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Dual Activation of TRIF and MyD88 Adaptor Proteins by Angiotensin II Evokes Opposing Effects on Pressure, Cardiac Hypertrophy and Inflammatory Gene Expression

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Abstract

Hypertension is recognized as an immune disorder whereby immune cells play a defining role in the genesis and progression of the disease. The innate immune system and its component toll-like receptors (TLRs) are key determinants of the immunological outcome through their proinflammatory response. TLR activated signaling pathways utilize several adaptor proteins of which adaptor proteins MyD88 and TRIF define two major inflammatory pathways. In this study, we compared the contributions of MyD88 and TRIF adaptor proteins to angiotensin II (Ang II)induced hypertension and cardiac hypertrophy in mice. Deletion of MyD88 did not prevent cardiac hypertrophy and the pressor response to Ang II tended to increase. Moreover, the increase in inflammatory gene expression (Tnfa, Nox4 and Agtr1a) was significantly greater in the heart and kidney of MyD88-deficient mice compared with wild type mice. Thus, pathways involving MyD88 may actually restrain the inflammatory responses. On the other hand, in mice with nonfunctional TRIF (Trif^{mut} mice), Ang II induced hypertension and cardiac hypertrophy were abrogated, and pro-inflammatory gene expression in heart and kidneys was unchanged or decreased. Our results indicate that Ang II induces activation of a pro-inflammatory innate immune response, causing hypertension, and cardiac hypertrophy. These effects require functional adaptor protein TRIF-mediated pathways. However, the common MyD88 dependent signaling pathway, which is also activated simultaneously by Ang II, paradoxically exerts a negative regulatory influence on these responses.

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Keywords

Hypertension; innate immune system; toll-like receptors; MyD88; TRIF; TICAM1; Angiotensin; gene expression

Introduction

In recent years, hypertension has increasingly been recognized as an immunological disorder whereby the components of the immune system play an important role in determining two key aspects of the disease, blood pressure and end organ damage ^{1–8}. Greater understanding of the immune system itself has accelerated the progress in defining the cell populations of the adaptive immune system that play a role in hypertension ^{9–15} and the neuro-immune axis that may activate these cells ^{1, 16–20}. More recently, the innate immune system, specifically the toll-like receptors (TLRs), has been shown to be the mediators of neuro-inflammatory response that can be modulated by angiotensin II (Ang II) ^{16, 17, 19}. The goal of this study was to determine the innate immune mechanisms involved in Ang II induced hypertension and cardiac hypertrophy.

Ang II is a major product of the renin-angiotensin system (RAS) that plays a crucial role in the pathology and treatment of hypertension. It causes vasoconstriction, promotes sodium retention in the kidneys, acts as a trophic factor in the myocardium, activates the sympathetic nervous system, and induces a pro-inflammatory state that results in organ damage $^{21-24}$. Ang II promotes inflammation by inducing cytokine release, immune cell infiltration in the kidney, expression of genes in kidneys and the heart that are pro-inflammatory, pro-hypertrophic and pro-fibrotic, such as Tumor Necrosis Factor- α (TNF- α), Interleukin-6 (IL-6), Interleukin-1 β (IL-1 β) and Transforming Growth Factor- β (TGF- β)^{1, 25}.

TLRs are a major class of innate immune receptors that constitute a diverse family of pattern recognizing receptor (PRR). They bind to exogenous pathogen associated molecular patterns (PAMP) or endogenous damage associated molecular patterns (DAMP) to initiate intracellular signaling cascades culminating in expression of pro-inflammatory genes as well as regulate the activity of adaptive immune cells ²⁶. TLR signaling is dependent on the association of TLRs with one of several adaptor proteins that determine the specificity of signaling ²⁷. Activation of TLRs often initiates two major adaptor-dependent signaling pathways: one is myeloid differentiation protein 88 (MyD88)-dependent, and the other is TICAM1 (Toll Interleukin Receptor-domain-containing adaptor molecule 1), also known as TRIF (TIR-domain containing adaptor-inducing interferon- β)-dependent ²⁸. Besides their role in induction of pro-inflammatory genes, signaling by different TLRs may be complementary, synergistic or even antagonistic ^{29–31} resulting in complex phenotypes and gene expression patterns.

