


Phenotypic switch of vascular smooth muscle cells in COVID-19: Role of cholesterol, calcium, and phosphate

Laura Ghanem¹ | Dina Essayli¹ | Jana Kotaich^{1,2} | Mohammad Al Zein¹ | Amirhossein Sahebkar^{3,4,5}  | Ali H. Eid⁶ 

¹Faculty of Medical Sciences, Lebanese University, Hadath, Lebanon

²MEDICA Research Investigation, Beirut, Lebanon

³Center for Global Health Research, Saveetha Medical College and Hospitals, Saveetha Institute of Medical and Technical Sciences, Saveetha University, Chennai, India

⁴Biotechnology Research Center, Pharmaceutical Technology Institute, Mashhad University of Medical Sciences, Mashhad, Iran

⁵Applied Biomedical Research Center, Mashhad University of Medical Sciences, Mashhad, Iran

⁶Department of Basic Medical Sciences, College of Medicine, QU Health, Qatar University, Doha, Qatar

Correspondence

Ali H. Eid, Department of Basic Medical Sciences, College of Medicine, QU Health, Qatar University, Doha, Qatar.
Email: ali.eid@qu.edu.qa

Abstract

Although the novel coronavirus disease 2019 (COVID-19), caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), primarily manifests as severe respiratory distress, its impact on the cardiovascular system is also notable. Studies reveal that COVID-19 patients often suffer from certain vascular diseases, partly attributed to increased proliferation or altered phenotype of vascular smooth muscle cells (VSMCs). Although the association between COVID-19 and VSMCs is recognized, the precise mechanism underlying SARS-CoV-2's influence on VSMC phenotype remains largely under-reviewed. In this context, while there is a consistent body of literature dissecting the effect of COVID-19 on the cardiovascular system, few reports delve into the potential role of VSMC switching in the pathophysiology associated with COVID-19 and the molecular mechanisms involved therein. This review dissects and critiques the link between COVID-19 and VSMCs, with particular attention to pathways involving cholesterol, calcium, and phosphate. These pathways underpin the interaction between the virus and VSMCs. Such interaction promotes VSMC proliferation, and eventually potentiates vascular calcification as well as worsens prognosis in patients with COVID-19.

KEYWORDS

atherosclerosis, calcification, cardiovascular disease, differentiation, SARS-CoV-2

Abbreviations: 25HC, 25-hydroxycholesterol; 27OHC, 27-hydroxycholesterol; ACAT, acyl-CoA cholesterol acyltransferase; ACE2, angiotensin-converting enzyme 2; ACTA2, smooth muscle aortic alpha-actin; ADAM17, disintegrin and metalloproteinase domain-containing protein 17; ALP, alkaline phosphatase; AMC, arterial medial calcification; ApoB, apolipoprotein B; ATP, adenosine 5'-triphosphate; Ca, calcium; CAC, coronary artery calcification; cAMP, cyclic adenosine monophosphate; CH25H, cholesterol 25-hydroxylase; ChemR23, chemerin receptor 23; CKD, chronic kidney disease; COL1, type 1 collagen; COVID-19, coronavirus disease of 2019; CREB, cAMP response element-binding protein; CVDs, cardiovascular diseases; ECM, extracellular matrix; ER, endoplasmic reticulum; Fe²⁺, ferrous ions; FN, fibronectin; GPI, glycosylphosphatidylinositol; HCoV-OC43, human coronavirus organ culture 43; HDL, high-density lipoprotein; HF, heart failure; HIF-1, hypoxia-inducible factor-1; HIPO⁴⁻, inorganic phosphate; HSPG, heparan sulfate proteoglycan; IL, interleukin; ISG, interferon-stimulated gene; LDL-c, low-density lipoprotein cholesterol; LDL-P, low-density lipoprotein particle; LOE, linderia obtusiloba; LXR, liver X receptor; MGP, matrix Gla protein; mtROS, mitochondrial reactive oxygen species; MYH11, myosin heavy chain; MβCD, methyl-β-cyclodextrin; NF-κB, nuclear factor kappa B; O₂, dioxygen; OPN, osteopontin; PDGF-BB, platelet-derived growth factor BB; PKA, protein kinase A; PMVs, platelet-derived microvesicles; Ppi, inhibitor pyrophosphate; RAAS, renin-angiotensin-aldosterone; RBD, receptor binding domain; Runx2, runt-related transcription factor 2; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; SM22α, smooth muscle 22 alpha; SMC, smooth muscle cell; SOD2, superoxide dismutase 2; Sox9, SRY-box transcription factor 9; SREBP2, sterol-regulatory element binding proteins-2; T3, triiodothyronine; TGF-β1, transforming growth factor beta 1; TLR4, toll-like receptor 4; TNF-α, tumor necrosis factor-alpha; TRPV4, transient receptor potential vanilloid 4; UBIAD1, UbiA prenyltransferase domain containing 1; UTP, uridine-5'-triphosphate; VEGF, vascular endothelial growth factor; VSMCs, vascular smooth muscle cells.

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1 | INTRODUCTION

Coronavirus disease 2019 (COVID-19) is a contagious disease caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), a Beta-coronavirus genus in the Coronaviridae family (Martinez-Salazar et al., 2022; SeyedAlinaghi et al., 2021). SARS-CoV-2 is an enveloped, positive-sense, single-stranded ribonucleic acid (RNA) virus (Harrison et al., 2020). Two-thirds of its genome encodes 16 nonstructural proteins (Harrison et al., 2020). Four structural proteins, namely the spike (S), envelope (E), membrane (M), and nucleocapsid (N), are encoded by the remaining one-third of the genome (Harrison et al., 2020). The spike S protein plays an essential role in mediating the entry of SARS-CoV-2 into the target cells (Harrison et al., 2020). In addition, another component that allows the virus to “enter” target cells is the angiotensin-converting enzyme 2 (ACE2) receptor (Salabei et al., 2022). This receptor is more ubiquitously expressed on myocytes, fibroblasts, endothelial cells, and smooth muscle cells (SMCs) (Salabei et al., 2022).

