



Role of Serum Endothelin-Converting Enzyme-1 (ECE-1), Adenosine Monophosphate-Activated Protein Kinase (AMPK), and Oxidized Low-Density Lipoprotein (Ox-LDL) in the Pathophysiology and Theoretical Diagnostic Framework of Type 2 Diabetic Mellitus Patients with Acute Myocardial Infarction: A Narrative Review (Article Review)

Rana Salah Noori Al-Saeagh^{1*}

¹College of Science for Women, ,Universiry of Babylon, sci184.rana.salah@uobabylon.edu.iq, Hilla, Babil, Iraq.

*Corresponding author email: sci184.rana.salah@uobabylon.edu.iq; mobile: 07725734988

دور كل من إنزيم محول الإندوثيلين-1 المصلي (ECE-1)، وبروتين كيناز المنشط بـ AMP (AMPK)، والبروتين الدهني منخفض الكثافة المؤكسد (Ox-LDL) في اطار تشخيصي نظري لمرضى السكري من النوع الثاني المصابين بالاحتشاء القلبي الحاد:مراجعة سردية (مراجعة مقالية)

رنا صلاح نوري الصائغ^{1*}

كلية العلوم للبنات، جامعة بابل، sci184.rana.salah@uobabylon.edu.iq ، ، الحلة، العراق

Accepted: 16/6/2026

Published: 30/6/2026

ABSTRACT

Background: Type 2 diabetes mellitus (T2DM) significantly increases the risk of acute myocardial infarction (AMI) through mechanisms including vascular dysfunction, oxidative stress, and inflammation. Conventional diagnostic tools, such as electrocardiography and troponins, often exhibit limited efficacy in patients with T2DM due to atypical symptom presentation.

Materials and Methods: This narrative review synthesizes published evidence from basic, preclinical, and clinical studies on the structure, functions, and interplay of a panel of serum biomarkers — endothelin-converting enzyme-1 (ECE-1), adenosine monophosphate-activated protein kinase (AMPK), and oxidized low-density lipoprotein (ox-LDL) — in the pathophysiology of AMI in the context of T2DM (T2DM-AMI). This narrative review synthesizes published evidence from basic, preclinical, and clinical studies on the structure, functions, and interplay of a panel of serum biomarkers — ECE-1, AMPK, and ox-LDL — in the pathophysiology of AMI in the context of T2DM (T2DM-AMI). No original patient data, clinical cohorts, or laboratory measurements are presented. The diagnostic potential discussed is derived from mechanistic and correlational studies and remains theoretical.

Results: ECE-1 promotes vasoconstriction via endothelin-1 (ET-1), AMPK serves as a master regulator of cellular energy balance, and ox-LDL is a key driver of atherosclerosis. Their interactions highlight a vicious cycle of metabolic and vascular damage. Based on published mechanistic and correlational evidence, combining these biomarkers may theoretically improve early diagnosis and personalized treatment; however, this remains hypothetical pending clinical validation in prospective cohorts containing original patient data. No quantitative diagnostic metrics (sensitivity, specificity, AUC-ROC, or cutoff thresholds) for the proposed biomarker panel are available from the current literature. Similarly, no quantitative comparison with existing diagnostic standards (high-sensitivity troponin or electrocardiography) is provided, as such validation studies have not yet been conducted.

Conclusion: The panel of ECE-1, AMPK, and ox-LDL shows theoretical diagnostic potential for T2DM-AMI based on existing mechanistic studies. However, because this is a narrative review without original patient data, clinical cohorts, or laboratory measurements, clinical validation in prospective studies is required before any diagnostic application can be recommended. However, without established diagnostic criteria, quantitative metrics (sensitivity, specificity, AUC-ROC, cutoff values), or direct quantitative comparison with current standards (hs-troponin, ECG), the term "diagnosis" remains aspirational. Clinical validation studies reporting these quantitative measures are urgently needed.

Keywords: Type 2 Diabetic Mellitus, Serum Endothelin, AMPK, ECE-1, Myocardial Infarction, Narrative review, Hypothesis-generating, Theoretical framework



INTRODUCTION

Cardiovascular disease (CVD) remains the leading cause of morbidity and mortality worldwide, and type 2 diabetes mellitus (T2DM) is a major contributor to its escalating prevalence. T2DM is characterized by chronic hyperglycemia, insulin resistance, and relative insulin deficiency. These metabolic derangements accelerate atherosclerosis and dramatically increase the risk of acute myocardial infarction (AMI). Patients with T2DM are more likely than non-diabetic individuals to suffer an AMI [1]. The atypical presentation of AMI in this cohort, often characterized by silent or unrecognized symptoms, frequently leads to delayed diagnosis and, consequently, more severe complications and poorer outcomes [1-3]. Given the heightened cardiovascular risk in this population, novel diagnostic strategies are urgently needed.

Currently, the diagnosis of AMI relies on clinical symptoms, electrocardiogram (ECG) findings, and elevated cardiac troponins. However, the diagnostic accuracy of these methods can be compromised in T2DM patients, due to atypical presentations and confounding comorbidities. For instance, diabetic neuropathy can mask typical anginal pain, while pre-existing ECG abnormalities and chronic kidney disease can complicate the interpretation of ECG and troponin results, respectively [4]. Furthermore, these traditional markers primarily indicate myocardial necrosis rather than the underlying pathophysiological processes. T2DM accelerates arterial stiffening and endothelial dysfunction through a confluence of metabolic stress, oxidative stress, and chronic inflammation. Therefore, biomarkers that reflect these upstream processes might theoretically enable earlier diagnosis and guide targeted therapies, but this remains hypothetical and requires validation in diagnostic accuracy studies [5].

This review examines ECE-1, AMPK and Ox-LDL as potential biomarkers for AMI in T2DM. ECE-1 catalyzes the production of endothelin-1 (ET-1), a potent vasoconstrictor that contributes to vascular fibrosis [5]. AMPK maintains energy balance and mitigates inflammation. Ox-LDL promotes endothelial damage and plaque formation. Their interplay—ox-LDL upregulating ECE-1 and down regulating AMPK—exacerbates T2DM-AMI pathology [6]. **The aim of the present review** is to elucidate their roles and interrelationships based on mechanistic evidence, and to explore their theoretical diagnostic utility, acknowledging that no diagnostic validation has yet been performed [6]. It is important to clarify that this narrative review does not provide quantitative diagnostic metrics (e.g., sensitivity, specificity, AUC-ROC, or cutoff thresholds) for the proposed biomarkers, nor does it offer a direct quantitative comparison with existing diagnostic standards such as high-sensitivity troponin or electrocardiography. Such data are not yet available in the published literature. Therefore, the use of the term "diagnosis" in the title reflects the theoretical potential of these biomarkers based on pathophysiological mechanisms, not an established clinical diagnostic utility.

