

# Elevated Transglutaminase Activity Triggers Angiotensin Receptor Activating Autoantibody Production and Pathophysiology of Preeclampsia

Chen Liu, PhD;\* Renna Luo, MD, PhD;\* Serra E. Elliott, PhD; Wei Wang, MD, PhD; Nicholas F. Parchim, PhD; Takayuki Iriyama, MD, PhD; Patrick S. Daugherty, PhD; Sean C. Blackwell, MD; Baha M. Sibai, MD; Rodney E. Kellems, PhD; Yang Xia, MD, PhD

**Background**—Preeclampsia (PE) is a life-threatening hypertensive disorder of pregnancy associated with autoantibodies, termed AT<sub>1</sub>-AA, that activate the AT<sub>1</sub> angiotensin receptor. Although the pathogenic nature of these autoantibodies has been extensively studied, little is known about the molecular cause of their generation.

**Methods and Results**—Here we show that tissue transglutaminase (TG2), an enzyme that conducts posttranslational modification of target proteins, directly modified the 7-amino acid (7-aa) epitope peptide that localizes to the second extracellular loop of the AT<sub>1</sub> receptor. These findings led us to further discover that plasma transglutaminase activity was induced and contributed to the production of AT<sub>1</sub>-AA and disease development in an experimental model of PE induced by injection of LIGHT, a tumor necrosis factor superfamily member. Key features of PE were regenerated by adoptive transfer of purified IgG from LIGHT-injected pregnant mice and blocked by the 7-amino acid epitope peptide. Translating our mouse research to humans, we found that plasma transglutaminase activity was significantly elevated in PE patients and was positively correlated with AT<sub>1</sub>-AA levels and PE features.

**Conclusions**—Overall, we provide compelling mouse and human evidence that elevated transglutaminase underlies AT<sub>1</sub>-AA production in PE and highlight novel pathogenic biomarkers and innovative therapeutic possibilities for the disease. (*J Am Heart Assoc.* 2015;4:e002323 doi: 10.1161/JAHA.115.002323)

**Key Words:** AT<sub>1</sub>-AA • autoimmunity • preeclampsia • transglutaminase

Preeclampsia (PE) is a life-threatening disorder of pregnancy with high morbidity and mortality for both mothers and babies, with symptoms of hypertension, renal dysfunction, abnormal fetal growth, and circulating antiangiogenic factors.<sup>1–4</sup> PE<sup>5–11</sup> and other hypertensive conditions<sup>12,13</sup> are

characterized by the presence of autoantibodies, termed AT<sub>1</sub>-AAs, that activate the AT<sub>1</sub> angiotensin receptor (AT<sub>1</sub>R). These autoantibodies are found in the maternal circulation of 70% to 90% of PE patients<sup>5,11,14</sup> and are likely to contribute to disease by activation of AT<sub>1</sub>Rs on various cell types.<sup>5,6,8,9</sup> These autoantibodies cause the defining clinical features of PE when introduced into pregnant mice and are likely to contribute to these features in the majority of women with PE who harbor these autoantibodies.<sup>10</sup> Available evidence suggests that the antibody titers correlate to the severity of the disease.<sup>11</sup> Because AT<sub>1</sub>-AAs are significant contributors to the pathophysiology of PE, it is important to define the molecular basis initiating their production. These findings will provide a better understanding of the pathogenesis of PE and will likely identify new therapeutic options to interfere with AT<sub>1</sub>-AA production and disease progression.

AT<sub>1</sub>-AAs recognize a specific seven amino acid (7-aa) epitope (AFHYESQ) found on the second extracellular loop of AT<sub>1</sub>Rs.<sup>5</sup> The immunological basis for the epitope specificity is not understood. Although previous studies have shown that infusion of inflammatory cytokines into pregnant rats results in the production of AT<sub>1</sub>-AA and many other PE features,<sup>15–17</sup> the mechanisms by which these cytokines cause the production of

From the Departments of Biochemistry and Molecular Biology (C.L., R.L., W.W., N.F.P., T.I., R.E.K., Y.X.) and Obstetrics, Gynecology and Reproductive Sciences (S.C.B., B.M.S.), The University of Texas Health Science Center at Houston, TX; The University of Texas Graduate School of Biomedical Sciences at Houston, TX (R.E.K., Y.X.); Department of Chemical Engineering, University of California, Santa Barbara, CA (S.E.E., P.S.D.); Nephrology Department, Xiangya Hospital, Hunan, China (R.L., W.W.); Department of Obstetrics and Gynecology, University of Tokyo, Japan (T.I.); Department of Nephrology, The First Affiliated Hospital of Dalian Medical University, Dalian, China (R.L.).