Respiratory involvement remains the primary manifestation of COVID-19. According to a systematic review and meta-analysis, more than half of patients with COVID-19 present with cough (da Rosa Mesquita et al., 2021). However, COVID-19 has also been shown to promote systemic symptoms. In essence, robust evidence suggests that significant portion of SARS-CoV-2 morbidity and mortality is attributed to cardiovascular complications, such as acute coronary syndrome, myocarditis, arrhythmia, and venous thromboembolism (Clerkin et al., 2020). A meta-analysis reported that the prevalence of cardiac injury in hospitalized patients with COVID-19 is 22%, and this proportion increases with disease severity, suggesting that cardiac injury is associated with worse prognosis and higher mortality (Fu et al., 2021). Furthermore, CVD is the most common comorbidity in patients with COVID-19, and COVID-19 patients with underlying cardiovascular disease demonstrate a worse prognosis (Li, Dong, et al., 2020). Among the mechanisms proposed to explain the bidirectional interaction between COVID-19 and the cardiovascular system, the ACE2 receptor appears to play a key role, as evidenced by both animal and human studies (Chilazi et al., 2021). The presence of ACE2 on the surface of endothelial cells and vascular smooth muscle cells (VSMCs) mediates cardiovascular injury and contributes to increased mortality (Liu et al., 2020).

VSMCs play integral roles in regulating vasotone and blood flow (Deng et al., 2021; Zhang et al., 2016). In healthy vasculature, VSMCs perform many of the physical homeostatic functions of arteries and arterioles, as well as the production and remodeling of the extracellular matrix (ECM) (Deng et al., 2021; Zhang et al., 2016). Under certain environmental factors, these highly specialized quiescent cells may undergo certain morphological, followed by functional, alterations that may lead to VSMC-driven vascular diseases (Deng et al., 2021; Zhang et al., 2016). This plasticity of VSMCs is most evident when they respond to various stimuli by switching from a differentiated to a dedifferentiated phenotype (Frismantiene et al., 2018; Zhang et al., 2016). In addition, when blood vessels are damaged or stimulated by growth factors, VSMCs can respond by

increased proliferation, migration, and synthesis of extracellular components, a phenomenon referred to as phenotypic switching (Frismantiene et al., 2018; Zhang et al., 2016). Many critical regulators can be implicated in the pathogenesis of this phenotypic switching, such as ions and molecules, including cholesterol, calcium, and phosphate.

The role of cholesterol, calcium, and phosphate in VSMC phenotypic switch has been extensively studied (Tang et al., 2022; Vengrenyuk et al., 2015). In patients with atherosclerosis, VSMC dedifferentiation plays a pivotal role in plaque formation. Indeed, it was shown that in atherosclerotic patients, approximately 40% of cells identified as macrophages were derived from VSMCs (Vengrenyuk et al., 2015). Interestingly, one molecule that plays a role in VSMCs transition from a contractile phenotype to a macrophage-like one is cholesterol (Vengrenyuk et al., 2015). Moreover, contractile VSMCs may exhibit resistance to vascular calcification, a process wherein calcium and phosphate are deposited in the form of hydroxyapatites (Furmanik et al., 2020). Intriguingly, phenotypic remodeling of VSMCs promotes vascular calcification (Furmanik et al., 2020), while calcification upregulates transcription factors that mediate differentiation of VSMCs to an osteoblastic phenotype (Bundy et al., 2021). Therefore, an imbalance of calcium and phosphate may also play an important role in the VSMC phenotype switching, and vice versa (Bundy et al., 2021; Furmanik et al., 2020).

Although the relationship between COVID-19 and arterial muscle cells has been suggested in recent studies, the exact mechanism of the interplay between SARS-CoV-2 and VSMC phenotypic switching remains largely unknown (Kar, 2022; Martínez-Salazar et al., 2022; Naeem et al., 2023). Thus, this review was undertaken to determine the molecular mechanism implicated in this switch while highlighting the active role of calcium, phosphate, and cholesterol in the pathophysiology of vascular outcomes and disease progression in COVID-19 patients. Moreover, pinpointing the dysregulation of VSMC modulation in patients with comorbidities may explain why COVID-19 is more serious in these individuals.

2 | COVID-19 AND THE CARDIOVASCULAR SYSTEM

As mentioned earlier, cardiovascular involvement is a common complication of COVID-19, and this is attributed to the powerful expression of ACE2 receptor on the surface of epithelial and vascular smooth muscle cells (Chatzis et al., 2022; Salabei et al., 2022; Wehbe, Hammoud, et al., 2021). The interaction between SARS-CoV-2 and ACE2 disrupts signal transduction pathways and cellular hemostasis, resulting in myocardial and vascular injury (Soumya et al., 2021). Furthermore, other mechanisms such as hypoxia and systemic inflammation put an additional burden on the heart and vessels, leading to a worse prognosis (Nishiga et al., 2020). Acute cardiac injury, defined by the elevation of serum cardiac biomarker levels >99th percentile of upper reference limit, is a common extra-pulmonary manifestation of COVID-19 (Tajbakhsh et al., 2021).

About 21% of patients hospitalized for COVID-19 demonstrate cardiac injury, and this proportion rises to almost 100% in critically ill patients (Li, Yang, et al., 2020). The pathophysiological mechanisms underlying myocardial injury in COVID-19 patients include direct viral damage, endothelial dysfunction, hypoxia, and systemic inflammation (Helms et al., 2022). Clinical evidence has also demonstrated that SARS-CoV-2 can trigger acute coronary syndrome, and this has been attributed to endothelial injury, thrombus formation, and plaque rupture (Schiavone et al., 2020). In addition, patients with COVID-19 are at increased risk of arrhythmias and sudden cardiac arrest. According to a systematic review, the incidence of arrhythmias in patients with COVID-19 is up to 18%, and this can be attributed to myocardial injury, electrolyte imbalances, fever, and sepsis (Hessami et al., 2021). Acute or decompensated heart failure, as evidenced by the development of cardiogenic pulmonary edema, has been reported in 6.5% of COVID-19 patients (Harrison et al., 2021). The development of systemic symptoms, such as fever, tachycardia, and hypoxia, in patients with underlying comorbidities might result in the decompensation of cardiac function. Furthermore, myocardial injury and acute coronary syndrome triggered by COVID-19 can also result in cardiac dysfunction (Mehra & Ruschitzka, 2020). Clinical observations showed that COVID-19 patients are also at increased risk of venous thromboembolism. Studies from China demonstrate that high levels of D-dimer (≥ 0.5 mg/l) were present in 46% of patients (Guan et al., 2020). In another study, elevated D-dimer was associated with greater odds of death (Zhou et al., 2020). It is suggested that the systemic inflammatory response and endothelial dysfunction promote a condition of hypercoagulable state in COVID-19 patients (Bikdeli et al., 2020).

