



Membrane and soluble endoglin role in cardiovascular and metabolic disorders related to metabolic syndrome

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Abstract

Membrane endoglin (Eng, CD105) is a transmembrane glycoprotein essential for the proper function of vascular endothelium. It might be cleaved by matrix metalloproteinases to form soluble endoglin (sEng), which is released into the circulation. Metabolic syndrome comprises conditions/symptoms that usually coincide (endothelial dysfunction, arterial hypertension, hyperglycemia, obesity-related insulin resistance, and hypercholesterolemia), and are considered risk factors for cardio-metabolic disorders such as atherosclerosis, type II diabetes mellitus, and liver disorders. The purpose of this review is to highlight current knowledge about the role of Eng and sEng in the disorders mentioned above, in vivo and in vitro extent, where we can find a wide range of contradictory results. We propose that reduced Eng expression is a hallmark of endothelial dysfunction development in chronic pathologies related to metabolic syndrome. Eng expression is also essential for leukocyte transmigration and acute inflammation, suggesting that Eng is crucial for the regulation of endothelial function during the acute phase of vascular defense reaction to harmful conditions. sEng was shown to be a circulating biomarker of preeclampsia, and we propose that it might be a biomarker of metabolic syndrome-related symptoms and pathologies, including hypercholesterolemia, hyperglycemia, arterial hypertension, and diabetes mellitus as well, despite the fact that some contradictory findings have been reported. Besides, sEng can participate in the development of endothelial dysfunction and promote the development of arterial hypertension, suggesting that high levels of sEng promote metabolic syndrome symptoms and complications. Therefore, we suggest that the treatment of metabolic syndrome should take into account the importance of Eng in the endothelial function and levels of sEng as a biomarker and risk factor of related pathologies.

Keywords Endoglin · Soluble endoglin · Endothelial dysfunction · Hyperglycemia · Metabolic syndrome

Introduction

Endoglin (Eng, CD105) is a 180 kDa transmembrane glycoprotein considered a co-receptor for ligands of the Transforming Growth Factor β (TGF β) superfamily, with an increasing research interest and currently over 2800 citations in PubMed. Eng is involved in the physiological function of the endothelium, but also plays an essential role in various

pathological conditions. There are two different isoforms of membrane Eng expressed by various cells and soluble endoglin (sEng) circulating in plasma or cell culture medium [1].

Eng is expressed by endothelial cells, vascular smooth muscle cells [2], fibroblasts [3], hepatic stellate cells [4], and activated monocytes and macrophages [5]. Changes in Eng expression and function are predominantly associated with several pathological conditions, including Hereditary hemorrhagic telangiectasia (HHT) [6], cancer, angiogenesis [7], heart development [8], fibrosis [9], endothelial dysfunction [10], and inflammation [11].

sEng is the N-terminal cleavage product of the extracellular domain of Eng formed by the activity of matrix metalloproteinases [12–14] that is released into the circulation. sEng can be detected and used as a biomarker in patients with various cardiovascular and metabolic disorders such as

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familial hypercholesterolemia [15, 16], arterial hypertension [17], preeclampsia [18], and diabetes mellitus [19].

The role of the membrane and soluble endoglin in cardiovascular diseases, atherosclerosis, and endothelial dysfunction, were reviewed previously [20, 21]. Furthermore, an excellent review focusing on the role of Eng in angiogenesis, cancer, and immune cells, was recently published by Schoonderwoerd et al. [22]. Indeed, sEng was demonstrated to be a circulating biomarker of development and progression of preeclampsia [18]. However, the elaborate summary of the Eng and sEng role in cardiovascular disorders associated with altered glucose and lipid metabolism was not summarized yet. Metabolic syndrome comprises conditions/symptoms that usually co-occur (endothelial dysfunction, arterial hypertension, hyperglycemia, obesity-related insulin resistance, and hypercholesterolemia or hypertriglyceridemia) and are considered as the risk factors for cardiometabolic disorders such as atherosclerosis, type II diabetes mellitus, and liver disorders. This review aims to critically evaluate recent results and update some information in the light of recent papers regarding membrane and soluble endoglin with respect to metabolic syndrome-related pathologies.

Membrane endoglin structure and signaling

Eng is composed of two disulfide-linked transmembrane monomers, as shown in Fig. 1. According to its structure, Eng belongs to the family of proteins containing the zona pellucida domain [23]. Eng contains an extracellular part, hydrophobic intramembrane part, and short serine/threonine-rich cytoplasmic region. The extracellular part of

Eng contains the orphan domain (which does not show any homology with other protein families) and the zona pellucida (ZP) part (responsible for protein–protein interaction). Part of the ZP domain is arginyl-glycyl-aspartic acid (RGD), a fundamental recognition structure for the binding with integrins and other RGD receptors [24, 25].

Two alternatively spliced isoforms of Eng have been identified [26]. Their structure differs in the number of amino acids in the cytoplasmic tail, level of phosphorylation, and affinity to receptors (Fig. 2) [27].

Long endoglin (L-Eng) is a predominantly expressed Eng isoform. L-Eng contains a cytoplasmic chain with 47 amino acids, and it is known to stimulate endothelial cell migration, proliferation, and angiogenesis through Eng/ALK1/SMAD1/5/8 pathway. Short endoglin (S-Eng) is a minor isoform of Eng with a cytoplasmic chain composed of 14 amino acids. S-Eng expression seems to inhibit endothelial cell proliferation, migration, and it was shown to induce endothelial cell senescence through Eng/ALK5/SMAD2/3 pathway [27]. However, the exact role of S-Eng was not widely explored [22, 26].

L-Eng isoform is the predominantly expressed isoform and plays a crucial role in pathological conditions. This review mainly focuses on L-Eng, which is mentioned as Eng in this paper.