\*Dr Liu and Dr Luo contributed equally to this work.

**Correspondence to:** Yang Xia, MD, PhD, or Rodney E. Kellems, PhD, Department of Biochemistry and Molecular Biology, The University of Texas Health Science Center at Houston, TX 77030. E-mail: yang.xia@uth.tmc.edu, rodney.e.kellems@uth.tmc.edu

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AT<sub>1</sub>-AAs that recognize the 7-aa epitope peptide of AT<sub>1</sub>Rs remain a mystery. Molecular mimicry with a homologous sequence on human parvovirus was originally considered as the molecular basis for the immunological origin of AT<sub>1</sub>-AA.<sup>18,19</sup> However, this possibility was not supported by epidemiological data that failed to show a correlation between PE and prior parvovirus infection.<sup>20</sup> A commonly considered mechanism for an autoimmune response is posttranslational modification (PTM), resulting in the creation of a neoantigen that is recognized as nonself by the immune system.<sup>21–26</sup> A large percentage of proteins in the body are targets of PTM, and it is now clear that some of these modifications create new antigens that stimulate an autoimmune response.<sup>21–26</sup> One of the best-studied autoimmune diseases associated with PTM is celiac disease, a condition affecting ≈1% of the population of developed countries.<sup>27</sup> Celiac disease is a chronic small bowel disorder caused by an abnormal immune response to a tissue transglutaminase (TG2)-modified dietary protein, termed gliadin (a component of gluteins that are prominent in wheat).<sup>28–30</sup> The TG2 modification of glutamine-rich gliadin creates a modified neoantigen peptide that stimulates an autoimmune response.<sup>28–30</sup> TG2 is the most ubiquitous member of the transglutaminase (TGase) family of enzymes that catalyze the PTM of glutamine residues on target proteins.<sup>31</sup> TGase-catalyzed posttranslational modifications include glutamine deamination, and isopeptide bond formation between a peptidyl glutamine residue and a peptidyl lysine residue or a primary amine.<sup>31</sup> PE is characterized by elevated pro-inflammatory cytokines<sup>32–38</sup> that promote TG2 gene expression.<sup>39</sup> We have recently reported that activated TG2 post-translationally modifies AT<sub>1</sub>Rs with isopeptide bond formation in placentas of women with PE and that circulating TGase activity is significantly elevated.<sup>40</sup> Moreover, in preeclampsia the elevated inflammatory condition favoring the activation and expression of TG2<sup>39</sup> has been demonstrated to give rise to the production of AT<sub>1</sub>-AA.<sup>15–17</sup> Thus, here we hypothesize that TGase activation is required for the production of AT<sub>1</sub>-AA induced by the pro-inflammatory environment of PE.

## Materials and Methods

### Human Subjects

Patients admitted to the Memorial Hermann Hospital were identified by the obstetrics faculty of the University of Texas Medical School at Houston. Preeclamptic patients were diagnosed with severe disease on the basis of the definition set forth by the National High Blood Pressure Education Working Group Report. Inclusion criteria, including no previous history of hypertension, have been reported previously and include blood pressure ≥160/110 mm Hg and urinary protein ≥300 mg in a 24-hour period or a dipstick value of ≥1. (Twenty-four-hour

urinary protein values were available for some of the PE patients. The rest of them had dipstick values.) Other criteria include the presence of persistent headache, visual disturbances, epigastric pain, or hemolysis, elevated liver enzymes, and low platelets syndrome in women with blood pressure ≥140/90 mm Hg. Control pregnant women were selected on the basis of having an uncomplicated, normotensive pregnancy with a normal term delivery. The research protocol was approved by the Institutional Committee for the Protection of Human Subjects. All participants gave informed consent. Clinical features of human subjects are presented in Table.