Several studies highlighted that the course of COVID-19 is not only dependent on the effect of SARS-CoV-2 infection on the host cells, but also on the downstream signaling pathways initiated by the binding of coronavirus' spike protein to its receptor on the endothelial cells and VSMCs (Parums, 2022; Suzuki et al., 2021). The endothelial cells, which constitute the innermost layer of the blood vessels, play a pivotal role in maintaining tissue homeostasis, through the regulation of systemic blood flow, immune system, coagulability state, and tissue perfusion, in accordance with other cells, such as pericytes and VSMCs (Pelisek et al., 2022; Soumya et al., 2021; Xu et al., 2023). However, disruptions in cellular homeostasis in the settings of COVID-19, such as the upregulation of reactive oxygen species production and subsequent imbalance in redox status, promote endothelial dysfunction and organ injury (Soumya et al., 2021). Furthermore, accumulating evidence suggests that SARS-CoV-2 infection causes various instances of VSMC dysfunction, including phenotypic switch, proliferation, and hypertrophy. These modifications are deemed to increase contractility and induce vascular remodeling. Essentially, a study highlighted that SARS-CoV-2 infection promotes vascular dysfunction characterized by enhanced vasoconstriction and impaired vasorelaxation by acting on the RhoA/Rho-kinase signaling pathway (Sykes et al., 2023). Furthermore, some studies focused on the involvement of SARS-CoV-2 in enhancing the systemic

inflammation mediated by the NLRP3 inflammasome in VSMCs via the SCAP-SREBP signaling pathway; which can put into relief the interconnectedness between COVID-19 and VSMCs (Liu et al., 2024a, 2024b). Another study elucidated the mechanisms underlying the increased cardiovascular risk after coronavirus infection by focusing on the IL18/IL18R1/HIF-1 signaling pathway (Zhang et al., 2021). Essentially, IL18 induction during SARS-CoV-2 infection results in the abnormal activation of HIF-1 signaling pathway, thereby promoting the synthetic phenotype of VSMC. Interestingly, HIF-1 overexpression has been demonstrated a key mechanism in the pathogenesis of vascular disease, such as atherosclerosis and aneurysm formation (Gao et al., 2012).

3 | COVID-19 AND VSMC PHENOTYPIC SWITCH: A FOCUS ON CHOLESTEROL, CALCIUM, AND PHOSPHATE

3.1 | Role of cholesterol and calcium in SARS-CoV-2 entry and replication

One of the factors that play a role in SARS-CoV-2 infection is lipid metabolism, particularly the one involving cholesterol. Indeed, it was demonstrated that the coronavirus entrance into the cell involves plasma fusion and endocytosis, a mechanism that primarily involves the lipid raft microdomains characterized by the presence of cholesterol, glycosphingolipids, and glycosylphosphatidylinositol (GPI)-anchored proteins (Casari et al., 2021; Kočar et al., 2021). These domains are considered the docking site of SARS-CoV-2 to pass through the cell membrane and release their genome (Casari et al., 2021; Kočar et al., 2021). In effect, it was shown that the cholesterol-rich lipid rafts are highly concentrated in receptors and co-receptors (Casari et al., 2021; Clausen et al., 2020; Li, Zhu, et al., 2021; Palacios-Rápalo et al., 2021). These receptors could have synergistic effects with the SARS-CoV-2 surface proteins and modulate the virus entry through the cell membrane (Figure 1) (Casari et al., 2021; Clausen et al., 2020; Li, Zhu, et al., 2021; Palacios-Rápalo et al., 2021). For example, one of these receptors, particularly the ACE2 receptor, binds to the S protein-receptor binding domain (RBD) of SARS-CoV-2 and facilitates its entry (Casari et al., 2021; Clausen et al., 2020; Palacios-Rápalo et al., 2021). Viral entry through this receptor also necessitates the interaction between the coronavirus spike protein and the heparan sulfate proteoglycan (HSPG) (Clausen et al., 2020; Palacios-Rápalo et al., 2021; Zhang et al., 2020). Another receptor, the toll-like receptor 4 (TLR4), which interacts with the S1 subunit of spike protein, facilitates SARS-CoV-2 entry even in the absence of the ACE2 receptor (Aboudounya et al., 2021; Butnariu et al., 2021; Palacios-Rápalo et al., 2021). To further elucidate the role of cholesterol in the life cycle of SARS-CoV-2, several studies have demonstrated that the manipulation of host membrane cholesterol alleviate virus entry into host cells. Indeed, it was demonstrated that depletion of membrane-bound cholesterol from

