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## Elevated Vasopressin in Pregnant Mice Induces T Helper Subset Alterations Consistent with Human Preeclampsia

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### Abstract

The pathogenesis of preeclampsia, a hypertensive disorder of pregnancy, involves imbalanced T helper (T<sub>H</sub>) cell populations and resultant changes in pro- and anti-inflammatory cytokine release. Elevated copeptin (an inert biomarker of arginine vasopressin (AVP)), secretion precedes the development of symptoms in preeclampsia in humans, and infusion of AVP proximal to and throughout gestation is sufficient to initiate cardiovascular and renal phenotypes of preeclampsia in wild-type C57BL/6J mice. We hypothesize that AVP infusion in wild-type mice is sufficient to induce the immune changes observed in human preeclampsia. AVP infusion throughout gestation

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#### Author Contribution

S.M.S., D.A.S., W.S.H., E.J.D., H.A.D., C.D.S., J.L.G., and M.K.S. designed the experiments, experiments and data collection were performed by S.M.S., D.A.S., J.M.L., J.A.S., L.K.K., W.S.H., E.J.D., H.A.D., K.N.G-C., J.L.G., and M.K.S., data analysis was performed by S.M.S., D.A.S., E.J.D., and M.K.S., data interpretation was conducted by S.M.S., D.A.S., E.J.D., G.L.P., C.D.S., J.L.G., and M.K.S., and all authors contributed to and approved the final manuscript.

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in mice resulted in increased pro-inflammatory interferon gamma (IFN $\gamma$ ) (T<sub>H</sub>1) in the maternal plasma. The T<sub>H</sub>17-associated cytokine IL-17 was elevated in the maternal plasma, amniotic fluid, and placenta following AVP infusion. Conversely, the T<sub>H</sub>2-associated anti-inflammatory cytokine interleukin (IL)-4 was decreased in the maternal and fetal kidneys from AVP-infused dams while IL-10 was decreased in the maternal kidney and all fetal tissues. Collectively, these results demonstrate the sufficiency of AVP to induce the immune changes typical of preeclampsia. We investigated if T cells can respond directly to AVP by evaluating the expression of AVP receptors (AVPRs) on mouse and human CD4<sup>+</sup> T cells. Mouse and human T cells expressed AVPR1a, AVPR1b, and AVPR2. The expression of AVPR1a was decreased in CD4<sup>+</sup> T cells obtained from preeclampsia-affected women. In total, our data are consistent with a potential initiating role for AVP in the immune dysfunction typical of preeclampsia and identifies putative signaling mechanism(s) for future investigation.

### Keywords

APCs= antigen presenting cells; AVP= arginine vasopressin; AVPR= arginine vasopressin receptor; BLD= below the limit of detection; CLII= major histocompatibility complex II; Ct= cycle threshold; DC= dendritic cell; ICS= intracellular staining; IFN $\gamma$ = interferon gamma; IL= interleukin; LAP= latency associated peptide; MFTB= maternal fetal tissue bank; MNC= mononuclear cell; PD-L1= programmed death ligand 1; pg/g= picogram per gram; PIR B= paired immunoglobulin receptor B; PreE= preeclampsia; TGF $\beta$ = transforming growth factor beta; T<sub>H</sub>= T helper cell; Treg= T regulatory cell

### Introduction

Preeclampsia, a hypertensive disorder in pregnancy, affects 5–7% of all U.S. pregnancies and yet, disproportionately comprises 15% of all maternal-fetal morbidity and mortality (1). Preeclampsia is known to cause immediate and long-term maternal and fetal morbidities (2–7). Current methods of early diagnosis and prevention of preeclampsia are limited and provide at best a few months of lead time before clinical symptoms appear (8). Aspirin can be used as a preventative agent with varied effectiveness (9, 10). This prevention is partially thought to be due to an early modulation of immune responses. If we can unravel the early immune pathogenic mechanism(s) involved in the development of preeclampsia, there is potential for improved prevention.

Human maternal plasma copeptin, a stable biomarker of arginine vasopressin (AVP) secretion, is elevated mid- to late-pregnancy in women who develop preeclampsia (11–16). Importantly, copeptin is a robust early-pregnancy (as early as the 6th week of gestation) predictor of the development of preeclampsia (17). Further, chronic infusion of AVP in wild-type, pregnant C57BL/6J females is sufficient to model human preeclampsia by inducing pregnancy-specific hypertension, proteinuria, glomerular endotheliosis, and fetal growth restriction (17). These data support a role for AVP in the early pathogenesis of preeclampsia, and provide key insight toward the future development of therapeutic interventions for preeclampsia

While the development of preeclampsia is multifactorial and involves many systems and processes including inadequate trophoblastic invasion and poor spiral artery remodeling, an altered inflammatory response is thought to be involved in the early pathogenesis of preeclampsia (18–23). In healthy pregnancies, pro-inflammatory CD4+ T helper (T<sub>H</sub>) 1 related activity is dominant early and later shifts to a more anti-inflammatory T<sub>H</sub>2 type of immune response. An imbalance of these T<sub>H</sub>1 and T<sub>H</sub>2 cells occurs during preeclampsia (24–27). This paradigm has expanded to include an increase in pro-inflammatory T<sub>H</sub>17 cells (24, 28–30). Although the precise mechanisms involved in the development of preeclampsia are complex and poorly understood, an aberrant pro-inflammatory T<sub>H</sub> cell response is clearly involved (18, 21, 31–34).

Roles for AVP in the regulation of blood pressure and fluid homeostasis are generally accepted, however, a role for AVP in immune system function is less well appreciated. AVP secretion stimulates pro-inflammatory cytokine secretion and activation of lymphocytes (35–38). The immune system is vital to the development of vascular dysfunction in hypertensive diseases. McMaster *et al.* review the evidence demonstrating T cells and T cell-derived cytokines, such as interferon gamma (IFN $\gamma$ ) and interleukin-17 (IL-17), play a role in the renal and vascular dysfunction present in hypertensive diseases (39). As in preeclampsia, hypertension is thought to skew T cells towards pro-inflammatory T<sub>H</sub>1 (IFN $\gamma$ ) and T<sub>H</sub>17 (IL-17) dominant phenotypes (40). T<sub>H</sub> IFN $\gamma$  production is a dynamic interaction that requires the presence of IL-2. Suppressor cells absorb IL-2, preventing induction of IFN $\gamma$  secretion and inhibiting T<sub>H</sub>1-associated responses (36, 37, 41). As reviewed by Chikanza *et al.*, AVP has been shown to boost T<sub>H</sub>1 responses *in vitro* and *in vivo* via enhancement of IFN $\gamma$  (42, 43). Further, AVP has been shown to replace the requirement of IL-2 for the production of IFN $\gamma$ , thus inhibiting appropriate down-regulation of T<sub>H</sub>1 responses. Further, Johnson *et al.* show that AVP induces proliferation of C57BL/6 thymocytes in culture (36, 37, 44). Taken together, these data support the potential for AVP to induce increased IFN $\gamma$  (and thus a T<sub>H</sub>1 response) in an IL-2 independent manner to circumvent appropriate conversion from an inflammatory T<sub>H</sub>1 response needed for placentation and spiral artery remodeling to a more anti-inflammatory T<sub>H</sub>2 environment. To our knowledge, the potential role of T<sub>H</sub>17 and IL-17 in the regulation of AVP production has not been investigated. Further, this immunologic role of AVP has not been assessed in pregnancy. Thus, AVP is uniquely positioned to potentially initiate known mechanisms of preeclampsia, and may therefore represent an early cause of preeclampsia. The objective of this study is to test the hypothesis that AVP infusion during pregnancy in wild-type C57BL/6J mice is sufficient to induce the immunologic alterations observed in human preeclampsia. These data may provide novel insights into immune mechanisms mediated by AVP in preeclampsia.

## Materials and Methods

### Animal Studies

The animal procedures used were approved by the University of Iowa Institutional Animal Care and Use Committee. Wild-type 12–16 week old C57BL/6J male and female mice were obtained from Jackson Laboratories and maintained on standard chow under standard care conditions. Virgin females were subcutaneously implanted with osmotic mini-pumps (Alzet

Model #1004, Cupertino, CA) infusing either saline or AVP (24 ng/hr, Sigma, St. Louis, MO) as previously described (17). Three days after implantation, dams were individually mated for a single overnight period. Pregnancies were timed by post-coital vaginal plugs indicative of gestational day (GD) 0.5. Maternal and fetal tissues were harvested on GD 18. Data were collected from multiple pregnancies from independent experiments. A subset of the cohort presented in the current study was previously published by Santillan *et al.* (17) and subsequent experiments have confirmed the preeclampsia phenotype (pregnancy-specific hypertension, proteinuria, fetal growth restriction, and kidney glomerular endotheliosis). Each pregnancy was considered N=1 for maternal tissues. Due to fetal tissue mass, 5 pairs of fetal kidneys and 5 fetal livers were pooled for an N=1 from a single pregnancy. Maternal tissues (plasma, kidney, and liver) and fetal tissues (amniotic fluid, kidney, liver, and placenta) had N = 5 per group from at least two independent experiments. All whole tissues (except spleen) and plasma were stored at -80°C until protein extraction and analysis. Maternal spleen was kept on ice in phosphate buffered saline supplemented with 2% fetal bovine serum for immediate dissociation of cells.