MATERIALS AND METHODS:

This narrative review synthesizes published evidence from basic, preclinical, and clinical studies on the structure, functions, and interplay of a panel of serum biomarkers — endothelin-converting enzyme-1 (ECE-1), adenosine monophosphate-activated protein kinase (AMPK), and oxidized low-density lipoprotein (ox-LDL) — in the pathophysiology of AMI in the context of



RESULTS

1. Endothelin Converting Enzyme -1 (ECE-1)

1.1 Structure of ECE-1

Endothelin-1 (ET-1) is a 21-amino-acid peptide essential for vascular homeostasis. Despite a short plasma half-life (1.5–4 minutes), ET-1 exerts prolonged effects due to strong receptor binding and downstream signaling. Its clearance is mediated by receptor- B (ETB) uptake, enzymatic degradation, and hepatic clearance. Among the endothelin peptides, ET-1 is the most potent isoform and a key mediator of vasoconstriction in T2DM and AMI. The biological effects of ET-1 are mediated by two receptor subtypes: ETA and ETB. ETA receptors, located on vascular smooth muscle cells, mediate vasoconstriction and cell proliferation. In contrast, ETB receptors on endothelial cells promote vasodilation through the release of nitric oxide (NO) and prostacyclin, while also facilitating ET-1 clearance. The balanced regulation of these receptors is essential for vascular health; an imbalance can precipitate disease [7].

ECE-1 is a zinc-activated metalloproteinase that converts the 38-amino-acid precursor, Big ET-1, into the active 21-amino-acid ET-1. There are four isoforms of ECE-1 (ECE-1a, ECE-1b, ECE-1c and ECE-1d), each with distinct subcellular localizations and functions. Notably, ECE-1c is responsible for significant ET-1 production in vascular tissues, linking it closely to T2DM and AMI. Alterations in ECE-1 activity can lead to endothelial dysfunction and exacerbate inflammation, processes that are central to these pathologies [8]. The isoforms feature zinc-containing domains that facilitate ET-1 creation at the plasma membrane, within intracellular vesicles, and in the Golgi apparatus. Pathophysiological states arise when the ECE-1/ET-1 axis is dysregulated in T2DM and AMI [9]. In T2DM, chronic hyperglycemia, oxidative stress, and inflammation elevate ECE-1 levels, promoting ET-1 release. This results in endothelial dysfunction, increased vascular stiffness, accelerated atherosclerosis, and diabetic microvascular complications. Similarly, in AMI, high levels of ET-1 worsen ischemia-reperfusion injury, promote adverse cardiac remodeling, and destabilize atherosclerotic plaques, contributing to heart failure [9].

The elevated action of ECE-1, specifically ECE-1a and ECE-1c at the endothelial membrane, is significant in increasing ET-1 production under these conditions. Thus, based on mechanistic evidence, ECE-1 has been suggested as a potential biomarker and a target for vascular and metabolic diseases. However, diagnostic accuracy studies have not yet been conducted [10]. The isoforms ECE-1a and ECE-1c are typically found at the cell surface and in early endosomes, enabling rapid, cell-surface production of ET-1 [11]. In contrast, ECE-1b resides in the trans-Golgi network, processing intracellular big ET-1 for secretion. ECE-1d is primarily located in early endosomes, where it is involved in intracellular signaling and the degradation of other neuropeptides. This differential localization allows for precise regulation of ET-1, which is critical for both normal physiology and disease states [12].

1.2. Physiological Functions of ECE-1

ECE-1 a protein of (753–770 amino acids) cleaves big ET-1 to generate active ET-1, thereby regulating vascular tone, proliferation, fibrosis, and inflammation [13]. In a healthy state,



ET-1 vasoconstrictive effects are balanced with NO. However, oxidative stress disrupts this equilibrium , favoring ET-1 dominance [12]. ETA receptor activation leads to vasoconstriction, while ETB receptors are primarily responsible for ET-1 clearance, mainly in the lungs [13].

1.3. Role of ECE-1 in the Pathophysiology of Type 2 Diabetes

The pathophysiological changes and vascular complications in T2DM are strongly associated with ECE-1, which catalyzes ET-1 production [14]. Diabetes causes endothelial dysfunction earlier than other complications which increases the risks for cardiovascular problems and microvascular problems such as retinopathy, nephropathy and neuropathy. The ECE-1-mediated increase in ET-1 is a major contributor to generalized endothelial dysfunction [15].

Chronic hyperglycemia, is a major driver of increased ET-1 production. The heightened activity of ECE-1 in high-glucose conditions is partly mediated by the activation of the protein kinase C (PKC)-delta isoform. Evidence indicates that dysglycemia leads to the activation of the endothelin system [15]. Researchers suggest that inhibiting this PKC isoform could prevent the hyperglycemia-associated rise in ET-1, highlighting ECE-1's crucial role in diabetic vascular damage. Consequently, ECE-1 is viewed not only as a marker of diabetes but also as a key enzyme in the development of its vascular complications. Monitoring this pathway's activity could help clinicians assess disease severity in diabetic patients [16].

Hyperglycemia in diabetic people causes oxidative stress, making NO less available and leading ET-1 to rise which stops the normal relaxation of blood vessels. As T2DM progresses, the decline in NO bioavailability, coupled with sustained ET system activation, promotes adverse vascular remodeling and growth of atherosclerotic plaques in the vessel walls [17]. The ability of ET-1 to induce vascular proliferation, fibrosis, and inflammation contributes significantly to many diabetic vascular problems. Though a direct action of ECE-1 on insulin resistance is lacking in the literature, it is clear that the role of ECE-1 in T2DM links it to problems that promote insulin resistance, including higher levels of free fatty acids and dysregulated production of adipokines. Therefore, the activity of ECE-1 in the body follows the metabolic abnormalities present in T2DM.[18-19]. All figures (1-4) in this review are conceptual/schematic diagrams created by the author for illustrative purposes. They are not derived from or adapted from any specific published figure unless explicitly stated. The mechanistic data underlying these diagrams are supported by the cited references[20].