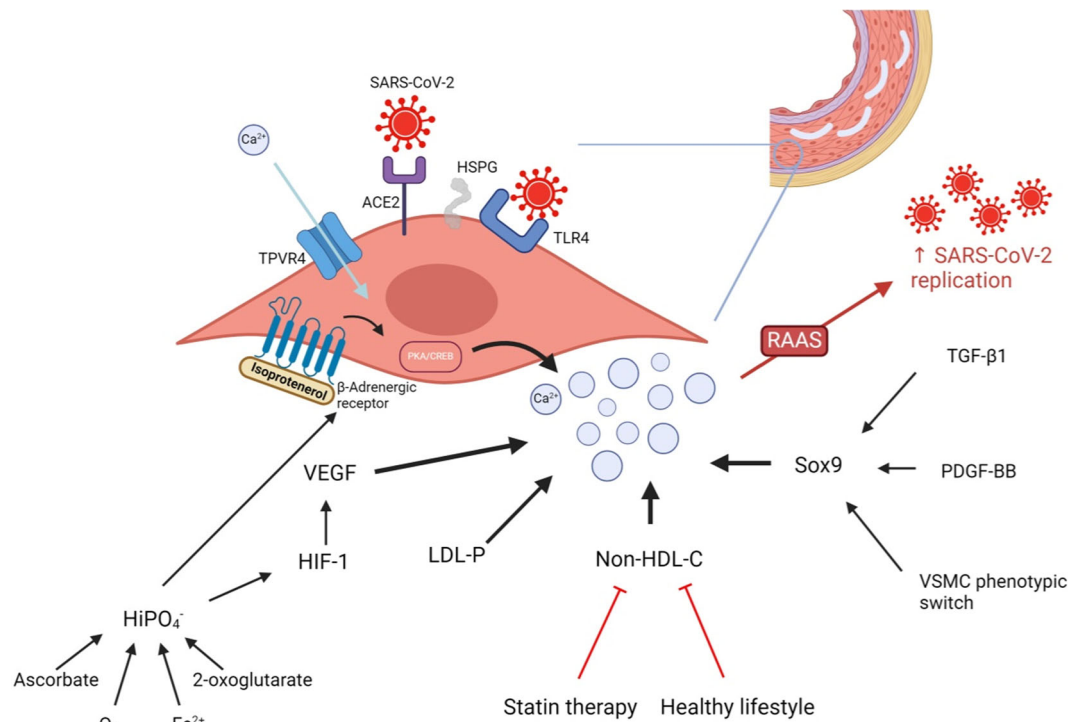


FIGURE 1 Potential mechanisms promoting vascular smooth muscle cells (VSMC) calcification and its phenotypic switch, resulting in increased severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) replication and entry. Calcification plays an important role in the VSMCs phenotypic switch. On one hand, an increased Ca^{2+} flow can be mediated through the transient receptor potential vanilloid 4 (TRPV4) on the VSMCs surface, resulting in their dedifferentiation. On the other hand, phenotypic switch can itself promote Ca^{2+} deposition. This calcification further enhances the SARS-CoV-2 replication via the renin-angiotensin-aldosterone (RAAS) activation. Moreover, lipid molecules can have a stimulatory effect on calcium, as well as on the coronavirus entry through toll-like receptor 4 (TLR4) and angiotensin-converting enzyme 2 (ACE2) receptor enhanced by heparan sulfate proteoglycan (HSPG). In fact, other mechanisms can also stimulate calcium deposits. For instance, ascorbate, O_2 , Fe^{2+} , and 2-oxoglutarate result in hyperphosphatemia, which activates the hypoxia-inducible factor-1 (HIF-1), responsible of stimulating the vascular endothelial growth factor (VEGF), an important factor for calcification. Moreover, high HiPO_4^- levels heighten beta-adrenergic receptor activity, thus stimulating the protein kinase A (PKA)/cAMP response element-binding protein (CREB) pathway and increasing Ca^{2+} levels. SRY-box transcription factor 9 (Sox9) expression is also involved in Ca^{2+} deposition. Created with [BioRender.com](https://www.biorender.com). Ca, calcium; Fe^{2+} , ferrous ions; HDL, high-density lipoprotein; HiPO_4^- , inorganic phosphate; LDL-P: low-density lipoprotein particle; O_2 , dioxygen; PDGF-BB, platelet-derived growth factor BB; TGF- β 1, transforming growth factor beta 1.

the lipid raft of ACE2-rich cells impaired SARS-CoV-2 entry into host cells (Sanders et al., 2021).

Another component contributing to the clinical manifestations of COVID-19 patients is calcium. A recent systematic review and meta-analysis showed that coronary calcification is associated with increased mortality in patients with COVID-19 (Cereda et al., 2022). This may be attributed to the greater activation, in these patients, of the renin-angiotensin-aldosterone system (RAAS), which increases the expression of ACE2 receptors on VSMC surface, thereby enhancing viral replication and vascular injury (Figure 1) (Beyerstedt et al., 2021; Cereda et al., 2022). This is in concordance with another study which showed that increased intracellular calcium load activates cathepsin L, which in turn aids in the cleavage of the spike S into S1 and S2 (Tang et al., 2020; Wei et al., 2022). S1 can bind to the ACE-2 receptor, while S2 facilitates the endocytosis of the virus into the host cells (Pizzato et al., 2022; Wei et al., 2022). The binding of the S2 to the lipid raft microdomains of the target cell is enhanced by elevated Ca^{2+} levels (Sultan et al., 2022).

3.2 | Role of calcium, cholesterol, and phosphate in VSMC phenotypic switch

Increasing evidence suggests that one of the major precipitating factors of the COVID-19 course is vascular calcification (Possari et al., 2021; Shabestari et al., 2022). Coronary artery calcification was associated with a worse prognosis in patients infected with SARS-CoV-2 (Dillinger et al., 2020; Luo et al., 2022; Shabestari et al., 2022). Moreover, many studies shed light on the higher mortality rate in COVID-19 patients who had vascular calcification, in comparison to those without calcified vessels (Gupta et al., 2021; Meyer et al., 2023; Slipczuk et al., 2021). It is important to mention that vascular calcification mainly emerges from the osteo-/chondrocyte phenotypic switch of VSMCs, which makes it highly essential to discuss the role of calcium, cholesterol, and phosphate on the VSMC phenotypic switch (Liu et al., 2022). Several studies show that the transient receptor potential vanilloid 4 (TRPV4) channel enables the entry of extracellular calcium (Figure 1) (Cao et al., 2018; Li, Gao, et al., 2021;

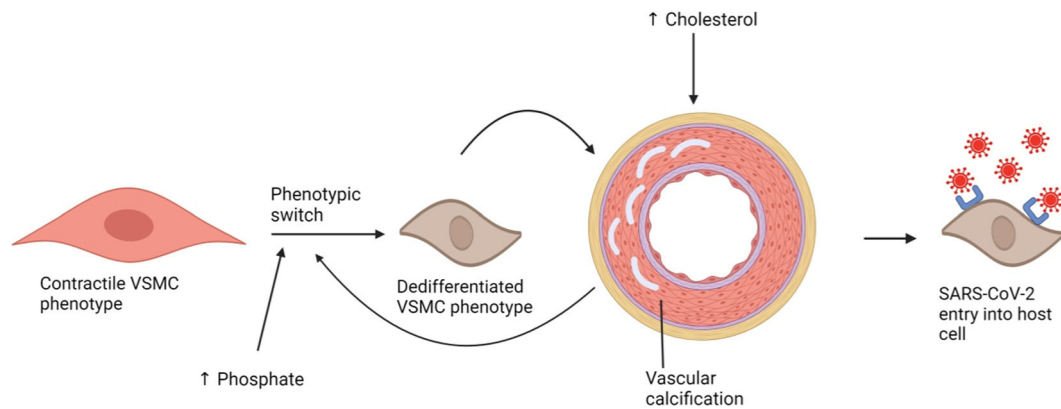


FIGURE 2 Summary of the interplay between vascular smooth muscle cell (VSMC) phenotypic switch, vascular calcification, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) entry into the cell. Contractile VSMCs can undergo a phenotypic switch under hyperphosphatemia, which results in dedifferentiated VSMCs. This can lead to vascular calcification, which can exert positive feedback on the phenotypic switch. Vascular calcification enhances the SARS-CoV-2 entry into the host cell, and can be stimulated by high cholesterol levels. Created with [BioRender.com](https://www.biorender.com).

