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Toll-Like Receptors Contribute to Sex Differences in Blood Pressure Regulation

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Abstract

Toll-like receptors (TLRs) play an important role in the innate immune system, and recently, they have been shown to be involved in the regulation of blood pressure. The incidence of hypertension is higher in men, and it increases in postmenopausal women. In fact, premenopausal women are protected from cardiovascular disease compared with age-matched men, and it is well established that this protective effect is lost with menopause. However, the molecular mechanisms underlying this protection in women are unknown. Whether or not it could be related to differential activation of the innate immune system remains to be elucidated. This review focuses on (1) the differences between men and women in TLR activation and (2) whether TLR activation may influence the regulation of blood pressure in a sex-dependent manner.

Keywords

TLR; sex difference; blood pressure; hypertension

INTRODUCTION

Toll-like receptors (TLRs) participate in innate immunity by recognizing pathogenassociated molecular patterns and endogenous damage-associated molecular patterns (DAMPs). Antigen-presenting cells, such as monocytes/macrophages, dendritic cells (DCs), and B cells, are the cell populations with the highest expression of TLRs to recognize ligands in the periphery.¹ Eleven types of TLRs (TLR1–TLR11) have been identified in humans. TLRs can be either localized on the cell surface or alternatively, some of the TLRs exist in intracellular compartments, localized in the endosome, lysosome, endolysosome, and endoplasmic reticulum (ER).²

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The TLRs expressed in the cell surface display 2 major domains: an intracellular portion that is called the Toll/IL-1 receptor (TIR) domain and an extracellular portion that contains leucine-rich repeats. After ligand-induced dimerization of the ectodomains of the TIR, a signal transduction pathway is initiated. Furthermore, the TIR domain of TLRs recruits downstream adaptor molecules, including MyD88 (myeloid differentiation primary response gene 88), TIRAP/Mal (TIR-domain–containing adaptor/MyD88 adaptor–like), TICAM1/ TRIF (TIR-domain–containing adaptor molecule 1/TIR-domain–containing adaptor-inducing interferon β), and TRAM (TRIF-related adaptor molecule).³ After the recruitment of distinct adaptor molecules, downstream signaling can be separated into 2 different pathways: MyD88-dependent and MyD88-independent signaling pathways or the so-called TRIF-dependent pathway. MyD88 is bound by all TLRs, except TLR3. Each signaling pathway results in the activation of inflammatory gene transcription factors [such as interferon-regulatory factors (IRFs), nuclear factor κ B (NF- κ B), and activator protein 1], thereby initiating an inflammatory response.^{2,3}

Intracellular TLRs (TLR3, TLR7, and TLR9) are activated only after acidification of endolysosomal compartments. In fact, receptors must traffic from the ER through the Golgi apparatus to take up residence in the endolysosomes before stimulation. The release of TLRs from the ER involves the posttranslational modification N-glycosylation, Ca⁺² stores, PRAT4A (also known as Cnpy3), and UNC93B1 (mediates the translocation of intracellular TLRs from the ER to the endolysosomes where they encounter and respond to their respective ligands). Furthermore, TLR7 and TLR9 are cleaved at endosomal compartments, which is essential for interaction with MyD88 and subsequent signaling. Cleavage of TLR3 has not been observed.^{4–9} See review¹⁰ for more details.

Hypertension is an important risk factor for cardiovascular disease (CVD) mortality. Between 2011 and 2014, the prevalence of hypertension among US adults was 45.6% (95% confidence interval, 43.6%–47.6%) using the new blood pressure (BP) thresholds from the 2017 American College of Cardiology/AHA guidelines versus 31.9% (95% confidence interval, 30.1%–33.7%) as predicted by the Seventh Report of the Joint National Committee on Prevention, Detection, Evaluation, and Treatment of High Blood Pressure.^{11,12}

Interestingly, the stimulation and activation of TLRs with host-derived molecules have given rise to a possible association between TLRs and CVD.¹³ Hypertension development has been associated with the infiltration of immune cells in the kidneys, the vasculature, and the heart.^{14–21} Furthermore, prevalence, absolute BP values, and molecular mechanisms have shown sex differences in this disease.²² In fact, sex differences in BP levels have also been demonstrated in experimental models of hypertension, including spontaneously hypertensive rats (SHRs),^{23–26} Dahl salt–sensitive rats,^{27,28} deoxycorticosterone acetate-salt,²⁹ angiotensin II (ANG II) administration,^{30–33} L-NAME–treated rats,³⁴ and others.

This review will briefly summarize sex differences in hypertension, with a focus on an examination of the evidence for sex differences in the activation of TLRs and explore whether this may play a role in the sexual dimorphism observed in BP regulation.

SEX DIFFERENCES IN HYPERTENSION

The current review focuses primarily on biological factors that differentiate male and female subjects. The term "sex" refers to the different biological and physiological characteristics of men and women, such as reproductive organs, chromosomes, and hormones, whereas "gender" refers to the socially constructed characteristics of women and men—such as norms, roles, and relationships of and between groups of women and men.³⁵ Consequently, the word "sex" will be used during this review.