### Tissue Protein Extraction and Protein Analysis

Total protein lysate was generated by homogenization of tissues in buffer containing 5M NaCl, 1M Tris, 0.5M EDTA, NP-40, protease inhibitor (Roche, Switzerland), and phosphatase inhibitor (Roche, Switzerland). A commercially available bicinchoninic acid (BCA) protein assay kit (Thermo Fisher Scientific, Waltham, MA) was utilized per protocol to assess total protein concentration. Plasma and tissue protein extracts were diluted where needed and commercially available ELISAs for mouse pro-inflammatory IFN $\gamma$  (T<sub>H</sub>1) and IL-17 (T<sub>H</sub>17) and anti-inflammatory IL-4 (T<sub>H</sub>2), IL-10 (T<sub>H</sub>2), and TGF $\beta$  cytokines were performed per protocol in duplicate (eBioscience, San Diego, CA). Cytokine concentrations were normalized to total protein and are presented as picograms/gram of total protein (pg/g).

### Cell Preparation and Flow Cytometric Analysis

Single cell suspensions were prepared by mechanical maceration from spleens of saline and AVP-infused dams followed by density gradient separation (FicoLite LM, Atlanta Biologicals, Lawrenceville, GA) to obtain MNCs. Cell suspensions were then stained with fluorochrome-conjugated or biotinylated antibodies (Supplemental Table 1), followed by streptavidin-conjugated fluorochromes. Fluorochrome-conjugated, purified rat immunoglobulins were used as isotype controls for background fluorescence. All cell samples were incubated with anti-CD16/32 (clone 2.4G2) and rat serum during staining to prevent background Fc $\gamma$ R binding. Intracellular Fixation and Permeabilization Buffer Set (eBioscience, San Diego, CA) was used per protocol for intracellular staining (ICS) of cytokines. Following staining, cells were fixed with either 0.1% formaldehyde or intracellular fixation buffer (eBioscience, San Diego, CA) where appropriate. Flow cytometric data were obtained within 24 hours using a Becton Dickinson LSR II (San Jose, CA) and analyzed using FlowJo software (Treestar Inc., Ashland, OR). Dead cells were excluded by forward/orthogonal light scatter characteristics. Single cells were identified via forward scatter-area (FSC-A) versus side scatter-width (SSC-W). Gating strategies of CD4+ T cells and dendritic cells (DCs) are shown in Supplemental Figure 2.

## Human Studies

Human mononuclear cells (MNCs) from control (n=27) and preeclampsia-affected (n=24) pregnancies were obtained from the University of Iowa Maternal Fetal Tissue Bank (MFTB, IRB# 200910784). The MFTB is a pregnancy focused human biorepository with clinically annotated biosamples collected throughout gestation with quality control measures for clinical data and biosample integrity. As a prospective, cross-sectionally collected biorepository and clinical data warehouse, early pregnancy samples are collected before the onset of diagnoses. Coded clinical data were obtained through the MFTB as previously published (45), which derives data from our Clinical Research Data Warehouse. The human control and preeclampsia-affected samples used in this study are a subset of a previously published cohort (17) and no differences in characteristics were observed between groups (Supplemental Table 2). The diagnosis and classification of PreE was based on the standard American College of Obstetrics and Gynecology (ACOG) definitions for analysis (46). PreE cases were identified by cross-referencing the MFTB database with the bioinformatics query of ICD-9 and ICD-10 codes of bank participants at the time of delivery. The electronic medical record of each potential case was evaluated by the MFTB to confirm the diagnosis of PreE by the ACOG definitions (46). Case-control verification training of the MFTB and verification of cases and controls was led by the senior author (MKS) who is a clinical maternal fetal medicine specialist. Maternal age-matched plasma samples and corresponding clinical data for the control population were obtained by querying the MFTB database. Control pregnancies were pregnant women who did not develop PreE. The gestational age at the time of sample collection was recorded. MNCs are all processed and stored for viability in liquid nitrogen as previously described (45).

## CD4+ T cell isolation

Human MNCs obtained from the MFTB. Splenic murine mononuclear cells were prepared as described above (cell preparation). Human and mouse CD4+ T cells were negatively selected using species-specific EasySep CD4+ T Cell Enrichment Kits per protocol (Stemcell Technologies Inc., Vancouver, BC). Human and mouse CD4+ T cell purity was determined to be 90% via flow cytometry.

## Quantitative PCR

Total cellular RNA was purified from cells using the mirVana miRNA isolation per protocol (ThermoFisher Scientific, Waltham, MA). RNA concentration and purity were determined using a NanoDrop 1000 spectrophotometer (ThermoFisher Scientific, Waltham, MA). Quantitative PCR (qPCR) gene expression assays of AVP receptors 1a (AVPR1a), 1b (AVPR1b), and 2 (AVPR2) were performed following SuperScript III (ThermoFisher Scientific, Waltham, MA) reverse transcription of fixed total RNA mass (500ng). Resulting cDNAs were amplified using primer sets shown in Table 1. All primer efficiencies were between 93–95%. Amplicons were detected in a PowerSYBR Green qPCR assay carried out on an Applied Biosystems 7900HT Real Time PCR System in the Genomics Division of the Iowa Institute of Human Genetics (IIHG). Raw cycle threshold (Ct) values were normalized (Ct) against the 18S rRNA endogenous control. The Ct and expression fold change were calculated as previously described (47, 48).

## Statistical Analysis

For continuous variables, a two sided Student's t test with unequal variance was utilized (GraphPad, Prism 7, La Jolla, CA). In addition, chi square was utilized for categorical variables. Statistical significance was designated at  $\alpha = 0.05$  or as determined by Bonferroni correction for multiple comparisons ANOVA.

## Results

### AVP infusion throughout gestation heavily skews maternal T<sub>H</sub> immunity toward a pro-inflammatory T<sub>H</sub>1 and T<sub>H</sub>17 phenotype

We previously demonstrated infusion of AVP into pregnant mice results in the hallmark features of human preeclampsia including, pregnancy-specific hypertension, proteinuria, renal glomerular endotheliosis, and fetal growth restriction (17). In the current study, we utilize this established mouse model of preeclampsia to investigate immune alterations caused by elevated AVP in pregnancy. The T<sub>H</sub>1- and T<sub>H</sub>17- associated pro-inflammatory cytokines, IFN $\gamma$  and IL-17, were significantly increased in the plasma of AVP-infused dams and unaffected by AVP in the maternal kidney and liver (Figure 1). T cells are activated in secondary lymphoid organs (eg. spleen); and consistent with circulating plasma IFN $\gamma$  and IL-17 elevations, we observed increases in both T<sub>H</sub>1 and T<sub>H</sub>17 CD4<sup>+</sup> T cells in the spleen of AVP-infused dams (Figure 1). These data suggest elevated AVP during pregnancy induces a pro-inflammatory T<sub>H</sub>1 and T<sub>H</sub>17 milieu in pregnant mice similar to that observed in human preeclampsia-affected women.

### Elevated AVP in pregnancy decreases anti-inflammatory T<sub>H</sub>2 and TGF $\beta$ cytokines in pregnant dams

Although we did not observe an AVP-induced change in the frequency of splenic CD4<sup>+</sup> T cells producing anti-inflammatory cytokines during pregnancy, there were tissue-specific changes in anti-inflammatory cytokines (Figure 2). The T<sub>H</sub>2-associated cytokines, IL-4 and IL-10, as well as the anti-inflammatory cytokine TGF $\beta$ , were all decreased in the maternal kidneys of AVP-infused dams. Additionally, there was a significant decrease in plasma TGF $\beta$  in these animals. The levels of IL-4, IL-10, and TGF $\beta$  were all comparable between the saline and AVP dams in the maternal liver (Figure 2). These data demonstrate AVP infusion during pregnancy results in an overall reduction in maternal anti-inflammatory cytokines.