We have previously shown a link between TLR/MyD88 signaling and myocardial infarction induced cardiac hypertrophy ³². Mice lacking MyD88 adaptor protein had significantly improved survival and reduced inflammation, cardiac fibrosis and cardiac hypertrophy after myocardial infarction. Prompted by these results we tested the hypothesis that TLR/MyD88

Our results revealed a selective and paradoxical dependence of Ang II hypertension responses on the two dominant adaptor proteins of the innate immune pathways TRIF and MyD88. Whereas TRIF-dependent pathways are essential for Ang II-mediated hypertension, cardiac hypertrophy and the inflammatory response, MyD88 pathways, in contrast, appear to be anti-inflammatory and restrain those responses.

Materials and methods

Animals

All experiments with animals were done in accordance with the regulations put forth by the Institutional Animal Care and Use Committee (IACUC) of University of Iowa. The wild type C57BL/6J (WT), and *Trif*^{mut} (C57BL/6J-*Ticam1Lps2*/J) ³³ mice were obtained from Jackson Labs. *MyD88^{-/-}* mice were obtained from Dr. Shizuo Akira's group ³⁴ and were bred in the University of Iowa animal facility in specific pathogen free environment. All mice were in C57BL/6 background.

Ang II infusion and tail-cuff recordings of blood pressure

Mini-osmotic pumps (Alzet model 1004, 0.11 μ l/h, 28 days), containing saline or Ang II (730, 1000 or 3000 ng/kg/min), were inserted subcutaneously in mice under anesthesia (2.0–2.5% isoflurane). Male mice of 8–12 weeks age were used for the experiments.

Systolic arterial blood pressure was measured by tail cuff using the Visitech-2000 system. A rigorous system of recording tail cuff pressures has been in effect in our laboratories over decades and found to be reliable for steady state blood pressure measurements over long periods of time.

Baseline blood pressure was measured prior to pump insertion. Tail-cuff recordings were performed at the same time of the day (before noon) to avoid diurnal changes in the pressure. To acclimatize the mice to the procedure, one week prior to the start of the experiments, tail-cuff pressures were measured on two days. During the experiments, twenty tail cuff measurements for each mouse were recorded for each session (3 times a week). Average measurements for each mouse were then used for the groups (n 3 mice per group). At the end of the experiments (3 weeks), mice were euthanized and organs were obtained for further analyses.

Cardiac hypertrophy

Cardiac hypertrophy was assessed by measurement of heart weight to body weight ratio (HW/BW) in hearts from mice after three weeks of saline or Ang II infusion. For measurement of cardiomyocyte diameter and cardiac fibrosis, heart tissues were examined from 4 separate cross-sections collected at different levels (each 250 μ m apart) from the center of the heart. In each tissue sample, a pathologist using post examination masking techniques ³⁵ screened each heart on low magnification to identify areas with the largest cardiomyocytes diameter. At these sites, the 5 largest cardiomyocyte diameters were

recorded. Thus for each heart 20 diameters were collected and the mean of these data calculated for each animal.

RNA isolation and RT-QPCR

Total RNA from mouse heart and kidneys were isolated using mirVana RNA isolation kit (Ambion) or RNeasy RNA Isolation Kit (Qiagen). A 2 µg aliquot of RNA sample was used to synthesize cDNA in 50 µl reactions using Oligo (dT) as primers and SuperScript III Reverse Transcriptase (Invitrogen). Quantitative real-time PCR was performed on QPCR cycler (ABI) using SYBR green based PCR reactions, as described ³². Quantifications were done using CT method where *Gapdh* was used as a reference gene for normalization of RNA expression ³⁶. The primers used in this study have been previously described ^{32, 37} and are listed in Supplemental Table-S1.

Statistical analyses

Statistical analyses were performed using an unpaired t-test or ANOVA as shown in respective figures. Post-hoc analyses on multiple group data were done using Tukey's test. A *p*-value of 0.05 or less was considered to be statistically significant. All results are presented as mean \pm SEM.

