated but inflammation, hormonal perturbation, metabolic disorders and oxidative stress can also lead to VC. Oxidative stress in VSMCs, in particular generated by hyperlipidemia and oxidized

lipoproteins or uremic milieu[30], causes the expression of runx2 [31], osterix and govers Wnt signaling [32], leading to osteogenic differentiation. Inflammatory cytokines, such as TNF- α , can also induce calcification via Msx2/Wnt/\(\beta\)-catenin pathway [33]. In support of that, calcium deposits colocalize with inflammatory cells in vitro [34, 35] and in vivo [36]. Moreover, it has been suggested that mineral crystals may themselves be pro-inflammatory, creating a vicious cycle of inflammation and calcification [37,38]. The receptor for advanced glycation endproducts (RAGE) endogenously expressed in endothelial cells and its ligands (in which S100 family proteins are found), are also known to be involved in atherosclerotic formation and VC. It has

been suggested that galectin-3 and RAGE modulate vascular osteogenesis in part via Wnt/\(\beta\)-catenin signaling [39]. Several trials have shown a raise in serum levels of \$100/calgranulins in vascular disease [40,41]. Thereby, S100 proteins could be a potential biomarker and therapeutic target to develop [3]. Involved in the control of both parathyroid hormone (PTH) and calcitonin secretion, the calcium-sensing receptor (CaSR) is a G protein-coupled cell surface receptor that is able to sense extracellular calcium ions. Evidences have been provided to demonstrate that a decrease in the CaSR protein expression in the vasculature is directly involved in the development of VC [42,43]. It is of particular interest to note that calcimimetics, which are allosteric drug compounds that selectively target the CaSR, decrease VC at least in part through local control of the CaSR expression in VSMC [44,45]. However, so far, the mechanism whereby the CaSR exert its protective effect remains largely unknown. Hormones have pleiotropic effects on calcific vasculopathy. For example, the adipose-derived factor, leptin, promotes VC in vitro [46] and in vivo [47]. Adiponectin-deficient mice have increased vascular calcification [48]. The influence of PTH is part of bone turnover process and is discussed further. A disruption between promoters and inhibitors can also generate VC. Moreover, similar to bone formation, there might exist a balance between VC and its resorption. Indeed, monocytes and macrophages contained in the calcified wall can differentiate into an osteoclast-like phenotype and counteracts the action of VSMCs that have undergone osteoblast differentiation [49]. Hyperphosphatemia would disadvantage osteoclast phenotype by down-regulating RANK ligand-induced signaling [50] but this is not clear whether osteoclast-like cells can really counteract VC or solely

All these modifications will favour for an optimal microenvironment for hydroxyapatite formation and calcification. Similar osteogenic differentiation is also observed, in vivo, in animal and human uremic models [12,16,51].

As calcium and phosphorus have a pivotal role in the pathogenesis of

2.3. The role of bone turnover

witness vascular remodeling process.

VC, it is not surprising that bone-turnover disturbances can enhance VC because bone is the most important reserve of calcium and phosphorus and could act as a buffer [52]. Some authors have written about the "bone-vascular" axis. The link between "low bone turnover" and risk of VCs is likely better illustrated by the current literature. However, the two "theories" of low and high turnover are not excluding each other in a context of "bone as a calcium-phosphorus buffer" [53].

2.3.1. Low turnover

In the first article measuring VCs, Braun et al. have already shown that coronary calcifications are inversely correlated with bone mass [54]. Probably the most interesting study on this topic has been published in 2004 by London et al. These authors have actually compared VCs by plain radiography with bone histomorphometry in 58 dialysis patients. They found that a high calcification score is independently associated with histomorphometry suggestive of low bone turnover (or a dynamic bone)[55]. One complementary study from the same

group especially underlined that calcium load is particularly deleterious,

in terms of VCs, for these patients with low bone turnover [56]. Studying

both coronary calcifications and bone histomorphometry in 101 dialysis patients. Barreto et al. also found an inverse correlation between calcium score and bone trabecular volume and trabecular thickness in an univariate analysis [57]. In 2009, Adragao et al. studied the relationship between bone histomorphometry and coronary calcifications in 38 dialysis patients. Contrary to precedent trials, low bone turnover was not associated with calcifications. However, lower bone volume was a risk factor for coronary calcification [58].

2.3.2. High turnover

The epidemiological link observed by some authors between PTH and CV mortality in dialysis patients has already been mentioned [59–62]. Coen et al. have demonstrated a relationship between high PTH levels (especially very high levels) and coronary calcifications [63,64]. Even if specific PTH fragments (1-34 PTH) have been shown to inhibit VCs in an animal model [65], Neves et al. have shown, in a model of parathyroidectomized rats with chronic kidney disease (CKD), that perfusion of PTH, per se, could induce VCs [66].

In this review, we aim to provide a profile of the various existing biomarkers of VC. We will try to determine the respective role they play in the physiopathology and how they can contribute to calcification risk assessment.

that requires the presence of the cofactor Klohto for its classical effects.