### AVP infusion in pregnancy decreases Treg and alters the ratio of Treg:T<sub>H</sub>17 cells

It has been observed in both humans and in animal models that Tregs are reduced and the ratio of Treg to T<sub>H</sub>17 cells is altered in preeclampsia (24, 28, 33). Given the changes observed in the anti-inflammatory cytokines IL-10 and TGF $\beta$  (Figure 2), we next evaluated Treg. Elevated AVP resulted in a decrease in anti-inflammatory Treg cells (Figure 3A) as well as an altered Treg:T<sub>H</sub>17 ratio (Figure 3B).

### **Dendritic cells from AVP-treated dams show enhanced co-stimulatory molecule expression**

Dendritic cells (DCs) are potent antigen presenting cells (APCs) that activate and influence  $T_H$  cell differentiation. This process involves not only cytokine production by both cell types, but also surface molecule interactions. The expression of co-stimulatory and inhibitory molecules by DCs was evaluated to determine if elevations in AVP during pregnancy alters the activation phenotype of DCs. The co-stimulatory surface molecules major histocompatibility complex class II (CLII), CD80, and CD86 are highly expressed on CD11c+ DCs from AVP-treated versus saline-treated dams (Figure 4A). Concomitantly, the expression of the inhibitory surface molecules programmed death ligand 1 (PD-L1) and paired immunoglobulin receptor B (PIR B) were significantly decreased on DCs from AVP-treated dams (Figure 4B). These data show that AVP programs DCs toward an activated, stimulatory phenotype and thus may enhance antigen presentation and pro-inflammatory T cell activation in preeclampsia.

### **Fetal exposure to elevated AVP in utero results in enhanced $T_H17$ and impaired $T_H2$ cytokine profiles**

Unlike maternal tissues, IFN $\gamma$  was not elevated in the fetal tissues from GD 18 AVP-infused dams (Figure 5A). Congruent with increased maternal IL-17, the amniotic fluid, placenta, and fetal kidney all showed increased IL-17 levels compared to tissues from saline infused dams, suggesting increased  $T_H17$  cells (Figure 5B). In addition to increases in pro-inflammatory IL-17, fetal tissues had marked decreases in the anti-inflammatory cytokines IL-4 ( $T_H2$ ), IL-10 ( $T_H2$ ), and TGF $\beta$ . IL-4 was significantly decreased in the amniotic fluid, placenta, and fetal kidney, while IL-10 was decreased in all fetal tissues, including fetal liver (Figure 6A and 6B). These data suggest a global alteration in  $T_H2$  cells in fetal tissues when exposed to elevated AVP during gestation. Further contributing to an elevated pro-inflammatory over an anti-inflammatory cytokine balance, the anti-inflammatory cytokine TGF $\beta$  was also decreased in the placenta and fetal kidney from AVP-infused pregnancies (Figure 6C). In humans, it is unknown if maternal AVP crosses the placenta. Radiolabeled  $I^{125}$ -AVP experiments in pregnant ewes suggest that AVP does not cross the placenta. Further, rat studies (49) suggest that the source of placental AVP is from the fetus. In total, our findings and others suggest that AVP is sufficient directly to cause preeclampsia related physiologic and immunologic phenotypes in the mother. Yet, the fetal physiologic and immunologic phenotypes may be due to a combination of direct stimulation at the maternal-fetal interface and/or a fetal response to the AVP induced maternal phenotypes.

### **Mouse and human CD4+ T cells express AVP receptors**

As  $T_H$  cells are altered in response to AVP during pregnancy in mice, CD4+ T cells were isolated and analyzed for expression of AVP receptors to determine if CD4+ T cells may be able to respond directly to elevated AVP. Interestingly, mouse and human CD4+ T cells expressed AVP receptors 1a, 1b, and 2 (Table 2 and 3). Similar to mouse CD4+ T cells, human CD4+ T cells highly expressed AVPR2, but AVPR1a was also highly expressed (Table 3). To focus on potential early and late pregnancy changes in circulating CD4+ T cells, AVPR expression was determined in the first and third trimester of human

pregnancies. CD4+ T cells isolated from human preeclampsia-affected pregnancies expressed significantly lower AVPR1a in the first trimester, with a 6.3 fold decrease in expression (Table 3). Additionally, although AVPR1b was not as highly expressed AVPR1a and 2, there was a 3.7 fold decrease in expression by CD4+ T cells isolated from preeclampsia-affected pregnancies during the first trimester. Interestingly, the changes in AVPR1a and AVPR1b expression observed normalized by the third trimester. Although AVPR2 expression was not altered in the preeclampsia-affected group in the first trimester, it was 1.7 fold increased the third trimester. These data not only demonstrate that CD4+ T cells express AVP receptors, but also that expression of AVPRs is differentially regulated throughout pregnancy in human preeclampsia-affected CD4+ T cells.

## Discussion

The early pathogenesis of preeclampsia involves the failure of the maternal immune system to normally tolerate the pregnancy (21). Previous studies have indicated that during a healthy human pregnancy, a shift occurs from a pro-inflammatory T<sub>H</sub>1 (IFN $\gamma$  dominant) CD4+ T cell response required early for appropriate placentation to an anti-inflammatory T<sub>H</sub>2 (IL-4 and IL-10 dominant) response required for fetal tolerance. The concept that persistent T<sub>H</sub>1 related fetoplacental intolerance is important in the pathogenesis of preeclampsia has been demonstrated by our lab and others (50–53). T<sub>H</sub>1 responses and related cytokines (IFN $\gamma$ ) directly inhibit the development of T<sub>H</sub>2 responses and vice versa (31, 54, 55). IFN $\gamma$  is elevated in human preeclampsia with a concurrent decrease in IL-10 and IL-4 production, driving a T<sub>H</sub>1 response (55–58). These T<sub>H</sub>1 cells further increase IFN $\gamma$  production, resulting in increased recruitment and activation of antigen presenting cells and cytotoxic CD8+ T cells. Recently, the production of IL-17 and the percent of T<sub>H</sub>17 cells have been shown to be increased in preeclampsia and potentially contribute to poor fetal tolerance (28, 33). Taken together, our data extend previous studies indicating there is both an increase in T<sub>H</sub>1 and T<sub>H</sub>17 cells, by identifying AVP as a potential, novel contributor to poor fetal tolerance and the development of preeclampsia.

For the first time, our data show that AVP-infusion during mouse pregnancy induces the T<sub>H</sub> subset changes and pro-inflammatory milieu similar to that observed in human preeclampsia. In mouse maternal circulation, there is a pro-inflammatory T<sub>H</sub>1 and T<sub>H</sub>17 milieu with elevated IFN $\gamma$  and IL-17 and a concomitant decrease in anti-inflammatory TGF $\beta$ . Correlating with previously observed AVP-induced renal changes (17), there were maternal kidney specific decreases in the T<sub>H</sub>2-associated anti-inflammatory cytokines IL-4 and IL-10, as well as anti-inflammatory TGF $\beta$ . Our observed differences in cytokines between the mouse maternal circulation and target tissues, underscores the concept that circulating immune responses may not completely represent tissue-specific immune responses. In addition to producing soluble proteins that are detectable in the circulation, T cells travel to the site of inflammation to exert their function. Although IFN $\gamma$  is elevated in the maternal plasma of our AVP-infused dams, but not in the tissues evaluated, it is likely IFN $\gamma$  is elevated in the spleen (given the observed increase in splenic CD4+ IFN $\gamma$ + T cells). As T cells are activated in the secondary lymphoid organs and also influence other immune responses, the observed changes in maternal plasma IFN $\gamma$  may be acting at the level of the spleen. In regards to IL-17, the elevation observed in the circulation does correlate with a



known target organ in preeclampsia, the placenta. Lastly, immune cells in the target organs may also locally produce cytokines and further contribute to tissue inflammation and disease progression. Collectively, our data expands what other groups have observed in mouse models and in human preeclampsia (18, 25, 27, 28, 33, 56, 59, 60), in that we observed changes in T<sub>H</sub>1/T<sub>H</sub>2/T<sub>H</sub>17 associated cytokines in our AVP-induced mouse model of preeclampsia, highlighting AVP as a novel, potential immunologic agent in the development of preeclampsia.

