

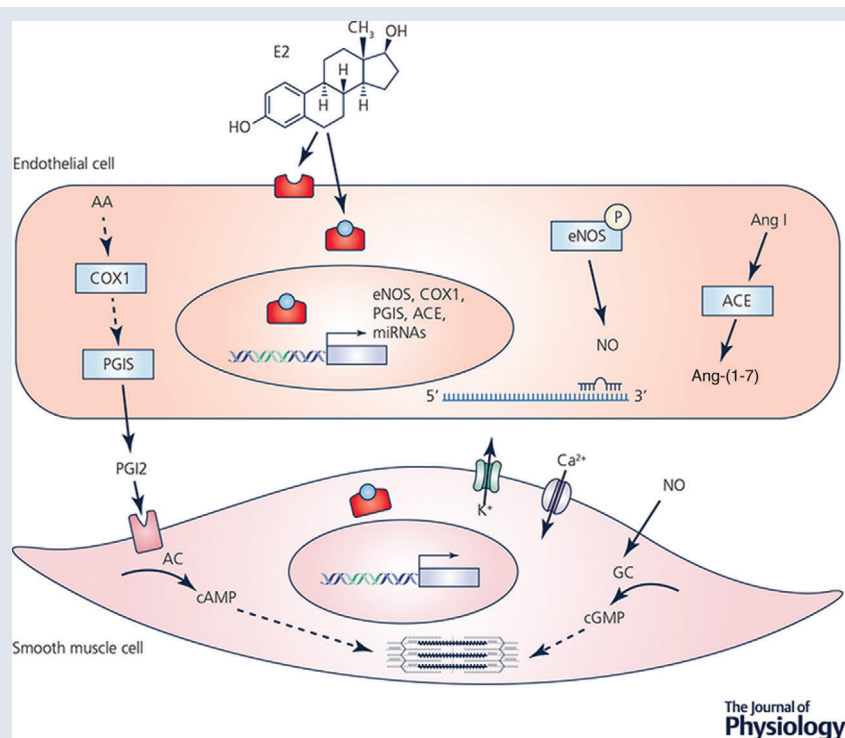
SYMPOSIUM REVIEW

Mechanisms underlying the influence of oestrogen on cardiovascular physiology in women

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Abstract Women show a lower incidence of cardiovascular diseases than age-matched men, but this benefit disappears after menopause. Oestrogen-mediated vascular actions are mainly attributed to oestradiol and exerted by oestrogen receptors ($ER\alpha$, $ER\beta$ and G protein-coupled oestrogen receptor), through rapid and/or genomic mechanisms, but these effects depend on ageing and inflammation. A cardiovascular approach in women's health has arisen due to

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controversy regarding oestrogen's beneficial impact as reported in experimental and observational studies and large randomized trials. These can be explained, in part, by two mutually non-exclusive hypotheses. On the one hand, the timing hypothesis, which states that oestrogen-mediated benefits occur before the detrimental effects of ageing are established in the vasculature; on the other hand, ageing and/or hormonal-associated changes in ER expression that could lead to a deleterious imbalance in favour of ER β over ER α , generally associated with higher inflammation and endothelial dysfunction. In experimental studies, oestradiol acting on ER α promotes the release of vasoactive compounds such as nitric oxide (NO) and prostacyclin, and shifts the angiotensin axis towards angiotensin 1–7 production. Mechanisms underlying oestradiol vascular function also include anti-inflammatory and epigenetic modifications. 17 β -Oestradiol changes the transcriptomic profile of endothelial cells, and the involvement of miRNA in the regulatory pathways of vascular function reinforces assumptions regarding the vascular actions of oestrogen. Thus, the present Symposium Review aims to postulate the role of ER α in oestrogen modulation of endothelium-derived mediators and vascular physiology, as well as its relationship with miRNA and inflammation, and elucidate how physiological changes in postmenopausal women counteract the observed effects.

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Abstract figure legend The beneficial effects mediated by oestrogen involve different intracellular signalling pathways, such as nitric oxide (NO), prostanoids and the renin–angiotensin system (RAS), towards a vasoprotective profile involving oestrogen receptors, mainly ER α . Physiological changes as ageing and menopause and in epigenomics affect the cardiovascular effects of oestrogen. (Created by Biorender.com.)

In-depth study of cardiovascular diseases (CVD) has led to a fuller understanding of sex differences in cardiovascular physiology. CVD is currently the leading cause of death in women from developed countries (WHO, 2016) although statistical data reveal that women develop CVD 10 years later than men (Burns & Korach, 2012), and incidence increases from menopause on (Deroo & Korach, 2006; Burns & Korach, 2012; Vrtacnik *et al.* 2014). Women's time-related advantage regarding CVD development has been attributed to hormonal status, and both clinical and experimental data have demonstrated the beneficial effects of oestrogen at the cardiovascular level (Hayward *et al.* 2000; Mendelsohn & Karas, 2005). However, hormonal replacement therapies (HRT) have been used in postmenopausal women with controversial findings (Mendelsohn & Karas, 1999; Mikkola *et al.* 2013). While the largest randomized controlled trial, Women's Health Initiative (WHI), initially reported no protective role against coronary heart disease risk (Rossouw *et al.* 2002), a reanalysis by age and years since menopause (Rossouw *et al.* 2007) demonstrated a significant benefit in healthy women initiating oestrogen therapy soon after menopause onset (Manson *et al.* 2003; Rossouw *et al.* 2007; Novella *et al.* 2012). In fact, age and years since menopause are important variables affecting the benefit/risk profile of HRT (Sood *et al.* 2014). The so-called timing hypothesis postulates that the beneficial impact

of hormonal replacement in CVD prevention can occur only when HRT is initiated before the detrimental effects of ageing on the cardiovascular system have become established (Clarkson *et al.* 2013). In this regard, it has been reported that age moderates oestrogen's vasodilatory (Sherwood *et al.* 2007) and anti-inflammatory (Novella *et al.* 2012) effect on vascular tissue in postmenopausal women. The current consensus on HRT is that the cardiovascular protective role of oestrogen depends on the timing of treatment after menopause (Lobo, 2017).