FGF-23 promotes phosphate excretion by reducing its proximal

re-absorption by reducing the expression of NPT2a and NPT2c mRNA,

sodium/phosphate transporters [67]. Although the administration of

1,25-dihydroxyvitamin D (1,25-(OH)2D) increases its circulating rate,

FGF-23 levels also reduce 1α -hydroxylase activity, which decreases

conversion of calcidiol into its active form, 1.25-(OH)2D and up-

3. Biomarkers

3.1. Fibroblast growth factor-23 (Fgf-23) and Klotho

Fig. 2. FGF-23 is an approximately 30 kDA protein released by bone

regulates 24-hydroxylase, which in turn converts 1,25-(OH)2D into more hydrophilic metabolites with lesser biological activity [68]. Thereby, gastrointestinal absorption of calcium and phosphate is reduced. In parathyroid glands, FGF23 decreases PTH secretion and parathyroid cell proliferation [69]. FGF-23 null mice develop hypercalcitriolemia and VC. The correction of hypercalcitriolemia improves survival but has no effect on VC. On the other side, poor phosphate diet leads to a complete normalization of the phenotype, suggesting that phosphate plays a major role in calcification abnormalities [67,70]. From this point of view, FGF-23 is more a hall marker of VC rather than a real actor.

However, it has been suggested that FGF-23 could actively participate in VC, perhaps by Klotho-independent functions [22,71]. Indeed, FGF-23 also induces left ventricular hypertrophy via FGFR-dependant, but Klotho independent, effects on cardiac myocytes [72]. This finding raises the possibility that elevated FGF-23 levels in CKD may also contribute to other forms of subclinical cardiovascular injury, such as VC [73]. A correlation between higher serum FGF-23 levels, mortality and VC, regardless of the serum phosphate levels, has been confirmed by several studies [74-82]. However, a recent study from Scialla et al. [73] has challenged this point of view. In this translational study, the association between FGF23, serum phosphate and calcification of the coronary arteries and thoracic aorta has been studied in a sample of 1501 individuals. In contrast to serum phosphate, FGF23 was not associated with arterial calcification. In hemodialysis (HD) patients, Lee et al. [83]

The elevation of serum FGF-23 is a common feature in CKD. It appears that the massive elevations in FGF-23 levels in CKD reflect both increased secretion and decreased removal from circulation. FGF-23 may be a compensatory phenomenon to the phosphate retention

have not found a correlation between calcification and FGF23 levels either. Further studies are required in order to examine whether

FGF23 may have a direct effect on progression of calcification.

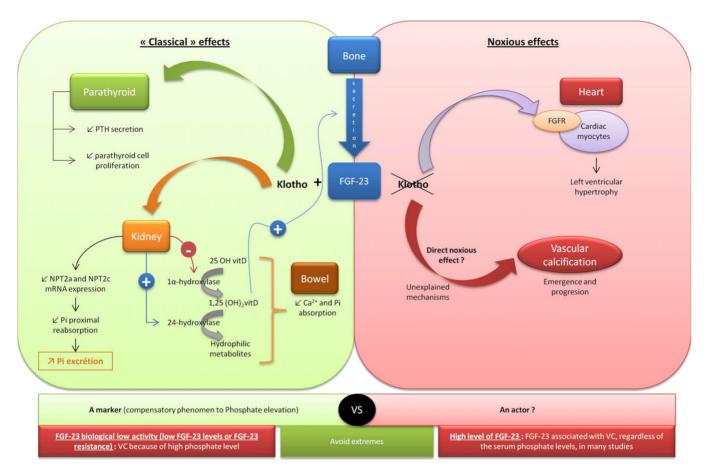


Fig. 2. Suspected duality of FGF23 effects.

seen in CKD [84]. In agreement with this hypothesis, FGF-23 was associated only with high coronary artery calcifications (CAC) (400) but not with moderate CAC (100-400) in a cohort of CKD patients[85]. Active vitamin D therapy may also contribute to the increased levels of FGF-23. Although the mechanistic link remains to be explained, FGF-23 may serve as a novel risk marker for the cardiovascular mortality in CKD [68]. In patients with coronary artery disease (CAD), the same independent link between FGF-23 and mortality has been demonstrated [84].

In contrast to FGF23, Klotho excess has never been shown to be noxious [86]. Klotho is a 1014 amino acids protein which is expressed on cell surface. It comprises a long extracellular queue, a transmembrane domain and a short intracellular portion [87]. Given the moderate FGF-23 affinity for its receptor, Klotho is needed to strengthen the link between the two entities. It thus increases FGF-23 ability to induce phosphorylation and signal transduction. This discovery emerged from the observation that either FGF-23 or Klotho knock-out mice resulted to the same phenotype [88]. As filtration rate decreases in CKD patients, Klotho expression is reduced and FGF-23 levels enhanced. This increase might represent an adaptation to Klotho diminution, also known as "FGF-23 resistance", or hyperphosphatemia [87]. Interestingly, Klotho levels are up-regulated by vitamin D receptor agonists (calcitriol or paricalcitol) in CKD mice submitted to a high phosphate diet. These mice show half less calcification than those who did not receive therapy. Phosphaturia is increased whereas phosphatemia and FGF-23 levels are lowered [89]. Klotho is expressed in a limited number of organs, including kidney and parathyroid gland. In addition to its membrane form, Klotho is also expressed in two soluble forms: one coming from cleavage of the membrane form and the other coming from an alternative splicing [87]. Membrane-anchored and soluble Klotho proteins seem to have distinct functions. Membrane Klotho mediates FGF-23 effects mentioned above whereas soluble Klotho is a pleiotropic protein functioning as an endocrine factor with a multitude of renal and extrarenal actions. Among these effects, anti-apoptotic, anti-oxidative, anti-Wnt signaling actions or modulation of renal ion channels are assumed [90]. In contrast, vascular Klotho deficiency favors the development of arterial calcification and mediates resistance to beneficial vascular effects of FGF23 [86].