We contend that AVP is an important initiating pathway in the early immunologic phenotype of preeclampsia. T<sub>H</sub>1 cells, seen in abundance in preeclampsia, are potent sources of IFN $\gamma$  which can induce AVP production (61). AVP secretion (via copeptin detection) is elevated very early in human preeclampsia (17) and is capable of inducing the early immunologic changes observed in the first trimester preeclampsia. Additionally, it has been demonstrated that the immune system, more specifically T<sub>H</sub>1 and T<sub>H</sub>17 cells, play a pivotal role in the development of vascular dysfunction (39, 40). We observed AVP-induced changes in both the maternal and fetal compartments. Although toward the end of mouse gestation, GD 18 mouse pups developmentally correspond to GD 58–60 in humans (62–64), which falls well within the first trimester of human gestation. Our mouse fetal immune findings may translate into early first trimester immunologic changes in the human fetus. This is consistent with the current understanding of a very early initiating cause of preeclampsia, as first trimester immunologic dysregulation lies upstream of the placental/vascular phenotype (21, 65). These observations also further the concept that AVP has immune effects beyond the canonical blood pressure/volume control. Together, these data suggest that AVP, whether via direct interaction with immune cells or indirectly via alterations in cardiovascular and renal function, may be an early cause of immune changes in preeclampsia.

Although the exact role of AVP in immune responses still requires further investigation, AVP is thought to stimulate immune cell activity in non-pregnant environments (35–38, 44, 66). As reviewed by Chikanza *et al.*, rats with chronic inflammatory disease also had elevated levels of plasma AVP. A reduction in circulating AVP decreased the pro-inflammatory response in these animals (67). *In vitro* experiments demonstrated a dose-dependent augmentation of autologous mixed lymphocyte reactions, including the requirement of IL-2 for IFN $\gamma$  production (36, 44). Further, addition of IL-1 beta (IL-1b) or IL-6 to *in vitro* cultures resulted in a dose-dependent increase in AVP production (68). Lastly, human cancer patients administered IL-6 had significantly higher circulating AVP levels within two hours of administration (69).

To our knowledge, the role of AVP in immune responses has not been investigated in pregnancy, and more specifically, not in preeclampsia-affected pregnancies. Here, we demonstrate that in pregnancy, mouse and human CD4<sup>+</sup> T cells express AVP receptors, and that AVPR1a and AVPR2 are highest expressed. Interestingly, AVPR1a and 1b expression by human CD4<sup>+</sup> T cells is significantly down-regulated in preeclampsia-affected pregnancies in the first trimester and this normalizes by the third trimester. AVPR1a is expressed in blood vessels and is known to play a role in baroreceptor reflexes and blood volume homeostasis (70–72). Although high placental AVPR1a expression has also been proposed as a potential mediator early in pregnancy for increased blood flow and appropriate placental growth (73),

these data suggest that AVPR1A may also mediate immune responses during preeclampsia and provide insight into potential therapeutic targets. In the third trimester, CD4<sup>+</sup> T cells from preeclampsia-affected pregnancies increased expression of AVPR2 compared to cells from normotensive control pregnancies. AVPR2 is highly expressed in the kidney and is pivotal in renal water reabsorption and thus urine concentration (74, 75) and under conditions of elevated AVP, it effects renal vasoconstriction (76). T cells have been shown to play a role in hypertension and renal dysfunction. More specifically, Increased CD4<sup>+</sup> IL-17 producing T cells are seen in hypertension and these cells traffic to the kidney and vasculature to cause dysfunction that leads to hypertension (77, 78). Renal dysfunction and hypertension are often observed in PreE in the third trimester. Our data showing an increase in inflammatory T<sub>H</sub>17 cells with altered AVPR2 expression suggests this receptor may play a currently unknown role in immune responses and PreE toward the end of pregnancy. Although the function of the expression of AVPRs on CD4<sup>+</sup> T cells in PreE is currently unknown, our data suggests these receptors may be a previously uncharacterized link between renal, cardiovascular, and immune dysfunction in PreE.

APCs are critical to the activation and programming of T<sub>H</sub> cells. AVP in pregnancy is sufficient to induce the up-regulation of co-stimulatory (CD80, CD86, and CLII) and down-regulation of inhibitory (PD-L1 and PIR B) molecule expression on DC, resulting in the generation of more T<sub>H</sub>1 and T<sub>H</sub>17 cells, which are necessary in the early pathogenesis of preeclampsia. Increased expression of co-stimulatory molecule CD86 has been shown to correlate highly with poor maternal-fetal immune balance (79). Blockade of CD86 in pregnancy allows for expansion of protective Treg cells and promotes the differentiation of a more tolerant T<sub>H</sub>2 environment (80). The expression of PD-L1 during pregnancy has been shown to be T cell dependent and required to confer fetal tolerance (81). Both, co-stimulatory and inhibitory molecules, play a key role in maternal-fetal tolerance. Our data show that elevated AVP during pregnancy results in the up-regulation of co-stimulatory and down-regulation of inhibitory molecules by DCs. These alterations in signals received by the DCs are a likely mechanism contributing to the altered T<sub>H</sub> phenotype induced by elevated AVP during pregnancy and the development of preeclampsia. These alterations in the co-stimulatory phenotype of DCs in the spleen leads to a differentiation of CD4<sup>+</sup> T cells toward a T<sub>H</sub>1 and T<sub>H</sub>17 phenotype. Peripheral T<sub>H</sub>1 and T<sub>H</sub>17 cells, as well as, decidual cell populations skew the placenta toward a milieu rich in pro-inflammatory cytokines that furthers contributes to the development and pathogenesis of preeclampsia (Figure 7).

Our data demonstrate that elevated AVP in mice during pregnancy induces maternal and fetal T<sub>H</sub> related changes; these alterations mimic those previously observed in human preeclampsia-affected pregnancies. Additionally, the expression of AVP receptors by CD4<sup>+</sup> T cells identifies a putative signaling mechanism and new therapeutic targets for future investigation. From our studies, we propose that manipulation of the AVP pathway may be a novel preventative and therapeutic target to address the vascular, renal, and immune causes of preeclampsia.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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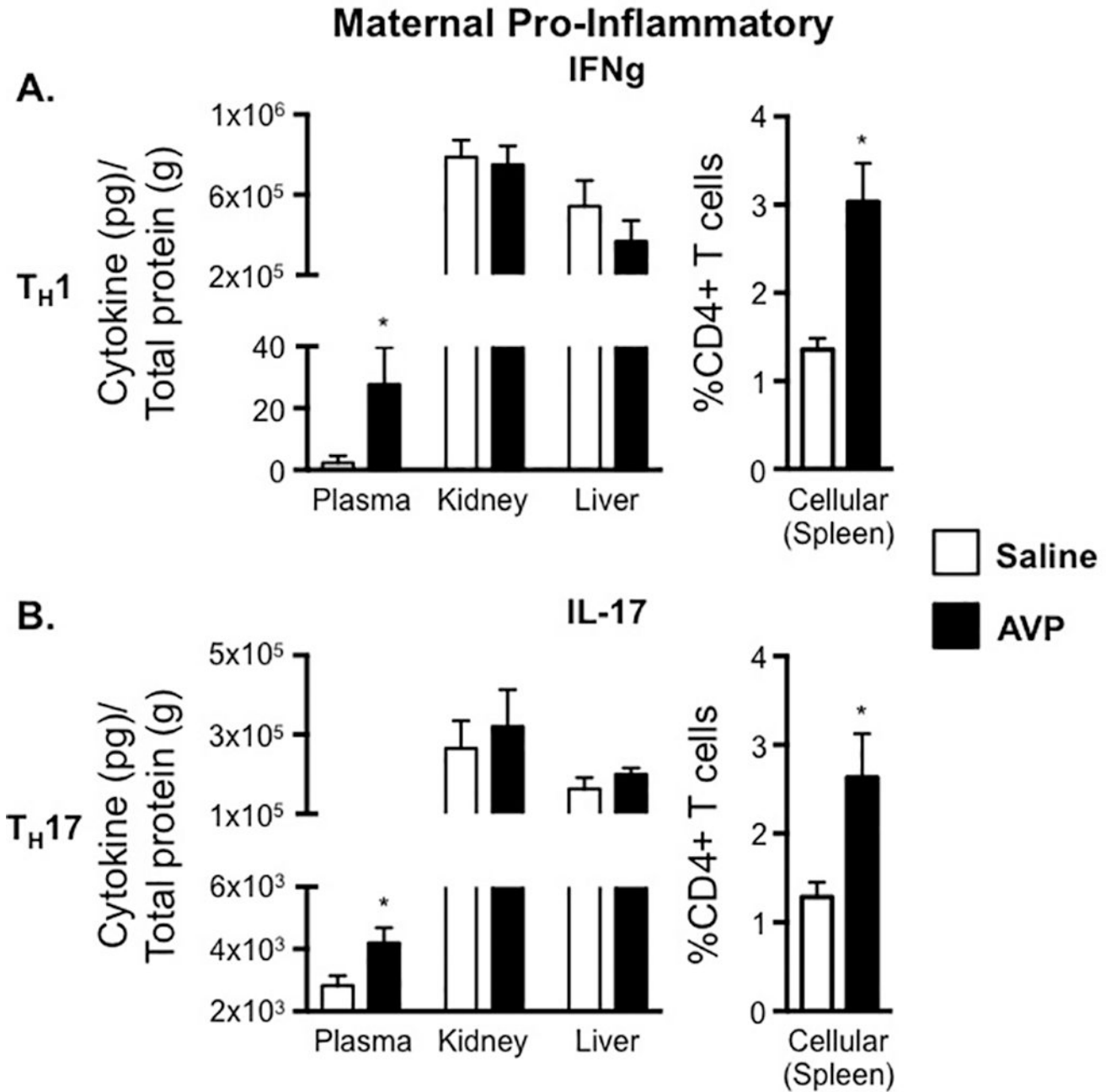
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### Clinical Perspectives