Since the publication of the WHI results in 2002, much has been learned, yet much controversy remains. The 2017 position statement of the North American Menopause Society (NAMS), which evaluates new literature and reaches consensus on recommendations for the use of HRT for the treatment of menopause-related symptoms, identified future research needs as the risks of HRT differ depending not only on timing of initiation but also on type, dose, duration of use, route of administration and whether a progestogen is needed (Hormone Therapy Position Statement Advisory Panel, 2017). In agreement with the timing hypothesis, the position statement of NAMS assessed that for women aged younger than 60 years or who are within 10 years of menopause onset, HRT appears favourable for treatment of some menopausal symptoms, but for those who initiate HRT more than 10 or 20 years from menopause onset or when aged

60 years or older, the benefit–risk ratio appears less favourable than for younger women, with greater absolute risks of coronary heart disease, stroke, venous thromboembolism and dementia.

Oestrogen receptors in the cardiovascular system

The most abundant form of circulating oestrogen is oestradiol, also termed 17β -oestradiol, which is predominantly synthesized and secreted by the ovaries during a woman's reproductive years. Vascular tissues, particularly endothelial cells, vascular smooth muscle cells (VSMCs) and cardiomyocytes, are oestradiol targets as they express different types of oestrogen receptors (ERs) (Khalil, 2013). This expression is also shared by monocytes, macrophages and dendritic cells, suggesting a modulatory role for oestradiol in inflammatory processes, a key event in onset and development of CVD (Harkonen & Vaananen, 2006; Kovats, 2015). Oestradiol binds to classical ERs including both $ER\alpha$ and $ER\beta$ in cytoplasm, to create homo- or heterodimers. They then bind to specific DNA motifs called oestrogen response elements (EREs) in the promoter region of oestrogen-responsive genes to regulate transcription (Klinge, 2001) and induce changes in gene expression. $ER\alpha$ and $ER\beta$ have different distributions, and selective activation of either the $ER\alpha$ or the $ER\beta$ isoform may involve contrasting biological effects, having opposing gene-expression regulatory effects (Lindberg *et al.* 2003; Tsutsumi *et al.* 2008) or alternatively having redundant mediatory roles (Arias-Loza *et al.* 2007; Lahm *et al.* 2008). Oestrogen signalling is thus selectively regulated by the relative balance between $ER\alpha$ and $ER\beta$ expression in target organs (Murphy & Steenbergen, 2014), although studies using $ER\alpha$ and $ER\beta$ knockout mice revealed that the beneficial effects oestrogen has on the vascular system are mainly mediated by $ER\alpha$ (Pare *et al.* 2002; Arnal *et al.* 2017).

Besides this classical genomic action, oestradiol also binds to membrane-bound $ER\alpha$ and $ER\beta$ receptors as well as to G protein-coupled ER (GPER) (Levin, 2009), rapidly activating nuclear transcription factors and triggering faster responses (within minutes). Many of the effects of oestrogen seen in human and animal models, such as reduced myocardial pro-inflammatory cytokine expression, inhibition of VSMC proliferation, and nitric oxide (NO)-dependent vasodilatation (Prossnitz & Barton, 2011), have recently been attributed to GPER expression in the cardiovascular system (Revankar *et al.* 2005).

The present overview is a Symposium Review presented in Europhysiology 2018, partially based on our own results, and aims to highlight the role of $ER\alpha$ in oestrogen modulation of endothelial-derived mediators and vascular physiology, and how physiological changes in postmenopausal women counteract the observed effects.

Vascular protective effects of oestrogen through $ER\alpha$

The vascular-protective impact of oestrogen has also been attributed to its effects on the vascular wall, in both endothelium and smooth muscle, releasing vasoactive-mediators which promote arterial vasodilatation, modulate inflammatory processes and regulate systemic lipid metabolism and oxidative-stress balance (Kondo *et al.* 2009; Barton, 2013; Usselman *et al.* 2016). Figure 1 summarizes the role of $ER\alpha$ in endothelium-derived mediator and vascular smooth muscle cell function. Next the effects on these mediators, in particular NO, prostacyclin and angiotensin 1–7 pathways, will be discussed (some of the actions are summarized in Table 1).