Several mechanisms have been described to contribute to hypertension and the resultant endorgan dysfunction in consequence of high BP. Sex-related differences have been consistently described in most of these mechanisms as observed in data provided from both clinical and experimental studies.^{1,11,14–16}

The prevalence of higher BP values in men relative to women starts at a young age, around the twenties, and this trend persists into the fifth–sixth decades of life. BP measurements collected from an observational cohort study comprising more than 20,000 subjects demonstrated that compared with men, women had lower values of systolic, diastolic, and mean arterial pressure at early ages (25 years old; 9, 6, and 7 mm Hg, respectively).³⁶

Studies have shown that premenopausal women display a cardiovascular protection from hypertension compared with men at the same age, and after which, the prevalence becomes higher in women.^{11,37} Indeed, in later life, hypertension is more prevalent in women than men.^{11,22,36,38–44} Interestingly, the increases in the prevalence of hypertension among women begin in the same period when menopause is established, suggesting that sex hormones play a role on the BP regulation, a fact that is supported by numerous studies.^{45–49}

Despite the amount of unquestionable evidence supporting sex differences in hypertension, no recommendations for sex-specific treatment strategies have been proposed, except the ones related to hypertensive disorders during pregnancy.⁵⁰ In fact, a large study including more than 9000 subjects suggested that more intensive lowering of systolic BP to ~120 mm Hg, compared with standard practice of targeting ~140 mm Hg, promotes heterogenic impacts on cardiovascular and renal outcomes, comparing men and women.⁵¹ Therefore, studies like this may turn our attention to the necessity to better evaluate BP mechanisms that are differently modulated between sexes.

Animal experimental models have been used as strategies to evaluate different mechanisms contributing to sex-associated differences in BP.⁵² In fact, several mechanisms have been shown to differently regulate BP between males and females, including sex steroids, the renin–angiotensin system, the endothelin system, the sympathetic nervous system, arachidonic acid metabolites, the nitric oxide system, oxidative stress, obesity and metabolic syndrome, and the contribution of the immune system.⁵² Some of these systems are briefly described below, and they will be further discussed during this review.

The renin–angiotensin system plays a major role in BP regulation, and sex differences have been demonstrated with several components of this system.⁵² Wild-type male mice infused with a low dose of ANG II developed hypertension, whereas female mice did not (increase

of BP around 30 vs. 7 mm Hg, respectively). Furthermore, gonadectomy and administration of an androgen receptor antagonist abolished ANG II–induced hypertension in male mice. By contrast, ovariectomy, estrogen receptor blockade, and estrogen receptor knockout increase the pressor effects of ANG II in female mice.³⁰

Rats infused with ANG II displayed a significant augmentation in BP in both males and females. However, males displayed a greater increase in the number of renal proinflammatory Th17 and IL-17+ T cells than females. Furthermore, hypertension in male SHRs is more dependent on increases of oxidative stress elements than in female SHRs.⁵³

SHRs display elevated levels of both ANG II and endothelin 1 (ET-1). Compared with female SHRs, male SHRs display accelerated progression of renal injury, blunted pressure–natriuresis relationship, and higher BP, with a mean difference of 22 mm Hg between the sexes.^{54–56} In fact, sex differences were demonstrated in the urinary excretion of prostanoids, subproducts from the arachidonic acid metabolism through COX activation, substances that are known to have antihypertensive and natriuretic effects. Conversely, female SHRs have enhanced prostaglandin E2 and thromboxane A2, and this effect is increased by orchidectomy but unchanged by ovariectomy (ovx), suggesting that PGE production is dependent on testosterone levels. In addition, orchidectomy abrogated the sexual dimorphism in BP and protein.⁵⁴

Lower BP values in female SHRs are similar to BP values in castrated male SHRs, indicating a central role for male sex hormones in mediating the sex difference in BP in SHRs. In addition, androgen receptor antagonism attenuates hypertension in male SHRs. Interestingly, treatment with testosterone in ovx females presents similar BP measurements to that in gonad-intact males.⁵⁵ By contrast, ovx does not change BP in female SHRs, although ovx female Dahl salt–sensitive rats showed augmented BP on normal salt diet and further increased on high salt diet compared with controls.⁵⁷ Therefore, the contribution of sex hormones to BP control is complex, and more studies are needed to fully understand how male versus female sex hormones regulate BP.

Sex differences have also been shown to occur in the ET-1 system.⁵⁸ Males display higher ET-1 levels, as well as a higher expression of ETA and ETB receptors in vascular smooth muscle cells (VSMCs), resulting in greater vascular contractility and remodelation. Alternatively, females display increased levels of ETB receptor expression in the endothelium, favoring the production of vasodilatory mediators, and therefore a lower BP.⁵⁸ Of importance, ET-1 elicits higher inflammatory and oxidative stress activation in males, compared with females, which as discussed below could be central to sex differences in BP control.