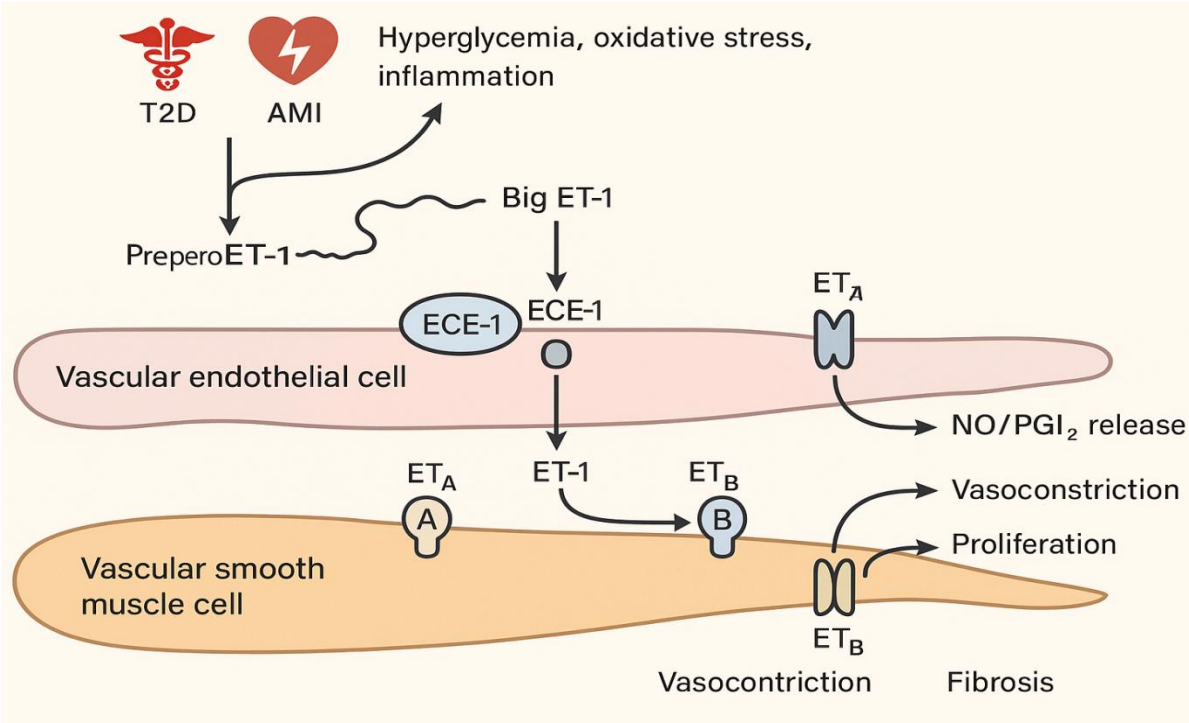


Figure 1: "Mechanistic Role of Endothelin-Converting Enzyme-1 (ECE-1) in Linking Type 2 Diabetes Mellitus and Acute Myocardial Infarction through Endothelin-1 Signaling" [18].

Figure 1 illustrates how hyperglycemia, oxidative stress, and inflammation in T2DM and AMI lead to increased production of preproET-1. This is converted to Big ET-1 and then to active ET-1 by ECE-1 on the vascular endothelium. ET-1 acts on ET_A and ET_B receptors on vascular smooth muscle cells to cause vasoconstriction, proliferation, and fibrosis, contributing to hypertension and increased cardiovascular risk.

2. Adenosine Monophosphate Activated Protein Kinase (AMPK)

AMPK is a highly conserved serine/threonine kinase that acts as a cellular energy sensor. It is activated by low energy states (e.g., elevated AMP:ATP or ADP:ATP ratios), triggering pathways to restore homeostasis. In addition to glucose uptake, fatty acid oxidation, protein synthesis, and mitochondrial function, AMPK plays a key role in metabolism, regulating glucose uptake, fatty acid oxidation, protein synthesis, and mitochondrial function.[20].

2.1. Structural Characteristics, Activation Mechanisms, and Physiological Roles in Metabolic Regulation

AMPK exists as a heterotrimeric complex composed of a catalytic α subunit and regulatory β and γ subunits. Gene diversity (α_1 , α_2 ; β_1 , β_2 ; γ_1 , γ_2 , γ_3) allows for twelve possible isoform combinations, leading to tissue-specific expression and function [20]. Complexes containing the α_2 subunit are predominantly found in tissues with high energy demand, such as skeletal muscle, heart, and liver. In contrast, α_1 -containing complexes are ubiquitously



expressed. This isoform diversity enables AMPK to fine-tune its response to metabolic stress across different tissues [21].

Most AMPK protein is present in the cytoplasm and at the plasma membrane, where it interacts with important energy-related molecules; however, some isoforms may be found elsewhere in the cell. If energy is limited (for example, there is high AMP/ADP or more free intracellular calcium), the binding of AMP/ADP to the γ -subunit activates Thr172 phosphorylation which raises AMPK activity. Further rules in the cell come from post-translational modifications, among them are ubiquitination and acetylation which control AMPK's functions inside the cell.[21].

Cells use and store energy in the body by using AMPK to increase metabolic reactions that produce ATP and inhibit those that use it. AMPK promotes glucose uptake in muscle and adipose tissue. At the same time, it helps cells use blood sugar, making the body respond better to insulin [22]. When AMPK is present, it stops the production of new fats by turning off the rate-limiting enzyme ACC and decreasing malonyl-CoA. It accomplishes this by boosting the work of the CPT1 which transports fats into the mitochondria. Protein production is stopped in the body when AMPK turns off mTOR, allowing energy conservation. The AMPK receptor improves mitochondrial function and helps create new mitochondria, increasing energy production capacity [23]. On top of that, it helps activate autophagy, whereby the body clears away damaged parts and proteins to keep cells healthy under challenging metabolic conditions. AMPK integrating energy-sensing functions with metabolic and homeostatic pathways indicates it is essential for energy balance and thus makes it an important target for therapeutics used in diabetes, obesity and cardiovascular diseases.

