

REVIEW

CD36, a scavenger receptor implicated in atherosclerosis

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CD36 is a membrane glycoprotein that is present on various types of cells, including monocytes, macrophages, microvascular endothelial cells, adipocytes and platelets. Macrophage CD36 participates in atherosclerotic arterial lesion formation through its interaction with oxidized low-density lipoprotein (oxLDL), which triggers signaling cascades for inflammatory responses. CD36 functions in oxLDL uptake and foam cell formation, which is the initial critical stage of atherosclerosis. In addition, oxLDL via CD36 inhibits macrophage migration, which may be a macrophage-trapping mechanism in atherosclerotic lesions. The role of CD36 was examined in *in vitro* studies and *in vivo* experiments, which investigated various functions of CD36 in atherosclerosis and revealed that CD36 deficiency reduces atherosclerotic lesion formation. Platelet CD36 also promotes atherosclerotic inflammatory processes and is involved in thrombus formation after atherosclerotic plaque rupture. Because CD36 is an essential component of atherosclerosis, defining the function of CD36 and its corresponding signaling pathway may lead to a new treatment strategy for atherosclerosis.

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INTRODUCTION

CD36 is a pattern recognition receptor that is expressed on various types of cells. CD36 binds multiple ligands and mediates different biological processes. Recent studies have demonstrated that CD36 is involved in the progression of atherosclerosis and that genetic deletion of CD36 or blockage of the CD36-induced signaling cascade reduces atherosclerotic lesion formation. CD36 has significant roles in different stages of the atherogenic process and consequently can be regarded as an important molecule in the progression of atherosclerosis. This paper provides an overview of reported functions of CD36, updates on the elucidation of CD36-mediated cell signaling, and its implications in atherosclerosis.

CD36 STRUCTURE AND LIGANDS

CD36, an 88-kDa transmembrane glycoprotein receptor, is expressed on various cell types, including monocytes and macrophages; platelets; microvascular endothelial cells; adipocytes; epithelial cells in the kidney and cardiac myocytes.¹ CD36 belongs to the class B scavenger receptor family, which also includes scavenger receptor B1 and lysosomal integral membrane protein 2.² CD36 is encoded by the human *Cd36*

gene, which is located on chromosome 7 (7q11.2) and consists of 15 exons. The CD36 protein is composed of a single peptide chain of 472 amino acids and is organized into two transmembrane domains: two very short cytoplasmic domains and a large glycosylated extracellular domain (Figure 1). The extensive glycosylation of CD36 is required for intracellular trafficking onto the cell membrane.³

CD36 binds many different ligands including thrombospondin-1, oxidized phospholipids (oxPL), oxidized low-density lipoprotein (oxLDL), hexarelin, fibrillar A β amyloid peptides and long-chain fatty acids. CD36 also binds *Plasmodium falciparum*-infected erythrocytes, bacterial cell wall components of *Staphylococcus* and *Mycobacterium*, cell-derived microparticles and apoptotic cells. The binding site for each ligand is different; thrombospondin-1 binds to the CLESH-1 domain (CD36 LIMP II Emp Structural Homology-1) that resides in amino acids 93–155. The binding site for oxPL is located in amino acids 157–171, and the binding site for oxLDL is nearby in the 155–183 sequence.⁴ Thrombospondin-1 binding to endothelial CD36 inhibits angiogenesis by inducing apoptosis.⁵ As a fatty acid translocase, CD36 binds long-chain free fatty acids and facilitates their