The immune system is another sex-dependent target that is differently activated during hypertension.^{59,60} The kidney is important for the control of BP, and experimental hypertension has been shown through renal immune cell infiltration. In fact, kidneys of female SHRs have more T regulatory cells (Tregs) and IL-10+ T cells in the kidney, whereas kidneys from male SHRs have more Th17 cells, suggesting a sex difference in the renal T-cell profile in hypertensive rats.^{34,61,62} In fact, it was demonstrated that premenopausal

females are protected from T-cell-mediated hypertension compared with male mice.⁶³ Moreover, systolic BP responses to ANG II are similar between premenopausal and postmenopausal mice (12 mm Hg); however, adoptive transfer of T cells significantly increased systolic BP in postmenopausal females (28 mm Hg).⁶³ Interestingly, male Rag^{-/-} mice (lacking T and B cells) receiving adoptive T cells of males displayed increased BP after infusion of ANG II compared with female Rag^{-/-} mice (37.7 vs. 13.7 mm Hg, respectively).⁶⁴ In the same way, peak mean arterial pressure was 13 mm Hg higher after adoptive T cells transfer of male versus female followed ANG II infusion.⁶⁵ In addition, male animals display increased CD4⁺ cells and Th17 cells, whereas female animals have more CD8⁺ cells and Tregs in hypertension models, and adoptive transfer of T cells from female mice to male Rag^{-/-} mice (lacking T and B cells) abrogated Ang II-induced increase in BP.^{22,61,64} Not all T cells are prohypertensive. Indeed, Tregs are antihypertensive, and females seem to be dependent on Tregs to maintain their lower BP relative to males. Depletion of Tregs in premenopausal mice induced a significant increase in BP. In fact, female rats are protected against deoxycorticosterone acetate-salt-induced hypertension through Tregs.⁶⁶ Taking together these data, a differential activation of the adaptive immune system between the sexes in hypertension is well established.

Therefore, the mechanism resulting in differential BP regulation between the sexes is complex and may be differently activated according to their age. Sexual dimorphisms in several pathways involved in the pathophysiology of hypertension have been described, and among then, the immune system may be central in mediating sex differences in BP control because activation of the renin–angiotensin system and the endothelin system have both been shown to induce an immune response.

The Relationship Between the TLRs and Hypertension

Hypertension, for many reasons, is considered a low-grade inflammatory disease. The relationship between the adaptive system and hypertension has been well established. However, the evidences linking the innate immune system and hypertension are still under investigation. Several pathogen-associated molecular patterns and DAMPs are recognized by TLRs, initiating an immune response. TLR activation leads to overproduction of cytokines and chemokines, among other signaling molecules. Although TLRs are primarily associated with innate immunity, they simultaneously contribute to a variety of clinical disorders, including CVD, where hypertension is the major risk factor. Of importance, the incidence of polymorphisms in genes that participate in TLR pathways may also contribute to CVD development.^{67–70}

TLR4 and Hypertension

TLR4 expression is enhanced in hypertension, suggesting a link with the increased BP. Studies have shown augmented TLR4 expression of mRNA and the protein level in mesenteric resistance arteries,⁷¹ VSMCs, and aortas from male SHRs⁷² compared with controls (WKY/Wistar).

In addition, L-NAME and ANG-II–induced hypertension models conducted in rats also showed enhanced TLR4 expression in the heart, VSMCs, and aorta.^{72–74} Inhibition of TLR4

using specific anti-TLR4 antibodies led to decreased BP and contractility improvement in resistance arteries from male SHRs,^{71,72} suggesting that TLR4 and the innate immune system play a role in hypertension. In addition, TLR4-deficient mice and rats treated with anti-TLR4 antibody have a blunted effect on the BP levels induced by ANG-II infusion.^{75,76}

After TLR4 is activated, its own expression is increased and a possible mechanism by which TLR activation can be controlled is through the reduction of its expression. In fact, anti-TLR4 treatment decreased TLR4 protein expression in mesenteric resistance arteries in adult SHRs.⁷¹ Moreover, TLR4 activation is a MyD88-dependent pathway that culminates in the activation of NF-κB, which mediates the transcription of COX-2 and proinflammatory cytokine genes.^{77–79} Anti-TLR4 treatment decreased the expression of MyD88 and NF-κB (p65) in mesenteric resistance arteries from SHRs.⁸⁰ Moreover, TLR4 antagonist (CLI-095) reduced p65 nuclear expression in VSMCs from and ANG II-induced hypertensive mice.⁷⁴ TLR4 inhibition also attenuates the activation of inflammatory markers such as IL-6 serum levels and COX-2 protein expression in mesenteric arteries from SHRs.⁷¹

In the same way, TLR4-deficient mice displayed reduced TNF-α and IL-6 in hypertensive kidneys.⁷⁵ Interestingly, endogenous TLR4 ligand heparin sulfate aggravated TNF-α and IL-6 mRNA response in cardiac tissue, which was significantly pronounced in SHRs.⁷³ The inhibition as well as the absence of TLR4 is also capable in preventing ROS formation in various tissues, in response to hypertension.^{72,75,76} Because proinflammatory cytokines and ROS formation are all linked to the development and exacerbation of hypertension, decreasing TLR4 inhibition will likely act to lower BP through multiple mechanisms.