Oestrogen and nitric oxide. In endothelial cells, which form the luminal cell monolayer of the vascular wall, oestradiol modulates the release of multiple vasoactive substances via both genomic and non-genomic action. Oestradiol increases NO bioavailability by either directly increasing NO generation or decreasing NO inactivation. Oestrogen increases NO bioavailability by mechanisms such as increasing endothelial NO synthase (eNOS) gene expression at the transcriptional level (Sumi & Ignarro, 2003); non-genomic and rapid activation of enzyme activity via cascades that activate kinases c-Src (Haynes *et al.* 2003), extracellular signal-regulated kinase (ERK) (Chen *et al.* 2004), phosphoinositide 3-kinase (PI3K) (Simoncini *et al.* 2003), and Akt, which leads to eNOS activation through phosphorylation at residue Ser1177 (Haynes *et al.* 2000; Meyer *et al.* 2009); increasing intracellular free Ca^{2+} concentration in endothelial cells (Rubio-Gayosso *et al.* 2000); regulating endogenous inhibitors and cellular location (Chambliss & Shaul, 2002; Monsalve *et al.* 2007; Novella *et al.* 2013); and attenuating superoxide anion (O_2^-) concentration, thereby decreasing O_2^- -mediated NO inactivation (Wassmann *et al.* 2001; Dantas *et al.* 2002; Ospina *et al.* 2002). Some of these rapid effects of oestradiol on the NO signalling pathway require no changes in gene expression and are mediated by different plasma membrane-associated ER subtypes. In addition to full-length $ER\alpha$ (ER66), an N-terminal truncated $ER\alpha$ isoform, ER46, plays a key role in these endothelial responses to oestradiol (Kim *et al.* 2014). Besides this, recent findings reveal that a GPER-mediated cascade acts as an alternative pathway in oestradiol-induced endothelium-dependent vasodilatation and NO formation via c-Src/PI3K signalling pathways (Fredette *et al.* 2018). Thus, in physiological conditions oestradiol stimulates vascular NO formation via GPER and mainly through $ER\alpha$, which acts at a vascular level as a potent vasodilator, but also conveys vasoprotection through antithrombotic mechanisms and modifies proliferation and migration of the underlying

VSMC (Förstermann & Sessa, 2012), thereby controlling the vascular tone.

In processes related to vascular injury associated with inflammation, $ER\beta$ expression in endothelium increases the expression of superoxide dismutase (SOD2) and eNOS, which altogether also raise NO bioavailability, ameliorating ischaemia–reperfusion-mediated vascular injury and minimizing reactive oxygen species generation (Zhan *et al.* 2016). On the other hand, pharmacological activation of the $ER\beta$ increases the expression of cytokine-driven inducible NO synthase (iNOS) in rat vascular smooth muscle (Panic *et al.* 2018), raising the hypothesis that $ER\beta$ can be induced by injuries and contributes to inflammation (Sartoretto *et al.* 2019). In non-vascular cells, $ER\beta$ activation also increases levels of phosphorylated neuronal NO synthase (nNOS) and NO

production through a Src/PI3K/Akt-dependent pathway in hypothalamic neurons (Gingerich & Krukoff, 2008).

The effect of oestradiol on the NO pathway observed in cultured cells has been confirmed in a large number of isolated blood vessel preparations including the rat aorta (Freay *et al.* 1997), rat femoral artery and rat portal vein (Kitazawa *et al.* 1997), rabbit coronary artery (Jiang *et al.* 1991) and porcine coronary artery (Teoh *et al.* 1999). Although oestradiol's mechanism of action differs according to the vascular bed and species studied, in general $ER\alpha$, $ER\beta$ and GPER all seem to contribute. Generally, oestradiol exposure in women increases vascular relaxation and endothelial-dependent vasodilatation, increasing blood flow in numerous vascular beds. In studies performed in healthy young women, oestradiol is also associated with increases in

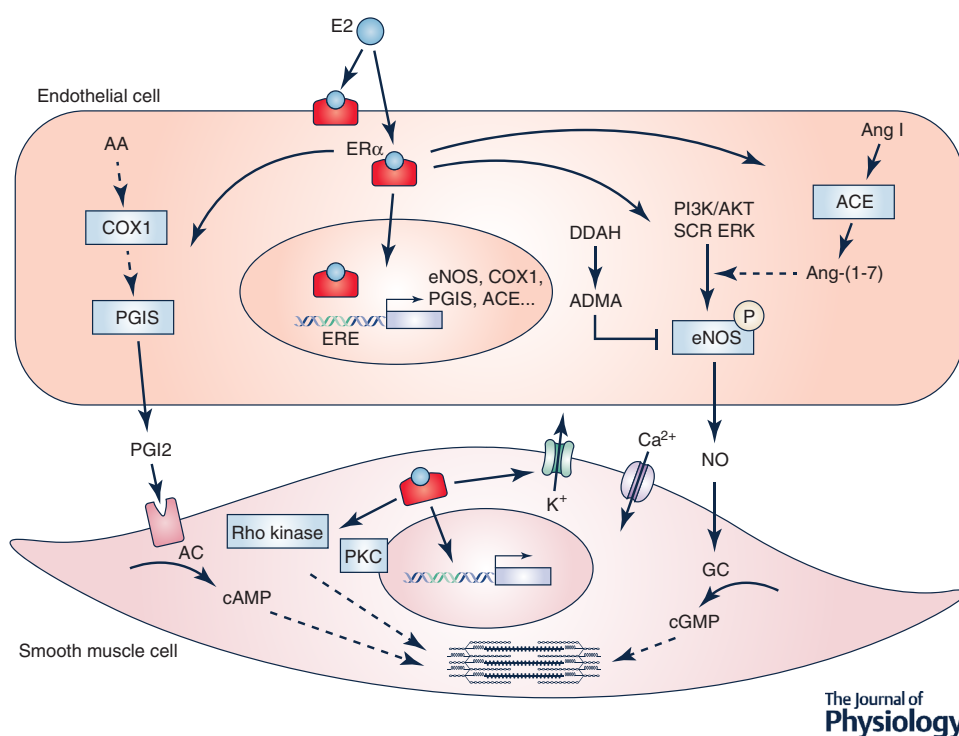


Figure 1. Role of $ER\alpha$ on endothelium-derived mediators and vascular smooth muscle cell function