- AVP secretion during pregnancy, via copeptin detection, has been shown to be elevated very early in women who later develop preeclampsia. AVP infusion into pregnant mice recapitulates the physiological aspects of human preeclampsia. It is unknown if elevated AVP during pregnancy is sufficient to cause the maternal and fetal immune phenotypes consistent with human preeclampsia
- Elevated AVP in pregnant mice results in increased  $T_H1$  and  $T_H17$  associated cytokines and T cells with a concomitant decrease in the  $T_H2$  associated cytokines. Additionally,  $CD4^+$  T cells express AVP receptors and expression of AVPR1a by human  $CD4^+$  T cells from preeclampsia-affected women is decreased.
- The data presented demonstrate a novel role for AVP in altering immune phenotypes in pregnancy. Additionally, the expression of AVP receptor expression by  $CD4^+$  T cells lends to the possibility of therapies utilizing AVP modulation to mitigate preeclampsia.





**Figure 1.** AVP infusion is sufficient to induce T<sub>H1</sub> and T<sub>H17</sub> maternal immune alterations. IFN $\gamma$  (A) and IL-17 (B) concentrations in maternal plasma, maternal kidney, and maternal liver were normalized to total protein and are represented as pg of cytokine per g of total protein. Lymphocytes were isolated from the spleen of saline and AVP infused dams. Following intracellular cytokine staining, CD3<sup>+</sup> CD4<sup>+</sup> cells were gated as shown in Supplemental Figure 1A and the frequency of cells producing the pro-inflammatory cytokines IFN $\gamma$  (A) and IL-17 (B) was determined. Open bars= saline-infused; Solid bars= AVP-infused. N = 5 per group from at least two independent experiments. Data are mean  $\pm$  SEM. Statistical

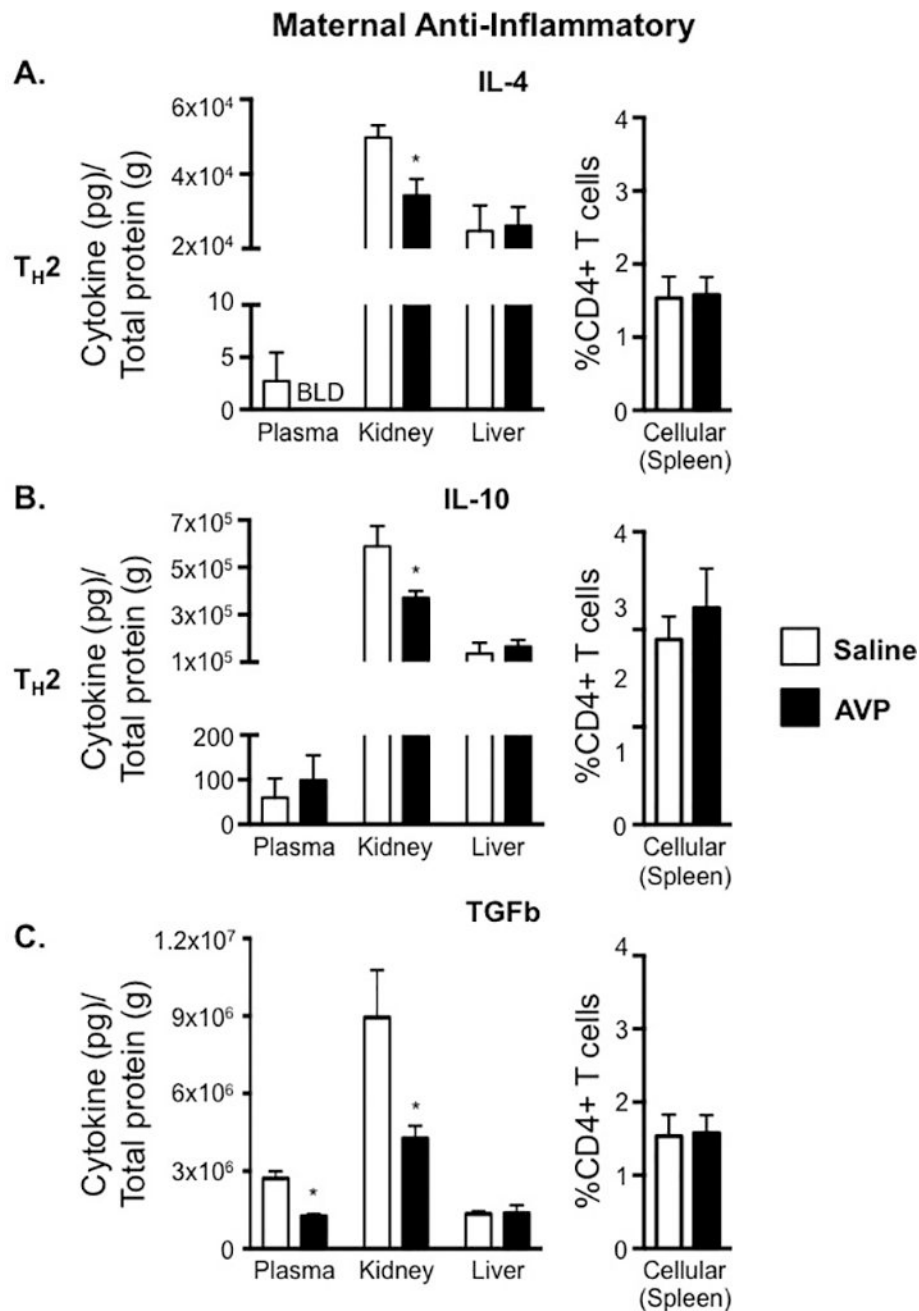
significance was determined using a Student t test and the minimal level of confidence deemed statistically significant was a p value  $<0.05$ .  $*=p<0.05$ .

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**Figure 2.**

Elevated AVP in pregnancy decreases anti-inflammatory cytokine production in maternal tissues. IL-4 (A), IL-10 (B), and TGFb (C) concentrations in maternal plasma, maternal kidney, and maternal liver were normalized to total protein and are represented as pg of cytokine per g of total protein. Lymphocytes were isolated from the spleen of saline and AVP infused dams. Following intracellular cytokine staining, CD3<sup>+</sup> CD4<sup>+</sup> cells were gated as shown in Supplemental Figure 1A and the frequency of cells producing the T<sub>H</sub>2-associated cytokines IL-4 (A) and IL-10 (B) as well as LAP/TGFb (C) was determined. Open bars= saline-infused; Solid bars= AVP-infused. BLD is below the limit of detection. N