TLR9 and Hypertension

DAMPs can activate TLRs, initiating an inflammatory response in hypertension. Mitochondrial DNA (mtDNA), a DAMP that is recognized by TLR9, is elevated in the circulation of SHRs, and it is suggested to be associated with increased BP.⁸¹ In fact, increased BP and endothelial dysfunction were observed in normotensive rats treated with a specific agonist for TLR9, CpG oligonucleotide (ODN2395), and administration of inhibitory oligo dinucleotide for TLR9 (ODN2088) lowered systolic BP in SHRs.⁸¹ In addition, patients with essential arterial hypertension display increased CpG-rich cell-free DNA.⁸²

Besides the canonical-inflammatory TLR9 signaling through MyD88, a newly established noncanonical stress tolerance TLR9 signaling pathway was demonstrated where on stimulation, sarcoplasmic reticulum/ER Ca²⁺ ATPase pump 2 (SERCA2) is reduced leading to activation of cell survival protein 5' AMP-activated protein kinase.⁸³ Acute in vitro incubation of mesenteric resistance arteries with ODN2395 resulted in augmented TLR9-induced inflammatory signaling (increased MyD88 and TRAF6 expression) and noncanonical TLR9-induced signaling (increased SERCA2 and phosphorylated AMPKa^{Thr172} expression). Interestingly, the presence of ODN2088 inhibited the augmented protein expression.⁸¹

Furthermore, it has been demonstrated that SHRs treated with chloroquine, a lysosomotropic agent able to disrupt endosomal TLRs—including TLR9, show improved nitric oxide

bioavailability as well as decreased BP, generation of reactive oxygen species, circulating T cells, and vascular infiltrating immune cells in SHRs.^{84,85} Of importance, mesenteric resistance arteries from young SHRs displayed decreased MyD88 and TRAF6 protein expression after chloroquine treatment, leading to lower NF-κB transcription, once its phosphorylation was reduced.⁸⁴ Moreover, treatment with hydroxychloroquine in patients with lupus lowers the prevalence of thromboembolic events⁸⁶ and reduced hypertension in mice with systemic lupus erythematosus.⁸⁷ Although little is known regarding the role of TLR9 in BP control in female experimental models of hypertension, activation of TLR9 by CpG oligonucleotides during gestation also induces maternal hypertension and increased contractility in mesenteric resistance arteries of pregnant rats.⁸⁸

TLR2, TLR3, and TLR7 and Hypertension TLR2

Inflammation of blood vessel plays a critical role during the initiation and maintenance of vascular diseases. In addition, vascular injury (through endothelial dysfunction) represents a link between cardiovascular risk factors and the increased incidence of hypertension. A study showed that vascular injury induced by cuff placement around the femoral artery in nontransgenic littermates results in increased TLR2 expression and mRNA expression of the proinflammatory cytokines TNF-a, IL-1β, and IL-6, as well as increased ROS production and neointimal proliferation through a TLR2-mediated signaling pathway. In fact, TLR2 knockout mice display blunted increases in vascular injury-mediated cytokine expression, ROS, and neointimal hyperplasia, suggesting that endogenous TLR2 activation might play a central role in the regulation of vascular inflammation.⁸⁹ High-density lipoprotein (HDL) is considered antiatherogenic and is known to prevent endothelial dysfunction. Abnormal HDL isolated from adult and children patients with chronic kidney disease injected into mice significantly increased their BP. Furthermore, human aortic endothelial cells incubated with HDL from chronic kidney disease strongly inhibited NO production. Of importance, selective inhibition or genetic deficiency of TLR2 almost completely abrogated the adverse effects of modified HDL on endothelial NO bioavailability, endothelial repair, and BP.90 Moreover, in a model of ischemia/reperfusion, TLR2 contributes to coronary endothelial dysfunction, inducing endothelial injury due to augmented neutrophil-induced oxidative stress formation.⁹¹ Interestingly, distal and proximal tubules in kidneys of rats infused with ANG II displayed increased TLR2 mRNA and protein expression as well as TNF-a and the number of infiltrated and matured DCs, which were abolished by treatment with losartan.92