Oestradiol (E2) binds to oestrogen receptor α ($ER\alpha$) triggering both genomic and cytoplasmic response. E2- ER complex is translocated to the nucleus and induces transcription of specific genes by binding to oestrogen response elements (ERE) in their promoter region. Endothelial nitric oxide synthase (eNOS), cyclooxygenase 1 (COX1), prostacyclin synthase (PGIS) and angiotensin converting enzymes (ECA) are regulated transcriptionally by $ER\alpha$. In addition to its genomic effect, E2- $ER\alpha$ also regulate eNOS activity by inducing phosphorylation through different kinase signalling pathways (PI3K/AKT, SCR, ERK), and reducing the endogenous inhibitor asymmetric dimethylarginine (ADMA) by regulating dimethylarginine dimethylaminohydrolase (DDAH). Moreover, E2- $ER\alpha$ enhances activity of angiotensin converting enzymes (ECA), increasing the production of angiotensin 1–7, and plays a role in NO-dependent vasodilatation through a mechanism that involves Mas receptor. As a result, NO diffuses into the vascular smooth muscle cells and binds to guanylate cyclase (GC), increasing cGMP that in turn cause relaxation. E2 also increase prostacyclin (PGI_2) production through the COX1–PGIS pathway. PGI_2 is released from endothelial cells and binds to specific receptors located in the membrane of smooth muscle cells, which leads to an increment of cAMP by adenylate cyclase (AC) and muscle relaxation. E2 can also interfere with different signalling pathways, such as protein kinase C (PKC) and Rho-kinase and membrane ion channel activity through non-genomic actions. Altogether, these mechanisms lead to oestrogen-mediated vascular relaxation. (Created by BioRender.com.)

Table 1. Vascular protective factors mediated by oestradiol

Vasoactive-mediator production	Mechanism	Experimental model	Reference
↑ NO bioavailability	↑ eNOS activation ER α	EA.hy926 cells	Haynes <i>et al.</i> (2003)
	↑ eNOS expression Oestrogen-related receptor α 1 (ERR α 1)	COS-7 cells and bovine pulmonary artery endothelial cells	Sumi & Ignarro, (2003)
	↑ NO production GPER	Human endothelial cells, TIVE cells	Fredette <i>et al.</i> (2018)
	↓ O $_2^-$ -mediated NO inactivation ↑ DDAH expression and ↓ the endogenous L-arginine analogue ADMA	Mesenteric arteries from SHR HUVECs	Dantas <i>et al.</i> (2002) Monsalve <i>et al.</i> (2007)
	↑ DDAH activity and ↓ the endogenous L-arginine analogue ADMA ER α	HUAECs	Novella <i>et al.</i> (2013)
↑ PGI $_2$	↑ COX-1 and PGIS expression ER α	HUVECs	Sobrino <i>et al.</i> (2010)
↓ ET-1	↓ ET-1 expression and secretion ER α	Mesenteric arteries from DOCA hypertensive rats	David <i>et al.</i> (2001)
	↓ ET-1 expression and secretion ER-independent mechanism	Porcine coronary artery endothelial cells	Dubey <i>et al.</i> (2001)
↑ Ang-(1–7)	↑ ACE2 activity and expression	Renal wrap model of hypertension	Ji <i>et al.</i> (2008)
	↑ ACE2 activity and expression ER α	HUVECs	Mompeón <i>et al.</i> (2016)

Oestradiol mediates the release of vasoactive mediators mainly from endothelium, some of which effects are summarized in the following table along with the oestrogen-receptor involved. ACE2, angiotensin-converting enzyme 2; ADMA, asymmetric dimethylarginine; Ang-(1–7), angiotensin 1–7; COX-1, cyclooxygenase 1; DDAH, dimethylarginine dimethylaminohydrolase; DOCA, deoxycorticosterone acetate; ER, oestrogen receptor; ET-1, endothelin-1; GPER, G protein-coupled oestrogen receptor; HUAEC, human umbilical artery endothelial cell; HUVEC, human umbilical vein endothelial cell; PGI $_2$, prostacyclin; PGIS, PGI $_2$ synthase; SHR, spontaneously hypertensive rat; TIVE, telomerase-immortalized human umbilical vein endothelial.

flow-mediated dilatation (FMD), a measure of conduit artery endothelial function mediated primarily by NO (Adler *et al.* 2018). When oestrogen levels declines in postmenopausal women, oestradiol administration improves also endothelial function (Hurtado *et al.* 2016) but the magnitude of improvement depends on the timing of when this treatment is initiated. The interactions of oestrogens on multiple pathways regulating vascular function, which also are involved in the ageing process, are complex, multifactorial and not completely understood. For example, the expression of ER α and eNOS in endothelial cells harvested from peripheral veins of women are lower in postmenopausal women than in young women (Gavin *et al.* 2009). Thus, not only does oestradiol decline with ageing, ER α receptor expression also declines, and an increase in oxidative stress is produced as well. How female and male sex hormones interact with the cardiovascular system, and in age-associated endothelial dysfunction in healthy woman and men has been recently reviewed in depth (Stanhewicz *et al.* 2018).