3.2. Fetuin-A (Fet-A)

Fig. 3. Fet-A, a 59-kDa glycoprotein, is a potent calcification inhibitor produced by the liver that possess a systemic action. Serum Fet-A concentrations are decreased in dialysis patients compared with healthy controls [91,92]. The accurate mechanism is still hypothetical, but is probably related to chronic inflammation, as Fet-A is a negative acute phase reactant [93].

Its calcification inhibitory action is powerful and illustrated by knock-out mice developing severe extraosseous calcifications [94]. In this murine model, the aorta remains free from calcifications. However, CKD and high phosphate diet induce severe aorta calcifications in these knock-out mice [95]. Fet-A is thought to inhibit calcification by binding early calcium phosphate crystals and by inhibiting crystal growth and mineral deposition. This could be facilitated by the formation of large calciprotein particles [96,97]. Indeed, the accumulation of naked calcium phosphate crystals is responsible for extraosseous calcification and causes inflammation. These crystals are usually digested by the cells of the reticuloendothelial system such as macrophages. In contact with the crystals, macrophages secrete proinflammatory cytokines and undergo more apoptosis. The formation of fetuin-A calciprotein particles (CPP) facilitates the clearance of these crystals and thereby reduces their negative impact. Fet-A likely plays a very important role in the stabilization of these complexes and reduces the inflammatory response.

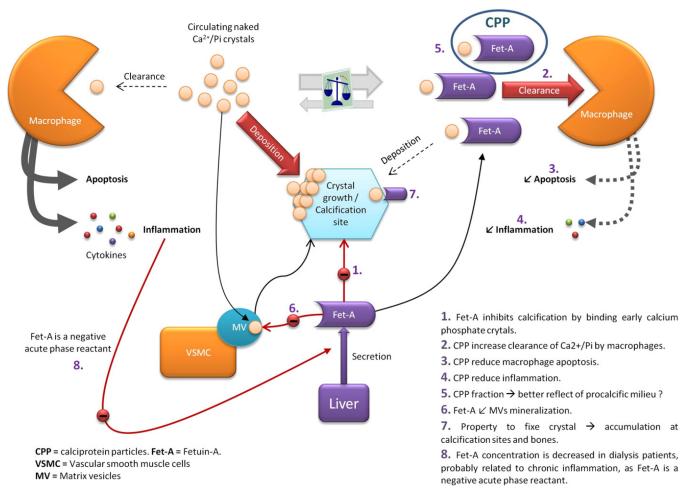


Fig. 3. Properties of Fetuin-A.

Fet-A binds and sequesters insoluble mineral nuclei, forming soluble colloidal CPP, thereby inhibiting crystal growth and aggregation. Macrophages secrete less cytokines and undergo less apoptosis phenomenon compared to reactions caused by naked crystals. This property of Fet-A to decrease inflammation may be influenced by the phosphorylation degree of the glycoprotein [98]. The above mechanism partially explains the inhibitory effect of Fet-A in VC. Fet-A could also prohibit MV calcification into the VSMCs which take up circulating Fet-A from an extracellular calcium- dependent way [99,100]. Because its ability to bind calcium apatite crystals, Fet-A accumulates at sites of VC as well as in bone [101]. Increasing Fet-A expression is found in calcified arteries from dialysis patients [102].

Several studies support the evidence that lower serum levels of Fet-A levels might result in VC. In these studies, lower serum Fet-A concentrations have been associated with increases in calcification scores, arterial stiffness, mortality and incidence of cardiovascular events [83,103–106]. Among recent studies, Lee et al [83] found lower serum Fet-A in HD patients with valvular calcification. Abdel-Wahab et al. [103] showed significant negative correlation between Fet-A and indices of vascular disease in systemic lupus erythematosus patients. In a 277 renal transplant recipients followed-up during 5 years, Marechal et al. [106] showed that low Fet-A levels are independently associated with aortic – but not coronary – calcifications and a higher risk of cardiovascular events and mortality.

However, contradictory results have also been published regarding the role of Fet-A. After adjustment for age, in a 58 patients study of Jung et al. [107], an inverse correlation between Fet-A and coronary artery calcification score became insignificant. Low Fet-A was nonetheless associated with a significant increase in all-cause mortality and the occurrence of a cardiovascular event. Manghat et al. [108] did not find significant correlation between Fet-A and arterial stiffness in 45 predialysis patients. Hermans et al. [109] reported that, after correction for covariates, Fet-A could not be identified as an independent predictor of aortic stiffness in a dialysis population.

Interestingly, Hamono et al. [110] have suggested that estimates of the specific CPP fraction of total circulating fetuin may provide a better measure of extraskeletal mineral stress in patients with CKD. A recent study of Smith et al. [38] showed that increased CPP Fet-A levels reflect an increasingly procalcific milieu and are associated with increased aortic stiffness in patients with pre-dialysis CKD.

Despite the discrepancy between these various results, it should be noted that, in the majority of reports, low Fet-A serum levels are actually related with VC and increased mortality in patients with end stage renal disease. Nonetheless, the exact role of Fet-A in VC and cardiovascular diseases needs to be further evaluated through additional experimental and clinical studies [111].