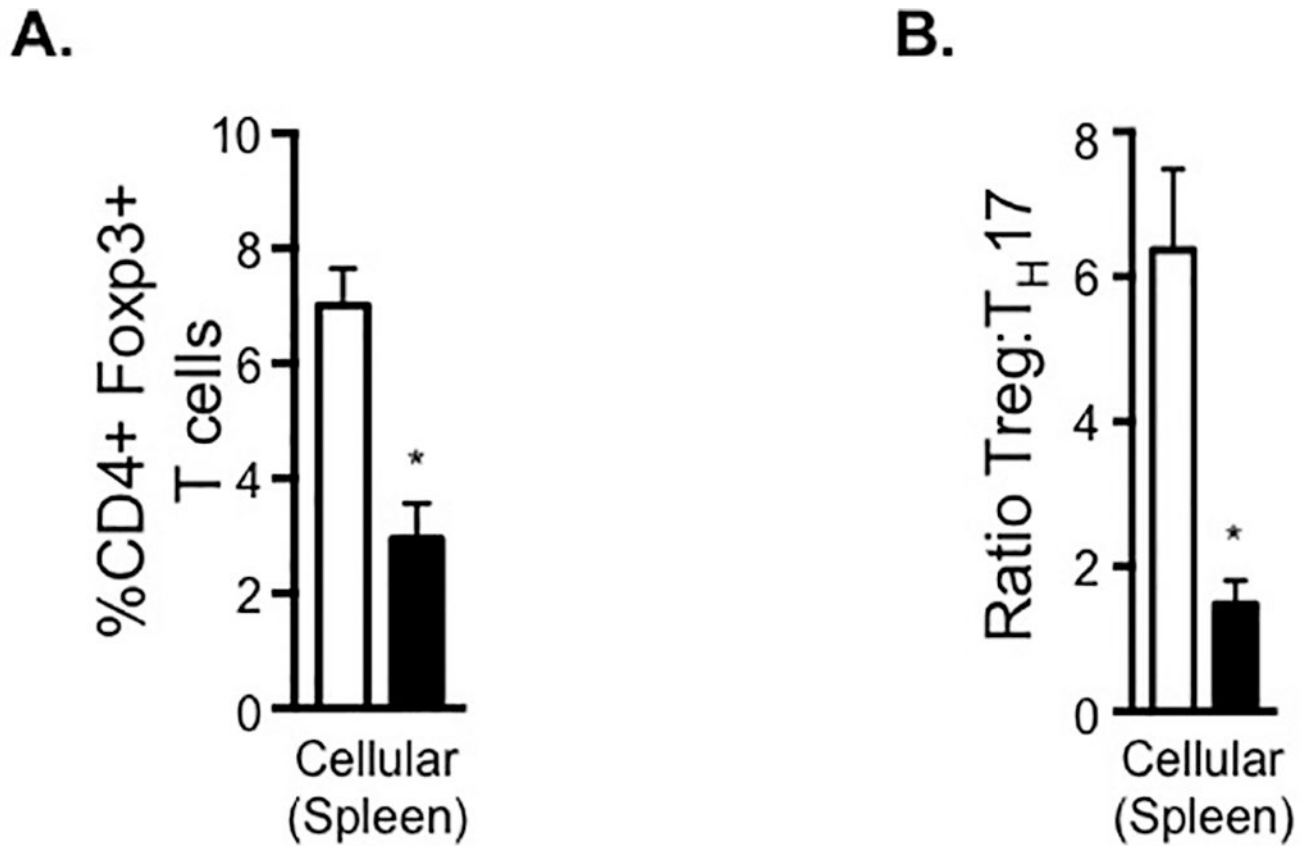
5 per group from at least two independent experiments. Data are mean  $\pm$  SEM. Statistical significance was determined using a Student t test and the minimal level of confidence deemed statistically significant was a p value  $<0.05$ .  $*=p<0.05$ .

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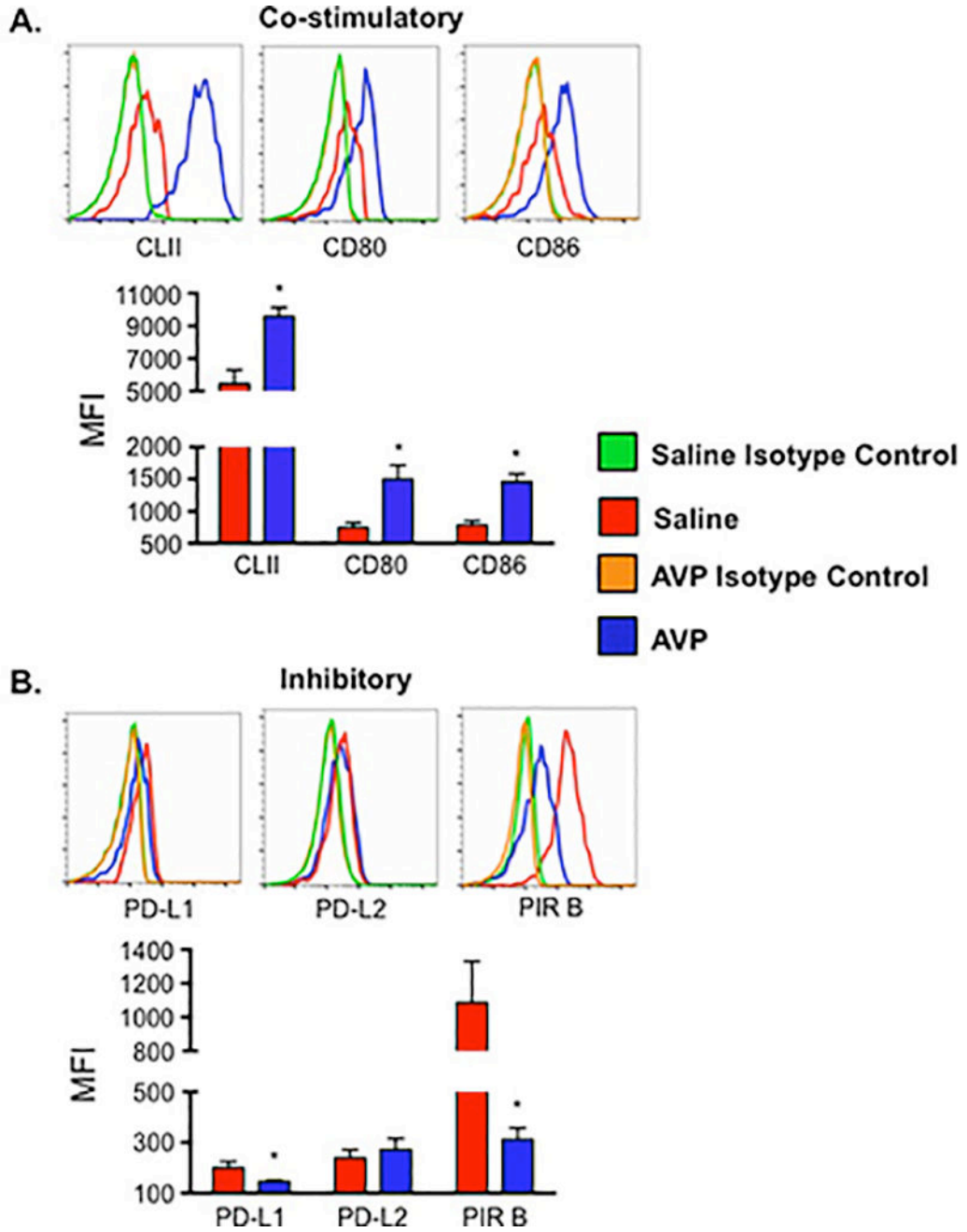
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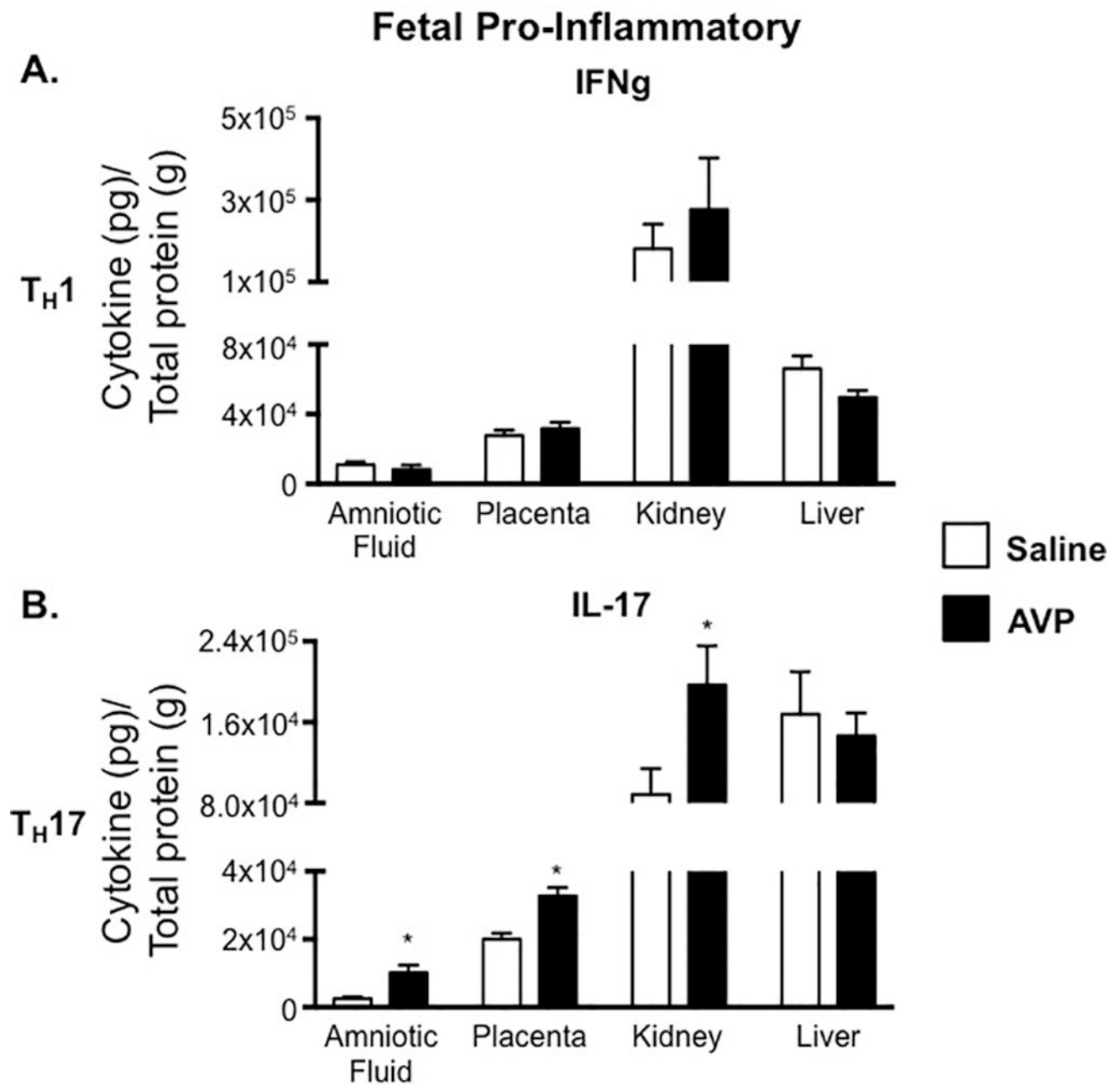
**Figure 3.**