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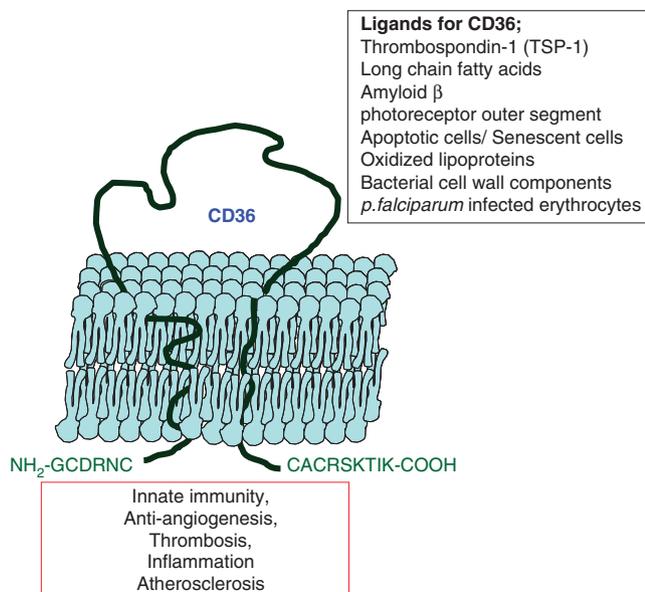


Figure 1 CD36 binds various ligands and functions in many biological processes. CD36 is a transmembrane receptor composed of two transmembrane domains, two very short cytoplasmic domains and a large glycosylated extracellular domain. CD36 is expressed in a variety of cell types, such as monocytes/macrophages, microvascular endothelial cells, adipocytes and microglia and platelets, and binds various ligands. CD36 is implicated in many biological processes including innate immunity, anti-angiogenesis, thrombosis, inflammation and atherosclerosis.

transport into cells. This function of CD36 provides an energy source for beta-oxidation to myocytes and lipid storage to adipocytes.⁶

oxPL are known to play an important role in atherosclerosis and are known to be ligands for CD36. Phospholipids are essential components of lipoproteins and cell membranes, and are composed of fatty acids bound to a glycerol backbone with a polar head group. Phospholipids, particularly those containing unsaturated fatty acids, are susceptible to free-radical or enzymatic oxidation by reactive oxygen and nitrogen species generated by myeloperoxidase, lipoxygenase and other enzymes present in the vessel wall.⁷ Modification of phospholipids produces conformational changes, including oxidized fatty acids protrusion from the hydrophobic membrane or lipoprotein interior into the more polar aqueous compartment. Thus, modified phospholipids gain access and are able to interact with the pattern recognition receptor CD36.⁸ oxPL are accumulated under conditions of oxidative stress, such as infection and inflammation, and are also generated by necrotic and apoptotic cells.⁹ Extensive studies have revealed the chemical and structural characteristics of oxidized, modified lipids on the surface of cell membranes and/or lipoprotein particles that serve as ligands for CD36. Early studies revealed that the lipid portions of oxidized lipoproteins and apoptotic cells retain a significant portion of CD36-binding activity and that oxidized phosphatidylcholine is the primary ligand.^{10–12}

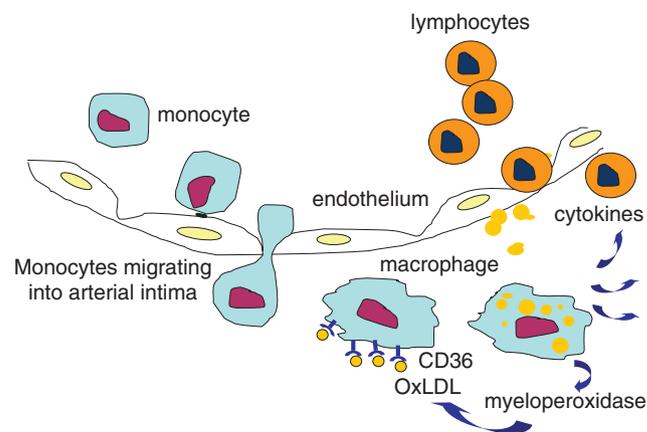


Figure 2 Macrophage trapping as a mechanism of atherosclerosis. The atherosclerotic process is initiated by monocyte entry into the arterial intima, followed by differentiation of these cells into macrophages. Macrophages internalize oxidized LDL (oxLDL) through scavenger receptors such as CD36 and are trapped in the arterial intima. They promote further inflammatory responses by secreting cytokines, which recruit other immune cells to the lesion. Progression of the atherosclerotic inflammation induces arterial lumen narrowing.