Toft-Bertelsen et al., 2019). This entry can result in neointimal hyperplasia and VSMC migration, while the L-type voltage-gated calcium channel has no effect on VSMC phenotype (Li, Gao, et al., 2021). Interestingly, medial calcification triggers changes in cell wall morphology and results in higher arterial stiffness (Jaminon et al., 2019), suggesting that this calcification is highly correlated with phenotypic switch of VSMCs where cells acquire some features of chondrocytes and osteoblasts (Jaminon et al., 2019) (Figure 2).

One of the factors affecting VSMC calcification, therefore promoting phenotypic switching, is cholesterol. A prospective study investigated this in 1980, with a follow-up period of over 28 years (Armstrong et al., 2021). In each stage of development, mainly in adolescence, non-HDL-c was associated with coronary artery calcification (Armstrong et al., 2021). Moreover, low-density lipoprotein particles (LDL-P) were highly associated with the presence of coronary artery calcification, independently of other lipid particles (Prado et al., 2011; Zaid et al., 2016). In addition, the low-density lipoprotein cholesterol (LDL-C) was shown to be dependent on other LDL-Ps and to be associated with coronary artery calcification, even if the association was weaker than that of LDL-P (Figure 1) (Prado et al., 2011; Zaid et al., 2016).

Levels of phosphate are yet another factor that affects VSMC differentiation. Interestingly, increased inorganic phosphate (HiPO^{4-}) levels rapidly activate the hypoxia-inducible factor-1 (HIF-1), a transcription factor implicated in VSMC phenotype (Figure 1) (Mokas et al., 2016). HiPO^{4-} -induced HIF-1 activation drives expression of various genes like including vascular endothelial growth factor (VEGF), a critical angiogenic factor (Mokas et al., 2016; Zimna & Kurpisz, 2015). Interestingly, VEGF is expressed in osteoblast precursor cells that boost VSMC calcification and osteogenic transdifferentiation (Figure 1) (Hu & Olsen, 2016; Mokas et al., 2016). Similarly, hyperphosphatemia in chronic kidney disease (CKD) increases the expression of β -adrenergic receptors (Moser et al., 2021). Stimulation of these receptors activates the protein

kinase A (PKA)/cyclic adenosine monophosphate (cAMP) response element-binding protein (CREB) signaling pathway, which then enhances the osteogenic transdifferentiation of VSMCs (Figure 1) (Moser et al., 2021).

3.3 | Effect of VSMC switch on calcification

Although the role of vascular calcification in promoting VSMC phenotypic switch is well established, some studies stated that these two processes may happen in a reversed order (Figure 2). For instance, VSMC phenotype switch was detected 4 weeks before arterial medial calcification (AMC) (Pai et al., 2011). Another study in $\text{ApoE}^{-/-}$ mice reported that vascular remodeling results in the transdifferentiation of VSMCs, evident by increased migration and proliferation (Augstein et al., 2018). This switch was concomitant with increased expression of the SRY-box transcription factor 9 (Sox9) (Augstein et al., 2018). Consequently, an increase in the number of apoptotic cells is noted, and ECM remodeling and increased calcium deposition ensue (Augstein et al., 2018; Faleeva et al., 2023) (Figure 1).

4 | DEFENSE MECHANISMS OF THE PATHWAYS INVOLVING CALCIUM, PHOSPHATE, AND CHOLESTEROL

4.1 | Inhibitors of SARS-CoV-2 entry into VSMCs

Inhibiting SARS-CoV-2 entry into host cells has been of much interest. A factor that plays an essential role in viral life cycle inhibition is lipid levels, especially at the stage of the virus fusion with the target cell membrane. While non-HDL-c is essential for VSMC calcification, allowing the SARS-CoV-2 to enter VSMCs more readily, HDL-c has a markedly different effect. Indeed, as

pertains to infection with this virus, multiple studies have shown that low levels of HDL-c are associated with an increased risk of hospitalization, more severe disease, and even mortality (Lahoz et al., 2022; Zang et al., 2020). Among other studies, a cohort study involving half a million participants demonstrated the existence of this inversely proportional relation between HDL-c level and the risk of SARS-CoV2 infection (Lahoz et al., 2022; Zang et al., 2020).

The role of cholesterol role in inhibiting coronavirus entry is also related to another pathway involving 25-hydroxycholesterol (25HC) (Koçar et al., 2021). This 25HC, the product of cholesterol-25-hydroxylase (CH25H), can block the sterol-regulatory element binding protein-2 (SREBP2), and can stimulate the liver X receptor (LXR), hence activating the Acyl-coenzyme A: cholesterol acyltransferase (Figure 3) (Koçar et al., 2021; Mao et al., 2022). These actions repress the cholesterol and the lipid raft microdomains present in the cell membrane, thus blocking membrane fusion and disrupting viral protein maturation (Figure 3) (Koçar et al., 2021; Mao et al., 2022; Zang et al., 2020). Supportably, 25HC was found to suppress SARS-CoV-2's infection in lung epithelial cells (Wang et al., 2020). Since 25HC is an endogenous particle with unrecognized toxicity at adequate concentrations, it is considered an effective therapeutic substance for COVID-19 (Wang et al., 2020). Multiple studies in recent years have demonstrated the broad-spectrum antiviral activity of 25HC by virtue of its ability to suppress

SARS-CoV-2 spike protein-catalyzed membrane fusion, thus viral entry into VSMCs.