AVP infusion in pregnancy alters the ratio of anti-inflammatory Tregs to pro-inflammatory TH17 cells. Lymphocytes were isolated from the spleen of saline and AVP infused dams. Following intracellular cytokine staining, CD3+ CD4+ cells were gated as shown in Supplemental Figure 1A and the frequency of cells producing the Treg transcription factor Foxp3 (A) or the pro-inflammatory cytokine IL-17 was determined. The ratio of Treg to TH17 cells was calculated (B). Open bars= saline-infused; Solid bars= AVP-infused. BLD is below the limit of detection. N = 5 per group from at least two independent experiments. Data are mean  $\pm$  SEM. Statistical significance was determined using a Student t test and the minimal level of confidence deemed statistically significant was a p value <0.05. \*= $p$ <0.05.



**Figure 4.** AVP induces altered surface receptor expression on DC. Inhibitory molecules paired immunoglobulin-like receptor B (PIR B) and programmed death ligand 1 (PD-L1) are decreased while co-stimulatory molecules MHC Class II (CLII), CD80, and CD86 were increased on dendritic cells from AVP infused dams. Lymphocytes were isolated from the spleen of saline and AVP infused dams. Following fluorescent antibody staining, CD11c+ DCs were gated as shown in Supplemental Figure 1B. (A) Representative histograms and mean fluorescence intensity (MFI) of co-stimulatory cell surface molecules. (B) Representative histograms and MFI of inhibitory cell surface molecules. Background

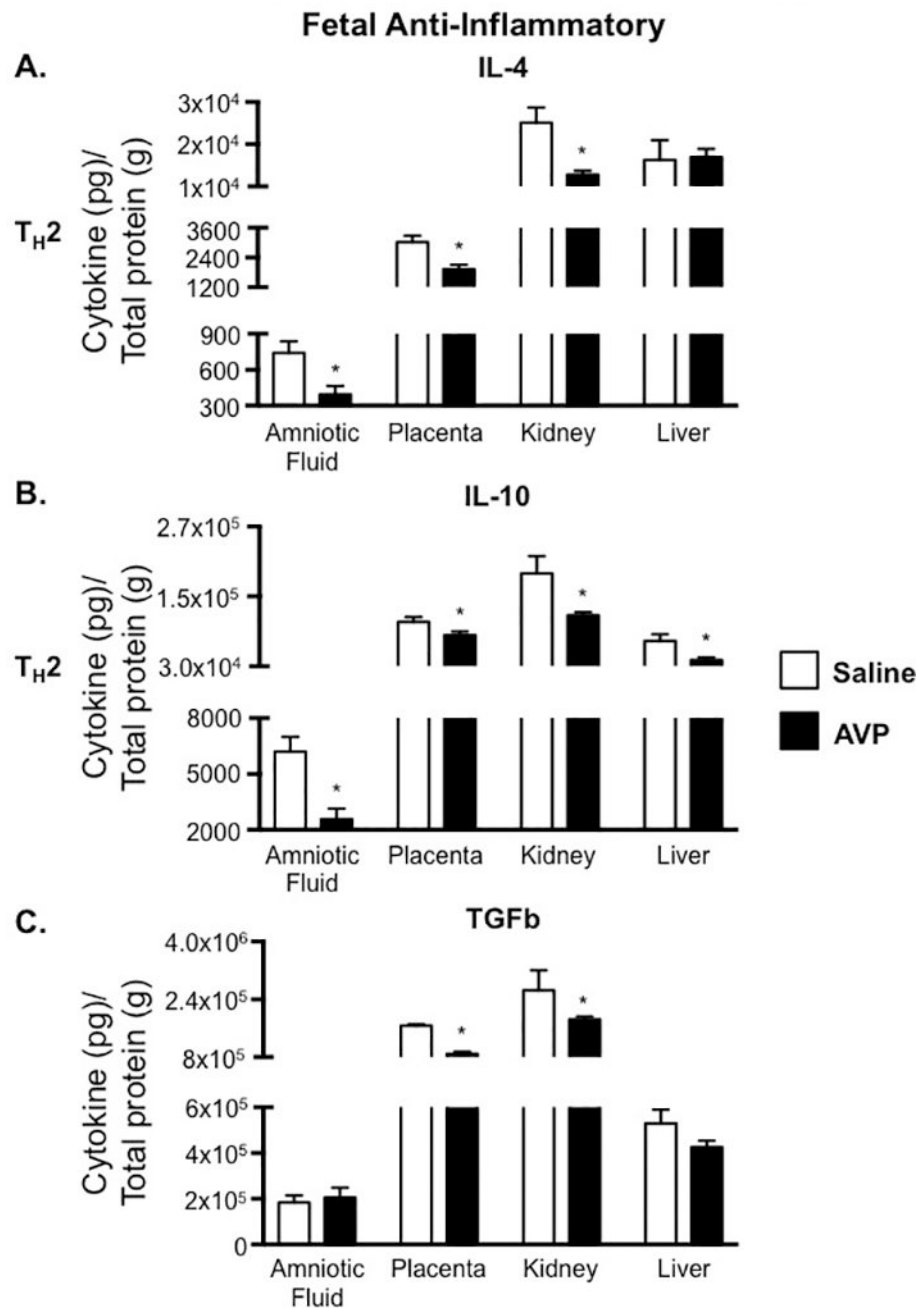
fluorescence was determined by staining cells with corresponding fluorochrome-conjugated rat immunoglobulin isotype control antibodies. Isotype control subtracted MFIs are shown. N = 5 per group from at least two independent experiments. MFIs are mean  $\pm$ SEM. Statistical significance was determined using a Student t test and the minimal level of confidence deemed statistically significant was a p value  $<0.05$ . \*= $p<0.05$ .



**Figure 5.**

Fetal tissue IL-17 cytokine production is altered by AVP. The T<sub>H17</sub>-associated cytokine IL-17 is increased in fetal tissues obtained from AVP-infused dams. IFN $\gamma$  (A) and IL-17 (B) concentrations in amniotic fluid, placenta, fetal kidney, and fetal liver were normalized to total protein and are represented as pg of cytokine per g of total protein. Open bars= saline-infused; Solid bars= AVP-infused. Due to fetal tissue mass, 5 pairs of fetal kidneys and 5 fetal livers were pooled for an N=1 from a single pregnancy. N = 5 per group from at least two independent experiments. Data are mean  $\pm$ SEM. Statistical significance was determined using a Student t test and the minimal level of confidence deemed statistically significant was a p value <0.05. \*=p<0.05.