Podrez *et al.*¹⁰ reported that the structural characteristics for binding CD36 require phospholipids with an sn-2 acyl group that incorporates a terminal γ -hydroxy (or oxo)- α , β -unsaturated carbonyl (oxidized phosphatidylcholine_{CD36}). In addition, oxPL with reactive groups covalently bind to proteins and form lipid–protein adducts. Both the free oxPL and the adducted forms are recognized by CD36.¹³

MACROPHAGE CD36 MEDIATES FOAM CELL FORMATION AND PROMOTES ATHEROSCLEROSIS

Foam cell formation is the initial critical step of atherosclerosis. The atherogenic process starts with the transmigration of blood-circulating monocytes into the arterial intima, where they differentiate into macrophages.¹⁴ Macrophages bind and internalize oxLDL through CD36.¹⁵ The internalized oxLDL provides its specific oxidized lipids as ligands for the nuclear hormone receptor PPAR- γ and upregulates the expression of CD36, which is called an ‘eat me signal.’ This facilitates further uptake of oxLDL.^{16,17} Activated macrophages secrete oxidants, including myeloperoxidase, which oxidizes LDL and thus, enlarges the pool of oxLDL.¹⁸ The interaction between CD36 and oxLDL also induces the secretion of cytokines that recruit immune cell infiltrates in the arterial intima,¹⁹ and the arterial inflammation provoked by foam cells induces arterial narrowing, establishing atherosclerotic vascular diseases (Figure 2).

Macrophages harvested from *Cd36*-null mice are defective in oxLDL uptake, and *Cd36*-null mice with atherosclerosis-prone background, including *ApoE* null or *LDL receptor*-null genotypes, showed less atherosclerotic lesion formation than *ApoE* null or *LDL receptor*-null mice on a high-fat diet.^{20–22} A bone marrow transplantation study also revealed that the

atherogenic mechanism is dependent on macrophage CD36. Mice that received *Cd36*-null macrophages had profoundly less atherosclerotic lesion formation, and re-introduction of macrophages with CD36 induced a twofold increase in the atherosclerotic lesion area.²³ In addition, treatment with a competitive peptide ligand (EP80317), derived from the growth hormone-releasing peptide family that blocks the oxLDL-binding site of CD36, also reduced atherosclerotic lesions by 51% in *ApoE*-null mice.²⁴ However, another study with a different *Cd36*-null mouse strain showed that *ApoE/Cd36* double-null mice that were fed a high-fat diet had a modest reduction or even an increase in some atherosclerotic lesions compared with *ApoE*-null mice. Moore *et al.*²⁵ also reported that a reduced lipid accumulation in peritoneal macrophages was coupled to significantly higher plasma cholesterol levels in male but not female *ApoE/Cd36* double-null mice in comparison with *ApoE*-null mice. These contradictory observations were originally highly controversial. As suggested by Witztum and Collot-Teixeira *et al.*,^{26,27} the differences may have been attributable to the use of two different mouse strains. The apparent controversy about the role of CD36 in murine atherosclerosis has been tempered by recent studies from Guy *et al.*,²¹ which showed significant atheroprotection in an additional *Cd36*-null strain crossed with the *ApoE*-null strain and in a different atherosclerosis model, the *LDL receptor*-deficient strain.²¹ Furthermore, two recent papers from Moore *et al.*²⁵ reported significant atheroprotection (approaching levels observed in Guy *et al.*²¹ reports) in the *Cd36/ApoE*-null strain developed in their lab.³⁰

The addressed studies, therefore, confirm that macrophage CD36 promotes atherosclerosis through the interaction with oxLDL.