AMPK is responsible for keeping the body's metabolism stable, serving as a kind of fuel gauge that checks AMP-ATP and ADP-ATP amounts. It decreases use of ATP in making fats and proteins, yet activates ATP-producing pathways such as fatty acid oxidation and glycolysis. Problems with this process are linked to the typical metabolic disturbances seen in type 2 diabetes (T2DM). AMPK works in glucose metabolism by allowing GLUT4 to move to cells, making insulin work better and preventing too much glucose production in the liver, protecting against both too high and too low blood sugar. Lipid metabolism allows for a rise in the rate of liver fatty acid breakdown and a decrease in cholesterol production, lipogenesis and triglyceride build-up, all of which suppresses basal adipocyte activity. Thus, AMPK enhances mitochondria production, aids in functioning under stressed conditions, decreases harmful oxidant species and lowers pro-inflammatory pathways such as TNF- α -mediated NF- κ B and therefore, helps stop chronic inflammation and oxidative damage in diabetes, as well as in CVD. In addition, cellular proliferation and senescence are controlled by AMPK, implying that changes in metabolism are linked to aging and the growth of chronic diseases. Therefore, AMPK has been proposed as a therapeutic target for metabolic illnesses such as T2DM, obesity and CVD, although this remains experimental. by helping balance the body's energy, adjusting the metabolism and extending the life of the cells. [25].

lowering the absorption of Ox-LDL and reducing both atherosclerosis and endothelial dysfunction.[27].

While a direct regulatory link between AMPK and ECE-1 has not been established, their functions are likely coordinated through shared anti-inflammatory and antioxidant pathways. Activation of AMPK may suppress the increased expression and production of ECE-1 and ET-1 by lowering inflammatory and oxidative stress through NF- κ B.[27]. As a result, excessive ET-1 in T2D and AMI conditions might decrease which can relax blood vessels, decrease inflammation and prevent excess fibrosis. Similarly, continuing high ET-1 signals, resulting from more ECE-1 activity, might lead to inflammation which decreases AMPK activity and can intensify both metabolism- and circulation-related issues.[28]

A decrease in AMPK points to poor regulation of metabolism, an increase in oxidative stress and problems with blood vessels in both T2D and AMI. A further rise in Ox-LDL and more ECE-1/ET-1 signaling emphasizes this disorder, causing harm to the vessel walls, increased atherosclerosis and cardiac changes. Activating AMPK with a therapeutic strategy might regulate metabolism, minimize the harmful effects from Ox-LDL and manage ECE-1, making it a compelling therapeutic target for patients with T2D and AMI, Adapted from [29].

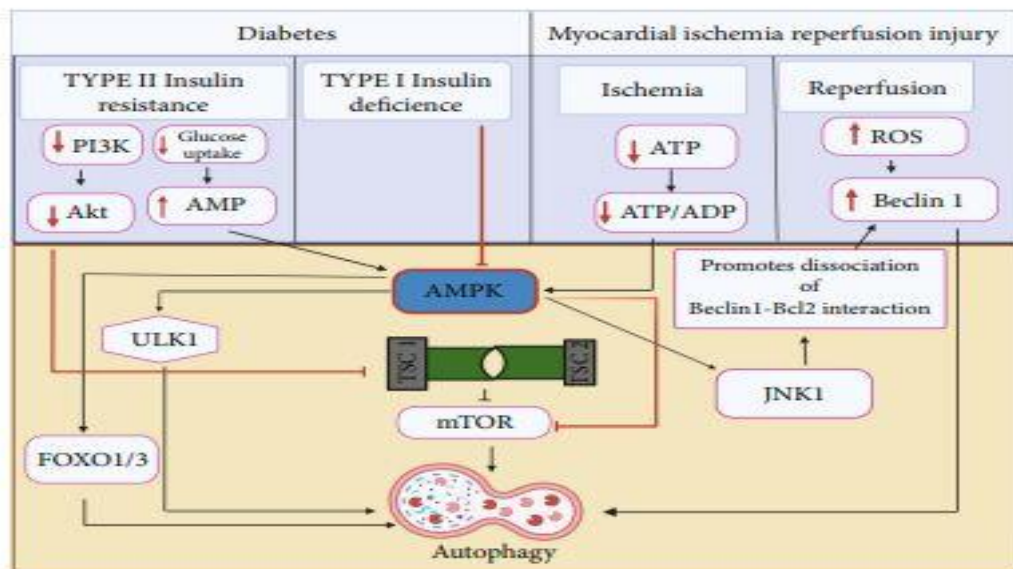


Figure 2: AMPK as a Central Regulator Linking Diabetes-Induced Insulin Signaling Dysfunction and Acute Myocardial. Adapted from [26]. This illustrates AMPK's role in autophagy during ischemia-reperfusion injury.



3.1. Structure and Formation of Ox-LDL

LDL particles are composed of a core of cholesterol esters surrounded by phospholipids and a single large protein, apolipoprotein B-100 (apoB-100). Oxidation by reactive oxygen species (ROS) and reactive nitrogen species (RNS) modifies both the lipid and protein components of LDL, generating lipid peroxides (e.g., malondialdehyde) and altering the structure of apoB-100 [30]. These modifications prevent Ox-LDL from being recognized by the native LDL receptor and instead promote its uptake by scavenger receptors on macrophages and endothelial cells [31].

3.2. Pathophysiological Significance in Type 2 Diabetes

Oxidized low-density lipoprotein (ox-LDL) is a pathologically modified form of LDL. Oxidation alters apolipoprotein B-100 and lipids, enabling binding to scavenger receptors such as SR-A, CD36, and LOX-1 on macrophages and endothelial cells. Under physiological conditions, Ox-LDL has no known function; instead, it is recognized as a key indicator of oxidative stress, suggesting that antioxidants are insufficient. Ox-LDL is essential in atherosclerosis because it leads macrophages to absorb cholesterol, causes the formation of "foam cells," creates problems in the endothelial layer, increases vascular smooth muscle cell activity, causes ongoing inflammation and saturates ways to remove excess cholesterol (ABCA1, ABCG1), so more cholesterol is maintained. Because of the fast-paced changes in glucose, cholesterol and oxidative stress seen in diabetics, LDL oxidation happens much faster. A rise in Ox-LDL in diabetic individuals contributes to heart and blood vessel disease by increasing dysfunction in veins, reducing nitric oxide production, increasing expression of adhesion molecules, accelerating the formation of foam cells, extending vascular inflammation and stimulating vascular smooth muscle cells to migrate and reproduce.[32].