### Animal Protocols

Pregnant C57Bl/6J mice (Harlan) were anesthetized with isoflurane, and 0.15 mL sterile PBS dissolved with or without recombinant mouse LIGHT (2 ng, R&D Systems) was introduced by retro-orbital sinus injection on gestation day (GD) 13.5 and 14.5. Some mice were also injected with 2.25 mg cystamine dihydrochloride (Sigma-Aldrich, MO) dissolved in the 0.15 mL sterile PBS to inhibit TGase activity. Since the retro-orbital injections of cystamine dihydrochloride were only performed on GD 13.5 and 14.5 together with LIGHT molecules, to ensure a constant and effective cystamine treatment afterwards, these pregnant mice were also administered drinking water containing 0.9 g/L cystamine dihydrochloride throughout the rest of their pregnancy. Both treatment routes were reported and validated in earlier publications.<sup>40–45</sup> All protocols involving animal research were reviewed and

**Table.** Clinical Features of Human Subjects

	NT	PE
N	24	55
Age, y	26.35±5.95	27.04±8.46
Race, %		
African American	50	51
White	35	33
Hispanic	15	16
Gravity	2.25±1.74	2.25±0
BMI	31.60±5.23	35.96±15.40*
Weeks gestational age	38.70±0.94	34.28±5.59*
Systolic BP, mm Hg	119.45±13.32	153.95±12.00*
Diastolic BP, mm Hg	70.15±9.97	90.81±0.68*
Proteinuria, mg/24 h	N/A	846.62±101.02

This table demonstrates that the blood pressure and proteinuria are significantly elevated in PE women vs NT pregnant women at term. BMI values are prepregnancy estimates. The value in each category is indicated as mean±SD. Clinical characteristic features of human subjects used in bacterial peptide display assay (Figure 2D) were reported before.<sup>48</sup> BMI indicates body mass index; BP, blood pressure; N/A, nonapplicable; NT, normotensive; PE, preeclampsia.

\**P*<0.001 vs normotensive pregnant women.

approved by the Institutional Animal Welfare committee of the University of Texas Health Science Center at Houston. All animal procedures were in accordance with institutional guidelines.

### In Vitro TG2 Function Assay, Western Blot, and Immunoprecipitation Followed by Mass Spectral Analysis

Approximately 100 µg TG2 purified from guinea pig liver (Sigma) was incubated with 0.1 mmol/L AT<sub>1</sub>-AA epitope peptide AFHYESQ (7-aa) for 30 minutes at 37°C in 100 µL reaction buffer containing 50 mmol/L Tris-HCl, pH 7.4 and 5 mmol/L CaCl<sub>2</sub>. Under the same conditions, a negative control assay was set up without the addition of the 7-aa epitope peptide. Following an incubation, a portion of the reaction mixture was examined by Western blot analysis using antibody raised against the 7-aa epitope peptide. A similar assay was reported before<sup>46</sup> to test whether TG2 reacts with the synthesized gliadin peptide in celiac disease. After the reaction, the assay mix was directly examined by Western blot using antibody directed against the 7-aa epitope peptide to assess the cross-linking efficiency.

To purify the cross-linking product for mass spectrometry analysis, the assay mix was also subject to immunoprecipitation using the 7-aa epitope antibody-bound to Protein G Sepharose High Performance beads (GE Healthcare Life Sciences) following manufacturer's instructions. The immunoprecipitated product was resolved by SDS-PAGE and subsequently stained with SimplyBlue™ SafeStain (Life Sciences). A band appearing near 80 kDa was excised and examined by mass spectrometry at the proteomics core facility of the MD Anderson Cancer Center.

### Blood Pressure Measurement and Quantification of Proteinuria

The systolic blood pressure was noninvasively measured by a volume pressure recording sensor and an occlusion tail-cuff (CODA System, Kent Scientific). This method shows good agreement with radiotelemetry measurements of blood pressure.<sup>47</sup> Blood pressure was measured at the same time daily (±1 hour) while the mice were kept warm using a warming pad. Twenty-four-hour urine was collected for analysis using metabolic cages (Nalgene). Mice were trained in metabolic cages for 2 days prior to urine collection. On GD18.5, the mean arterial pressure was determined by cannulating the right carotid artery with a mouse jugular catheter connected to a pressure transducer and an amplifier unit, just prior to euthanasia. The amplifier was connected to a data acquisition module and mean arterial pressure was recorded on a personal computer by Chart 5 Software (AD Instruments, Inc).