TLR3

Preeclampsia, a pregnancy-specific hypertensive syndrome, may result from overactivation of the maternal immune system and is characterized by endothelial dysfunction and excessive inflammation. It was demonstrated that pregnant rats treated with a TLR3 agonist (Poly I:C) displayed preeclampsia-like symptoms, including elevated systolic BPs, increased urinary protein excretion, and malformed pups/litter, which are exacerbated by the absence of interleukin 10.^{93,94} Interestingly, the TLR3 pathway is required for ANG II–induced hypertension; TLR3-deficient (*Tlr3*^{-/-}) mice do not display increased systolic BP with ANG II infusion. Moreover, cardiac hypertrophy and increases in IL-6 and TNF- α gene expression in the kidney and heart caused by ANG II infusion were attenuated in *Tlr3*^{-/-} mice.⁹⁵ Otherwise, TLR3 activation improved mechanical and hypercholesterolemia-

induced arterial injury. In fact, systemic administration of Poly I:C reduced neointima formation in an arterial injury model involving the placement of a perivascular collar, and this protection was lost in *Tlr3*^{-/-} mice. In addition, deficiency of TLR3 accelerated the onset of atherosclerosis in hypercholesterolemic ApoE^{-/-} mice, demonstrating a protective role of TLR3 signaling in the vessel wall.⁹⁶

TLR7

MicroRNAs (miRNAs) are short (18–22 nt), single-stranded, noncoding RNAs involved in the regulation of physiological and pathological processes. Specifically, miR-17, miR-21, miR-34a, miR-92a, miR-126, miR-145, miR-146a, and miR-150 have been linked with the development of CVD. In fact, a study with normotensive and hypertensive middle-aged adults (with both men and women; female subjects were at least 1 year postmenopausal) showed increased circulating expression of miR-34a in hypertensive individuals, whereas miR-21, miR-146a, and miR-126 were lower. Moreover, these miRNAs were related with BP.⁹⁷ Of importance, ssRNAs are ligands for TLR7. TLR activation leads to adaptive immune system activation, and inflammation through increased lymphocytes contributes to the development of hypertension. In this regard, transgenic mice that overexpress TLR7 have a massive expansion of inflammatory DCs and increased activation of B and T cells, which triggers autoimmunity.⁹⁸

Therefore, evidence currently available supports the notion that TLRs, in particular TLR4 and TLR9, play a role in the BP control and could contribute to the development of arterial hypertension. However, in hypertension, it is unclear whether TLR4 and TLR9 are differently modulated in males compared with females. The lack of mechanistic data comparing expression, activity, and availability of endogenous ligands with TLRs between the sexes needs to be better investigated based on the well-accepted central role for the immune system in modulating both the development of hypertension and sex differences in hypertension. Table 1 summarizes the role of TLRs in hypertension.

SEX DIFFERENCES IN ACTIVATION/EXPRESSION OF TLRs

A sex bias in TLR activation has already been shown. In fact, women exhibit increased cellular-mediated and humoral-mediated immune responses and a higher risk of autoimmune disease compared with men.^{1,99–102}

TLR4 RNA expression was evaluated in the postmortem fresh-frozen cerebellum sample from children who experienced inflammation before death. Inflammation, in this case, included exacerbated infection, asthma, asphyxia, inflammatory tumors, or that were treated with antibiotics or nonsteroidal anti-inflammatory drugs around death. TLR variants 1, 3, and 4 were augmented in cerebellums from children categorized with inflammatory disease. However, TLR4 variant 1 and variant 4, but not variant 3, were associated with sex and age. ¹⁰³ The limitation in this study was the experimental size of the sample.

A clinical study demonstrated that TLRs expressed in peripheral blood mononuclear cells (PBMCs) are differently activated between sexes. In healthy women, TLR4, TLR3, and TLR7 stimulated with lipopolysaccharide, Poly I:C, and loxoribine, respectively, displayed

increased TLR7 ligand–induced type 1 interferon (IFN) responses compared with men. ^{104,105} However, TLR3 and TLR4 activations, evaluated by IL-6 production after their stimulation, were not different between the sexes.¹⁰⁵

Yet, PBMCs from men produced less IFN-a after TLR7 stimulation, as well as greater amounts of interleukin (IL)-10 in response to TLR8 and TLR9 ligands.^{106,107} By contrast, it was demonstrated that men presented higher levels of inflammatory cytokines compared with women after stimulating whole blood with various TLR ligands.