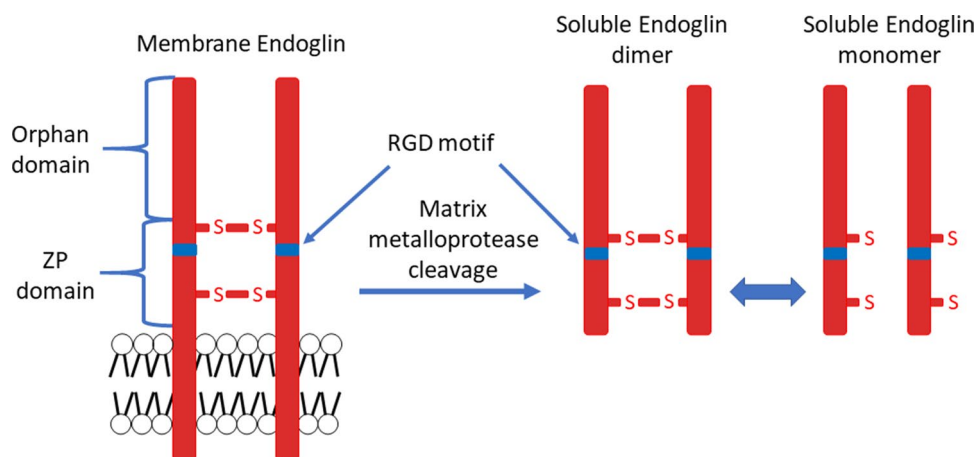
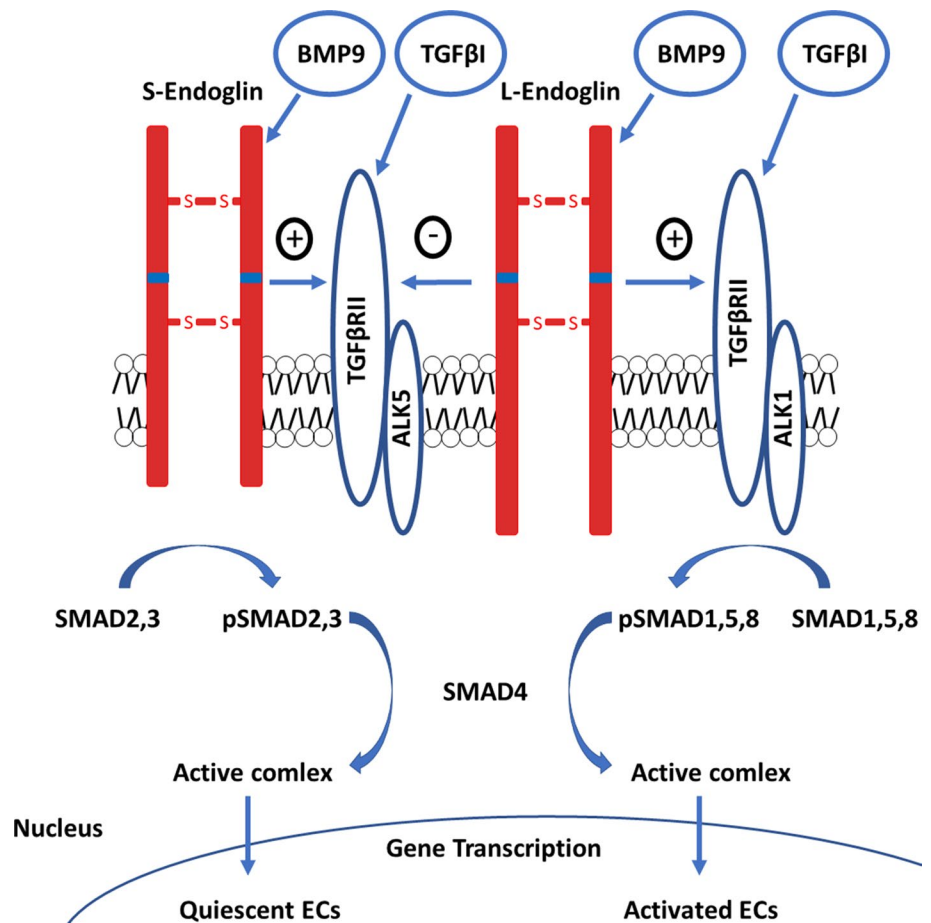


Fig. 1 Schematic of Eng and its cleavage product sEng. The extracellular part of Eng consists of the orphan domain and the ZP domain. ZP domain contains the RGD motif, which is a crucial recognition structure for the binding with integrins and other RGD receptors.

Matrix metalloproteinases can cleave the whole extracellular part of Eng and produce sEng, which is released into the circulation or cell culture media. Up to date, both soluble forms of endoglin, including monomer and dimer, were proposed to be detected in the circulation

Fig. 2 Role of different Eng isoforms in the endothelium. S-Eng participates in TGFβRII/ALK5-mediated signaling, where it promotes SMAD2/3 phosphorylation. Phosphorylated SMAD2/3 (pSMAD2/3) binds with common mediator SMAD4 (co-SMAD4) and creates an active complex. This complex translocates to the nucleus and induces transcription of genes related to the quiescent endothelial cell phenotype. On the other hand, L-Eng participates in BMP9 and in both TGFβRII/ALK1 and TGFβRII/ALK5-mediated signaling, as well. L-Eng promotes TGFβRII/ALK1—SMAD1/5/8 pathway and inhibits TGFβRII/ALK5—SMAD 2/3 pathway. Binding of phosphorylated SMAD1/5/8 (pSMAD1/5/8) with co-SMAD4 creates an active complex. Translocation of this complex to the nucleus induces transcription of genes related to activated endothelial cells phenotype



Membrane endoglin expression and function in cardiometabolic disorders in vivo and in vitro

Eng expression is predominant in the cells found in the vessel wall, including endothelial cells, monocytes/macrophages, fibroblasts, and vascular smooth muscle cells (SMCs) [28]. It is also expressed, in lesser amounts, in other cell types, for instance, in hepatic stellate cells [4]. We reviewed the role of Eng in atherosclerosis and endothelial dysfunction, where we suggested the importance of proper Eng expression and signaling in healthy endothelium and its potential involvement in the development/progression of atherogenesis [21]. In the current review, we significantly extend that information and try to highlight novel findings in the field, regarding the importance of Eng expression and function in metabolic syndrome-related pathologies.