All of the mice were euthanized on GD18.5 before delivery when their serum and organs were collected. We quantified urinary albumin by ELISA (Exocell) and measured urinary creatinine by a picric acid colorimetric assay (Exocell). We used the ratio of urinary albumin to urinary creatinine as an index of urinary protein.

### Measurement of AT<sub>1</sub>-AA

#### ELISA

AT<sub>1</sub>-AA was measured using a commercially available sandwich ELISA (CellTrend GmbH, Luckenwalde, Germany, catalog #12000), which is now marketed by One Lambda (catalog number EIA-AT2RX). The human plasma samples were diluted 10-fold rather than 100-fold as suggested by the manufacturer's instructions. The kit uses straightforward ELISA-based procedures on 96-well plates for the quantitative determination of AT<sub>1</sub>-AA in human plasma or serum. The kit was developed using membrane preparations from mammalian cells that overproduce human AT<sub>1</sub>R as the bait. The kit has intra-assay and interassay coefficients of variation of ≈7% and 12% according to the manufacturer. For measurement of murine AT<sub>1</sub>-AA, the HRP-linked anti-mouse IgG secondary antibody (Jackson ImmunoResearch Labs, Catalog#715-035-150) was employed. Other steps in this measurement were carried out according to vendor instructions.

#### Peptide display

Plasma samples from humans and mice were assayed for binding to the 7-aa epitope peptide using an assay in which the epitope sequence AFHYESQ is displayed in a high-avidity format on the bacterial cell surface as reported before.<sup>48</sup> In this assay, AT<sub>1</sub>-AA detection can be performed with the 7-mer and does not require the entire second extracellular loop. This peptide display assay is important for showing the epitope specificity of the AT<sub>1</sub>-AA produced in the LIGHT-injected mice.

### Plasma TG Activity

TGase activity in human and mouse samples were determined using in vitro TGase assay kits (Sigma-Aldrich, MO) following the manufacturer's instructions.

### Purification of Total Mouse IgG and Adoptive Transfer Method

#### Affinity purification of IgG from pregnant mice injected with saline or LIGHT

Total IgG was purified by GammaBind G Sepharose chromatography according to the manufacturer's instructions from sera of mice injected either with saline or LIGHT (Saline-IgG or

LIGHT-IgG). Briefly, 400  $\mu$ L serum from mice with either saline or LIGHT injection were loaded on the columns and incubated for 3 hours. The flow-through was collected by centrifugation at 1000g for 1 minute. The eluted bound IgG was further collected by centrifugation at 1000g for 1 minute after incubation with 100  $\mu$ L 100 mmol/L glycine (pH 2.7) for 10 minutes. The low pH of the eluted fraction was neutralized by adding 5  $\mu$ L of 1 mol/L Tris, pH 9.0. We performed all steps at 4°C. The levels of IgG in the elution fraction (affinity-purified total IgG) were quantified by ELISA (Jackson ImmunoResearch Labs, Catalog#715-035-150).

### Introduction of antibody into mice

Nonpregnant C57BL/6J mice were used (18–22 g; Harlan) in our study. Mice were anesthetized with isoflurane. Purified total IgG ( $\approx$ 800  $\mu$ g) from 200  $\mu$ L sera were concentrated by lyophilization and resuspended in  $\approx$ 200  $\mu$ L saline and then introduced into nonpregnant mice by osmotic minipump. Some mice also received a 7-aa peptide corresponding to an epitope on the second extracellular loop of the AT<sub>1</sub> receptor (1.5 mg/mice). This peptide effectively neutralizes AT<sub>1</sub>-AA. Mouse systolic blood pressure, mean arterial pressure, and proteinuria were measured as described above. Plasma AT<sub>1</sub>-AA concentrations on day 18.5 were quantified by ELISA (CellTrend GmbH, catalog #12000).