TLR expression in PBMCs seems to be selectively modulated by sex hormones. Estradiol stimulation increased RNA and protein levels of TLR8 in PBMCs from healthy premenopausal females, compared with age-matched males. On the other hand, incubation with testosterone had no effect in either sex.¹⁰⁸

PBMCs from healthy women exposed to TLR1–8 stimulation display variations in IL-1β, IL-6, IL-8, and TNF-α production during different phases of the menstrual cycle, as demonstrated by in vitro experiments.¹⁰⁹ After stimulation, all TLRs evaluated displayed a different pattern of cytokine release across the menstrual cycle, except TLR3 and TLR7. Interestingly, use of hormonal contraception was associated with a reduced capacity for circulating plasmacytoid DCs (pDCs) to produce IFN-α and TNF-α in response to TLR9 stimulation in the mucosa of the lower female genital tract.¹¹⁰ Moreover, there was a demonstrated association between the use of hormonal contraception and activation of migrating genital tract–derived DCs after stimulation with LPS, PAM3, and R848 (TLR4, TLR2, and TLR7/8, respectively).¹¹¹

Sex hormones have been shown to modulate the immune system during pregnancy. On TLR7/8 stimulation, PBMCs from first-trimester pregnant women displayed decreased IFNa production, compared with nonpregnant controls. By contrast, the production of TNF-a is increased after TLR4 stimulation in the beginning of pregnancy, followed by a significant decrease during pregnancy. In addition, TLR7-induced IFN-a production by pDCs is augmented, whereas TLR4-induced TNF-a production by monocytes is lower during pregnancy.¹¹²

Incubation with estradiol generated augmented TLR7 and TLR9 expression in pDCs from postmenopausal women, with markedly increased IFN- α production.¹¹³ Moreover, stimulation of pDCs by a TLR7 agonist also results in increased IFN production^{104,114} and higher mRNA expression of all IFN- α subtypes in pDCs extracted from women,¹¹⁵ even before puberty. However, puberty itself is associated with increased production of type 1 IFN,¹¹⁶ which could be explained by TLR7 localization into X chromosome, and the differential TLR expression between men and women.¹⁰⁴ In fact, the presence of two X chromosomes may be responsible for the greater production of IFN- α in pDCs after TLR7 stimulation.¹¹⁶

Data from experimental models also evaluated TLRs in PBMCs. A greater mRNA expression of TLR2, TLR3, and TLR4 was shown in female peritoneal leukocytes of mice and rats compared with male mice and rats, as well as higher TLR2 and TLR4 expression in macrophages. By contrast, ovarian removal led to significantly reduced leukocyte mRNA

expression of TLRs and protein expression of TLR2 and TLR4 on female resident macrophages.¹¹⁷ TLRs play an important role in the immune control of viruses, including mouse cytomegalovirus. A sex difference was demonstrated to cytomegalovirus infection, where female animals showed decreased expression of TLR9 accompanied by a lower activation of the innate immune system, compared with males.¹¹⁸ On the other hand, activation of TLR2 and TLR4 in splenic lymphocytes from mice with myocarditis induced by coxsackievirus B3 infection led to decreased expression of FoxP3+ regulatory T cells in male mice. The suppression of Treg cells by TLR signaling in males but not females suggests a correlation with increased myocarditis susceptibility in males.¹¹⁹

In accordance with human data, mouse pDCs have been demonstrated to be positively regulated by estradiol, where humanized mice present increased TLR7–mediated responses to human pDCs in female host mice relative to males. In agreement, exogenous and endogenous estrogen promoted enhanced TNF-a production by mouse pDCs after TLR7/ TLR9 stimulation through the estrogen receptor, whereas inhibition and genetic ablation of estrogen receptors in DC lineage blunted the TLR7-mediated IFN response.^{1,113,120,121}

It has been shown that macrophages from male mice generate a higher mRNA expression to TLR4 and its coreceptor CD14, as well as higher levels of IL-1 β , following in vivo LPS exposition.¹²² By contrast, macrophages from mice treated with testosterone, generated in the absence of androgen, elicit lower TLR4 expression and activation on stimulation. In addition, in vivo removal of endogenous testosterone (orchiectomy) led to higher TLR4 cell surface expression and increased susceptibility to endotoxic shock.¹²³

Nonetheless, data may follow a different regulatory pathway in different tissues because the pattern of TLR expression does not change during preovulation, menstruation, and implantation in the human fallopian tube epithelial cell line.¹²⁴ In fact, the expression of TLR1–6 in the fallopian tube epithelial cells was demonstrated and was not altered by incubation with different concentrations of estradiol and progesterone. However, the simultaneous combination of both hormones was able to change TLR1, TLR4, TLR5, and TLR6 RNA expression.¹²⁴ In addition, these hormones, in combination, suppressed the production of IL-6 induced by Poly I:C, a TLR3 agonist, suggesting that TLR3 function is somehow mediated by different concentrations of sex hormones, resulting in an inhibitory effect on the TLR3-induced pathway.¹²⁵

Conversely, expression of IFN- or NF- κ B-inducible genes (IFN- α/β or TNF- α) in the spleen and hypothalamus of rats by acute TLR3 activation through Poly I:C did not show sex differences, with the exception of IFN- α expression 8 hours after the activation, exclusively in the spleen from female rats.¹²⁶

TLR7/8 encoded by the X chromosome is expressed in the X-sperm but not the Y-sperm. Activation of TLR7/8 selectively suppressed the mobility of X-sperm but not the Y-sperm without changing sperm viability or acrosome formation. Interestingly, in vitro fertilization using the ligand-selected high-mobility sperm resulted in 83% male pups, whereas activation of TLR7/8 resulted in slow mobility sperm and 81% female pups, suggesting that TLR7/8 are specific receptors that differentially impact the functions of X-sperm but not Y-sperm.¹²⁷

THE ROLE OF THE TLRs IN SEX DIFFERENCES IN BP

Despite several studies demonstrating differences in BP control between males and females, there are no studies to date that have directly addressed the potential interaction between sex, TLRs, and hypertension.