3.3. Matrix Gla protein (MGP)

MGP is a vitamin K, 14-kDa γ-carboxylated protein expressed by chondrocytes, VSMCs, endothelial cells and fibroblasts. Its role as a calcification inhibitor has been illustrated by MGP knock-out mice who develop extensive arterial calcifications [112,113]. In 2002, Moe et al. demonstrated a correlation between vascular MGP expression and the calcification of epigastric arteries in dialysis patients [51,102]. MGP-deficiency in humans leads to Keutel syndrome, a rare genetic disease hallmarked by abnormal soft tissue calcification [113].

MGP binds calcium crystals, inhibits crystal growth and plays a role in the normal phenotype of VSMCs in preventing the osteoblastic differentiation [17,114]. MGP also binds and inactivates a pro-mineralization factor, bone morphogenetic protein-2 (BMP-2) [115]. Among other effects, BMP-2 promotes osteogenic conversion of VSMCs via MSX2 transcription factor. Nevertheless, BMP-2 inhibition may not play the leading role in MGP preventive action on calcification. Lomashvili et al. [116] used warfarin as MGP inhibitor in cultured aortas but did not observe a decrease in calcification by adding BMP-2 antagonist. In this study, the lack of involvement of BMP-2 supports the possibility that MGP prevents medial calcification by directly binding hydroxyapatite and inhibiting its formation, which is consistent with the fact that the content of MGP in calcified aortas is proportional to hydroxyapatite content. MGP could also play a role in Fet-A calciprotein complexes.

tent. MGP could also play a role in Fet-A calciprotein complexes. Recently, Khavandgar et al. [113] hypothesized that deleterious effect of MGP deficiency could be linked to an extracellular matrix alteration. Indeed, in their study on MGP knock-out mice, they found that VSMCs differentiation did not precede the calcification phenomenon while MGP —/— aortas showed altered extracellular matrix remodeling. Furthermore, they showed that elastin haploin sufficiency reduced arterial calcification in MGP -/- elastin +/- mice. Elastin is a major elastic lamina protein and has been shown to act as a mineral nucleator. MGP could protect mineral nucleation sites on elastin and thereby prevent spontaneous calcification of the elastic laminae. In support of that, the irregular calcification of the thoracic and abdominal aorta segments in MGP - / - mice correlates with the local variations of the elastin content [113]. Parallel to this study, other authors hypothesized a mineralization process by size exclusion, in which MGP proves to be essential to prevent mineralization within fibrils. Indeed, most inhibitors are bigger

than MGP and cannot access nucleation sites located on fibrils, generat-

carboxylation and serine phosphorylation. It thus exists in four different

forms. Whereas the accurate role of phosphorylation is not fully

MGP may undergo two post-translational modifications: glutamate

ing mineralization in the absence of MGP [117,118].

elucidated, it is well established that carboxylation is needed for MGP activity. Balance between carboxylated (c) MGP and uncarboxylated (uc) MGP is thereby very important [117]. Using conformation specific antibodies, it has been reported that uc MGP is significantly present in the calcified vasculature whereas c MGP is nearly absent. It has been shown that non fully γ -carboxylated (but not γ -carboxylated MGP) is associated with VC [5,119,120]. However, total uc MGP seems to be inversely correlated with VC [121]. As total uc MGP includes phosphorylated and non-phosphorylated fractions and given the fact that phosphorylation may be involved in MGP activity, it is important to selectively dose the dephosphorylated-uncarboxylated (dp-uc) MGP inactive fraction. In a multivariate linear regression analysis, Schurgers et al. [122] have evidenced an independent association between plasma dp-uc MGP levels and aortic calcification whereas Schlieper et al. [123] did not find any correlation with the extent of VC. The discrepancy between these results could perhaps be explained by the use of different assays, as there is no validated analytic process yet.

Activation of MGP by γ -carboxylation is vitamin K dependant

[5,20,112,120]. In the case of vitamin K deficiency, VC phenomenon

can therefore be magnified. Treatment with Vitamin K antagonists

(AVK) also lead to the same consequence [117]. This is the actual expla-

nation for the observations that warfarin leads to extensive VCs in

animal and human studies [124-126]. It is interesting to note that vita-

min K therapy partially reverses warfarin-induced VCs in rats [126]. As AVK are the most prescribed anticoagulant medication to date, it is nec-

essary to establish whether the benefit outweighs the long-term risk to

develop increased VC, especially in diabetic or CKD patients [117].

3.4. Bone Morphogenetic Protein – 2 (BMP-2)

Bone morphogenetic proteins (BMPs) belong to a subdivision of TGF- β like growth factors family. BMPs regulate growth, differentiation

and development in the embryo as well as during tissue remodeling processes in the adult organism. BMP-2 is an important molecule in the regulation of bone formation as well as in VC. In bone, it promotes osteoblast differentiation and mineralization [127]. Inhibition of BMP-2 inhibits osteoblast differentiation and bone formation in vivo and in vitro [128] and protects against atherosclerosis and VC [129]. BMP-

nohistochemistry. The change in gene expression of these cells might be considerably regulated by BMP-2. It appears that a novel mechanism of BMP-2 activation would have been discovered recently. A protease from the coagulation called factor VII-activating protease (FSAP) would mediate BMP-2 cleavage and generation of truncated BMP-2, which is characterized by enhanced activity [130].