### Statistical Analysis

Data were expressed as the mean $\pm$ SEM. GraphPad Prism software was employed to run the statistical programs. Measurements of blood pressure, urinary protein concentration, and plasma concentrations of autoantibodies and enzymes typically approximated a normal distribution. Student *t* tests (paired or unpaired as appropriate) were applied in 2-group analysis. The means of multiple groups were compared by the 1-way ANOVA, followed by a Tukey multiple comparisons test. Data presented in Figure 2A were analyzed using a repeated-measures ANOVA followed by a Tukey multiple comparisons test. For the bacterial epitope peptide display assay, statistical significance was determined by 1- or 2-tail Mann–Whitney *U* test. A value of *P*<0.05 was chosen as the threshold of statistical significance. Pearson product-moment correlation coefficient *r* was employed to determine the correlation between the 2 variables in Figure 4.

## Results

### The 7-aa Epitope Peptide of the AT<sub>1</sub>R is Modified by TG2 In Vitro

TG2 catalyzes the PTM of glutamine residues on target proteins by deamination or by isopeptide bond formation with

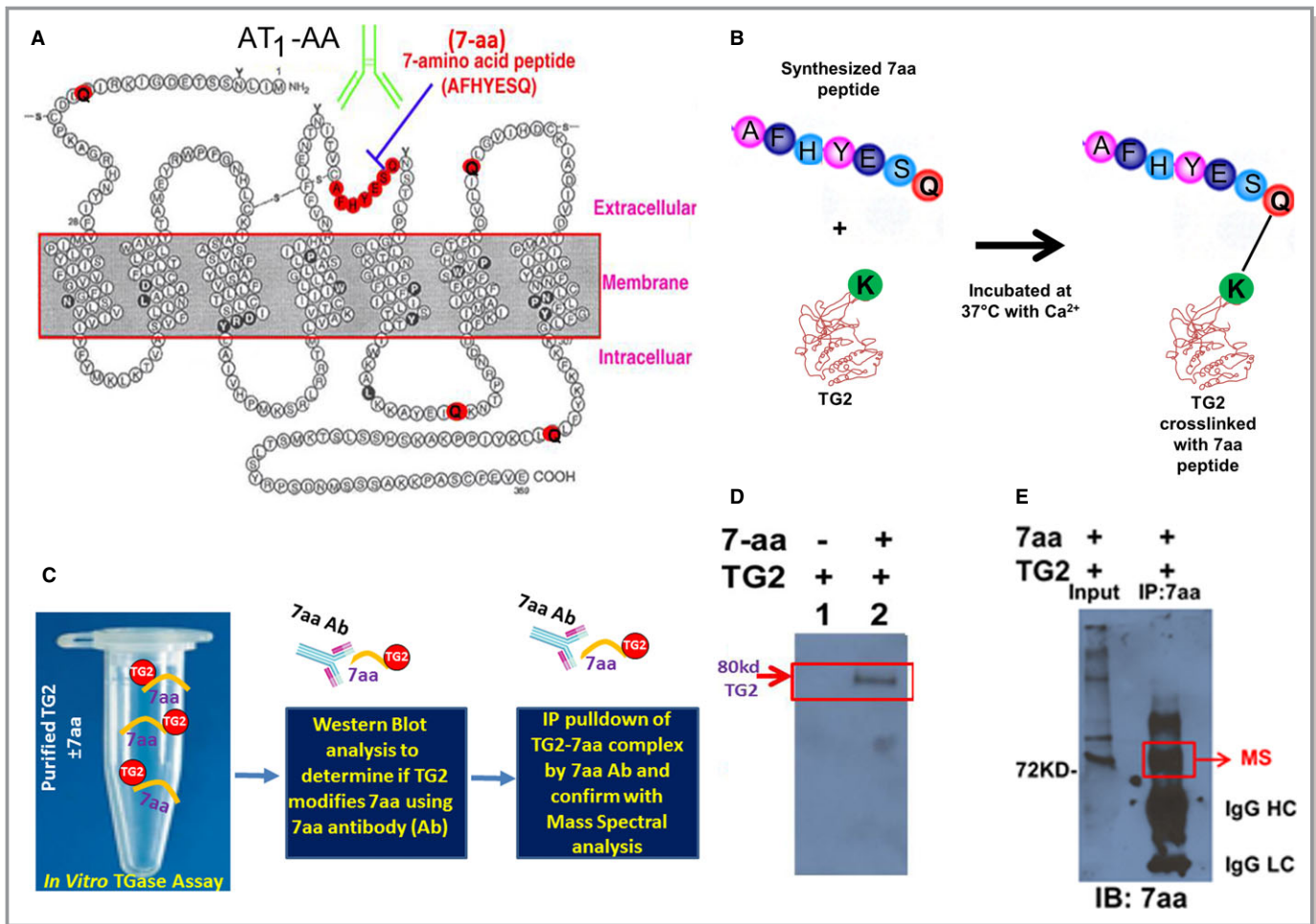
a lysine residue or a primary amine. Of the 359 amino acids that constitute the AT<sub>1</sub>R, only 5 are glutamine residues (marked red in Figure 1A), and represent potential sites of TGase modification. One glutamine (Q187) resides at the end of the 7-aa epitope sequence (AFHYESQ) in the second extracellular loop that is uniformly recognized by AT<sub>1</sub>-AA in women with PE.