Oestrogen and cyclooxygenases. Cyclooxygenase (COX)-derived factors are particularly important in regulating vascular tone as they can induce both vascular relaxation (through prostacyclin (PGI $_2$) production) and contraction (through thromboxane A $_2$ (TXA $_2$) and prostaglandin H $_2$ (PGH $_2$) production). Oestradiol has been implicated in the modulation of peripheral vascular synthesis of vasodilatory mediators, including prostanoids through COX, as the rate-limiting step in the formation of vasoactive prostanoids (Sobrino *et al.* 2009). In human endothelial cells, oestradiol, acting through ER α , induces stimulation of the vasodilator and antiaggregatory PGI $_2$ production by up-regulating COX-1 and PGI $_2$ synthase (PGIS) expression without altering vasoconstrictor TXA $_2$ production (Sobrino *et al.* 2010). This mechanism supports the hypothesis that oestradiol is able to maintain vascular health and protect endothelial cells against vascular disorders (Mikkola *et al.* 2013). The beneficial effects of oestrogen on the endothelium can also be partially explained by an inhibitory effect

on production of the COX-derived vasoconstrictor agents PGH₂ and TXA₂ (Davidge & Zhang, 1998; Dantas *et al.* 1999; Vidal-Gómez *et al.* 2016), and endothelin-1 (David *et al.* 2001; Dubey *et al.* 2001), tipping prostanoid balance toward increased PGI₂ production. However, in the absence of oestrogen, arachidonic acid is actively converted to a COX-1-dependent constrictor, indicating that oestrogen-mediated elevation in COX-1 and PGI₂ synthase appears to shift the balance of prostanoid products from constrictor to dilator (Ospina *et al.* 2002). These effects observed in cultured endothelial cells have been also observed in cerebral blood vessels from ovariectomized rats, where oestradiol increases protein levels in both COX-1 and PGIS and up-regulates the production of PGI₂, promoting increased cerebral perfusion and conferring resistance against thrombotic events (Ospina *et al.* 2002). Apart from this, ER β has also been associated with COX-2 expression and both PGI₂ and TXA₂ concentrations at basal state, which suggests the possibility of a ligand-independent regulation of COX-2 activity and PGH₂ substrate availability (Su *et al.* 2009). GPER also mediates oestrogen-dependent inhibition of endothelium-derived vasoconstrictor prostanoid production and activity under pro-inflammatory conditions, providing evidence for a novel mechanism through which GPER could inhibit vascular tone and inflammation (Meyer *et al.* 2015).

In addition to regulating endothelium-derived factors, oestradiol directly regulates the smooth muscle layer by inhibiting VSMC proliferation, migration and vascular contraction (Suzuki *et al.* 1996). Indeed, oestrogen-mediated relaxation can also occur in endothelium-denuded segments (Mugge *et al.* 1993). Several mechanisms, involving among others ion channels and kinase cascades, have been proposed to explain this vasorelaxant effect. Oestrogen can interfere with ion channels through non-genomic actions and decrease smooth muscle constriction by interfering with Ca²⁺ mobilization and Ca²⁺ entry responses (Crews & Khalil, 1999) and activating K⁺ channels (White *et al.* 2002), leading to membrane hyperpolarization and vascular relaxation. The role of ER has been studied in female rat mesenteric microvessels, where ER subtypes mediate distinct vasodilatation and decreased intracellular Ca²⁺ (mainly through ER α , with both ER β and GPER being also implicated) through endothelium- and K⁺ channel-independent inhibition of Ca²⁺ entry mechanisms of VSMC contraction (Mazzuca *et al.* 2015). Direct interaction of oestradiol with voltage-gated Maxi-K channel subunit β , which confers higher Ca²⁺ sensitivity, may modulate vascular smooth muscle (Valverde *et al.* 1999). Oestrogen can also modulate vasoconstriction by interfering with protein kinase C (Kanashiro & Khalil, 2001) and Rho-kinase signalling in VSMC (Hiroki *et al.* 2005).

Oestrogen and angiotensin 1–7. Additionally, oestradiol is able to modulate the renin–angiotensin system (RAS) (Farhat *et al.* 1996; Alvarez *et al.* 2002), which plays a pivotal role in physiological regulation of blood volume and blood pressure and is involved in controlling vascular contractility. Renin released from the kidney converts angiotensinogen from the liver to the decapeptide angiotensin-I (Ang I), which undergoes proteolytic cleavage, through activating angiotensin-converting enzyme (ACE) to generate angiotensin-II (Ang II). The discovery of angiotensin-converting enzyme 2 (ACE2), which cleaves COOH-terminal residues from Ang I and II, producing primarily vasoprotective angiotensin 1–7 (Ang-(1–7)), suggested that RAS involves two axes: (1) Ang II, which mediates vasoconstriction and remodelling effects through receptor type 1 (AT1R) while exert opposing effects through Ang II receptor type 2 (AT2R), and (2) Ang-(1–7), which acts as a protective and vasodilator pathway acting on the Mas receptor. Changes in Ang II/Ang-(1–7) balance are therefore essential to maintain cardiovascular homeostasis (Jiang *et al.* 2014).

Evidence indicates that components of the RAS are markedly affected by oestrogen (Sullivan, 2008; Hilliard *et al.* 2013b) shifting the balance towards the ACE2/Ang-(1–7)/Mas and AT2R pathways in females. In general, oestrogen increases the synthesis of circulating angiotensinogen, while decreasing the synthesis of the RAS enzymes renin and ACE (Fischer *et al.* 2002; Komukai *et al.* 2010). Accordingly, sex differences in vascular RAS mechanisms have commonly been assumed to play a role in the relative protection against CVD in premenopausal women. Circulating plasma Ang-(1–7) concentrations have been reported to be higher in healthy premenopausal women than in healthy men of a similar age (Sullivan *et al.* 2015), and its relationship with oestrogen is underscored in studies showing an increase of urinary Ang 1–7 levels among pregnant women (Valdes *et al.* 2001). Vascular AT1R expression in ovariectomized rats treated with oestradiol is down-regulated (Nickenig *et al.* 1998; Rogers *et al.* 2007). There is also evidence that AT2R plays a protective role by regulating blood pressure in female mice (Armando *et al.* 2002; Brown *et al.* 2012), rats (Sampson *et al.* 2012) and women due to its up-regulation by oestrogen (Hilliard *et al.* 2013a). Furthermore, there is evidence that ER α is involved in oestradiol-mediated effects on RAS as primarily responsible for oestrogen regulation of kidney ACE2, AT1R and AT2R genes in ovariectomized mice (Brosnihan *et al.* 2008), which reinforces the central role of ER α in oestrogen's beneficial impact on cardiovascular physiology.