Ligand-bound receptor signaling results in smad 1/5/8 phosphoryla-

tion. The phosphorylated smads are then able to enter nucleus and lead

2 was detected in calcified human VSMCs by mRNA levels and by immu-

to transcriptional modification [130]. BMP-2 would also be involved in microRNA-30b and microRNA-30c decreasing, likely through a smadindependent pathway. MicroRNAs are small noncoding RNAs which link specific complementary mRNA sequences. This binding drives to translation inhibition and promotes mRNA degradation, so that this one cannot produce protein anymore. MicroRNA-30b and microRNA-30c are plausible candidates to regulate runx2 expression. Runx2 is an important factor in VSMCs transition into osteoblast-like cells. Thus, BMP-2 would perhaps promote VSMCs calcification by increasing runx2 expression through microRNA-30b and microRNA-30c down-regulation [131].

mal cells. OPG plays a pivotal role in the regulation of the bone turnover.

inhibiting osteoclast differentiation and acting like a decoy receptor for

the receptor activator of NF-KB ligand (RANKL system) [132]. It inter-

feres with the interaction between RANK (expressed by osteoclast-like

cells) and RANKL (expressed by osteoblast-like cells). OPG is also

3.5. Osteoprotegerin (O) and RANKL/Rank axis OPG is a regulatory factor produced by bone marrow derived stro-

thought to inhibit ALP activity [133]. OPG levels are significantly higher in CKD patients, in relation with the severity of renal failure. Although OPG is known to impede osteoclast differentiation in bone, OPG is usually considered as a protective factor against VC as it blocks the bone remodeling process in the vascular tissue whereas RANKL is thought to have a negative action. OPG is also a neutralizer of the proapoptotic actions of TRAIL (TNF-related apoptosis-inducing ligand), which strongly activates vascular cells apoptosis [134]. Apoptotic bodies can also lead to mineralization. In support of that, it has been observed that OPG deficient mice do develop both severe aortic calcifications and osteoporosis [135] and that calcified arteries are known to have higher levels of RANKL expression and lower levels of OPG [136]. Zhou et al. [137] isolated VSMCs from thoracic aorta of rats and cultured them in osteogenic medium. They found osteoblastic differentiation was markedly reduced by addition of OPG in a dose responsive way. They noticed that this addition also lead to decrease in Notch1 and RBP-Jk mRNA,

However, other findings suggest a more complex picture. Morena et al.[138] found a strong relationship between OPG high plasma levels and CAC in CKD patients. In this study, the cut-off value best predicting CAC score was 757.7 pg/ml (sensitivity = 91.7%; specificity = 59.0%). To improve diagnostic performance, two threshold values have been proposed: the lower one below which CAC is excluded (<709.9 pg/ml) and the highest one above which CAC is accepted (>963.7 pg/ml) with a grey area within which the diagnostic remains uncertain. This compromise resulted in a reduced 12.8% error rate. It is important to consider that these values were established in CKD pa-

tients before dialysis initiation. Indeed, OPG is not removed correctly

by HD and therefore accumulates. Interestingly, OPG seems to be a mark-

er of VC onset rather than a severity or progression predictor. This was an

which are upstream contributors of the Msx2 pathway, known as an

osteoblastic conversion path.

observation of a study which investigated the relation between OPG. FGF-23 and CAC in a ND-CKD population [85]. Despite the fact that OPG levels were significantly increased in patients with moderate vs. mild CAC, an additional increase of OPG levels in patients with severe cal-

cification was not observed. In contrast, FGF-23 better reflected severe calcifications vs. milder ones. Many other trials have also shown that VC as well as arterial stiffness and cardiovascular events are inversely related to serum RANKL [139–141] and positively related to serum OPG

[83.136.138.139.141–149]. Although some convincing assumptions are

proposed to explain why OPG levels correlate with calcification, it has still to be decided whether OPG is rather a marker, a protector or an ac-

tive player in calcification process. Position of RANKL in the physiopathology is not perfectly clear

either. Some in vitro studies have showed that RANKL treatment induces VC, whereas others found no effect, albeit in different species and culture conditions [136]. Many studies used transfected RANKL mice models to assess impact of RANKL overexpression on vascular system. The overexpression was global or the distribution was systemic (released from the liver for example). These methods do not permit to distinguish RANKL direct effects on the vasculature from indirect effects induced by bone remodeling. So Morony et al. [136] have recently conceived a transfected mice model in which RANKL was solely expressed in VSMCs. To do so, they linked RANKL gene to a promoter whose expression was highly restricted to VSMCs in adult tissues. No calcium deposition in the aortic roots has been revealed. Although SM22 α -Rankl^{tg}

mice do not develop spontaneous VC, it is possible that these mice develop greater VC than normal mice when exposed to procalcific factors.

The team has not been able to demonstrate an increase in calcification with cultured SM22 α -Rankl^{tg} cells either. Furthermore, Olesen et al.