Results

Ang II infusion increases blood pressure in *MyD88^{-/-}* but not in *Trif^{mut}* mice

Compared to saline infusions, which did not alter systolic blood pressure (SBP) over three weeks, Ang II infusion (3000 ng/kg/min) during the same period increased SBP in both WT and in $MyD88^{-/-}$ mice (Figure 1A and 1B). In WT, blood pressure peaked at 147 ± 4 mm Hg (Figure 1A) whereas in Ang II-infused $MyD88^{-/-}$ mice SBP reached a peak value of 163 ± 6 mm Hg (Figure 1B). The increase in SBP during the third week of infusion was significantly greater in $MyD88^{-/-}$ mice than in WT mice (p < 0.05). In contrast to the WT and $MyD88^{-/-}$ mice, Ang II did not increase blood pressure in $Trif^{mut}$ and the response to Ang II was identical to the response to saline during the last two weeks of infusion averaging 116 ± 3 mm Hg with saline and 118 ± 4 mm Hg with Ang II (Figure 1C).

In addition, at lower Ang II dosage (730–1000 ng/kg/min) in a separate group of studies the average SBP over the last two weeks of infusion of Ang II was significantly lower in *Trif^{mut}* (129 ± 3) vs. WT 135 ± 5) and highest in MyD88 $^{-/-}$ (142 ± 5 mm Hg; Suppl. Figure 1).

These results indicate that Ang II-induced hypertension is MyD88-independent and that the presence of MyD88 restrains the hypertensive response to Ang II, which is TRIF-dependent.

Ang II-induced cardiac hypertrophy is abrogated in Trif^{mut} mice

The heart weight to body weight ratio (HW/BW) in WT mice infused with saline, low dose (730 ng/kg/min), and high dose (3000 ng/kg/min) of Ang II were 4.88 ± 0.16 , 5.23 ± 0.15 and 6.83 ± 0.21 mg/g respectively (Figure 2A). Corresponding values in $MyD88^{-/-}$ mice were 3.93 ± 0.15 , 4.70 ± 0.06 and 6.30 ± 0.86 (Figure 2B) and in *Trif^{mut}* mice they were 4.39 ± 0.19 , 4.72 ± 0.11 , and 5.10 ± 0.29 (Figure 2c). The increase in HW/BW ratio with the

high dose Ang II vs. saline was greater in MyD88^{-/-} than WT but lowest in *Trif^{mut}*. The high dose of Ang II infusion induced a 60% increase in HW/BW over saline infusions in $MyD88^{-/-}$ mice, a 40% increase in WT, and a 22% increase in *Trif^{mut}* (Figure 2). Analyses of variance and linear regressions confirming the differences in the three genotypes are in Supplemental Table-S2). Cardiomyocyte diameter and fibrosis confirmed Ang II-induced cardiac structural changes (Supplemental Table-S3).

Cardiac gene expression in Ang II-infused mice

We compared the expression of inflammation related genes in the hearts from saline and Ang II-infused WT, $MyD88^{-/-}$, and $Trif^{mut}$ mice. Tumor necrosis factor- α (*Tnfa*) expression was significantly increased by Ang II in WT and $MyD88^{-/-}$ mice, but the increase was much greater in the $MyD88^{-/-}$ hearts than in WT hearts (Figure 3A and B). However, Ang II-infusion had no effect on *Tnfa* expression in *Trif^{mut}* hearts (Figure 3C).

Expression of NADPH oxidase-4 (*Nox4*) was significantly increased to a similar extent in hearts from WT and $MyD88^{-/-}$ mice with Ang II-infusions (Figure 3D and E). However, *Nox4* expression was unaffected in *Trif^{mut}* hearts upon Ang II-infusion (Figure 3F).

The angiotensin type 1-receptor (AT1 receptor, *Agtr1a*) expression in heart was unaffected by Ang II infusion in WT mice (Figure 3G), was significantly increased by Ang II in $MyD88^{-/-}$ mice (Figure 3H) and was slightly decreased by Ang II in *Trif^{mut}* mice (Figure 3I). Similarly the expression of *Cxcl10* a gene that is selectively induced by the TRIF/IFN pathway was enhanced in $MyD88^{-/-}$ supporting a negative regulatory effect of MyD88 on the TRIF pathway. Expression of *Cxcl10* was actually decreased in *Trif^{mut}* mice (Suppl. Fig. 2A).