Membrane endoglin, hypercholesterolemia, endothelial nitric oxide synthase, endothelial dysfunction, and inflammation

Endothelial nitric oxide synthase (eNOS) is a crucial enzyme maintaining the proper function of vascular

endothelium. For instance, when the activity of eNOS is reduced by hypercholesterolemia and/or inflammation, it may result in the development of endothelial dysfunction [29].

Several studies showed a mutual relationship between Eng, eNOS, and the proper function of the endothelium. It was demonstrated that changes in Eng levels result in the alteration of blood vessels function and possibly to the development of endothelial dysfunction [10, 30].

For instance, decreased expression of eNOS in Eng haploinsufficient mice (Eng[±]) resulted in impaired endothelium-dependent vasodilation in mice [30]. In addition, Toporsian et al. found that Eng is an essential component of the eNOS activation complex and that it stabilizes eNOS protein, which indicates the crucial role of Eng in the regulation of local vascular tone [31]. Another in vitro study, published by Santibanez et al., showed that Eng increases SMAD2 protein levels, phosphorylation status, and stability, which results in an increased eNOS expression in either absence or presence of exogenous TGF-β1 in endothelial cells [32]. Last but not least, reduced Eng/eNOS expression was also demonstrated in vitro in human umbilical vein endothelial cells (HUVECs) after the simulation of inflammation by the treatment with TNF-α [33].

Also, Jerkic et al. used mouse endothelial cells derived from the yolk sac of Eng homozygous Eng^{-/-}, heterozygous Eng[±] and control Eng^{+/+} mice during embryonic day 8.5 to demonstrate the role of Eng in endothelial function [34]. In Eng[±] deficient cells, they showed decreased levels of protein stabilizing factors VEGFR2, PAK-1, VE-cadherin, and Rac-2 and increased levels of endothelial barrier destabilizing factors TSP-1 and CD148. Destabilization of the endothelial barrier was related to increased RhoA activation and increased permeability of neutrophils through the endothelial barrier. They suggested that reduced Eng expression results in increased endothelial permeability and impairment of endothelial barrier function, which are general hallmarks of endothelial dysfunction [34]. Similarly, Anderberg et al. showed significantly weakened endothelial cell barrier to tumor cell intravasation and extravasation enhanced ALK5 signaling and increased endothelial-to-mesenchymal transition in endoglin deficient mice suggesting the importance of Eng for the proper barrier function of the endothelium [35]. Supporting that, Rossi et al. nicely demonstrated the role of Eng in adhesion of mural and endothelial cells (human aortic endothelial cells (HAECs), HUVECs, and umbilical artery smooth muscle cells) [36]. In general, they showed that adhesion of mural cells with endothelial cells depends at least partially on Eng. Once Eng expression was reduced (by siRNA of Eng) or blocked by sEng treatment, integrin-mediated adhesion of mural cells was altered. In line with that, reduced expression of Eng resulted in increased endothelial permeability, suggesting a critical role of Eng in endothelial function [36].

In another experiment, Rossi et al. used rat myoblast transfectants expressing human Eng and corresponding mock transfectants to demonstrate the role of Eng in the adhesion and transmigration of leukocytes through cell monolayer [11]. For the first time, they demonstrated the interaction of leukocyte integrin $\alpha 5 \beta$ and platelet $\alpha \text{IIb} \beta 3$ with Eng expressed on endothelial cells via RGD motif, suggesting that Eng might be considered as an adhesion molecule that promotes the development of endothelial dysfunction. They also showed that reduced Eng expression resulted in a decreased inflammation-induced transendothelial migration of leukocytes, suggesting the crucial role of Eng in leukocyte transmigration, which is the hallmark of endothelial dysfunction [11]. Moreover, Ojeda-Fernandez demonstrated an essential role of Eng for the immune system function in mice macrophages. They compared the immune response of Eng myeloid lineage-specific knock out mice (Eng KO mice) with the immune response of wild-type mice. Eng KO mice developed spontaneous infections in soft tissues, impaired leukocyte transmigration, phagocytosis, and altered gene expression in macrophages, suggesting the important role of Eng in macrophage function and inflammation [37].

Altogether, these studies might suggest the potential proinflammatory role of Eng.

However, it is interesting to mention that these in vitro experiments show the effect of Eng in macrophages in the microvasculature, not in large arteries, where we usually study endothelial dysfunction and atherosclerosis. Thus, we propose the different roles of Eng in various parts of the macrovasculature and microvasculature.