[150] studied the effect of RANKL recombinant treatment on human

OPG knockdown VSMCs and they did not find any modification in

calcification-associated genes expression pattern. Nonetheless, a different approach has been tempted by Deuell et al. [151]. As macrophages are able to induce apoptosis and differentiation of VSMCs through an inflammatory process, the hypothesis that RANKL regulates VSMCs calcification by modulating macrophage production of pro-calcific cytokines was tested. In order to explore this, they used a bone marrow-derived macrophage (BMDM)/VSMCs co-culture system. Treatment of BMDMs with RANKL resulted in expression of IL-6 and TNF- α . According to this team, increased expression of these pro-calcific cytokines in macrophages might then mediate RANKL-induced VSMCs calcification in a

greater calcification, but this may be a reflection of a noxious OPG activity on the vasculature or it also could be the witness of a compensatory mechanism setting. Indeed, OPG levels could rise in response to a vascular injury and would then attempt to control, in part, vascular remodeling. This hypothesis is supported by OPG-/- animal experiences in which they developed severe calcifications, unless calcification has been an indirect repercussion of an excessive bone resorption. Some publications have shown a decrease or even an inhibition in VC, using recombinant OPG treatment or denosumab (a RANKL inhibitor) in

mice [152,153], but not in clinical study with post menopausal women [154]. As OPG can be secreted by VSMCs which underwent osteoblastic

differentiation, OPG could merely be a bystander too.

Those are the reasons why it is difficult to establish the accurate role

of OPG-RANK/RANKL axis. High serum levels of OPG are associated with

3.6. Osteopontin (OPN)

paracrine way.

OPN is a phosphoprotein expressed in the mineral tissues which inhibits mineralization by blocking hydroxyapatite formation and activating osteoclast function [155]. OPN is present in calcified vessels. OPN knock-out mice do not develop VC but, when these mice are bred with MGP knock-out mice, the VCs are more important than in simple MGP knock-out mice [156]. OPN must be phosphorylated to act as a

calcification inhibitor [157]. OPN inhibits mineralization of VSMC by

binding to the mineralized crystal surface [158]. On the contrary to the fully phosphorylated OPN, cleaved OPN could act as a proinflammatory cytokine and a proangiogenic factor facilitating vascular mineralization [155,159]. The possibility that OPN could serve as a calcification serum marker

is controversial. Berezin et al. [160] showed that OPN was a good predictor of coronary calcification in type two diabetes mellitus patients. Tousilis et al. [149] found a positive association between OPN and arterial stiffness in coronary artery disease. However, all trials do not come to the same conclusions: although OPN serum levels were considerably increased in patients with abdominal aortic aneurysms, no independent correlation between arterial stiffness (assessed by pulse wave velocity) and OPN could be highlighted after multiple regression analysis [161]. Albu et al. [142] did not find any correlation between OPN levels and arterial stiffness in postmenopausal women. However, it is interesting to notice that OPN correlated positively with hsCRP and negatively with HDL-C. Indeed, the discrepancy between the different studies may perhaps be explained by the differences in patient populations. It is thought OPN plays a key role in inflammatory process. Its relation with diseases related to inflammation such as atherosclerosis, obesity and auto-immune diseases has already been shown [142,155,162–164].

It has also been suggested that hyperglycemia could up-regulate OPN

and thereby lead to VSMCs proliferation [165].

3.7. Osteonectin (On)

in bone development that demonstrates affinity for hydroxyapatite and collagen [166]. ON has been found in association with large calcifications in atherosclerotic plaques [167]. The lack of knowledge on osteonectin doesn't allow to define its specific role. Further investigations are needed to precise its utility in calcification assessment.

ON, also called SPARC or BM40, is a calcium binding protein involved

3.8. Osteocalcin (Oc) OC, a vitamin-K dependent matrix protein that inhibits calcium salt precipitation in vitro [168], shows a strong affinity for hydroxyapatite. OC has been found in calcified atherosclerotic plagues and calcified aortic valves [169]. It was generally thought that OC inhibits crystal growth

[170] and limits bone formation [171], but different studies challenge

this point of view and suggest the opposite idea [172,173]. Hence, a metabolic role for OC has newly been suggested. As OC has been

shown to stimulate insulin secretion by pancreas and glucose uptake

in adipose tissue, Ideleviech et al. [174] wanted to consider direct met-

abolic effects of OC on VSMCs and chondrocyte cells with an in vitro OC

overexpression model. They found that OC functions as a stimulator of

differentiation and mineralization, up-regulating Sox9, Runx2, collagen

type X, ALP, proteoglycans, and mineral content in both of these cells. In

particular, OC increases hypoxia-inductible factor 1 α (HIF-1 α) activa-

tion, which promotes glucose uptake and modulate its metabolism.

HIF-1 α is hydroxylated under normal conditions but in hypoxic situa-

tions, hydroxylation of HIF-1 α does not occur and α subunit translocates to the nucleus to bind β counterpart. The heterodimer triggers a modification of genes expression. These genes are involved, among other effects, in glucose metabolism. This mechanism is likely used in bone tissue as an adaptive way to poorly vascularized and hypoxic environment. The group also showed that OC siRNA silencing counteracts 1,25(OH)₂D induced model of VC in young rats [174]. In an interesting way, Flammer et al. [175] selected CD133 +/CD34-/KDR + endothelial progenitor cells (EPCs) and demonstrated that expression of OC on circulating early EPCs is strongly associated with unstable coronary artery disease. Fadini et al. [176] also showed a procalcific polarization of circulating progenitor cells in type 2 diabetes mellitus patients, especially those who suffer from coronary artery disease. OC levels are known to increase in HD patients [177,178]. OC

regulation clearly undergoes disruption during mineralization process.