Entry of SARS-CoV-2 may also be inhibited by 27-hydroxycholesterol (27OHC). 27OHC exerts an antiviral effect against many viruses, including two human CoVs belonging to the β -coronavirus genus: SARS-CoV-2 and *Human Coronavirus Organ Culture 43* (HCoV-OC43) (Mao et al., 2022; Marcello et al., 2020). It has been suggested that 27OHC acts by modifying the cell structure rather than by targeting viral components (Marcello et al., 2020). This mode of action makes the 27OHC able to impede SARS-CoV-2 entry into cells including VSMCs, and hence rendering it as a potentially potent inhibitor.

Another process involved in inhibiting coronavirus entry is ACE2 shedding. As mentioned earlier, the ACE2 catalytic ectodomain is a required entry receptor for SARS-CoV-2 infection (García-Escobar et al., 2022; Glende et al., 2008). The liberation process of the ACE2 catalytic domain is facilitated by disintegrin and metalloproteinase domain-containing protein 17 (ADAM17) through the calcium signaling pathway field (García-Escobar et al., 2022; Zipeto et al., 2020). This mechanism "releases" a soluble catalytic ectodomain of ACE2 (García-Escobar et al., 2022). Such soluble forms of ACE2 block SARS-CoV-2 infection, thus offering a prognosis improvement in COVID-19 patients (García-Escobar et al., 2022; Glende et al., 2008; Haga et al., 2010; Zipeto et al., 2020). The shedding of the ACE2 catalytic ectodomain is a predictor of all-cause death, including

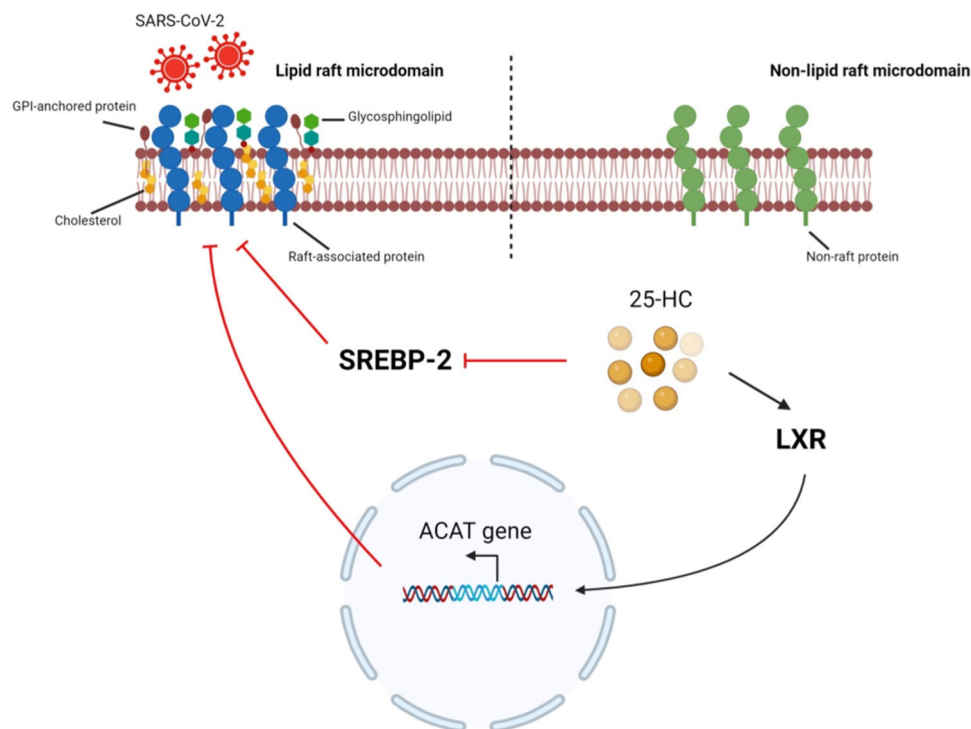


FIGURE 3 Mechanism by which 25HC blocks severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) fusion with the vascular smooth muscle cell (VSMC) lipid raft microdomain. 25-hydroxycholesterol (25HC) inhibits sterol-regulatory element binding proteins-2 (SREBP-2) and activates liver X receptor (LXR), which in turn stimulates acyl-CoA cholesterol acyltransferase. This results in the depletion of cholesterol, glycosphingolipids, and the glycosylphosphatidylinositol (GPI)-anchored proteins from the lipid raft microdomain, which then inhibits the entry of the SARS-CoV-2 into the host cell. Created with [BioRender.com](https://www.biorender.com). ACAT, acyl-CoA cholesterol acyltransferase.

cardiovascular mortality, cardiac remodeling, and endothelial dysfunction (García-Escobar et al., 2022; Haga et al., 2010).

4.2 | Inhibitors of calcification or VSMC switch

As stated above, vascular calcification and VSMC phenotypic switch affect SARS-CoV-2 entry (Figure 2). Therefore, it is not surprising that efforts are being expended to determine whether inhibitors of calcification could be efficacious against COVID-19. Contextually, patients with low calcium scores are more resilient to the SARS-CoV-2 infection (Cereda et al., 2022). Similarly, other studies confirm the antiviral effect of calcium channel blockers especially against coronaviruses, and further suggest their potential in suppressing viral entry as well as treating SARS-CoV-2-infected patients (Crespi & Alcock, 2021; Fani et al., 2023; Straus et al., 2021).

Calcification and phenotypic switch of VSMCs can be modulated by various enzymes. One such enzyme is superoxide dismutase 2 (SOD2), which downregulates mitochondrial reactive oxygen species, resulting in the downregulation of Ca^{2+} -sensitive intracellular cysteine protease calpain-1 (Roman-Garcia et al., 2011; Tsai et al., 2021). This leads to lower expressions of alkaline phosphatase and increases the expression of adenosine 5'-triphosphate (ATP) synthases, as well as calcification inhibitors (Roman-Garcia et al., 2011; Tsai et al., 2021). This, in turn, leads to diminished VSMC apoptosis and a low phosphate-induced VSMC calcification, thereby inhibiting VSMC phenotypic switch (Roman-Garcia et al., 2011; Tsai et al., 2021).