Ox-LDL plays a role in cardiac problems by forming weak plaques that can rupture, increasing matrix metalloproteinase secretion that destroys their protective caps, bringing about vascular smooth muscle cell death that weakens the plaques, and damaging heart muscle cells with oxidative stress working as a sign of higher risk for cardiovascular events, particularly in patients with diabetes.[33]. When Ox-LDL interacts with LOX-1 receptors at the molecular level, it has been suggested to potentially upregulate ECE-1, leading to higher endothelin-1 production and increased vasoconstriction, oxidative stress, and inflammation. However, direct evidence linking Ox-LDL to ECE-1 upregulation remains limited and requires further investigation. This process is opposed by AMP-activated protein kinase which increases the movement of cholesterol out of cells, reduces damaging ROS and decreases inflammatory NF- κ B activity. In Type 2 diabetes, having too many nutrients for a long time makes AMPK less active which favors ECE-1/endothelin-1 and leads to more oxidized LDL-related damage to blood vessels. Ox-LDL appears to play a central role linking abnormal metabolism, vascular inflammation, and cardiac complications in diabetes, suggesting several potential therapeutic targets, all of which require experimental validation. [30].

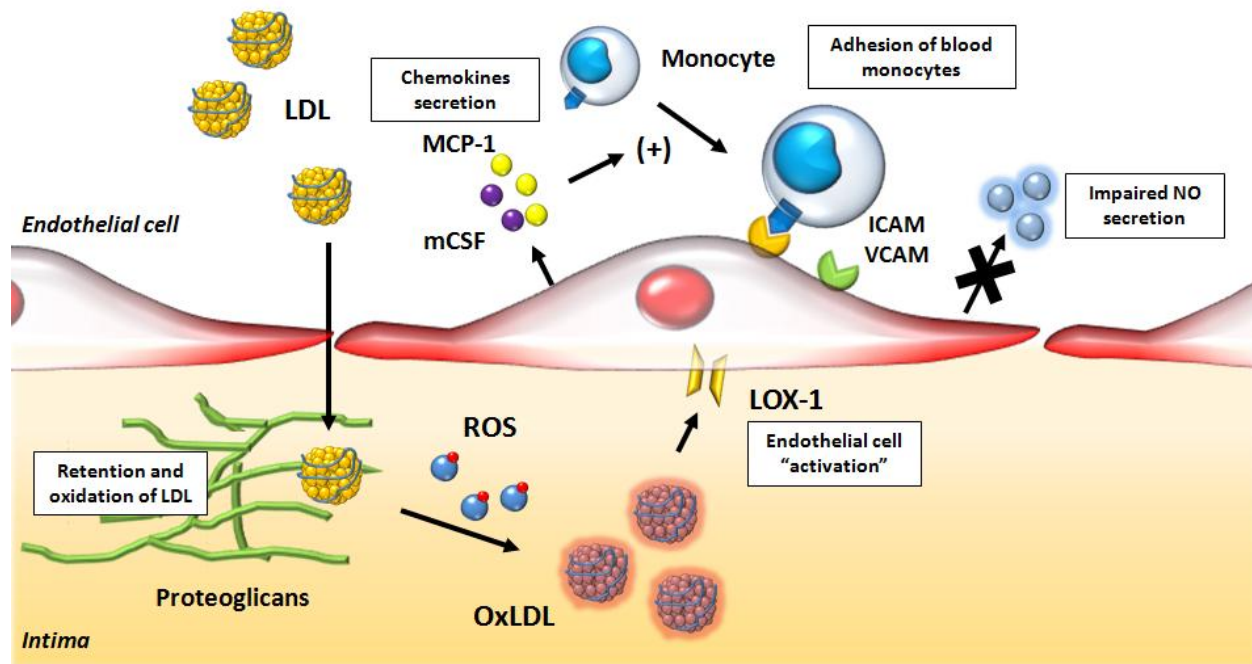


Figure 4: Role of Oxidized LDL in Atherosclerosis [30].

Figure 4 shows that Oxidized LDL (Ox-LDL) is taken up by endothelial cells (ECs) mainly through the action of overexpressed LOX-1 in human atherosclerotic lesions, leading to important changes in the ECs. The MCP-1 chemokine is triggered by LOX-1 and the MAPK pathway, while ICAM-1 and VCAM-1 adhesion molecules are turned on by LOX-1 through NF- κ B activation. In addition, OxLDL-LOX-1 pathways block production of nitric oxide and encourage EC death by turning on NF- κ B and AP- pathways. Overall, they increase endothelial dysfunction, attract more inflammatory cells and allow plaques inside the arteries to grow faster [33].

DISCUSSION

4. Clarification on the Nature of Claims Made in This Review

Before discussing the interrelationships between ECE-1, AMPK, and ox-LDL, it is important to clarify the evidentiary basis of this review. All claims regarding the diagnostic or therapeutic potential of these biomarkers are based exclusively on mechanistic and correlational evidence from basic science, preclinical, and observational studies. No diagnostic accuracy studies (e.g., prospective cohorts reporting sensitivity, specificity, or AUC-ROC values) have been published for this biomarker panel in T2DM-AMI patients. Therefore, statements such as "could serve as a biomarker," "may improve diagnosis," or "shows diagnostic potential" are speculative and hypothesis-generating. They should not be interpreted as evidence-based clinical recommendations. The primary value of this review is to synthesize current mechanistic knowledge and to identify gaps for future research, not to claim established diagnostic utility.



for observational studies were not applied. Consequently, the strength of the evidence underlying the reported findings has not been formally evaluated. ***Third**, no grading of evidence strength (e.g., GRADE criteria) was conducted. Statements regarding the diagnostic potential of ECE-1, AMPK, and ox-LDL are based on mechanistic and correlational evidence, not on high-quality diagnostic accuracy studies. ***Fourth**, as noted in previous sections, this review contains no original patient data, clinical cohorts, or laboratory measurements. All conclusions are derived from published literature. Given these methodological limitations, the findings of this narrative review should be considered hypothesis-generating rather than conclusive. Future research should employ systematic reviews, meta-analyses, and prospective diagnostic accuracy studies to validate the theoretical framework proposed here.