Nonetheless, its utility as serum marker is still discussed in conflicting studies. Aoki et al. [148] did not show any relationship between OC and VC in type 2 diabetes mellitus patients whereas Kim et al. [179] found an inverse correlation between OC and Agatston calcification

score in Asian women, even after adjusting for age. In order to define if OC can be used as a diagnostic or a screening tool, the role of OC in the pathogenesis of VC clearly remains to be clarified.

3.9. Pyrophosphate (Ppi)

hydrolysis [186].

PPi is a small molecule made of two phosphate ions. It acts as a calcification inhibitor by inhibiting hydroxyapatite crystal formation [180]. Once again, knock-out mice (in fact, knock-out mice for a precursor) develop VCs [181]. Absence of PPi would promote VSMC differentiation but the mechanism is not fully understood [182,183]. Absence of PPi combined with high phosphate levels and presence of type 1 collagen could facilitate the development of calcification [184,185]. Two ectonucleotidases have been implicated in extracellular PPi (ePPi) homeostasis during VSMCs calcification: tissue-non specific ALP hydrolyzes ePPi to phosphate, and nucleotide pyrophosphatase/ phosphodiesterase-1 (eNPP1) generates ePPi by extracellular ATP

It has been shown that dialysis patients exhibit low serum PPi and that these are further lowered during a HD session. PPi plasma levels were not affected by the mode of dialysis or by nutritional or inflammatory status [187]. O'Neill et al. [188] demonstrated the negative association between PPi and VC in CKD.

Although the short half life of PPi limits the possibility for improving

VC by bolus injections, daily peritoneal dialysis achieved with a solution

which contains PPi in CKD mouse model do succeed in inhibiting calcification [189]. O'Neill et al [190] demonstrated that daily intraperitoneal injections in rats could also reduce both incidence and amount of calcification. PPi has been shown to inhibit mineralization on rat aortic VSMCs cultures too [191]. Furthermore, biphosphonates, non hydrolysable analogs of PPi, have also proved their ability to inhibit aortic calcifications in experimental renal failure rats. Calcification was stopped in cultures of rat aortas as well as in vivo model. It supports the idea that biphosphonates have direct effects on VC, independent of bone [192], maybe via a down regulation of Notch1-RBP-Jk signaling pathway and MsX2 gene induction [193]. ATP, which is a polyphosphate associ-

ated with nucleoside, might also act as calcium phosphate deposition

inhibitor, not only as the source of PPi but also as a direct inhibitor

[186]. Even if PPi seems to be a promising marker, its determination

has been performed in a single center only and the transferability to

Sclerostin is an osteocyte-specific glycoprotein and is considered as a

potent inhibitor of bone formation [194,195]. It inhibits specific co-

3.10. Sclerostin

other centers should be validated.

receptors needed for β-catenin-dependant signaling activation [196]. This pathway is involved in osteoblast-mediated bone formation [197] . It is thought that sclerostin plays a role in bone mechanosensibilisation. When bone undergoes a substantial strain, sclerostin production would be decreased and bone could thus increase its formation in response to mechanical stress [198]. As β-catenin belongs to Wnt cascade signaling and as Wnt pathway is thought to be implicated in development of VC, it is interesting to investigate a potential association between sclerostin

levels and VCs. In non-CKD patients, some studies have demonstrated

a positive association between sclerostin levels and VC [199,200]

whereas in other ones, there was not a significant correlation between

the two parameters [201,202]. Results of similar investigations seem

quite different in specific population of HD patients. Indeed, in their

study including 154 CKD patients, Claes et al. [203] highlighted

absence of calcifications as an independent predictor of higher level of

sclerostin. Further studies are needed for confirming a possible sclerostin

involvement in VC genesis and for defining its potential usefulness in VC assessment.

Today, the ideal marker of VC does not exist. The pathophysiological

4. Discussion

mechanisms underlying this phenomenon are still poorly understood. As explained in the introduction, calcification can be induced by various situations. Etiologies that induce VC in diabetes mellitus patients are likely different from those which lead to the same result in CKD patients or postmenopausal women. Signaling pathways that are involved in VC may then depend on patient's status. A perfect marker would be ideally located on a hypothetical convergence point of all these pathological conditions. Thus, it could reflect reliably calcification emergence and progression in any situation. But this view is maybe too utopian and simplistic. Over the years, study of biomarkers showed a large variety of conditions that can modulate vascular microenvironment composition, such as bone turnover, inflammation, vitamin D status or even oxidative stress. Within this vascular microenvironment itself, a dense and interconnected network of calcification inhibitors and promoters was highlighted, as shown by Fig. 4. (1) VSMCs undergo differentiation into osteoblast-like cells, in

tion, likely mediated by the co-transporter Pit-1, in response to extracellular hyperphosphatemia. (2) Renal failure is one of the major hyperphospatemia origin whereas (3) FGF-23 is a factor which tend to moderate it by increasing phosphate renal excretion. This FGF-23 action is achieved with Klotho's help. FGF-23 has other effects described in Fig. 2, among which noxious ones are also suspected. (4) Other factors such as BMP-2, absence of PPi (in part, due to ALP activity), oxidative stress, inflammatory process or metabolic disorders are also known to be responsive to VSMCs conversion. (5) Calcium enhancement also proved to be deleterious, especially by its ability to induce Pit-1 overexpression and also by alteration of MGP and Fetuin-A actions, two VC inhibitors.