The muted effect of Ang II infusion on cardiac gene expressions was not due to a general unresponsiveness of *Trif^{nut}* mice because Ang II induced significant over expression of skeletal muscle actin- $\alpha 1$ (*Acta1*) in all three strains of mice and particularly in *Trif^{nut}* (Suppl. Figure 2B). Thus, the intact MyD88 pathway in WT mice has a restraining inhibitory effect on Ang II-induced inflammatory gene expression, whereas the intact TRIF pathway is required for the inflammatory effect of Ang II.

Renal gene expression in Ang II-infused mice

We compared gene expression in the kidneys of saline and Ang II-infused mice. Ang II infusion resulted in an insignificant increase in expression of *Tnfa* in the kidneys of WT mice (Figure 4A), however, there was a clear and significant increase in *Tnfa* in kidneys of $MyD88^{-/-}$ mice (Figure 4B), and a much smaller negligible increase in kidneys of *Trif^{mut}* mice (Figure 4C).

Ang II-infusion did not increase expression of *Nox4* in the kidneys of WT mice but it significantly increased the expression in kidney of $MyD88^{-/-}$ mice (Figure 4D and 4E) and, conversely, significantly decreased *Nox4* in kidney of *Trif^{mut}* mice (Figure 4F).

The renal expression of *Agtr1a* was unaltered in kidney of Ang II-infused WT (Figure 4G), significantly increased in kidney of $MyD88^{-/-}$ mice (Figure 4H), but significantly decreased in kidney of *Trif^{mut}* mice (Figure 4I).

Thus, the absence of MyD88 caused a greater magnitude of pro-inflammatory gene expression with Ang II in both heart and kidney. In contrast, in the absence of TRIF, the inflammatory gene overexpression was eliminated or even reduced when compared to WT, confirming the inhibitory effect of the MyD88 pathway and the TRIF dependence of the inflammatory response.

Changes in Cardiac TLR and adaptor protein expressions in MyD88^{-/-} and Trif^{mut} mice

Since TLR3 and TLR4 are likely to be involved in signaling pathways that use the two adaptor proteins we measured their cardiac expressions in $MyD88^{-/-}$ and $Trif^{mut}$. The results in Figure 5 indicate that deletion of MyD88 was associated with an increase in Trif expression, which along with increases in both Tlr3 and Tlr4 expression, must have contributed to the enhanced inflammatory gene expressions observed in $MyD88^{-/-}$ mice. In $Trif^{mut}$ mice, expression of MyD88 was increased, whereas Tlr4 expression was decreased and Tlr3 expression was unchanged. Together, the negative regulatory effect of MyD88 and decreased Tlr4 expression must have contributed to the suppressed inflammatory gene expression in $Trif^{mut}$.

Discussion

We tested the contribution of the innate immune signaling by the two major adaptor proteins of TLRs, MyD88 and TRIF, in Ang II-induced hypertension. Consistent with our earlier demonstration that the innate immune system plays a major role in a genetic model of hypertension, our results show that pathways involving both adaptor proteins contributed to Ang II pressor responses in WT mice albeit in a mutually opposite manner.

Our results show that: (a) MyD88-dependent innate immune signaling pathways are not responsible for Ang II-induced hypertension, cardiac hypertrophy, and pro-inflammatory gene expression, (b) TRIF-mediated pathways, on the other hand, are the essential determinants of Ang II-induced responses, and (c) the absence of MyD88 unmasks exaggerated Ang II responses indicating that MyD88 signaling is simultaneously activated by Ang II and functions as a negative regulator of pro-inflammatory pathological responses mediated by TRIF. These results are summarized in the schematic in Figure 6. We arrived at an original and provocative finding that TRIF-dependent pathways are essential and pro-inflammatory in evoking Ang II hypertension, whereas MyD88-dependent pathways are negative regulators of TRIF pathways, and therefore may be considered anti-inflammatory. These results provide novel mechanistic insight in one of the most commonly used models of hypertension and identify putative targets of pathologic and therapeutic significance.