CD36: A SIGNALING MOLECULE THAT COMPRISES MULTIMOLECULAR COMPLEXES

Many studies have revealed that CD36 transduces signals, although it lacks known intracellular signaling domains, such as kinase, phosphatase, g-protein binding or scaffolding domains. CD36 is known to physically associate with src family non-receptor tyrosine kinases, such as *fyn*, *lyn* and *yes*.^{31–33} In macrophages, the interaction between CD36 and oxLDL induces the phosphorylation of *lyn* and the subsequent activation of the mitogen-activated kinases Jun-kinase (JNK) 1 and 2. JNK activation mediates the uptake of oxLDL; the treatment of oxLDL-loaded macrophages with a JNK inhibitor was associated with a reduced oxLDL uptake. Atherosclerotic arterial lesions from mice had significantly increased levels of activated JNK compared with arterial tissue from non-atherosclerotic mice.³⁴ Recently, a guanine nucleotide exchange factor, *Vav*, was revealed to be in the CD36-mediated signaling cascade. *Vav* is activated by src family kinases and mediates the activation of the small-molecular weight G proteins *Rac* and *Rho*. In addition, *Vav* is a scaffold protein for various signaling molecules including phospholipase C and dynamin. A recent paper showed that oxLDL induces *Vav* activation in macrophages, and *Vav*-null macrophages have reduced oxLDL uptake. Inhibition of dynamin, a *Vav*-interacting protein that functions in endocytic vesicle fission, blocked the uptake of oxLDL (Figure 3).³⁵ Consequently, *ApoE/vav1* double-null mice showed reduced atherosclerotic lesion formation compared with *ApoE*-null mice.³⁶

IRGM1, another member of the small GTPase family that is highly expressed on macrophages, also mediates the internalization of CD36 and oxLDL. Macrophages from IRGM1-null mice exhibited impaired CD36 internalization and oxLDL uptake, and IRGM1-null mice had reduced atherosclerotic

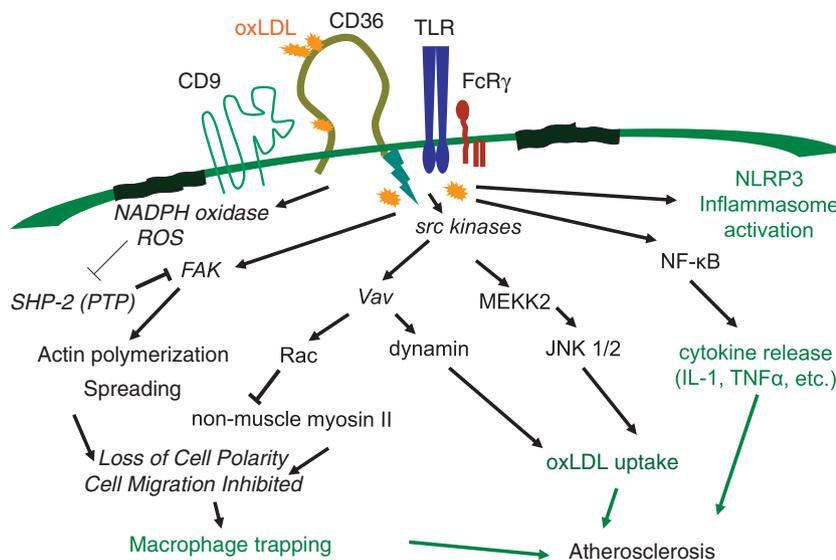


Figure 3 CD36-mediated signaling promotes atherosclerosis. The interaction between CD36 and oxidized LDL (oxLDL) provokes signaling cascades that mediate the uptake of oxLDL, cytokine production and macrophage trapping, thus promoting the atherosclerotic inflammatory process.

lesion formation.³⁷ OxLDL via CD36 induces the activation of other mitogen-activated kinases such as p38 mitogen-activated kinase and ERK 1/2.³⁸ Signaling mediated by CD36 activates nuclear factor kappa beta. CD36-deficient macrophages have reduced nuclear factor kappa beta activation and reduced the release of interleukin (IL)-1 β and tumor necrosis factor- α .³⁹