<sup>5</sup> Thus, TG2-mediated PTM of Q187 may generate a neoantigen that is recognized by AT<sub>1</sub>-AA. Previously, we demonstrated that AT<sub>1</sub>Rs are modified by TGase in PE placental trophoblasts, but due to the low cellular abundance of AT<sub>1</sub>R as a membrane bound GPCR, we were unable to purify enough receptor from PE placentas or trophoblasts to thoroughly determine the endogenous isopeptide modification sites and the chemical nature of their amine donor. Thus, to overcome this obstacle, we conducted in vitro TGase assays to determine whether TG2 can directly modify the 7-aa epitope peptide of AT<sub>1</sub>R (illustrated in Figure 1B). This was done by incubating purified TG2 with or without the specific 7-aa epitope peptide. If the epitope peptide is recognized as a substrate by TG2, then it should become covalently attached to the enzyme, either at the active site as a covalent intermediate, or crosslinked to available lysine residues on other TG2 enzymes in the reaction mix.<sup>31,49</sup> To determine whether the 7-aa epitope peptide was covalently attached to TG2 enzymes, we examined the reaction products<sup>39</sup> by denaturing gel electrophoresis and Western blot analysis using an antibody generated specifically to the 7-aa peptide of the AT<sub>1</sub>R.<sup>32</sup> We also used this antibody in pull-down experiments followed by mass spectral analysis, to determine whether the epitope peptide became covalently associated with TG2 during the in vitro reaction (illustrated in Figure 1C).