Cellular necrosis after cell death or tissue injury results in the release of DAMPs (for instance, mtDNA, histone and S100 proteins, and heat shock proteins), which are recognized by TLRs leading to their activation.^{128–131} The susceptibility of cells to undergo cell death has been demonstrated to have sex differences.^{132–135} In fact, a recent study demonstrated that mtDNA released in hypertensive rats differently modulates vascular responses in males and females. In this work, male SHRs displayed increased mtDNA, proinflammatory cytokines, and higher contractility to phenylephrine in the aorta incubated with mtDNA, which was attenuated by TLR9 inhibitor. On the other hand, female SHRs did not show changes in the contractile response, however, displayed decreased proinflammatory cytokines and prevented oxidative stress.¹³⁶

Consistent with TLR-sex difference, genetic differential expression of TLRs is observed in the kidney and mesenteric arteries from SHRs. TLR3, TLR5, TLR6, and TLR7 mRNAs are more highly expressed in the renal cortex of male versus female SHRs. TLR6 mRNA is also more highly expressed in the mesenteric arterial bed from male versus female SHRs. Conversely, TLR4 mRNA is more highly expressed in the mesenteric arterial bed of female versus male SHRs.¹³⁷

Consistent with animal models, a cardiovascular cohort study at Framingham showed that platelets from women display increased TLR mRNA expression compared with men, and it is associated with different cardiovascular risk factors. In women, TLR1, TLR3, TLR6, and TLR7 were associated with body mass index, and TLR5, TLR7, and TLR10 were associated with the total cholesterol to HDL ratio. In men, TLR1, TLR2, and TLR3 were associated with the lipid profile and TLR8 with hypertension treatment. Similarly, TLR expression in men was more commonly associated with circulating inflammatory markers (TNFR1 and ICAM1), whereas in women TLR expression was associated with P-selectin levels.¹³⁸

In addition, polymorphism TLR6 Ser249Pro, a single-nucleotide polymorphism, when observed in hypertensive women, resulted in lower left ventricular hypertrophy and reduced proinflammatory response. Conversely, hypertensive men carrying the TLR6 variant do not show echocardiographic features.¹³⁹ Indeed, TLR4 knockout mice developed less-severe left ventricular dysfunction,¹⁴⁰ and blockade of MyD88 attenuated load-induced cardiac growth.¹⁴¹ In the same way, an investigation of 2774 subjects reported that hyporesponsive TLR4 polymorphisms (Asp299Gly and Thr399Ile; single-nucleotide polymorphism [SNP]) were associated with an increased risk of myocardial infarction in men, but not women,¹⁴² further supporting sex-specific effects of TLR on cardiovascular function.

In pregnant women, hormonal changes lead to differences in response to TLR activation and BP regulation. Preeclampsia (characterized by high BP and proteinuria) shows irregularities in the immune system that could be linked to the immunological changes related to placental environments.¹⁴³ Trophoblast surface as well as placentas from preeclamptic women

overexpress TLR2, TLR4, TLR3, TLR7, TLR8, and TLR9 when compared with healthy pregnancies.^{144–146}

Similar to data in human, in pregnant mice, intraperitoneal injections of TLR3, TLR7, and TLR8 agonists led to a significant increase in protein expression and mRNA for TLR3/7/8 in placenta and an increase in systolic BP at gestational day 17 compared with vehicle-treated controls.¹⁴⁶ TLR3 and TLR9 in rodents, using Poly I:C and synthetic CpG oligonucleotide (respectively), led to the development of the characteristic markers of preeclampsia, including hypertension, endothelial dysfunction, decreased fetal weight, and proteinuria. ^{88,93,94,147–150}

These observations led to the inference that activation of TLRs is somehow sensitive to sex hormones and this could be one of the mechanisms related to sex differences in the BP control. However, it is imperative to make clear that most of the current available data on this field are limited, if immunity from these individuals is evaluated only by a single time point observations, because the innate immunity is a result of previous immunogenic expositions.

CONCLUSIONS

The present review summarizes recent studies that suggest a link between TLRs and hypertension. It is well known that the BP control in men and women is different, and exciting evidence demonstrates that expression/activation of TLRs are sex specific. However, the sex differences in TLR activation mediating hypertension remain to be elucidated. Figure 1 and Table 2 summarize sex differences and TLRs in hypertension.

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FIGURE 1.