Aside from the classical circulating RAS pathway, the intracellular RAS described as the 'non-classical' RAS pathway has gained attention for its ability to antagonize classical RAS signalling. RAS components are also expressed in the heart and vascular wall, and control

vascular tone and arterial structure (Nguyen Dinh Cat & Touyz, 2011). Oestrogen also regulates tissue RAS, in that oestradiol diminishes cardiac ACE expression in human atrial tissue, while simultaneously inducing ACE2, which counteracts the classical RAS activity towards the vasodilator pathway. This ACE2 induction is prevented by the ER α antagonist, suggesting a role for ER α in mediating the cardiovascular protective effects of oestrogen (Bukowska *et al.* 2017). Enhanced ACE2 activity and expression have also been reported in the kidney and uterus of experimental animals during pregnancy (Joyner *et al.* 2007; Neves *et al.* 2008) and in different models of hypertensive rats, where both ACE (Gallagher *et al.* 1999; Dean *et al.* 2005) and ACE2 tissue expression are decreased by ovariectomy and restored by oestrogen replacement (Ji *et al.* 2008; Shenoy *et al.* 2009). In endothelial cells we reported that oestradiol stimulates the production of Ang-(1–7) via ER α by increasing ACE and ACE2 expression and activity (Mompeón *et al.* 2016), and demonstrated that the Mas receptor plays an essential role in NO-dependent vasodilatation mediated by oestradiol (Sobrino *et al.* 2017). In this regard, the blockade of Mas receptor is equivalent to ER blockade in preventing the effects of oestradiol, indicating crosstalk between oestradiol and the Ang-(1–7)–Mas axis (Sobrino *et al.* 2017). Thus, the loss of cardiovascular protection observed in postmenopausal women could also partly result from the change from the ACE2–RAS protective axis to the classic ACE–RAS pathway (Komukai *et al.* 2010; Hilliard *et al.* 2013b; Stanhewicz *et al.* 2018).

Oestrogen-regulated miRNA

Besides oestrogen-mediated regulation of important cardiovascular pathways through a direct gene transcription mechanism, oestrogen has recently been posited as a modulator of cardiovascular physiology by modifying another group of important gene expression regulators based on epigenetic mechanisms. Among them, miRNAs are small non-coding RNAs that can inhibit gene expression post-transcriptionally via sequence-specific interactions with target genes. In addition, circulating miRNAs found in the blood stream have been proposed as non-invasive biomarkers in CVD (Fichtlscherer *et al.* 2011) and changes in the circulating miRNA profile have been linked to oestrogen levels in women (Pérez-Cremades *et al.* 2018a).

Changes in miRNA levels induced by sex hormones, sex chromosome expression and regulation of key components of miRNA biosynthesis machinery have been described as possible underlying mechanisms of miRNA-mediated regulation of cardiovascular function in women. In this regard, ERs have an important role in the regulation of the miRNA-mediated oestrogen effects. First, they act as a transcription factor, as demonstrated

by differences in miRNA profile between ER⁺ and ER[−] breast cancer cells (Bailey *et al.* 2015; Cizeron-Clairac *et al.* 2015). Indeed, down-regulated miRNAs in ER[−] breast cancer compared to ER⁺ lose their ER binding sites in the promoter region near the miRNA sequence (Bailey *et al.* 2015). At a cardiovascular level, ER binding sites near oestrogen-regulated miRNAs have also been found in VSMC (Zhao *et al.* 2013; Deng *et al.* 2015) and endothelial cells (Vidal-Gomez *et al.* 2018). We recently demonstrated the involvement of different ERs in the expression of oestradiol-regulated miRNAs by using specific ER agonists and antagonists. Although most of the analysed miRNA were regulated by ER α , ER β and GPER were also found to be involved in oestradiol-regulated miRNA expression (Vidal-Gomez *et al.* 2018). In addition, oestradiol can regulate miRNA expression by acting directly on its biosynthesis machinery (Gupta *et al.* 2012). Although oestradiol regulation of miRNA biosynthesis components has been reported mainly in reproductive tissues, for example in differences observed between ER⁺ and ER[−] breast cancer cells (Cheng *et al.* 2009; Cizeron-Clairac *et al.* 2015), transcriptomic data in human endothelial cells treated with physiological concentrations of oestradiol highlight the changes in the expression of specific genes involved in miRNA synthesis (Pérez-Cremades *et al.* 2018b).

The role of specific miRNAs in the regulatory mechanisms of oestrogen in cardiovascular function has recently been reviewed in depth elsewhere (Pérez-Cremades *et al.* 2018a). However, the role of ERs in this effect has been addressed only for certain miRNAs (Fig. 2). For example, miR-203 is one of the dysregulated miRNAs in cultured VSMCs after oestradiol exposure (Zhao *et al.* 2013). It is up-regulated by an ER α -dependent mechanism, but not by ER β , through a transcription activation mechanism mediated by the transcription factors activator protein 1 (AP-1) and Zinc finger E-box-binding homeobox 1 (Zeb-1). Moreover, the authors demonstrated the role of miR-203 in regulating VSMC proliferation, showing that inhibition of miR-203 expression cancelled out the oestradiol-mediated effect on VSMC proliferation through targeting Abl and p63 (Zhao *et al.* 2013). These results suggest that this miRNA is involved in the antiproliferative action of oestrogens on VSMCs and could explain oestrogen-induced inhibition of neointimal formation after vascular damage (Mori *et al.* 2000; Xing *et al.* 2009). In addition, the ER-regulated miRNA, miR-22, contributes to the antioxidant effect of oestrogen on cardiovascular tissues (Wang *et al.* 2015). miR-22 activity is related to cardiac remodelling and hypertrophy (Huang & Wang, 2014). However, oestradiol treatment decreases miR-22 expression in *in vitro* cardiomyocytes and *in vivo* myocardium of ovariectomized mice via ER α -mediated mechanisms. miR-22 down-regulation increases the expression of its