Dominance of the Immune System in Ang II Responses

Ang II is the major product of the renin-angiotensin system (RAS). It plays a major role in cardiovascular regulation of arterial pressure and blood volume, salt retention, and

sympathetic activation. It adversely alters the course of several pathological states including hypertension, heart failure, atherosclerosis, and the metabolic syndrome.

The contribution of Ang II to hypertension has long been ascribed to its powerful vasoconstrictor action, its salt retaining properties and its central and peripheral sympathoexcitatory effects. A seminal notion was advanced by the work of Harrison and colleagues that Ang II induced hypertension is dependent on the immune system ¹³. A complementary finding has been our previous results that in a genetic hypertension model, the spontaneously hypertensive rat (SHR), Ang II dramatically enhances cytokine release (IL-6 and IL-1 β) by splenocytes upon their activation by ligands of TLR 7, 8 and 9 ¹⁷. That the pro-inflammatory effect of Ang II was evident in SHR splenocytes prior to the onset of hypertension suggests a genetic abnormality of the innate immune cells that links their AT1 receptors to the TLR signaling pathways and potentiates pro-inflammatory responses. More recently, Harrison's group has also identified an important role of the innate immune system, specifically the dendritic cells, in Ang II hypertension ²⁵.

The processes that mediate cardiac hypertrophy in response to Ang II have been ascribed to the mechanical afterload of a raised arterial pressure and also to trophic effects on cardiac myocytes with overexpression of contractile proteins by direct activation of AT1 receptors or indirectly by the adrenergic stimulus of excessive sympathetic activation. However, in a manner similar to the Ang II pressor response the trophic cardiac effect seems to be also mediated by immunologic pathways ^{24, 38, 39}

The results of this study provide strong confirmation that Ang II hypertension, cardiac hypertrophy, and the pro-inflammatory gene expressions in both the heart and kidney, depend on innate immune signaling pathways that require the presence of the specific adaptor protein TRIF.

Ang II hypertension and cardiac hypertrophy are MyD88-independent but TRIF-dependent

An unexpected finding of this study was that MyD88, a widely recognized essential adaptor protein for the activation of pro-inflammatory TLR pathways ^{32, 34, 40}, negatively regulated the pro-inflammatory TRIF-mediated response to Ang II. In fact, we had previously reported a beneficial reduction of cardiac hypertrophy in the $MyD88^{-/-}$ mice after myocardial infarction ³². Similarly Wang, et al. ⁴¹ reported a reduction in hypertrophy in $MyD88^{-/-}$ mice upon Ang II-infusion. However, they used a subpressor dose of Ang II (~400ng/kg/min). In our present study our model was primarily a model of hypertension using Ang II infusions at high pressor rates (3000ng/kg/min) and the resulting hypertension, cardiac hypertrophy and particularly the pro-inflammatory gene expressions were exaggerated.

These doses of Ang II cause significant vasoconstriction, neuro-humoral and central nervous system (CNS) activation with increased sympathetic activity in addition to the renal and cardiac effects. Thus, the differences in results may represent the release of different endogenous ligands that may engage different adaptor mediated pathways. Furthermore, a recent abstract ⁴², indicates that the hypertension and hypertrophy caused by Ang II infusions (1000ng/kg/min) were preserved in $MyD88^{-/-}$ mice. This study's conclusion was in accordance with ours that MyD88 does not contribute to hypertension.

Contrasting Effects of Signaling through TRIF vs. MyD88 Adaptors on Gene Expression in Heart and Kidney

We observed contrasting changes in pro-inflammatory gene expression in MyD88^{-/-} and $Trif^{mut}$ mice compared to WT mice. Tnfa is a key cytokine in Ang II induced increases in blood pressure and cardiac hypertrophy ^{38, 39, 43, 44}. We found that in both heart and kidney the magnitude of induction of Tnfa expression was much higher in MyD88^{-/-} than in WT and $Trif^{mut}$.

Similarly reactive oxygen species (ROS) production resulting from activation of RAS system in renal tissue leads to hypertension ¹⁶ and the expression and activation of *Nox4* RNA ^{45, 46} promotes cardiac hypertrophy ^{47–49}. We found that *Nox4* expression was slightly increased in heart of WT mice after three weeks of Ang II but significantly enhanced in both kidney and heart of MyD88^{-/-} and, in contrast, markedly downregulated in kidney and unchanged in heart of *Trif^{mut}* mice.