CD36 has diverse downstream signaling pathways and functions by interacting with other membrane receptors such as integrins, toll-like receptors (TLRs) and tetraspanins, including CD9.^{40,41} Macrophage CD36 responds to lipoteichoic acid on the membrane of *Staphylococcus aureus* and diacylated lipoproteins, depending on interactions with TLR2/6.^{42,43} The CD36-mediated uptake of oxLDL, the CD36-mediated phagocytosis of *P. falciparum* malaria-parasitized erythrocytes and the endocytosis of CD36 are all independent of TLR2, while the activation of TLR2 enhances CD36-mediated uptake.^{34,44} A recent study revealed that CD36 induces the assembly of the TLR4/6 heterodimer, which is responsible for the responses to oxLDL and amyloid- β , including inflammatory gene expression, IL-1 β release and nuclear factor kappa beta activation.³⁰ Sheedy *et al.*⁴⁵ showed that CD36, via the binding and uptake of endogenous soluble ligands, including oxLDL, amyloid- β peptide and amylin peptide, delivers two signals to promote the assembly of the cytoplasmic protein complex NLRP3 inflammasome, during sterile inflammation (as in atherosclerosis and Alzheimer's disease). The signals for NLRP3 inflammasome activation include 'signal 1,' which upregulates expression of IL-1 family members and some Nod-like receptors, and 'signal 2,' which leads to inflammasome assembly and secretion of IL-1 family members.⁴⁶ CD36 directs soluble cargo, including oxLDL and soluble amyloid- β , to the lysosomal compartment where it is crystalized. The crystals destabilize the lysosomal membranes and induce inflammasome activation.⁴⁵ In accordance, CD36-deficient *ApoE*-null mice have less cholesterol crystal accumulation in atherosclerotic plaque and lower serum concentrations of IL-1 compared with *ApoE*-null mice.⁴⁵ These studies reveal a mechanism by which CD36 promotes sterile inflammation.

The interaction between platelet-activating factor receptor and CD36 also mediates oxLDL uptake and IL-10 production.⁴⁷ Rios *et al.*⁴⁷ showed that oxLDL promotes colocalization of platelet-activating factor receptor and CD36 in lipid rafts, and this colocalization is required for oxLDL uptake and oxLDL-induced IL-10 production.

A recent paper verified a mechanism by which the CD36-containing multimolecular signaling complex enables internalization of CD36 and its ligands. Heit *et al.*⁴⁸ demonstrated that CD36 resides in heterogeneous membrane receptor complexes that contain integrin β 1/ β 2 and the tetraspanins CD9 and CD81, and that this receptor complex links CD36 to the immunoreceptor tyrosine activation motif-bearing adaptor Fc γ (FcER1G). Coupling to an immunoreceptor tyrosine activation motif adaptor allows CD36 to engage Src and Syk kinases and triggers internalization of CD36 and its bound ligands.

Therefore, macrophage CD36 coordinates with many other signaling partners and functions as a signaling molecule that promotes atherosclerosis by mediating foam cell formation and the release of inflammatory mediators.