Western blot analysis showed that antibody to the 7-aa epitope peptide recognized an 80-kDa protein the size expected for TG2 (Figure 1D). No cross-reactivity was observed in control reactions that lacked the 7-aa epitope peptide (Figure 1D). Mass spectral analysis confirmed that the 80-kDa protein immunoprecipitated by the 7-aa epitope peptide antibody was TG2 (Figure 1E). Thus, Western blot analysis following denaturing gel electrophoresis and mass spectral analysis of co-immunoprecipitated protein indicated that the 7-aa epitope peptide was modified and covalently cross-linked to TG2.

### Plasma Transglutaminase Activity and AT<sub>1</sub>-AA Specifically Recognizing the 7-aa Epitope Peptide Are Induced in an Experimental Model of PE Based on LIGHT Injection

Although we demonstrated the direct modification of the 7-aa epitope peptide by TG2 in vitro, the importance of TG activation in AT<sub>1</sub>-AA production in vivo is undetermined. To

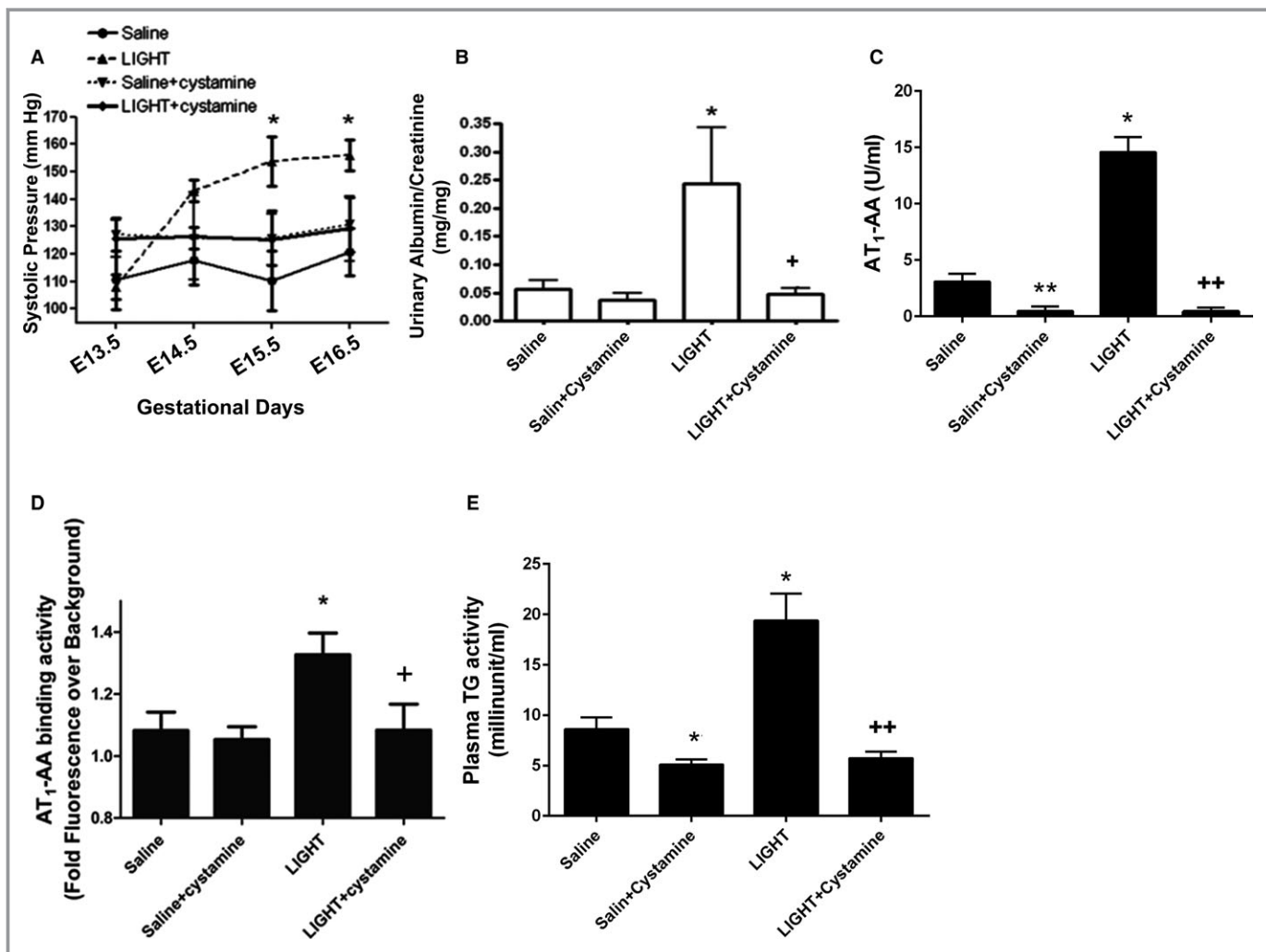




**Figure 1.** AT<sub>1</sub> receptor seven amino acid (7-aa) epitope peptide is modified by TG2 in vitro. A, Two dimensional illustration of AT<sub>1</sub> receptor showing location of the 7-aa epitope peptide and 5 glutamines (Q). B and C, Illustration of experimental strategy to determine whether TG2 modifies the 7-aa epitope peptide in vitro. D, 7-aa epitope peptide was covalently cross-linked to TG2 in the in vitro assay. TG2 was incubated in the presence or absence of the 7-aa epitope peptide (AFHYESQ). The reaction mix was subsequently fractionated by denaturing gel electrophoresis and transferred to a blot. Antibody specific for the 7-aa epitope peptide recognized an 80-kDa protein, the size expected for TG2, indicating that the 7-aa peptide was covalently cross-linked to TG2. E, Mass spectral characterization of TG2 cross-linked with the 7-aa epitope peptide. After the in vitro reaction, proteins covalently linked with 7-aa epitope peptide in the reaction mix were immunoprecipitated with the antibody directed against this peptide and resolved in the denaturing gel electrophoresis. The 80-kDa protein in the immunoprecipitation products was characterized as TG2 in mass spectral analysis. The presence of 7-aa epitope in the immunoprecipitation products was also confirmed in Western blot using the antibody. AT receptor indicates angiotensin receptor 1; IgG HC, immunoglobulin G heavy chain; IgG LC, immunoglobulin G light chain; TG2, transglutaminase 2; MS, mass spectral analysis.