Increased expression of Ang II type 1 receptors (*Agtr1a*) was also significantly increased during the 3 week infusion of Ang II in both heart and kidney of MyD88^{-/-} and conversely reduced significantly in $Trif^{onut}$ mice whereas no significant changes were observed in WT mice.

Taken together these expressions of pro-inflammatory genes correlate with the development of hypertension and cardiac hypertrophy, and demonstrate that the TRIF adaptor protein is the essential mediator of those responses whereas MyD88 is an effective negative regulator.

TLR Mediated Dual Activation of MyD88 Dependent and TRIF-dependent Signaling Pathways

Multiple TLRs interact selectively and specifically with ligands to initiate signaling cascades through adaptor protein molecules resulting in pro- or anti-inflammatory responses ^{50, 51}. The most common adaptor protein is the MyD88 that was initially recognized as a component of the IL-1 receptor signaling complex ²⁷. All TLRs except TLR3 and a subset of TLR4 signaling events depend on MyD88 ^{27, 50, 51}. MyD88-independent TLR4 pathways are activated by LPS in macrophages causing induction of several genes, including CXCL10, RANTES and macrophage colony stimulating factor ⁵². The adaptor molecule of the MyD88 independent signaling events is TRIF ^{27, 51}. TRIF induction of IRF-3 activation and IFN- β and IFN-inducible genes are MyD88 independent ^{53–55}.

Our results indicate that Ang II hypertension induces simultaneous activation of TLR pathways that are dependent on both adaptor proteins and have opposing effects in the WT mice with a net increase in arterial pressure, cardiac hypertrophy, and modest increases in cardiac and renal pro-inflammatory gene expression. The hypertension, cardiac hypertrophy, and the pro-inflammatory gene expression caused by Ang II were predominantly TRIF mediated. The expression of a TRIF-dependent chemokine *Cxcl10*, which was enhanced in the hearts of $MyD88^{-/-}$ mice compared to WT but decreased in *Trif^{mut}* mice, further supports the induction of TRIF dependent inflammatory signaling, which is negatively regulated by MyD88-dependent signaling.

Since TLRs are selectively activated by a large number of specific ligands of either exogenous or endogenous origin, the determination of which ligands are generated during the prolonged infusion of Ang II is challenging. One might suggest that the early rise in pressure during the first days of Ang II infusion may provoke cellular damage which releases DAMPs into the circulation which then activate TLRs on innate immune cells initiating the inflammatory cascade. An important question is how MyD88 exerts its anti-inflammatory effects.

Negative regulation of TRIF responses by MyD88 is anti-inflammatory

TLR signaling via different adaptor proteins may function in context-specific ^{56, 57} as well as tissue-specific manner ⁵⁸. Moreover, multiple mechanisms of negative regulation of these adaptor pathways have been described ⁵⁹. It is not clear whether the MyD88 anti-inflammatory effect results from direct stimulation of specific inhibitors or from gene repression effects of transcription factor NF- κ B ^{60–62}. Our results, however, suggest that deletion of MyD88 results in increased cardiac expression of TLR3 and TLR4 as well as TRIF, all of which would contribute to exaggerated responses and support a negative regulatory influence of MyD88. In contrast, the TRIF mutation decreased TLR3 without altering TLR4 expression but increased MyD88 expression. The latter would contribute to the observed suppression of responses to Ang II.

Another conceptually attractive mechanism that would explain the enhanced TRIFdependent inflammatory responses in *MyD88^{-/-}* mice is "Signaling Flux Redistribution" (SFR) ⁶³ whereby removal of MyD88 enhances the availability and flux of signaling substrates through the alternative TRAM/TRIF dependent pathway. Enhanced inflammatory response in MyD88-deficiency has also been reported in other models as well ^{57, 64}. Indeed, MyD88 functions as a negative regulator of the TRIF-dependent pathway in a corneal inflammation model ⁶⁵. In this study, TLR3 ligand-induced inflammatory cellular infiltration in cornea was exacerbated in *MyD88^{-/-}* mice. It is also noteworthy that MyD88 signaling also produces anti-inflammatory molecules, such as IL-10 ⁶⁶, which requires intact MyD88 for its anti-inflammatory action ⁶⁷. The constitutive expression by MyD88 of A20, IRAK-M and PPAR gamma, which result in downstream repression of NF-kB transcription factor but not TRIF-dependent interferon regulatory factor 3 (IRF3), would also result in enhancement of pro-inflammatory responses in MyD88 deficiency ^{68, 69}.