Interestingly, we found that exposure to similar stimuli results in opposite outcomes in vitro and in in vivo experiments with respect to Eng. Our study focused on how hypercholesterolemia in vivo or its simulation in vitro can affect the expression and function of Eng. We explored the differences between the acute effect of 7-ketocholesterol (7 K) [simulating oxidized LDL (oxLDL) effects in atherogenesis] in HAECs and the chronic effect of hypercholesterolemia in apolipoprotein E-deficient/LDL receptor-deficient (ApoE^{-/-}/LDLR^{-/-}) mice [10]. It was demonstrated that 7 K induces Eng expression in HAECs via simultaneous activation of transcription factors regulating Eng expression, including Krüppel like factor 6 (KLF6) [38, 39] nuclear factor kappa B p65—hypoxia-inducible factor 1 (RELA-HIF-1) [40, 41] and liver X receptor (LXR) nuclear receptor subfamily 1 group H member 3 (NR1H3) [14, 42]. Increased expression of Eng was associated with increased expression of eNOS and phosphorylated form of eNOS (p-eNOS) protein levels, which may suggest activation of potential protective mechanisms of Eng after oxidized cholesterol treatment in HAECs. On the other hand, significantly increased protein levels of Eng after 7 K treatment were associated with significantly increased levels of cell adhesion proinflammatory molecules (E/P-selectins, VCAM-1, and ICAM-1). Consequent increased adhesion and transmigration of monocytes through 7 K pre-treated endothelial monolayer confirmed the development of endothelial dysfunction. After the silencing of Eng, induction of adhesion and transmigration were prevented [10], suggesting a crucial role of Eng in endothelial dysfunction in acute (12 h) hypercholesterolemic condition in vitro.

In vivo part of the study was focused on the chronic effect of hypercholesterolemia in ApoE^{-/-}/LDLR^{-/-} mouse model of spontaneous hypercholesterolemia, endothelial dysfunction, and atherogenesis [43]. Hypercholesterolemia (in the time frame of two months) resulted in the development of vascular and endothelial dysfunction in the aorta, with reduced Eng/eNOS/pSMAD2/3 expression in the aorta and decreased nitric oxide (NO) production, induction of inflammation, and increase of sEng levels in plasma. These results show that reduced Eng expression is related to the alteration of NO production and vascular function (endothelial dysfunction) even before the formation of atherosclerotic lesions, which suggests that proper Eng expression is crucial for the prevention of endothelial dysfunction. This is in line

with our previous papers, showing that the progression of atherogenesis (increased size of atherosclerotic lesions) is accompanied by reduced Eng expression in the aorta and increased levels of sEng in blood [44–46]. Thus, we suggest that reduced Eng expression and increased levels of sEng are hallmarks of endothelial dysfunction development and atherogenesis, as shown in Fig. 3.

To summarize, acute exposure to oxLDL in vitro results in the development of endothelial dysfunction, which is strongly supported by increased expression Eng. In contrast, chronic exposure to cholesterol in vivo shows reduced Eng expression during the development of endothelial dysfunction and atherogenesis.

Membrane endoglin, hyperglycemia, diabetes mellitus, and arterial hypertension

Hyperglycemia is one of the metabolic syndrome-related symptoms and a hallmark of diabetes mellitus. Even though the role of Eng in hyperglycemia and diabetes is not widely investigated, some studies have attempted to establish a possible relationship between them.

Alvarez-Munoz et al. evaluated Eng expression in cultured skin fibroblasts from patients with Type 1 diabetes mellitus (DM1) with and without diabetic nephropathy,

because skin fibroblast behaviors in these patients are related to diabetic nephropathy risk. They showed increased Eng mRNA expression and protein levels in fibroblasts isolated from patients with DM1 with a lower risk of nephropathy development (“slow-track”) when compared to the patients with a higher risk of nephropathy development (“fast-track”). Therefore, they suggested a potential protective role of Eng in patients with DM1 for the development of diabetic nephropathy (fibrotic changes). However, there might be some limitations of that study, related to the fact that only four subjects from each group were tested for Eng protein expression studies, and no representative Western blots were shown in the article [47].

Another in vitro study performed on HUVECs treated with high glucose (25 mmol/l) or oscillating glucose (5–25 mmol/l) resulted in increased mRNA expression of Eng, HIF-1 α , and KLF6, which could be prevented by antioxidant alpha-lipoic acid [48]. However, no significant changes in Eng protein levels were found by Wang et al. when studying human blood outgrowth endothelial cells exposed to high glucose levels (25 mmol/l) [49].

Systemic arterial hypertension represents another symptom of metabolic syndrome, affecting blood vessels, and promoting endothelial dysfunction, suggesting that it may affect Eng expression and function. Nonetheless, to the best