**Figure 6.**

Fetal concentrations of anti-inflammatory cytokines are decreased in the presence of elevated AVP during gestation. IL-4 (A), IL-10 (B), and TGFb (C) concentrations in amniotic fluid, placenta, fetal kidney, and fetal liver were normalized to total protein and are represented as pg of cytokine per g of total protein. Open bars= saline-infused; Solid bars= AVP-infused. Due to fetal tissue mass, 5 pairs of fetal kidneys and 5 fetal livers were pooled for an N=1 from a single pregnancy. N = 5 per group from at least two independent experiments. Data are mean  $\pm$  SEM. Statistical significance was determined using a Student

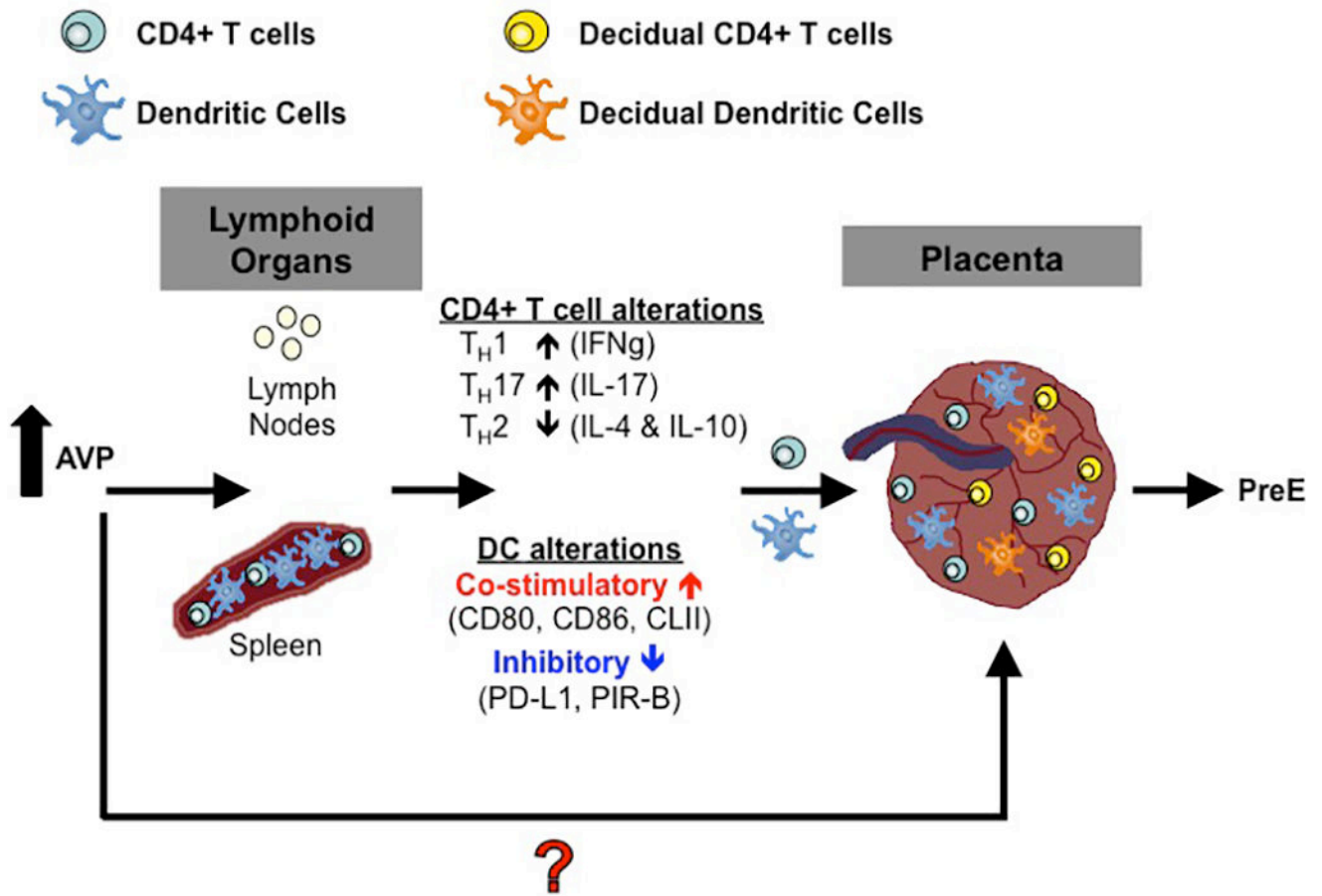
t test and the minimal level of confidence deemed statistically significant was a p value <0.05. \*=p<0.05.

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**Figure 7.** Proposed model of AVP action in pregnancy. Elevated AVP during pregnancy induces DCs in secondary lymphoid organs to upregulate co-stimulatory molecules and down-regulate inhibitory molecules. These DCs activate T cells and the resultant helper T cells secrete more pro-inflammatory  $T_H1$  and  $T_H17$  and less anti-inflammatory  $T_H2$  cytokines. These cells traffic to the placenta and along with resident decidual cell populations alter cytokine profiles in the placenta and fetal tissues. The red question mark indicates possibility of AVP acting directly on the placenta and tissue resident cell populations. PreE=preeclampsia.

**Table 1**

## qPCR Primer Sequences \*

Target#		Sequence	T <sub>m</sub>	Amplicon
hAVPR1a	S	GTGCAGAGCAAGCGGGTGTG	61.8°C	438bp
	AS	CGAGTCCTTCCACATAACCCGT	58.7°C	
mAVPR1a	S	CTCTGCTGGACACCTTCTTC	55.5°C	218bp
	AS	GTTGGGCTTCGGTTGTAGA	55.2°C	
hAVPR1b	S	CCAAGATCCGAACAGTGAAGAT	54.6°C	206bp
	AS	GCTGTTGAAGCCCATGTAGA	55.0°C	
mAVPR1b	S	AAGATCCGAACCGTGAAGATG	54.8°C	320bp
	AS	TGGGTCAGCAGTGTGTG	55.4°C	
hAVPR2	S	GGCCAAGACTGTGAGGATGA	56.9°C	200bp
	S	ACACGCTGCTGCTGAAAGAT	57.5°C	
mAVPR2	S	AGGACACCGGACAGGAA	55.7°C	275bp
	AS	AAAGCAGGCTACGCAACT	54.8°C	
18S rRNA	S	AAC TTTCGATGGTAGTCGCCG	57.3°C	104bp
	AS	CTTGGATGTGGTAGCCGTTT	57.6°C	

\* qPCR primers were designed using the Integrated DNA Technologies PrimerQuest online tool ([www.idtdna.com](http://www.idtdna.com)) against GenBank mRNA sequences.

#h = human-specific; m = mouse-specific; S = sense strand; AS = anti-sense strand; All primer efficiencies were between 93–95%.

**Table 2**

Murine CD4+ T cell expression of AVPRs

	Saline	AVP	Expression Fold Change	p-value
	Average CT			
AVPR1a	20.7 ±0.9	21.0 ±0.5	-1.3	0.72
AVPR1b	20.6 ±0.7	21.6 ±0.3	-2.0	0.19
AVPR2	9.7 ±0.6	10.5 ±0.4	-1.7	0.26

Mouse splenic CD4+ T cells were negatively purified from saline and AVP-infused dams. Expression of AVP receptors was determined via qPCR. Raw cycle threshold (Ct) values were normalized ( - Ct) against the 18S rRNA endogenous control. The lower the Ct value of a specific target, the higher the expression. The Ct and expression fold change of AVP versus saline and preeclampsia-affected versus control pregnancy were calculated as previously described (47, 48). Data are mean ±SEM. Statistical significance was determined using a Student's t test and the minimal level of confidence deemed statistically significant was a p value <0.05.

CD4+ T cells isolated from human preeclampsia-affected women have trimester-specific alterations in AVPR expression

**Table 3**

	1 <sup>st</sup> Trimester				3 <sup>rd</sup> Trimester			
	Control		PreE		Control		PreE	
	Average	CT	Fold Change	p-value	Average	CT	Fold Change	p-value
AVPR1a	9.4±0.6	12.1±0.3	-6.3	0.0009	10.2±0.4	10.5±0.3	-1.1	0.56
AVPR1b	20.4±0.7	22.3±0.5	-3.7	0.04	21.2±0.6	21.6±0.4	-1.3	0.62
AVPR2	8.0±0.3	8.7±0.3	-1.7	0.14	8.8±0.2	8.0±0.1	1.7	0.01

Human CD4+ T cells were negatively selected from peripheral blood mononuclear cells of control (1<sup>st</sup> trimester N=6; 3<sup>rd</sup> trimester N=15) and preeclampsia-affected (PreE; 1<sup>st</sup> trimester N=9; 3<sup>rd</sup> trimester N=12-14) pregnancies. Expression of AVPR receptors was determined via qPCR. Raw cycle threshold (Ct) values were normalized (Ct) against the 18S rRNA endogenous control. The lower the Ct value of a specific target, the higher the expression. The Ct and expression fold change of AVPR versus saline and preeclampsia-affected versus control pregnancy were calculated as previously described (47, 48). Data are mean ±SEM. Statistical significance was determined using a Student's t test and the minimal level of confidence deemed statistically significant was a p value <0.05.