4.4. Assay Availability, Measurement Methods, and Standardization Challenges

For any biomarker to be clinically useful, reliable and standardized assays must be available. Currently, the measurement of ECE-1, AMPK, and ox-LDL in serum presents several challenges.

ECE-1: Commercial ELISA kits for human ECE-1 are available from several suppliers (e.g., Cloud-Clone Corp, MyBioSource, Cusabio). However, these kits are primarily research-use-only (RUO) and not FDA-approved or CE-marked for diagnostic purposes. No standardized reference ranges or cutoff values have been established for ECE-1 in T2DM-AMI patients. Furthermore, ECE-1 exists in four isoforms with different subcellular localizations, and most commercial kits do not distinguish between them.

AMPK: AMPK is an intracellular kinase that is typically measured in tissue lysates rather than serum. While serum AMPK levels can be detected by ELISA, the physiological significance of circulating AMPK is poorly understood. Most published studies measure AMPK phosphorylation (activation state) in platelets, mononuclear cells, or cardiac tissue rather than serum levels. Therefore, serum AMPK as a diagnostic biomarker for T2DM-AMI lacks established biological plausibility and requires further mechanistic validation.

Ox-LDL: Ox-LDL is the most clinically advanced of the three biomarkers. Several commercial ELISA kits are available (e.g., Mercodia, Cloud-Clone Corp, Abcam). Some studies have reported reference values, and ox-LDL has been investigated in multiple cardiovascular cohorts. However, standardization remains problematic because different assays use different antibodies (e.g., against malondialdehyde-modified apoB-100 versus other epitopes), leading to variable results across studies.

4.5. Cost-Effectiveness and Feasibility in Resource-Limited Settings

Even if technical assay challenges are overcome, the clinical adoption of any new biomarker panel depends on cost-effectiveness and feasibility, particularly in resource-limited settings where T2DM and AMI burdens are highest.

Estimated Costs: Commercial ELISA kits for ox-LDL typically cost 300-600 \$ per kit (96 wells), translating to 3-6 \$ per sample plus labor and equipment costs. ECE-1 kits are similarly priced. AMPK kits range from 400-800 \$ per kit. A panel combining all three biomarkers would cost approximately 15-25 \$ per patient for consumables alone, not including instrument



maintenance, personnel, and quality control. In comparison, high-sensitivity troponin assays cost approximately 2-5 \$ per sample and are widely available on automated clinical chemistry platforms.

Feasibility in Resource-Limited Settings: ELISA-based measurements require specialized equipment (plate readers, washers, incubators), trained personnel, stable electricity, and cold chain storage for reagents. These requirements are often unavailable in low- and middle-income countries, particularly in rural areas. Point-of-care testing for troponin is increasingly available, but no point-of-care tests exist for ECE-1, AMPK, or ox-LDL.

Cost-Effectiveness: No cost-effectiveness studies have been published for this biomarker panel. Given the modest additional diagnostic information that might be provided beyond existing standards (hs-troponin, ECG), it is unlikely that a three-biomarker ELISA panel would be cost-effective unless it demonstrates superior diagnostic accuracy in prospective trials. Future studies should include formal cost-effectiveness analyses using incremental cost-effectiveness ratios (ICERs) and quality-adjusted life years (QALYs).

4.6. Confounding Factors Affecting ECE-1, AMPK, and Ox-LDL Levels

Several demographic, clinical, and pharmacological factors may influence serum levels of ECE-1, AMPK, and ox-LDL, potentially confounding their diagnostic utility in T2DM-AMI.

Age: Age is a major determinant of vascular function and oxidative stress. Endothelin-1 levels increase with age in healthy individuals, and ECE-1 activity may be similarly affected. Ox-LDL levels also rise with age due to accumulated oxidative damage. Age-matched control groups are essential in diagnostic validation studies.

Renal Function: Chronic kidney disease (CKD) is common in T2DM and affects biomarker levels through multiple mechanisms. ET-1 and its precursors are cleared by the kidneys, and ECE-1 levels may be elevated in CKD independent of cardiac status. Ox-LDL levels are also increased in CKD due to enhanced oxidative stress and reduced clearance. AMPK signaling is dysregulated in CKD. Therefore, renal function (eGFR, creatinine) must be measured and adjusted for in any diagnostic study.

Other Comorbidities: Hypertension, dyslipidemia, obesity, and heart failure independently affect all three biomarkers. Hypertension increases ET-1 and ECE-1 expression. Dyslipidemia directly affects LDL oxidation. Obesity is associated with reduced AMPK activity. Studies must either exclude patients with these comorbidities or adjust for them statistically.

Medications: Several commonly used medications affect these biomarker pathways:

- **Metformin** activates AMPK, potentially altering serum AMPK levels or activity
- **Statins** reduce LDL cholesterol and may decrease ox-LDL formation
- **ACE inhibitors and ARBs** interact with the endothelin system
- **Insulin and oral hypoglycemics** affect metabolic pathways that regulate AMPK.

4.7. Comparison with Emerging Biomarkers Further Along in Validation

The diagnostic potential of ECE-1, AMPK, and ox-LDL for T2DM-AMI must be evaluated in the context of other emerging biomarkers that have progressed further in clinical validation.



cohorts, or laboratory validation, the clinical utility of these biomarkers remains unproven. Key limitations of the current literature include its reliance on correlative data from observational studies and a lack of prospective, interventional trials. Future research should focus on validating these biomarkers in large, prospective cohorts of T2DM patients with direct, original measurements to establish their diagnostic and prognostic utility for AMI. Furthermore, exploring therapeutic strategies that modulate these pathways — such as AMPK activators or ECE-1 inhibitors — could pave the way for more personalized interventions, but such strategies remain experimental at this stage.

Acknowledgments: The authors are grateful to all the researchers who support us to achieve this study.

Conflict of interests. The authors declare that they have no conflict of interests regarding the publication of this work.

Methodological Note : This paper is a narrative (traditional) review. Readers should be aware of the following methodological features:

- No systematic search protocol was registered.
- No PRISMA checklist was followed.
- No quality assessment tools (e.g., QUADAS-2, Cochrane Risk of Bias, Newcastle-Ottawa Scale) were applied.
- No GRADE assessment of evidence strength was performed.
- No original patient data, clinical cohorts, or laboratory measurements are included.