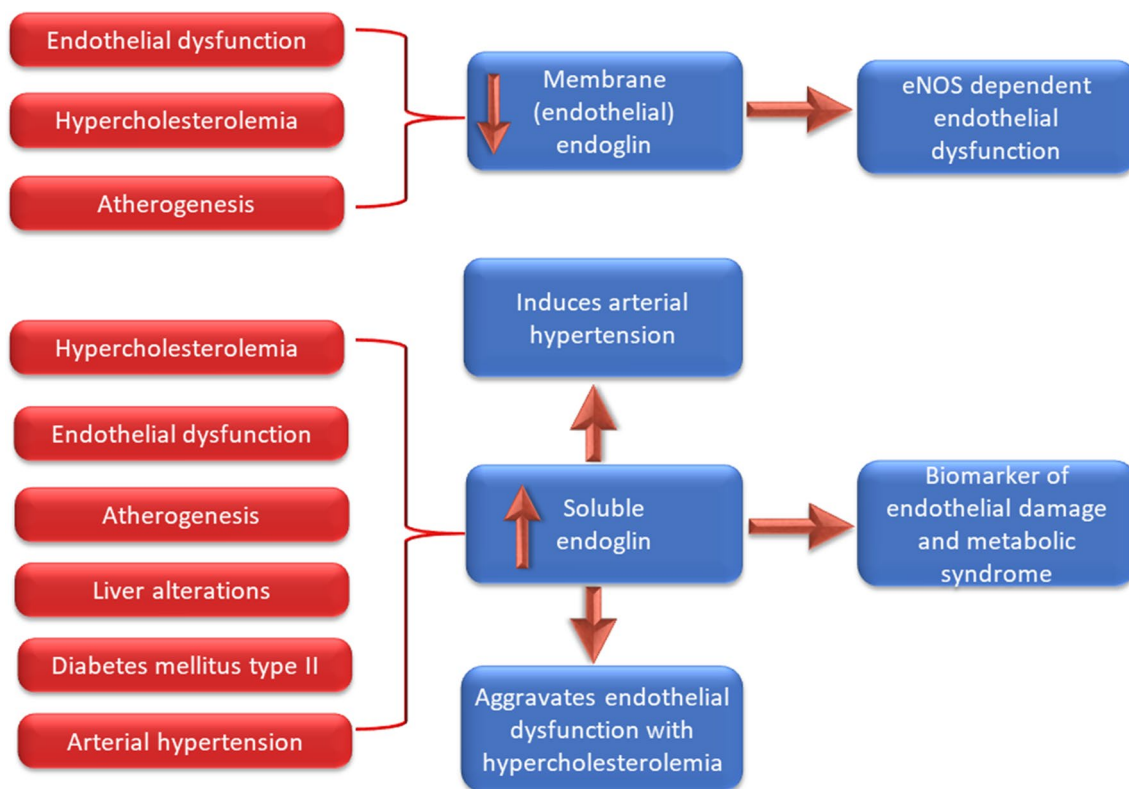


Fig. 3 Potential role of Eng and sEng in the metabolic syndrome

of our knowledge, no studies are focusing on the role of Eng in the systemic arterial hypertension up to date.

Membrane endoglin, liver alteration, and obesity-related to metabolic syndrome

Profibrotic changes in the liver are a part of the pathophysiology of diseases such as nonalcoholic fatty liver disease (NAFLD) and nonalcoholic steatohepatitis (NASH). NAFLD is often recognized as a hepatic manifestation of metabolic syndrome-related to obesity and fat accumulation in the liver. Some NAFLD patients may develop a more severe form termed NASH, which may progress to cirrhosis and permanent liver damage [50].

Eng is known to play an important role in fibrosis development in the liver [51, 52]. Meurer et al. reported high protein expression of Eng in isolated liver hepatic stellate cells, myofibroblast-like cells, liver macrophages (Kupffer cells), and liver sinusoidal endothelial cells; however, no expression of Eng in hepatocytes neither in HepG2 cell line was found. Up-regulation of Eng expression was observed during hepatic stellate cell activation and transdifferentiation to myofibroblast-like cells in cell culture and both mice and rat experimental models of liver injury (CCl₄ application or bile duct ligation) [53] [54]. However, it is of interest to mention that role of endoglin in liver fibrosis is still not well documented and lacks in-depth studies; thus, it might be too early to review these findings now to provide any meaningful insights.

Obesity (visceral) represents an essential step for the development of insulin resistance, and it plays a crucial role in the pathophysiology of type II diabetes mellitus and metabolic syndrome [55]. Kurki et al. focused on the expression profile of adipose tissue cytokines and angiogenesis-related proteins in obese and lean mice. They found significantly increased expression of Eng in adipose tissue of obese mice when compared to lean mice [56]. When exposed to calorie restriction for 50 days, the expression of Eng in obese mice did not decrease, even though they lost approximately 15.6% of their weight. On the contrary, energy restriction further increased the expression of Eng in obese mice. Surprisingly, the protein expression of Eng induced by energy restriction in lean mice was superior to the Eng expression in obese mice without energy restriction [56]. These data suggest that stress and catabolic processes during fast weight reduction can increase Eng expression as well as chronic inflammation during obesity [56].

Jilkova et al. further investigated the expression of Eng in white adipose tissue (WAT) in obese mice focusing on the relation between obesity, inflammation, and Eng expression in mice. Diet-induced low-grade inflammation in C57BL/6 J obese male mice (fed by corn oil-based high-fat diet) resulted in increased expression of Eng,

suggesting promotion of the angiogenesis during the early stages of WAT inflammation. To prove these results, long-chain polyunsaturated fatty acids or rosiglitazone were added to the diet to inhibit inflammation. Significantly reduced VCAM-1 and Eng expression was found in obese mice fed with a combination of long-chain polyunsaturated fatty acids and rosiglitazone when compared to mice fed high-fat diet [57]. The authors suggested the potential involvement of Eng in WAT inflammation and remodeling of adipose tissue (probably via promoting inflammation-related angiogenesis), which can be prevented by long-chain polyunsaturated fatty acids or rosiglitazone.

Taken together, the increase of Eng expression in adipose tissue during obesity is probably related to angiogenesis, inflammation, and/or adipogenesis. Despite that, there is no direct evidence of Eng playing a crucial role in the inflammation induced by visceral obesity development.

Soluble endoglin generation and the role of matrix metalloproteinases

Matrix metalloproteinases (MMPs) are zinc-dependent endopeptidases with various and often opposing effects in the organism. They can cleave many non-matrix targets such as Eng, chemokines, cytokines, cell adhesion molecules, and other proteinases [58]. MMPs may be divided according to the structure in the four main groups. Gelatinases (MMP2, MMP9), matrilysins (MMP7, MMP26), archetypal MMPs (MMP1, MMP8, MMP13, MMP3, MMP10, MMP12, MMP19, MMP20, MMP27), and furin-activable MMPs (MMP11, MMP21, MMP28, MMP14, MMP15, MMP16, MMP24, MMP17, MMP25, MMP23) [59]. Moreover, MMPs may also be divided into soluble forms circulating in the blood or cultured medium in experiments (MMP1, MMP2, MMP3, MMP7, MMP8, MMP9, MMP10, MMP11, MMP12, MMP13, MMP26) and membrane-bound forms expressed in various tissues (MMP14, MMP15, MMP16, MMP17, MMP24, MMP25) [60].