target SP1, a transcription factor that regulates the cytoprotective enzyme cystathionine γ -lyase, as well as H₂O production and antioxidant defence (Wang *et al.* 2015). Taken together, these results may explain in part the female cardioprotection against oxidative stress (Wang *et al.* 2010). Furthermore, miR-22 inhibits oestrogen signalling by targeting ER α , suggesting a reciprocal regulation (Pandey & Picard, 2009). Finally, miR-21 is regulated by oestradiol via ER β -dependent mechanisms in female cardiac tissue (Queiros *et al.* 2013). This miRNA has been implicated in myocardial hypertrophy by regulating mitogen-activated protein kinase (MAPK) signalling in fibroblasts (Thum *et al.* 2008). In this regard, miR-21 is down-regulated in female cardiac fibroblasts exposed to oestradiol and to a specific ER β agonist, and its expression is up-regulated in the left ventricle of ER β knockout female mice (Queiros *et al.* 2013). Oestradiol regulates MAPK signalling through targeting three specific negative regulators of this pro-fibrotic pathway, the negative regulator sprouty homologue 1 (Spry1), RAS p21 protein activator 1/GTPase activating protein 1 (Rasa1) and

Gap1m/RAS p21 protein activator 2 (Rasa2), results that may explain the mechanisms underlying the protective effect of oestrogen on cardiac remodelling.

Effects of oestrogen on inflammation

Female cardiovascular health is a complex issue, since at menopause women face a decrease in oestrogen levels together with an active vascular ageing process. Coupled with known risk factors, findings from epidemiological and experimental studies have closely linked inflammatory processes with vascular ageing (Seals *et al.* 2011). Predominant features of the ageing process are chronic progressive increase in pro-inflammatory status (Najjar *et al.* 2005) and development of a more adhesive endothelium (Csiszar *et al.* 2008), and the process *per se* is known to have a positive association with levels of inflammation biomarkers and increased risk of CVD (Singh & Newman, 2011).

Data regarding the effects of oestrogen on the inflammatory process are contradictory, with both

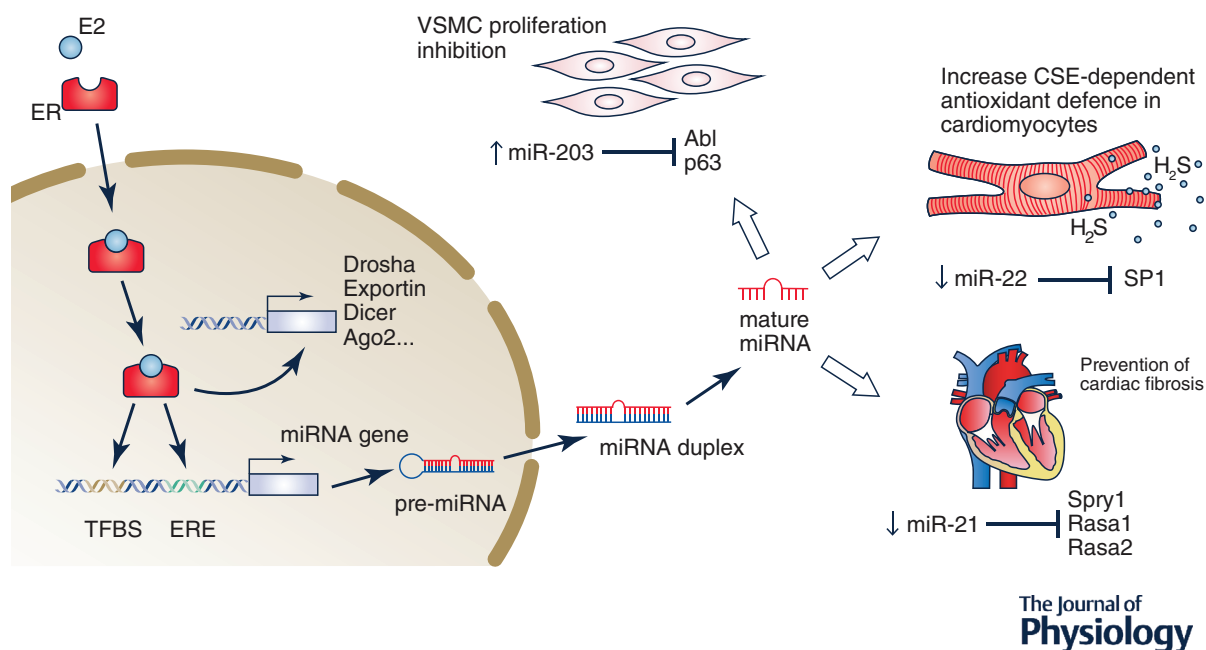


Figure 2. Role of oestrogen receptors in miRNA-dependent regulation of cardiovascular physiology in women

Oestradiol (E2) binds to oestrogen receptor (ER) and regulates miRNA expression directly by binding to specific oestrogen responsive elements (ERE) in the promoter region of the miRNA genes, or indirectly by recruiting different transcription factors that regulate miRNA expression via their transcription factor binding sites (TFBS). E2 can also impact miRNA expression by regulating transcription activity of miRNA biosynthesis components. The role of ER in E2-dependent miRNA regulation in cardiovascular physiology has been described for different miRNA: E2-dependent up-regulation of miR-203 is implicated in vascular smooth muscle cell (VSMC) proliferation through targeting Abl and p63 (Zhao *et al.* 2013); miR-22 down-regulation mediated by E2 in cardiomyocytes induces increased expression of SP1 transcription factor that increase cystathionine γ -lyase (CSE)-dependent H₂S production and antioxidant defence (Wang *et al.* 2015); miR-21 is regulated by E2 via ER β in cardiac tissue, preventing cardiac fibrosis through direct inhibition of three negative regulators of the MAPK signalling pathway (Spry1, Rasa1, Rasa2) (Queiros *et al.* 2013). (Created by BioRender.com.)