CD36-MEDIATED CYTOSKELETAL MODULATION INDUCES MACROPHAGE TRAPPING IN ATHEROSCLEROTIC PLAQUE

Characteristically, atherosclerotic inflammation does not spontaneously resolve, which leads to irreversible arterial remodeling. Macrophage activation is a typical attribute of atherogenic inflammation. In contrast with acute inflammation, in which inflammatory responses are resolved by the emigration of infiltrated immune cells to draining lymph nodes, during atherogenic inflammation, macrophages are trapped in atherosclerotic lesions and continue propagating the inflammatory response without resolution. However, previous studies revealed that, under certain conditions, atherosclerotic lesions regress concurrently with macrophage emigration.^{49–51} Llodra *et al.*⁵¹ induced regression of atherosclerotic plaque by transplanting arterial segments with atherosclerotic plaque from hypercholesterolemic mice into normal mice. This macrophage-trapping mechanism suggests novel approaches to the re-mobilization of lipid-laden macrophages from atherosclerotic lesions to facilitate disease regression and treatment. My work in Silverstein's laboratory reproduced findings reported nearly 26 years ago by Quinn *et al.*⁵² that oxLDL inhibits macrophage migration. We attempted to define the mechanism and show that CD36 mediates oxLDL-induced inhibition of macrophage migration. *In vitro* and *in vivo* migration assays indicated that oxLDL inhibited wild-type macrophage migration but not *Cd36*-null macrophages. OxLDL facilitates cell spreading by activating focal adhesion kinase through src-kinases. However, the inactivation of src homology 2-containing phosphotyrosine phosphatase (SHP-2) leads to sustained focal adhesion kinase activation, which results in the dysregulation of cytoskeletal assembly and disassembly and the loss of macrophage mobility. Inactivation of SHP-2 occurs through reactive oxygen species-mediated oxidative modification of the enzyme. Therefore, the blockage of the CD36-mediated signaling cascade by antioxidants and nicotinamide adenine dinucleotide phosphate oxidase inhibitors blocks the effect of oxLDL and restores macrophage migration in the presence of oxLDL.⁵³ Loss of macrophage cell polarity is another mechanism of oxLDL macrophage trapping. Our recent study revealed that oxLDL, through CD36, induces front-end lamellipodia retraction and the loss of cell polarity. The mechanism is caused by the oxLDL-induced activation of the Vav/Rac pathway and the subsequent inactivation of non-muscle myosin II. Thus, Vav deletion or Rac inhibition blocks the effect of oxLDL on macrophage cell polarity and migration (Figure 3).⁵⁴ These studies identified and suggested molecules in CD36-mediated signaling as new targets for reversing atherosclerosis.

CD36 ACTIVATES PLATELETS, WHICH EXERT PRO-ATHEROGENIC ACTIONS

Platelets promote thrombus formation after atherosclerotic plaque rupture, and are also involved in the initiation and development of the inflammatory process of atherosclerosis.⁵⁵

Podrez *et al.*⁵⁶ showed that oxLDL binding to CD36 activates platelets, inducing the expression of P-selectin and the activation of integrin $\alpha_{IIb}\beta_3$. Recent studies from Silverstein's lab revealed that oxLDL binding to CD36 provokes platelet signaling and drives the platelets to become hyperactive.^{57,58} Platelets cultured with oxLDL were more sensitive to a low dose of ADP, the classic platelet activator. This process was mediated by the CD36-induced activation of JNK and Vav.^{57,58} A phosphoproteomic analysis of platelets revealed additional signaling pathways that mediate oxidized phospholipid/CD36-induced platelet activation, including Src family kinases, Syk, and phospholipase C- γ .⁵⁹ These data support the previous observation by Eitzman *et al.*⁶⁰ that high-fat diet-fed *ApoE*-null mice had faster thrombi formation at the site of vascular injury. Accordingly, *Cd36*-null mice with an *ApoE*-null background showed slower thrombotic response than *ApoE*-null mice when challenged with vascular injury.⁵⁷ Similarly, oxPL on the endothelial cell-derived microparticles also bind to platelet CD36 and promote thrombosis in mice with ferric chloride (FeCl_3)-induced vascular injury.⁶¹ Advanced glycation end product, which is abundant in the blood of diabetic patients, also functions as a CD36 ligand and augments the response of platelets to ADP, promoting faster thrombosis in an animal model.⁶²