Some endogenous molecules may also be involved in this phenotypic switch. For instance, ATP, uridine-5'-triphosphate (UTP), and the ubiquitous mineralization inhibitor pyrophosphate (Ppi) can protect VSMCs from apoptosis (Opdebeeck et al., 2020; Patel et al., 2018). As such, those cells will no longer create a nucleation site for the hydroxyapatite crystal formation (Patel et al., 2018), hence preventing VSMC calcification from taking place (Patel et al., 2018). In addition, calcium and phosphate play an essential role in the VSMC phenotypic switch. These two minerals induce vascular calcification (Freise et al., 2015), which triggers VSMC dedifferentiation into a synthetic osteoblast-like phenotype (Houben et al., 2016; Opdebeeck et al., 2020). However, the coexistence of calcium and phosphate in large quantities yields a feedback loop; they upregulate Matrix Gla protein (MGP) levels, and in turn block calcification (Houben et al., 2016).

5 | COMORBIDITIES, VSMC, AND COVID-19

5.1 | CKD and COVID-19

Several studies highlight the notion that CKD may increase the risk of COVID-19 (Schiff & Lang, 2023). In patients with renal failure, the immune system is suboptimal, and a decrease in important immunologic mediators such as antibodies and complement is noted

(Schiff & Lang, 2023). This implies a consequent decline in the innate and adaptive immune system efficiency, resulting in more susceptibility to infections.

Because lower calcium levels may put a brake on SARS-CoV-2 entry, abating vascular calcification in CKD patient can potentially reduce their risk of being infected with SARS-CoV-2 (Jdiaa et al., 2022). In this context, interactions between VSMCs and chemerin, a biomarker of declined renal function, have been noted to play a role in calcification (Figure 4) (Carracedo et al., 2019; Su et al., 2021). Indeed, chemerin binds to its G-protein coupled receptor on VSMCs. This triggers a signaling pathway that culminates in increased expression of two calcification inhibitors, namely fetuin-A and MGP (Carracedo et al., 2019; Sun et al., 2021). Both these proteins suppress osteogenic differentiation of VSMCs, and hence reduce vascular calcification (Figure 4) (Carracedo et al., 2019).

5.2 | Atherosclerosis and COVID-19

Atherosclerosis, a major cardiovascular disease, is the most common cause of worldwide mortalities. It is characterized by the formation of vascular plaques made up of fats, cholesterol, fibrin, among others, and it also involves VSMC phenotypic switching (Bennett et al., 2016; Ibrahim et al., 2023). Indeed, contractile phenotype markers like smooth muscle cell myosin heavy chain (MYH11), and smooth muscle aortic alpha-actin (ACTA2), SMC lineage-restricted protein are reduced in atherosclerosis (Bennett et al., 2016). This downregulation is concomitant with increased VSMC-derived secretion of exosomes carrying molecules, like phosphatidylserine or annexin A6, that

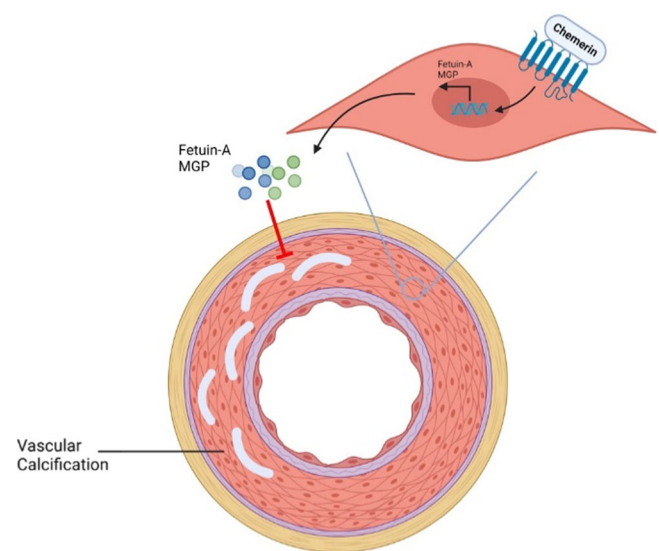


FIGURE 4 The mechanism of inhibiting calcification through chemerin's effect on vascular smooth muscle cells (VSMC). Chemerin, an adipokine found in patients with impaired renal function binds to its receptor on VSMC surface, promoting in the expression of two calcification inhibitors, the fetuin A and matrix Gla protein (MGP). Created with [BioRender.com](https://www.biorender.com).

induce calcification in the atherosclerotic plaque (Bennett et al., 2016; Grootaert & Bennett, 2021). Furthermore, production of membrane-bound apoptotic bodies in the plaque ensue (Bennett et al., 2016; Grootaert & Bennett, 2021). These bodies induce the recruitment of macrophages to this plaque, making it a nucleation site for the calcification (Figure 5) (Bennett et al., 2016; Grootaert & Bennett, 2021).

In the context of atherosclerosis, VSMCs may follow another dedifferentiation path. For instance, they may switch to an osteochondrogenic phenotype (Sanyour et al., 2020) by virtue of a notable suppression of smooth muscle 22 alpha (SM22 α) along with the upregulation of osteochondrogenic markers such as runt-related transcription factor 2 (RUNX2), osteopontin, osteocalcin, Type II and X collagen, alkaline phosphatase, and Sox9, hence precipitating calcification (Grootaert & Bennett, 2021). Another cause for this calcification in atherosclerotic patients is the deposition of calcifying vesicles, and the downregulation of mineralization inhibitory molecules such as vit K-dependent MGP and fetuin-A (Figure 5) (Durham et al., 2018; Grootaert &

Bennett, 2021). Consequently, microcalcification in the fibrous cap and macrocalcification in the necrotic core of the plaque are induced (Grootaert & Bennett, 2021). Given that vascular calcification is a player in the pathogenesis of COVID-19, patients suffering from atherosclerosis, especially ones with calcified plaques, are at higher risk of suffering from COVID-19 or acquiring its severe symptoms (Poznyak et al., 2021).