Sex differences in TLRs during hypertension: TLRs are differently modulated and expressed between the sexes, during hypertension. In fact, male SHRs display increased mtDNA, augmented TLR9 activation, and augmented expression of TLR4, compared with female SHRs, features that favor hypertension. When TLR9 and TLR3 are overactivated during pregnancy, preeclamptic-like symptoms are observed. In females, polymorphisms at TLR6 increase cardiac protection, whereas polymorphisms in TLR4 favor infarction in males. ROS, reactive oxygen species.

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TABLE 1.

Role of TLRs in Hypertension

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Observation	Local	Characteristics	References
Increased expression or activity of TLR			
TLR2	Femoral artery	Vascular injury but basal BP levels	89
TLR2	Kidney	II-9NG-II	92
TLR3	Placenta	Preeclampsia	93 and 94
TLR4	MRA	SHR	71
TLR4	Cardiac	L-NAME	73
TLR4	CSMC	SHR	76
TLR4	VSMCs	SHR/ANG-II	72
TLR4	Aorta	SHR/ANG-II	72 and 74
Increased expression or activity of the TLR ligand			
miR-34a (TLR7)	Plasma	Patients with essential arterial hypertension	76
mtDNA (TLR9)	Serum	SHR	81
CpG-rich cell-free DNA (TLR9)	Serum	Patients with essential arterial hypertension	82
Genetic or therapeutic interventions targeting to TLRs resulting in reduced blood pressure			
TLR2-deficient mice	Coronary artery	Cardiac ischemia/reperfusion (basal BP levels)	91
TLR3-deficient mice	Heart and kidneys	II-9NG	95
Chloroquine (endosomal TLRs)	MRA	SHR	84 and 85
Hydroxychloroquine (endosomal TLRs)	Systemic lupus erythematosus	Thromboembolic events	86
Hydroxychloroquine (endosomal TLRs)	Aorta and kidney	Systemic lupus erythematosus mice	87
Anti-TLR4	MRA and aorta	SHR	71 and 72
TLR4-deficient mice	Kidney	II-9NA	75

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CSMC, cavernosal smooth muscle cells; L-NAME, N-nitro-L-arginine methyl ester; MRA, mesenteric resistance artery.

TABLE 2.

Sex Differences and TLRs in Hypertension

Observation	Local	Hypertension Classification	Role on Hypertension	References
Female				
Increased expression or activity of TLR				
TLR3/TLR5/TLR6	Renal cortex	SHR	Immune response	137
TLR4/TLR6	MRA	SHR	Immune response	137
TLR1/TLR3/TLR6/TLR7	Platelets	Basal BP levels	Associated cardiovascular risk	138
TLR5/TLR7/TLR10	Platelets	Basal BP levels	factors with body mass index Associated cardiovascular risk factors with the total cholesterol to HDL ratio	138
Increased expression or activity of the TLR ligand				
mtDNA (TLR9)	Aorta	SHR	Decreased proinflammatory cytokines and prevented oxidative stress	136
Polymorphism				
TLR6—Ser249Pro	Serum/heart	Hypertension	Lower LV and reduced proinflammatory response	139
TLR4—Asp299Gly and Thr399Ile	Serum	Basal BP levels	No association with myocardial infarction	142
Male				
Increased expression or activity of TLR				
TLR7	Renal cortex	SHR	Immune response	137
TLR1/TLR2/TLR3	Platelets	Basal BP levels	Associated cardiovascular risk factors with lipid treatment	138
TLR8	Platelets	Basal BP levels	Associated cardiovascular risk factors with hypertensive treatment	138
Increased expression or activity of the TLR ligand				
mtDNA (TLR9)	Aorta	SHR	Increased proinflammatory cytokines and higher contractility	136
Polymorphism				
TLR6—Ser249Pro	Serum/heart	Hypertension	No changes in echocardiographic features	139
TLR4—Asp299Gly and Thr399Ile	Serum	Basal BP levels	Increased risk of myocardial infarction	142
TLR4 knockout mice	Heart	Cardiac hypertrophy	Less-severe LV dysfunction	140 and 141
Pregnancy				
Increased expression or activity of TLR				
TLR2/TLR4/TLR3/TLR7/TLR8/TLR9	Trophoblast and placenta surface	Women with preeclampsia	Classical markers of preeclampsia	144–146
Increased expression or activity of the TLR ligand				

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	Local	Hypertension Classification	Role on Hypertension	References
t-837/CL097 (TLR3/TLR7/TLR8)	Placenta	Preeclampsia (animal model)	Classical markers of preeclampsia: Impaired relaxation, elevated systolic blood pressures, increased urinary protein concentrations, and malformed pups/litter	93 and 146
nucleotide (TLR9)	MRA	Preeclampsia (animal model)	Classical markers of preeclampsia: Increased systolic blood pressure and resistance artery contraction and decreased fetal weight	88 and 150

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LV, left ventricle; MRA, mesenteric resistance artery.