Up to date, only two MMPs demonstrated to be able to cleave Eng to release sEng, namely MMP12 [12] and MMP14 [13]. Interestingly, both MMP12 [61] and MMP14 [62] are known to be involved in the development of vascular diseases such as atherosclerosis.

Several papers were published in recent years concerning the role of sEng as a biomarker of various cardio-metabolic disorders. Besides, some studies pointed out the possible role of sEng in the pathophysiology of cardio-metabolic disorders/metabolic syndrome, which will be discussed below.

Soluble endoglin as a biomarker of cardiometabolic disorders

Our last review was focused on the importance of sEng as a biomarker of various cardiometabolic disorders [63]. Briefly, sEng levels are increased in patients with hypercholesterolemia [16] and Familial hypercholesterolemia [15]. In addition, sEng levels were also increased in the experimental mouse model of atherosclerosis with hypercholesterolemia, before the formation of visible atherosclerosis in aorta, and during the formation of advanced lesions [10, 45, 46]. Moreover, a correlation of sEng levels with cholesterol levels in hypercholesterolemic mice was demonstrated [63].

Hypercholesterolemia is a crucial risk factor for the development of atherosclerosis, and oxidative modification of cholesterol (oxLDL) plays an important role in the pathogenesis. It was demonstrated that sEng levels are higher in mice with more prominent atherosclerotic lesions [45, 46]. However, no correlation between atherosclerotic plaque size and sEng levels was confirmed [63]. Interestingly, sEng showed a positive linear correlation with carotid intima-media thickness in 978 patients, suggesting that sEng might be an interesting biomarker of subclinical carotid atherosclerosis [64].

Li et al. showed an association between sEng concentration and atherosclerotic cardiovascular disease risk [64]. On the other hand, Saita et al. found an inverse association of sEng concentration and severity of coronary atherosclerosis in patients with coronary artery disease [65]. Finally, Charytan et al. did not find a significant correlation between sEng levels and atherosclerotic burden in 122 patients' study [66]. According to these data, sEng could be considered a biomarker related to hypercholesterolemia, but there is no conclusive evidence that sEng reflects the development of atherosclerotic lesions.

Soluble endoglin, diabetes mellitus, and obesity

Hyperglycemia is considered as a risk factor for the development of cardiometabolic disorders, especially type II diabetes mellitus (DM). Plasma concentration of sEng was increased in patients with advanced DM, and the concentration of sEng positively correlates with the severity of diabetic vascular alterations such as retinopathy [67], diabetic peripheral neuropathy [68], and nephropathy [19, 69]. Ceriello et al. showed that both hypoglycemia and hyperglycemia increased sEng and MMP14 levels, and this effect could be reversed with glucagon like peptid-1 (GLP-1) in patients with type I diabetes mellitus. They suggested that an increase of sEng and MMP14 is related to oxidative stress development both in hypoglycemia and hyperglycemia [70].

Cawyer et al. focused on the effects of high glucose in the human extravillous cytotrophoblast cell line Sw.71 to evaluate the effects of high glucose on angiogenic (VEGF, PIGF) and antiangiogenic (sEng, sFLT-1) factors. They showed that high glucose levels ≥ 8.3 mmol/l significantly decreased the concentration of angiogenic and increased concentration of antiangiogenic factors, including sEng. This effect of high glucose could be reversed using a p-38 inhibitor (SB203580) or rosiglitazone [71, 72].

Lappas et al. investigated the role of sEng in gestational diabetes mellitus (GDM) and maternal obesity. They focused on the expression of sEng, Eng, and adhesion molecules in the placenta and adipose tissue. They examined healthy women, women with obesity, women with GDM, and women with a combination of obesity and GDM. Interestingly, no changes in the placenta were found, except significantly increased sEng and cell adhesion molecules in samples of adipose tissue of women with GDM [73]. However, Vieira et al. found the opposite trend of sEng concentration changes in obese women. They found a significantly decreased blood plasma concentration of sEng in the group of 834 obese women compared to the group of 3106 non-obese women [74].

These data demonstrate that sEng levels reflect hyperglycemia, and we might propose that sEng could be considered an important biomarker of developing diabetic changes; however, its relation to obesity (especially visceral obesity) must be further investigated.

Soluble endoglin as a biomarker of arterial hypertension

As mentioned above, sEng levels are increased in hypercholesterolemia and hyperglycemia, both risk factors for the development of systemic arterial hypertension. The only paper describing the sEng levels and systemic arterial hypertension showed that sEng levels correlated with systolic blood pressure, left-ventricular hypertrophy, and endothelial dysfunction [67]. In addition, several papers described the relation between sEng and Pulmonary arterial hypertension (PAH). Indeed, PAH is not a subtype of arterial hypertension, and PAH etiopathogenesis is very different from systemic arterial hypertension. Increased sEng concentration in the blood plasma of patients correlated with the New York Heart Association (NYHA) classification of PAH. High sEng concentrations were significantly increased in patients with idiopathic or hereditary PAH, PAH associated with connective tissue disease, drug, or toxin-related PAH, but not with PAH related to congenital heart disease [17]. Also, Coral-Alvarado et al. found significantly increased sEng levels in the plasma of patients with systemic sclerosis combined with PAH compared to the healthy controls [75]. Moreover, Bakouboula et al. tested peripheral venous

blood from healthy donors, blood from the jugular vein, and occluded pulmonary artery blood from patients with PAH. They found a significantly increased concentration of sEng in circulating procoagulant microparticles from patients with PAH compared to healthy donors. The concentration of sEng in occluded artery blood was even higher than the concentration of sEng in microparticles from the jugular vein of patients with PAH, suggesting the production of sEng in impaired pulmonary endothelial cells [76]. These data suggest that sEng might be associated with the changes of blood pressure; however, the precise mechanism remains to be elucidated.