Recent studies suggested that oxLDL, bound by platelets via scavenger receptors, exerts additional pro-atherogenic effects. Platelets interact with oxLDL and release atherosclerosis-promoting chemokines.⁶³ Furthermore, platelets internalize oxLDL, and these lipid-laden platelets activate the endothelium and inhibit endothelial regeneration.⁶⁴ Platelets also mediate the conversion of monocytes into foam cells. Curtiss *et al.*⁶⁵ showed that platelets facilitate cholesteryl ester accumulation in peripheral blood mononuclear cell-derived macrophages. A recent study by Badrnya *et al.*⁶⁶ reinforces this observation by elucidating the role of CD36 in this process. In their study, they found that platelets form aggregates with monocytes in response to oxLDL, and this aggregate facilitates oxLDL uptake by monocytes in a CD36-dependent manner. Platelets activated by oxLDL promote monocyte extravasation in the thioglycollate-elicited peritonitis model. In addition, an *in vitro* experiment using transwell-migration assays also showed that oxLDL/CD36 platelet interactions enhanced monocyte migration across a human umbilical vein endothelial cell monolayer.⁶⁶

GENETIC VARIATIONS OF CD36

The role of CD36 has been extensively studied since the generation of mice with CD36 genetic deletion. The genetic deletion of CD36 led to a reduction in the generation of atherosclerotic plaque in mice. In humans, several *Cd36* gene mutations have been reported to result in a diminished

expression of CD36 protein.^{67–70} The two mutation types are categorized into type I and type II CD36 deficiencies. Type I deficiency is characterized by the lack of expression of CD36 in many cell types, including monocytes, macrophages and platelets. In type II deficiency, CD36 is deficient only in platelets.^{68–75} However, the physiological consequences of the CD36 deficiency have not been clearly verified. Patients with type II CD36 deficiency have normal platelet functions,⁷⁶ while the incidence of cardiomyopathy, hyperlipidemia and insulin resistance are relatively high in these patients.^{77–80} It is not clear if the effects of human CD36 deficiency are caused by the lack of CD36 in certain cell types or if there are aberrant functions of mutant CD36 or any compensatory effects. Therefore, additional studies are warranted to clarify the effects of CD36 deficiency in humans.

SOLUBLE CD36: A MARKER FOR METABOLIC SYNDROME INCLUDING ATHEROSCLEROSIS

Non-cell bound CD36 has been isolated from human plasma and is referred to as soluble CD36 (sCD36). The level of sCD36 represents the expression level of CD36 in various cell types and tissues in human and rodent models of insulin resistance and type 2 diabetes. The concentration of sCD36 is approximately fivefold higher in plasma of obese diabetic subjects than in lean healthy subjects.⁸¹ In concordance with the role of CD36 in atherogenic inflammation, the sCD36 level was correlated with cardiovascular mortality in a cohort of chronic renal disease patients⁸² and with intima-media thickness measured by ultrasound.⁸³ HMG-CoA reductase inhibitor (statin) treatment reduced the serum concentration of sCD36.⁸² However, whether sCD36 is composed of full-length CD36 or a proteolytic fragment of the CD36 extracellular domain has not been determined. Therefore, additional studies that investigate the structure of sCD36, the mechanism of sCD36 release, and the role of sCD36 are needed.

CONCLUSION

As a signaling molecule, CD36 plays significant roles in atherogenic processes, including foam cell formation, release of inflammatory mediators, macrophage trapping and thrombosis. Therefore, verification of the CD36-mediated signaling pathway may suggest novel therapeutic targets and a new strategy for the treatment of atherosclerosis. Recent studies of multimolecular complex formation, induced by CD36, revealed details about CD36 signal regulation and the manner in which CD36 drives various functions. Because CD36 binds a variety of ligands and functions in many different biological processes, a ligand-specific signaling pathway for CD36 should be examined. The atherogenic function of CD36 has been demonstrated in animal studies and *in vitro* experiments. However, only a few studies have examined CD36 in humans. Moreover, previous reports about type II CD36 deficiency in humans contradicted observations from animal studies. Therefore, additional studies that focus on the function of CD36 in humans are warranted to target this molecule for the treatment of atherosclerosis.

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