For definitive diagnostic validation of ECE-1, AMPK, and ox-LDL in T2DM-AMI, prospective cohort studies with quantitative performance metrics are required.

References

- [1] Adams, J. A., Uryash, A., Lopez, J. R., & Sackner, M. A. (2021). The endothelium as a therapeutic target in diabetes: A narrative review and perspective. *Frontiers in Physiology*, *12*, 638491. <https://doi.org/10.3389/fphys.2021.638491>
- [2] Akhtar, M., Khan, R. U., Saadia, S., Dawood, M. H., Khan, M., Collins, P., Sauer, A. J., & Ahmed, R. (2025). Trends and disparities in acute myocardial infarction and type 2 diabetes mellitus-related mortality in the United States from 1999 to 2022. *Nutrition, Metabolism and Cardiovascular Diseases*, *35*(6), 104063. <https://doi.org/10.1016/j.numecd.2025.104063>
- [3] Babakr, A. T. (2024). Scavenger receptors: Different classes and their role in the uptake of oxidized low-density lipoproteins. *Biomedicine & Pharmacotherapy Journal*, *17*(2). <https://dx.doi.org/10.13005/bpj/2897>
- [4] Babakr, A. T. (2025). Oxidized low-density lipoproteins and their contribution to atherosclerosis. *Exploration of Cardiology*, *3*, 101246. <https://doi.org/10.37349/ec.2025.101246>
- [5] Böhm, F., & Pernow, J. (2007). The importance of endothelin-1 for vascular dysfunction in cardiovascular disease. *Cardiovascular Research*, *76*(1), 8–18. <https://doi.org/10.1016/j.cardiores.2007.06.004>
- [6] Bonilha, I., Hajdich, E., Luchiar, B., Nadruz, W., Le Goff, W., & Sposito, A. C. (2021). The reciprocal relationship between LDL metabolism and type 2 diabetes mellitus. *Metabolites*, *11*(12), 807. <https://doi.org/10.3390/metabo11120807>



- [7] Butler, J., Hammonds, K., Talha, K. M., Alhamdow, A., Bennett, M. M., Bomar, J. V. A., Ettlinger, J. A., Martinez Traba, M., Priest, E. L., Schmedt, N., Zeballos, C., Shaver, C. N., Afzal, A., Widmer, R. J., Gottlieb, R. L., Mack, M. J., & Packer, M. (2025). Incident heart failure and recurrent coronary events following acute myocardial infarction: A multicenter study. *European Heart Journal*, *46*(16), 1540–1550. <https://doi.org/10.1093/eurheartj/ehae885>
- [8] Caturano, A., Rocco, M., Tagliaferri, G., Piacevole, A., Nilo, D., Di Lorenzo, G., Iadicco, I., Donnarumma, M., Galiero, R., Acierno, C., et al. (2025). Oxidative stress and cardiovascular complications in type 2 diabetes: From pathophysiology to lifestyle modifications. *Antioxidants*, *14*(1), 72. <https://doi.org/10.3390/antiox14010072>
- [9] Leon, B. M., & Maddox, T. M. (2015). Diabetes and cardiovascular disease: Epidemiology, biological mechanisms, treatment recommendations and future research. *World Journal of Diabetes*, *6*(13), 1246–1258. <https://doi.org/10.4239/wjd.v6.i13.1246>
- [10] Dong, Y., Zhang, M., Wang, S., Liang, B., Zhao, Z., Liu, C., Wu, M., Choi, H. C., Lyons, T. J., & Zou, M.-H. (2010). Activation of AMP-activated protein kinase inhibits oxidized LDL-triggered endoplasmic reticulum stress in vivo. *Diabetes*, *59*(6), 1386–1396. <https://doi.org/10.2337/db09-1637>
- [11] Dyck, J. R., & Lopaschuk, G. D. (2006). AMPK alterations in cardiac physiology and pathology. *The Journal of Physiology*, *574*(1), 95–112. <https://doi.org/10.1113/jphysiol.2006.109389>
- [12] Evandiar, M., Airlangga, P. S., Setiawan, P., Santoso, K. H., Kriswidyatomo, P., & Utomo, B. (2025). Role of endothelin-1 on myocardial ischemia-reperfusion injury: A comprehensive review. *Journal of Neonatal Surgery*, *14*(9S), 394–399. <https://doi.org/10.52783/jns.v14.2686>
- [13] Golino, M., Marazzato, J., Blasi, F., Morello, M., Chierchia, V., Cadonati, C., Matteo, F., Licciardello, C., Zappa, M., Ageno, W., Passi, A., Angeli, F., & De Ponti, R. (2022). High-sensitivity cardiac troponin T and the diagnosis of cardiovascular disease in the emergency room: The importance of combining cardiovascular biomarkers with clinical data. *Journal of Clinical Medicine*, *11*(13), 3798. <https://doi.org/10.3390/jcm11133798>
- [14] Minamino, T., Kurihara, H., Takahashi, M., Shimada, K., Maemura, K., Oda, H., Ishikawa, T., Uchiyama, T., Tanzawa, K., & Yazaki, Y. (1997). Endothelin-converting enzyme expression in the rat vascular injury model and human coronary atherosclerosis. *Circulation*, *95*(1), 221–230. <https://doi.org/10.1161/01.CIR.95.1.221>
- [15] Jeon, S.-M. (2016). Regulation and function of AMPK in physiology and diseases. *Experimental & Molecular Medicine*, *48*(7), e245. <https://doi.org/10.1038/emm.2016.81>
- [16] Martín-Timón, I., Sevillano-Collantes, C., Segura-Galindo, A., & del Cañizo-Gómez, F. J. (2014). Type 2 diabetes and cardiovascular disease: Have all risk factors the same strength? *World Journal of Diabetes*, *5*(4), 444–470. <https://doi.org/10.4239/wjd.v5.i4.444>
- [17] Kufazvinei, T. T. J., Chai, J., Boden, K. A., Channon, K. M., & Choudhury, R. P. (2024). Emerging opportunities to target inflammation: Myocardial infarction and type 2 diabetes. *Cardiovascular Research*, *120*(11), 1241–1252. <https://doi.org/10.1093/cvr/cvae142>
- [18] Lee, H., Fernandes, M., Lee, J., Merino, J., & Kwak, S. H. (2025). Exploring the shared genetic landscape of diabetes and cardiovascular disease: Findings and future implications. *Diabetologia*, 68(6), 1087–1100. <https://doi.org/10.1007/s00125-025-06403-9>
- [19] Li, J., Miller, E. J., Ninomiya-Tsuji, J., Russell, R. R., III, & Young, L. H. (2005). AMP-activated protein kinase activates p38 mitogen-activated protein kinase by increasing recruitment of p38 MAPK to TAB1 in the ischemic heart. *Circulation Research*, *97*(9), 872–879. <https://doi.org/10.1161/01.RES.0000187458.77026.10>
- [20] Low Wang, C. C., Hess, C. N., Hiatt, W. R., & Goldfine, A. B. (2016). Clinical update: Cardiovascular disease in diabetes mellitus—Atherosclerotic cardiovascular disease and heart failure in type 2