Soluble endoglin as an inducer of cardiometabolic disorders

Venkatesha et al. showed that placenta-derived sEng from pregnant preeclamptic women is able to inhibit capillary tube formation *in vitro* and increase vascular permeability and induce arterial hypertension *in vivo*. sEng was able to inhibit TGF β 1 binding and signaling in endothelial cells, which resulted in decreased activation of eNOS and impaired vasodilatation in isolated rat renal microvessels and mesenteric vessels. Authors concluded that sEng plays an essential role in the pathogenesis of arterial hypertension, proteinuria, glomerular endotheliosis, and HELLP syndrome [77]. Walshe et al. demonstrated that sEng (using adenovirus expression of sEng) increased the expression of P-selectin, promoted leukocyte rolling on endothelium, and elevated levels of soluble E-selectin, soluble VCAM-1, and impaired endothelial dependence vasodilation [78]. Besides, in an experimental model of preeclampsia, transgenic mice overexpressing human sEng (sEng levels higher than 2000 ng/ml) had higher systolic blood pressure when compared to wild-type littermates [14]. Furthermore, we showed that sEng treatment induced inflammation (represented by increased NF- κ B and IL6) in HUVECs, suggesting that sEng has a proinflammatory potential, as well [79].

Based on these studies, we might propose that sEng promotes the development of endothelial dysfunction, which might contribute to an increase of systemic blood pressure, suggesting that high levels of sEng promote metabolic syndrome (Fig. 3). However, how exactly sEng contributes to the development of metabolic syndrome is still unknown.

Some studies are showing that sEng might have beneficial effects on fibrosis development. Kapur et al. showed that sEng was able to prevent cardiac fibrosis by decreasing collagen synthesis in human cardiac fibroblasts. They demonstrated *in vivo* the ability of sEng injection to decrease cardiac fibrosis via pSmad 2/3 inhibition in mice after thoracic aortic constriction [80]. However, it is of interest to mention that fibrotic process with respect to ALK1/Smad 1/5

and ALK5/Smad 2/3 pathways is controversial. Most studies suggest the protective antifibrotic role of ALK1/Smad 1/5 pathway, which counteracts the profibrotic ALK5/Smad 2/3 pathway [81].

On the other hand, Pannu et al. demonstrated the profibrotic response of systemic sclerosis fibroblasts on the persistent activation of SMAD1 and Erk-1/2 pathway [82]. Also, in hepatic stellate cells, renal fibroblasts, and mesangial cells, increased activation of Smad 1/5 resulted in increased production of extracellular matrix proteins and progression of fibrosis [83]. Indeed, this topic was nicely revised by Muñoz-Félix et al. [81] and Dituri et al. [84] recently.

In addition, it is of interest to mention that dimeric sEng has been considered to be an inhibitory ligand trap for BMP9. Moreover, Breikopf-Heinlein et al. showed profibrogenic actions of BMP9, suggesting that BMP9 promotes liver fibrogenesis under damage, whereas the absence or inhibition of BMP9 promotes wound healing and liver repairment. This might suggest sEng involvement in liver fibrosis and/or regeneration [85]. On the contrary, Lawera et al. demonstrated that majority of sEng in blood plasma is in monomeric form, and this one does not act as an inhibitory ligand trap for BMP9 signaling [86]. Thus, the interaction of sEng and BMP9 with respect to fibrosis remains to be elucidated.

Reduced inflammatory response after proinflammatory stimuli (LPS, Carrageenan) and ischemia *in vivo* was demonstrated in transgenic mice model expressing human sEng (high sEng mice) by Ruiz-Remolina et al. In addition, they showed that morphological changes after ischemia–reperfusion of lung and kidney were decreased, as well as leukocyte recruitment in high sEng mice [87]. Moreover, it was shown that decreased leukocyte recruitment in high sEng mice might suggest binding of leukocytes integrin α 5 β 1 with sEng RGD motif. This should result in limited availability of leukocyte integrins for the binding of Eng, thus resulting in decreased transmigration of these cells through endothelium in high sEng mice [11]. Thus, these data show the potential anti-inflammatory effects of sEng. However, it is of interest to mention that the above-mentioned studies were performed in various experimental conditions, which most likely contribute to these contradictory results with respect to sEng.

Our experimental group focused on the role of sEng effects in the aorta, a blood vessel prone to the development of endothelial dysfunction and atherosclerosis, which is part of metabolic syndrome.

It was demonstrated that high levels of sEng did not affect aortic endothelial function either at the protein or at the functional level, suggesting possibly no contribution to endothelial dysfunction when sEng is operating as a single factor [88]. In general, it is accepted that endothelial dysfunction and atherosclerosis “requires” hypercholesterolemia. Thus,

we combined high sEng levels with high-fat diet (HFD) administration for 3 months. The results showed that this combination induced proinflammatory (increased expression of P-selectin, ICAM-1, phosphorylated NF- κ B and COX-2) and oxidative stress (increased expression of HO-1, NOX-1, and NOX-2) phenotype in aortic endothelium (without any visible atherosclerosis). Surprisingly, endothelium-dependent vasodilatation induced by acetylcholine was preserved better in mice with high sEng levels, suggesting activation of a compensatory mechanism in endothelium [89]. Once we increased the exposure to sEng and HFD to 6 months, we showed significant aggravation of endothelial dysfunction characterized by reduced Eng, p-eNOS/eNOS, pSMAD2/3/SMAD2/3 signaling pathway only in mice with high sEng levels.

Thus, we might suggest that sEng, especially combined with other risk factors of metabolic syndrome, might be considered as a risk factor for the development of endothelial dysfunction/inflammation and possibly atherosclerosis (Fig. 3).