6 | CONCLUSION

The interplay between VSMCs phenotypic switch and cholesterol, calcium, or phosphate is crucial for the pathogenesis of SARS-CoV2 or its remission, especially in patients with CKD and atherosclerosis. That would be especially important, as it represents a potential for targeting viral entry inhibition, thus antiviral therapy, since currently, many studies aim at targeting pathways that mediate viral entry. Furthermore, despite the currently available therapeutic options, more studies are needed to identify molecules other than cholesterol,

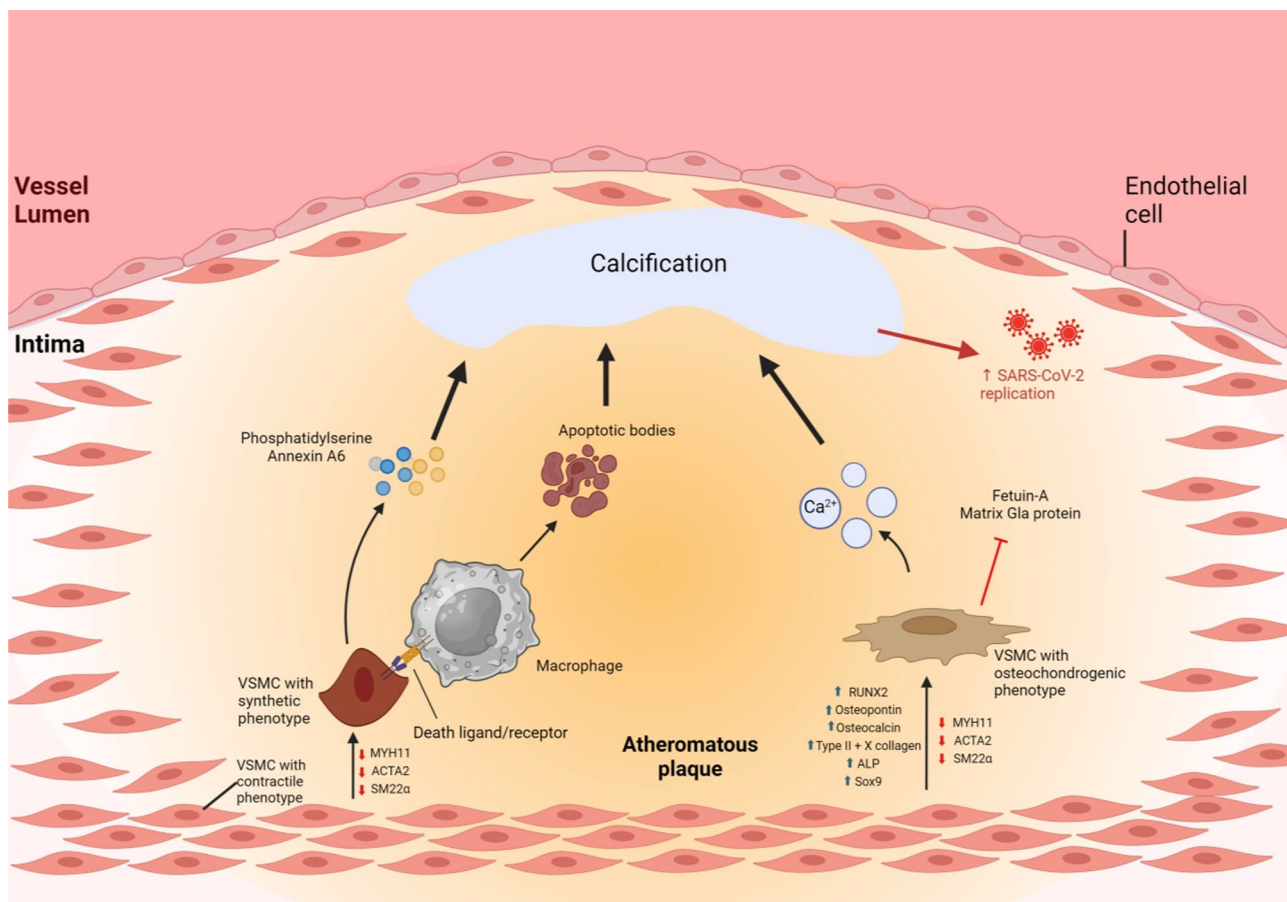


FIGURE 5 The effect of various vascular smooth muscle cell (VSMC) phenotypes on the calcification state of the atherosclerotic plaque. Some contractile VSMCs' markers, such as myosin heavy chain (MYH11), smooth muscle aortic alpha-actin (ACTA2), and smooth muscle 22 alpha (SM22 α) are downregulated in atherosclerosis; however, others like runt-related transcription factor 2 (RUNX2), osteopontin, osteocalcin, type II and X collagen, alkaline phosphatase (ALP), and SRY-box transcription factor 9 (Sox9) are upregulated. This prompts VSMCs to assume another phenotype. Synthetic VSMCs secrete PS and annexin A6, recruit macrophages to the plaque, and decrease calcification inhibitors like matrix Gla protein (MGP) and fetuin-A, therefore inducing in calcium deposits. Created with [BioRender.com](https://www.biorender.com).

calcium, and phosphate that could prove to be tractable targets in the fight against this virus (Bhimraj et al., 2024; Giordo et al., 2021; Issa et al., 2021; Kaddoura et al., 2020; Wehbe, Wehbe, et al., 2021; Younis et al., 2020, 2021; Zareef et al., 2020). This will aid in developing new drugs that can treat patients who are resistant to the current standard of care.

AUTHOR CONTRIBUTIONS

Laura Ghanem: Writing—original draft. **Dina Essayli:** Writing—original draft. **Jana Kotaich:** Writing—original draft. **Mohammad Al Zein:** Writing—original draft. **Amirhossein Sahebkar:** Writing—original draft; writing—review and editing. **Ali H. Eid:** Writing—review and editing; supervision; resources; conceptualization.

CONFLICT OF INTEREST STATEMENT

The authors declare no conflict of interest.

ORCID

Amirhossein Sahebkar  <http://orcid.org/0000-0002-8656-1444>

Ali H. Eid  <http://orcid.org/0000-0003-3004-5675>

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How to cite this article: Ghanem, L., Essayli, D., Kotaich, J., Zein, M. A., Sahebkar, A., & Eid, A. H. (2024). Phenotypic switch of vascular smooth muscle cells in COVID-19: Role of cholesterol, calcium, and phosphate. *Journal of Cellular Physiology*, e31424. <https://doi.org/10.1002/jcp.31424>