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anti-inflammatory (Straub, 2007) and pro-inflammatory (Cutolo *et al.* 2006) effects reported. The inflammatory pathway is downstream of many vascular signalling mechanisms that are affected by sex and ageing, further obscuring distinct effects of sex hormones on inflammation. On one hand, oestrogen has been reported to suppress vascular inflammation by down-regulation of pro-inflammatory molecules, including cytokines and adhesion molecules (Stork *et al.* 2002; Kip *et al.* 2005). On the other hand, several clinical studies have described oestrogen as a pro-inflammatory modulator in autoimmune diseases (Cutolo *et al.* 2006). Numerous experimental studies report that oestradiol down-regulates tumor necrosis factor α and interleukin 1β in different cell types and suggest an anti-inflammatory and vasculoprotective action for oestrogens (Straub, 2007; Novella *et al.* 2012). Whether the conversion of vasoprotective/anti-inflammatory effects of oestrogen to vasotoxic/pro-inflammatory effects in ageing subjects is a function of prolonged oestrogen deficiency *per se* or is related to the ageing process and/or the development of vascular disease remains unresolved.

As previously expressed, ER α mediates a great number of oestradiol effects that can be beneficial to cardiovascular physiology: it produces vasodilatation and prevents vasoconstrictive and proaggregating factors, reduces VSMC proliferation and induces a beneficial lipid profile. ER β (and as far as is known, GPER) exerts different effects and, in some conditions, counteracts the beneficial profile of oestradiol through ER α . The balance, or imbalance, between ER α and ER β is therefore an important factor when analysing the cardiovascular effects of oestradiol. In fact, ER α activation has been shown to attenuate injury-induced vascular remodelling (Brouchet *et al.* 2001), but *in vitro* studies have also shown that ER β also plays a protective role in injured arteries (Xing *et al.* 2007), leading us to posit that both ER subtypes contribute to vasoprotection.

Note that ER β is more highly expressed than ER α in oxidative stress, hypoxia and inflammation (Rider *et al.* 2006). In these cases, ER β modulation can be important in regulating pathophysiological ER α -stimulated processes. This link between ERs seems to be more evident in the vascular response to oestrogen, which appears to change during ageing and depend on years since menopause. In previous studies, we observed a gradual increase in ER β expression in uterine arteries of postmenopausal women in line with age, even 10 years after menopause onset, while there was only a slight increase in ER α expression (Novella *et al.* 2012). This age-related increase in ER β expression was positively associated with a pro-inflammatory profile of oestradiol. Likewise, in an experimental murine model of menopause, an increased ratio of ER β /ER α in both vascular endothelium and smooth muscle in aged female mice caused a reversal of the antioxidant effect of oestrogen

to a pro-oxidant profile responsible for increased oxidative stress during ageing (Novensa *et al.* 2011); also, in bone marrow-derived macrophages, ER α expression is greatly diminished with age (Bowling *et al.* 2014). Thus, evidence suggests that vasoprotective effects of oestradiol are age-dependent and this could explain the high cardiovascular risk of HRT seen in clinical trials in postmenopausal women. While the role of ER α has been extensively studied, the actions of ER β on the cardiovascular system and the age- and menopause-related changes of vascular ER β actions remain unclear.

Conclusion

The beneficial effects conferred by oestrogen involve a precise balance of different intracellular signalling pathways, such as NO, prostanoids and RAS, towards a vasodilator and vasoprotective profile involving oestrogen receptors, mainly ER α . Changes in vascular oestrogen receptor expression, age- and menopause-related endothelial injury and epigenomics could also affect the cardiovascular effects of oestrogen. More research is therefore warranted to elucidate these important topics that are probably closely related to the sex differences observed in cardiovascular physiology and pathophysiology.

Future perspectives

Recent studies have provided compelling evidence that the sex of the endothelial cells will influence the responses to not just the sex hormones, but the host of vasoactive agents (Hermenegildo *et al.* 2013; Addis *et al.* 2014; Cattaneo *et al.* 2017). Even more, it is important to note that not only the sex of the subject but the location of the endothelial cells in the body have profound influence (Huxley *et al.* 2018).

The majority of the studies performed so far, including those reviewed in the present article and Table 1, do not take into account those factors. Sex is as a fundamental variable that should be considered when designing and analysing basic and clinical research. Cells of males and females have many basic biochemical differences, and many of these stem from genetic and also hormonal differences. Thus, including female subjects or female-derived specimens in research would lead to a better understanding of cardiovascular physiology in both women and men.

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Additional information

Competing Interests

None of the authors has any conflicts of interests.

Author Contributions

All authors worked together to conceive the topic of the review, and contributed to writing, editing and revising the text and figures. All authors have read and approved the final version of this manuscript and agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. All persons designated as authors qualify for authorship, and all those who qualify for authorship are listed.

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Keywords

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