Recently, Gallardo-Vara et al. demonstrated that sEng induces the expression of BMP4 in endothelial cells in various organs, suggesting BMP4 is a downstream mediator of sEng, responsible for the development of systemic arterial hypertension in these mice [90] (Fig. 3).

Hypercholesterolemia-induced endothelial dysfunction and atherogenesis reduce Eng expression, which is related to the development of endothelial dysfunction. Risk factors that determine the development of metabolic syndrome are associated with increased levels of sEng levels, which might be considered as biomarkers of metabolic syndrome-related pathological disorders. In addition, sEng was shown to aggravate existing endothelial dysfunction and induce the development of systemic arterial hypertension. Adapted from [10, 14, 15, 21, 30, 31, 63, 67, 77, 90, 91]

During this review, we noticed that sEng levels are increased in metabolic syndrome, mostly related to hypercholesterolemia. As the liver plays an important role in cholesterol metabolism *in vivo*, it became interesting to explore the potential impact of high sEng levels on cholesterol and bile acid metabolism in the liver [92].

An experiment in healthy mice showed that the presence of high levels of sEng resulted in increased cholesterol content in the liver due to increased hepatic import of LDL and total cholesterol in these mice. Despite that, it is important to mention that this modulation occurred within physiological levels, since all mice were normocholesterolemic. The presence of sEng also resulted in increased conversion

of cholesterol into bile acids via upregulation of Cyp7a1, increased Mdr1 expression, and additionally increased biliary elimination of bile acids coupled with choleric activity [92].

Taken together, we may speculate that the potential impact of sEng on bile acid metabolism should be considered in patients with metabolic syndrome when they are treated with pharmaceuticals that require hepatic metabolism.

Membrane and soluble endoglin changes after the treatment

So far, we focused on the possible role of Eng and sEng in the pathological conditions related to metabolic syndrome. We showed that both Eng and sEng are involved in the process, so now we would like to focus only on studies mentioning drugs that are used for the treatment of metabolic syndrome-related pathologies such as hypercholesterolemia, diabetes mellitus, and arterial hypertension with respect to Eng and sEng. The summary of all papers is demonstrated in Table 1.

Conclusion

In conclusion, we propose that reduced Eng expression is a hallmark of endothelial dysfunction development in chronic disorders associated with metabolic syndrome. On the other hand, Eng also plays an essential role in leukocyte transmigration and acute inflammation, suggesting that Eng is crucial for the regulation of endothelial function during the acute phase of vascular defense reaction to harmful conditions. sEng is a circulating biomarker of metabolic syndrome-related symptoms and pathologies, including hypercholesterolemia, hyperglycemia, diabetes mellitus, arterial hypertension, and liver damage. In addition, sEng is able to participate in the aggravation of endothelial dysfunction and promote the development of arterial hypertension, suggesting that high levels of sEng promote metabolic syndrome symptoms and its complications. Therefore, we suggest that the treatment of metabolic syndrome should take into account the importance of Eng in the endothelial function and levels of sEng as a biomarker and risk factor of related pathologies.

Table 1 Effect of selected drugs on Eng expression and sEng levels

Drug	Effect/impact	References
Atorvastatin	Decreased cholesterol levels, plaque size, and sEng plasma concentration and simultaneously increased the expression of Eng and eNOS in the aorta of hypercholesterolemic mice with atherosclerosis, suggesting a beneficial effect of Eng during atherogenesis	[46, 93]
	Increased Eng, SMAD2, eNOS expression, and decrease plaque size, without lipid-lowering effects suggesting a beneficial effect of Eng during atherogenesis	[44]
	Increased expression of Eng in HUVECs	[94]
	Prevented inflammation-induced decrease of Eng and eNOS expression in endothelial cells, suggesting a beneficial effect of Eng against the development of endothelial dysfunction	[95]
	Inhibited Eng and collagen type I formation and protein expression induced by TGF β 1 in the heart, suggesting endoglin involvement in pathological fibrosis in myocardium	[96]
Simvastatin, Rosuvastatin, Pravastatin	Increased sEng secretion from HUVECs but not from placental explants showing potential statin effect on release of sEng	[97]
Rosiglitazone	Reversed the effect of high glucose treatment on sEng levels, suggesting that improvement of insulin resistance in diabetes mellitus type II also results in reduced levels of sEng	[71, 72]
GLP-1 agonists	Were demonstrated to reduce sEng and MMP14 levels in patients with diabetes mellitus type I, suggesting that sEng-level changes reflect diabetic changes that can be affected by anti-diabetic drugs	[70]
Metformin	Decreased sEng release from endothelial cells and villous cytotrophoblast cells, but does not change sEng concentration from preterm preeclamptic placental villous explants suggesting the potential effect of the drug on sEng levels	[97, 98]
Perindopril	Significantly lower the concentration of sEng compared to patients treated with different antihypertensive agents and healthy volunteers, suggesting sEng levels might be increased during arterial hypertension	[99]
Resveratrol	Decreased sEng release from endothelial cells by decreasing MMP14 activity together with decreased endothelin-1 expression and increases phosphorylation of eNOS, suggesting that beneficial effects on endothelial dysfunction can be related to reduced sEng levels	[100]
Carotuximab	Affected Eng signaling by activation of SMAD2/3 signaling, while SMAD1/5/8 signaling was reduced, suggesting a capacity to affect cardiometabolic effects of Eng in various conditions	[101]
	Increased release of sEng in the conditioned media of HUVECs and HMEC-1, most likely via activation of MMP14 suggesting a potential influence on sEng levels	[102]

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Compliance with ethical standards

Conflict of interest No conflicts of interest, financial or otherwise, are declared by the authors.

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