Our results showing exacerbation of Ang II hypertension and inflammatory gene expression in kidney and heart including Cxcl10 in the $MyD88^{-/-}$ heart supports a similar antiinflammatory effect of MyD88. Thus, in the context of Ang II-induced hypertension and cardiac hypertrophy it is clear that MyD88 exerts a novel negative regulatory role.

Interaction of Ang II with TLRs

Exactly how the Ang II pathway interacts with the TLR pathways is not understood. Ang IIinduced hypertension involves the adaptive immune system ¹³, which in turn is regulated by the activation of the innate immune system by TLRs that detect specific ligands (DAMPs) ⁷⁰. Cytokines released by the innate immune system, such as pro-inflammatory cytokine IL-6 are essential for Ang II-induced hypertension in mice ⁷¹. Moreover, as

mentioned above, we have recently shown that Ang II is capable of markedly enhancing specific pro-inflammatory TLR responses in the immune cells of SHRs but not in the immune cells of normotensive Wistar-Kyoto rats ¹⁷. The components of TLR signaling are widely expressed in multiple tissues and cell types. Although systemic Ang II infusion has wide-ranging effects on various tissues, Ang II is not known to be an agonist of the TLRs. It is possible that a downstream product of Ang II-AT1 receptor binding may generate a TLR agonist or the AT1R signaling pathway might interact with the components of the TLRs pathway to activate it.

Perspective

Inflammation is a hallmark of the etiology of hypertension and end organ damage including cardiac hypertrophy. TLRs and their adaptor proteins are integral components of innate immunity whose role in hypertension and end organ damage is now fully recognized. Our finding that Ang II-responses result from activation of different pathways with pro- and anti-inflammatory adaptor proteins opens up opportunities for new molecular targets to treat recalcitrant hypertension.

Hypertension represents an abnormal immunological state that involves the adaptive as well as the innate immune system. Our finding indicates that a hypertensive state may be exacerbated as a result of suppression or impairment of MyD88-dependent pathways. Conversely, targeting the TRIF pathway may be therapeutic by blocking the inflammatory response and enhancing the negative regulatory effects of MyD88.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

Ang II	Angiotensin II
AT1R	Angiotensin II type 1 receptor
DAMP	Damage associated molecular patterns
PAMP	Pathogen associated molecular patterns
PRR	Pattern recognizing receptors
MyD88	Myeloid differentiation protein 88

SBP	Systolic blood pressure
TICAM1	TIR domain-containing adaptor molecule 1
TLR	Toll-like receptor
TRIF	Toll-interleukin receptor-domain containing adaptor protein inducing interferon- $\!\beta$
Trif ^{mut}	C57BL/6J- <i>Ticam1Lps2</i> /J mice

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Novelty and Significance

What is new?

We discovered a dual simultaneous activation by Ang II of two innate immune signaling pathways. One that includes TRIF, as its adaptor protein, is essential for Ang II-induced hypertension, cardiac hypertrophy, and pro-inflammatory gene expression, whereas the other that is dependent on adaptor protein MyD88 mediates an anti-inflammatory response that restrains Ang II hypertension, hypertrophy and inflammatory gene expression.

What is relevant?

Ang II-induced hypertension and end organ damage depend on inflammatory immune responses. However, the mechanisms of Ang II-induced inflammatory responses are not fully understood. Now, the knowledge of the specific adaptor proteins that regulate innate immune inflammatory pathways provides opportunities for novel interventions to treat hypertension and end organ damage.