great part because of an intracellular phosphate increased concentra-

(6) The VSMCs conversion will favor excretion of bone-associated proteins, such as OPN, collagen type 1, BMP-2 and OC and (7) VSMCs will release mineralization-competent MVs. In turn, (8) BMP-2 can promote osteoblast differentiation and is a potent calcification inducer. (9) MGP, expressed by chondrocytes and VSMCs under normal conditions, inactivates BMP-2. It also binds calcium crystals and inhibits crystal growth. Finally, it prevents osteoblastic differentiation too. As osteocalcin, its activity is vitamin-K dependent and can be countered by vitamin-K antagonists. (10) Osteocalcin and (11) osteonectin are known to bind Ca/Pi

crystal but their accurate actions as inducers or inhibitors need to be specified, whereas the (12) PPi inhibiting effect on crystal formation is well known. The recent discovery of OC metabolic effects might suggest OC is a promoter of VSMCs differentiation into osteoblast-like cells. (13) OPN activity would depend on its phosphorylation state. Fully phosphorylated OPN would inhibit mineralization by blocking hydroxyapatite formation and activating osteoclast function while the cleaved one could act as a proinflammatory cytokine and proangiogenic factor facilitating vascular mineralization. (14) OPG is considered as a regulatory factor. On one hand, it can

prevent VC by blocking bone remodeling process in vascular tissue and by neutralizing the pro-apoptotic actions of TRAIL. It might also inhibit ALP activity. On the other hand, the inhibition of bone remodeling process by OPG could induce a calcium shift into vascular cells. (15) Fet-A released by the liver inhibits mineralization, perhaps through calciprotein particles formation while (16) PTH secreted by parathyroid enhances calcification phenomenon, as both low and high bone turnover might lead to VC. Under normal conditions, there is a balance between all these

parameters. It is possible that each pathological condition disrupts the balance with its own approach. Nevertheless, it seems that all the

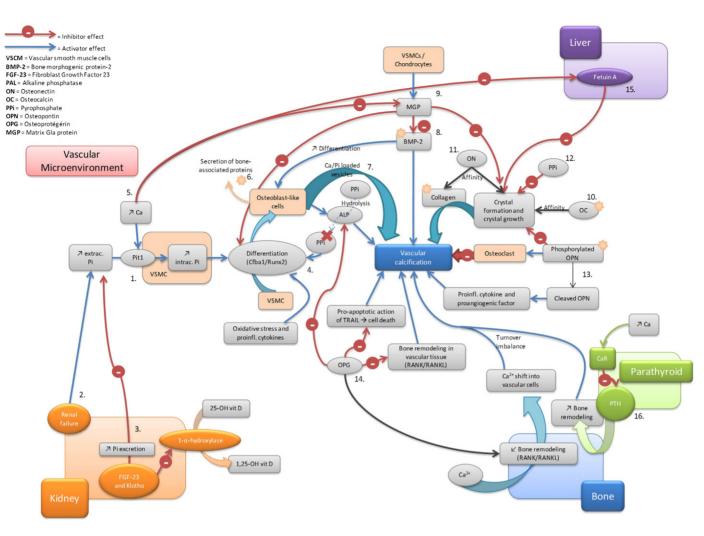


Fig. 4. Vascular microenvironment: inhibitors and promoters of vascular calcification.

calcification inhibitors do not possess the same potential. OPG serum levels appear to be correlated repeatedly to calcification in many pathological conditions. However, OPG would not reflect the severity of damages as would FGF-23 do. FGF-23 levels should be followed up, either upwards or downwards, given the suspected duality of the effects of FGF-23. Low serum Fet-A levels are usually associated with VC but an increased CPP fraction of Fet-A would also be useful to reflect a procalcific milieu. Non fully γ -carboxylated MGP is associated with VC too and might be an interesting marker to monitor patients under AVK treatment. OPN would be an attractive marker in diseases with inflammatory component such as diabetes or autoimmune diseases. Within this category of disease, it might help to reclassify asymptomatic subjects with classical risk factors into high risk group for further examination.

In population with renal deficiency particularly susceptible to develop VC, the problem becomes even more complex. Whether patients undergo dialysis or not, reference values will need to be adaptated, depending on renal failure severity and marker ability to be removed by dialysis. In addition, as described earlier, complex interactions exist between different actors (pro or anti-calcifications) such as PTH, vitamin D, FGF-23, OPG, Sclerostin, acting sometimes via redundant signaling pathways such as Wnt/ β -catenin, RunX2/cbfa1, Notch1-RBP-Jk. Furthermore, these interactions could be different according to the stage of CKD [204]. Thus, the stage of disease should be taken into account in the interpretation of biomarker and/or the combination of biomarkers.

As evidenced by the present discussion, VC physiopathology is still far from being fully elucidated. The role of each biomarker needs to be

clarified and many studies are still leading to contradictory results. In vitro observations are sometimes very different from conclusions observed in in vivo studies. Direct effects on vasculature and indirect effects mediated by bone turnover are not easy to discriminate. When a correlation between the serum levels of a calcification marker and calcification is clearly showing up, it still remains to determinate whether level fluctuations attest a noxious effect of the biomarker or if they highlight a compensatory process or even solely reflect phenomenon as bystander.

Qualities that would be appreciated for selecting a good marker depend on its capacities to achieve clinical goals, particularly its ability to select high risk patients for further investigation, to make a reliable calcification assessment, to provide a prognostic, to help in treatment choice or to follow up the treatment efficiency. Given importance to assess and control mineralization process, it is essential to keep going on building-up more and more knowledge.

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