الخلاصة

المقدمة: يُعد داء السكري من النوع الثاني عاملاً رئيسياً يرفع بشكل كبير احتمالية الإصابة باحتشاء عضلة القلب الحاد، وذلك عبر مسارات مرضية تشمل اختلال الوظيفة البطانية، والإجهاد التأكسدي، والعمليات الالتهابية. أما الوسائل التشخيصية التقليدية المتمثلة في تخطيط كهربية القلب وقياسات التروبونين، فغالباً ما تكون فعاليتها محدودة لدى مرضى السكري من النوع الثاني، ويعود ذلك إلى أن الأعراض تظهر لديهم بشكل غير نمطي مقارنة بغيرهم من المرضى.

طرق العمل: تعتمد هذه المراجعة السردية على تجميع المعطيات المنشورة في الأدبيات العلمية المستمدة من الدراسات الأساسية وما قبل السريرية وكذلك الدراسات السريرية، وذلك فيما يتعلق ببنية ووظائف وتداخلات ثلاث مؤشرات حيوية موجودة في المصل، وهي: إنزيم المحول للإنديوثيلين-1 (ECE-1)، وكيناز البروتين المنشط بـ (AMPK) AMP، والبروتين الدهني منخفض الكثافة المؤكسد (ox-LDL). ويركز البحث على دور هذه المؤشرات في الآليات المرضية المرتبطة باحتشاء عضلة القلب الحاد لدى المصابين بداء السكري من النوع الثاني. من المهم التنبيه إلى أن هذا العمل لا يتضمن أي بيانات أصلية مستمدة من مرضى، ولا يعرض أي نتائج من مجموعات سريرية، ولا يقدم قياسات مخبرية جديدة. إن الإمكانية التشخيصية التي يناقشها البحث تستند فقط إلى أدلة مستخلصة من دراسات آلية ودراسات ارتباطية، ولذلك تبقى هذه الإمكانية في إطارها النظري غير المثبت سريرياً حتى الآن.

النتائج: يؤدي إنزيم ECE-1 إلى تضيق الأوعية الدموية من خلال تحفيزه لإنتاج الإندوثيلين-1 (ET-1)، بينما يعمل بروتين AMPK كمُنظم مركزي لتوازن الطاقة داخل الخلايا، ويُعد الـ ox-LDL محركاً أساسياً في نشوء تصلب الشرايين. وتُظهر التداخلات بين هذه المؤشرات الثلاثة وجود حلقة مرضية مغلقة تتفاقم فيها الأضرار الاستقلابية والوعائية. وبناءً على الأدلة المنشورة المستمدة من الدراسات الآلية والارتباطية، فإن دمج هذه المؤشرات الحيوية قد يؤدي - من الناحية النظرية - إلى تحسين التشخيص المبكر وتخصيص العلاج لكل مريض على حدة، إلا أن هذا الافتراض لا يزال مجرد فرضية نظرية تنتظر التحقق السريري من خلال دراسات مستقبلية جماعية تتضمن بيانات أصلية مستمدة من مرضى حقيقيين. تجدر الإشارة إلى أنه لا توجد في الأدبيات العلمية الحالية أي مقاييس كمية تشخيصية (مثل معايير الحساسية، أو النوعية، أو المساحة تحت المنحنى التشغيلية، أو قيم القطع التشخيصية) للمجموعة المقترحة من المؤشرات الحيوية. كما أنه لا تتوفر أي مقارنة كمية بين هذه المؤشرات والمعايير التشخيصية الحالية (كالتروبونين عالي الحساسية أو تخطيط كهربية القلب)، وذلك ببساطة لأن دراسات التحقق السريري اللازمة لم تجر بعد.

الاستنتاجات: استناداً إلى ما توصلت إليه الدراسات الآلية المتاحة حالياً، فإن مجموعة المؤشرات الحيوية المكونة من ECE-1 و AMPK و ox-LDL تمتلك إمكانات تشخيصية نظرية لمرضى السكري من النوع الثاني المصابين باحتشاء عضلة القلب الحاد. لكن يجدر التأكيد على أن هذا العمل هو مراجعة سردية لا تحتوي على بيانات أصلية مستمدة من مرضى، ولا تشمل مجموعات سريرية، ولا تقدم قياسات مخبرية جديدة. ولهذا السبب، لا يمكن التوصية بأي تطبيق تشخيصي لهذه المؤشرات إلا بعد إجراء دراسات تحقق سريري مستقبلية جماعية. كما أن مصطلح "تشخيص" الوارد في عنوان هذا البحث لا يزال طموحاً نظرياً في غياب معايير تشخيصية محددة، وغياب المقاييس الكمية (مثل الحساسية، والنوعية، والمساحة تحت المنحنى، وقيم القطع)، وعدم وجود مقارنة كمية مباشرة مع المعايير التشخيصية الحالية كالتروبونين عالي الحساسية وتخطيط كهربية القلب. إن الحاجة ملحة وماسة إلى إجراء دراسات تحقق سريري توفر هذه المقاييس الكمية بشكل موثوق.

الكلمات المفتاحية: داء السكري من النوع الثاني، الإنديوثيلين المصلي، وكيناز البروتين المنشط بـ (AMPK) AMP، إنزيم المحول للإنديوثيلين-1، احتشاء عضلة القلب. مراجعة سردية، توليد الفرضيات، إطار نظري.