Summary

We have discovered that a TRIF adaptor-dependent inflammatory pathway of the innate immune system rather than the common MyD88 adaptor-dependent pathway mediates Ang II-induced hypertension, cardiac hypertrophy, and pro-inflammatory gene expression. Paradoxically the MyD88 adaptor signaling activated by Ang II is antiinflammatory. This study provides valuable insight into the mechanisms of hypertension and the potential for developing new therapies.



Figure 1.

Systolic blood pressure (SBP) measured during infusion of saline (black lines) or Ang II (3000 ng/kg/min, red lines) for 3 weeks in (A) WT, (B) $MyD88^{-/-}$ and (C) $Trif^{mut}$ mice. Values represent means \pm SEM (n 3 mice in each group). The pressor response was significantly greater in the last week of Ang II infusion in $MyD88^{-/-}$ than in WT (*p <0.05). Conversely, the pressor response during Ang II infusion was abrogated in the $Trif^{mut}$ mice; the SBP was unchanged during infusion of saline and Ang II for the last two weeks (n.s.= p > 0.05).



Figure 2.

Measurement of heart weight to body weight ratios (HW/BW ratio) after 3 weeks of saline or Ang II infusions were determined in (A) WT, (B) $MyD88^{-/-}$, and (C) $Trif^{mut}$ hearts. Low (730 or 1000 ng/kg/min) or high (3000 ng/kg/min) doses of Ang II were used (ANOVA with post-hoc Tukey's test; n 3; n.s. = not significant, asterisks *= P < 0.05, ***= P < 0.001). Increase in heart weight to body weight ratios with Ang II vs. saline were greater in $MyD88^{-/-}$ and lesser in $Trif^{mut}$ compared to WT. Linear regression analyses and slopes of dose response curves are in supplemental Table S2.

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Figure 3.

Comparison of pro-inflammatory gene expression in hearts from WT, $MyD88^{-/-}$ and $Trif^{mut}$ mice after chronic infusion (3 weeks) of saline or low dose Ang II. Fold change in RNA expression was measured by C_T method using *Gapdh* RNA as a reference. Values in graphs are normalized to measurements in saline-infused mice. *Tnfa*= tumor necrosis factor-alpha, *Nox4*= NADPH oxidase 4, and *Agtr1a*= angiotensin receptor type 1a. (n 3 samples each group; Asterisks show P 0.05, n.s. = not significant). In general Ang II induced increase in gene expressions were enhanced in $MyD88^{-/-}$ and reduced in $Trif^{mut}$ compared to WT.



Figure 4.

Comparison of pro-inflammatory gene expression in kidneys of WT, $MyD88^{-/-}$ and $Trif^{mut}$ after chronic infusion (3 weeks) of saline or low dose Ang II. Fold change in RNA expression was measured by C_T method using *Gapdh* RNA as a reference. Values in graphs were normalized to measurements in saline-infused mice (n 3 each group; Asterisks show P 0.05, n.s. = not significant). Ang II induced increase in gene expressions were enhanced in $MyD88^{-/-}$ and reduced in $Trif^{mut}$ compared to WT.



Figure 5.

Baseline RNA expression of TLR3, TLR4 and their adaptor proteins MyD88 and TRIF in the hearts of WT vs. $MyD88^{-/-}$ and WT vs. $Trif^{mut}$ mice. Comparisons of RNA expressions between WT and $MyD88^{-/-}$ or WT and $Trif^{mut}$ hearts were done by QPCR and C_T method. Statistical significance was tested by unpaired t-test. Asterisks show significant differences from WT (P < 0.05) whereas n.s. denotes no significant change. In $MyD88^{-/-}$ the expressions of Trif, Tlr3, and Tlr4 were increased. In $Trif^{mut}$, the expression of MyD88 was increased, Tlr4 was decreased and Tlr3 unchanged.



Figure 6.

Schematic portrays Ang II induced hypertension, cardiac hypertrophy and pro-inflammatory gene expression in innate immune cells of WT, $MyD88^{-/-}$ and $Trif^{mut}$ mice. The dual activation of MyD88 and TRIF pathways through TLR and AT1 receptors is shown in WT cells to induce NF- κ B and interferon (IFN), respectively. The TRIF-dependent and negative regulatory influence of MyD88 are highlighted.