address this question, we used a mouse model of PE based on injection of the inflammatory cytokine LIGHT (tumor necrosis factor [TNF] superfamily member 14) into pregnant mice on GD 13.5 and 14.5.<sup>32</sup> We have previously shown that the injection of LIGHT into pregnant mice induced key features of PE including hypertension, proteinuria, reduced fetal and placental weight, elevated sFlt-1, and elevated endothelin-1, thus making this a convenient and relevant experimental model of PE.<sup>44</sup> We confirmed previously published results showing that LIGHT infusion induced hypertension and proteinuria (Figure 2A and 2B). To determine whether LIGHT also induced AT<sub>1</sub>-AA production, we used 2 assays to quantify AT<sub>1</sub>-AA in the circulation of the pregnant mice with or without

LIGHT injection. Using a cell-based ELISA to quantify the presence of AT<sub>1</sub>-AA, we found that autoantibody titers in LIGHT-injected pregnant mice were confidently identified and significantly higher than the saline controls (Figure 2C). The cell-based ELISA measures antibodies that recognize native AT<sub>1</sub>Rs, and is not specific for binding to the 7-aa epitope peptide of the AT<sub>1</sub>R. Thus, to determine whether the AT<sub>1</sub>-AA produced in LIGHT injected pregnant mice recognized the same 7-aa epitope that is recognized by AT<sub>1</sub>-AA produced in women with PE, we used a recently developed flow cytometry-based bacterial peptide display assay<sup>48</sup> to measure binding to the epitope peptide (AFHYESQ). Similar to the cell-based ELISA, we found that LIGHT injection induced specific



**Figure 2.** Elevated TGase is required for LIGHT-induced production of AT<sub>1</sub>-AA and PE features in pregnant mice. Pregnant mice were injected with LIGHT on GD 13.5 and 14.5 in the presence or absence of cystamine, a competitive inhibitor of TGase. Cystamine treated animals continued to receive the TGase inhibitor in the drinking water. Animals were sacrificed on GD 18.5. A and B, Blood pressure and proteinuria were induced in the pregnant mice with LIGHT injection. C, Plasma AT<sub>1</sub>-AA levels—Results shown represent specific binding to AT<sub>1</sub>Rs in a cell-based ELISA assay. D, Plasma AT<sub>1</sub>-AA levels—Results shown represent binding to the 7-aa epitope peptide presented in a bacterial peptide display format. E, Plasma TGase activity. \**P*<0.05, \*\**P*<0.01 vs saline injected mice; +*P*<0.05, ++*P*<0.01 vs LIGHT injected mice. Saline (n=8), Saline + cystamine (n=3), LIGHT (n=13), and LIGHT + cystamine (n=4). AT<sub>1</sub>-AA indicates angiotensin II receptor 1 agonistic autoantibody; GD, gestation days; PE, preeclampsia; TGase, transglutaminase.

autoantibody directed to the 7-aa epitope peptide of AT<sub>1</sub>Rs. Thus, 2 independent assays confirmed that LIGHT infusion into pregnant mice leads to the generation of autoantibody directed against the AT<sub>1</sub>R (Figure 2D).

### Essential Role of Elevated TGase Activity in LIGHT-Induced PE Features and AT<sub>1</sub>-AA Production in Pregnant Mice

In addition to inducing the production of AT<sub>1</sub>-AA, we found that LIGHT injection significantly increased plasma TGase activity in pregnant mice (Figure 2E). To determine the functional role of elevated plasma TGase in LIGHT-induced

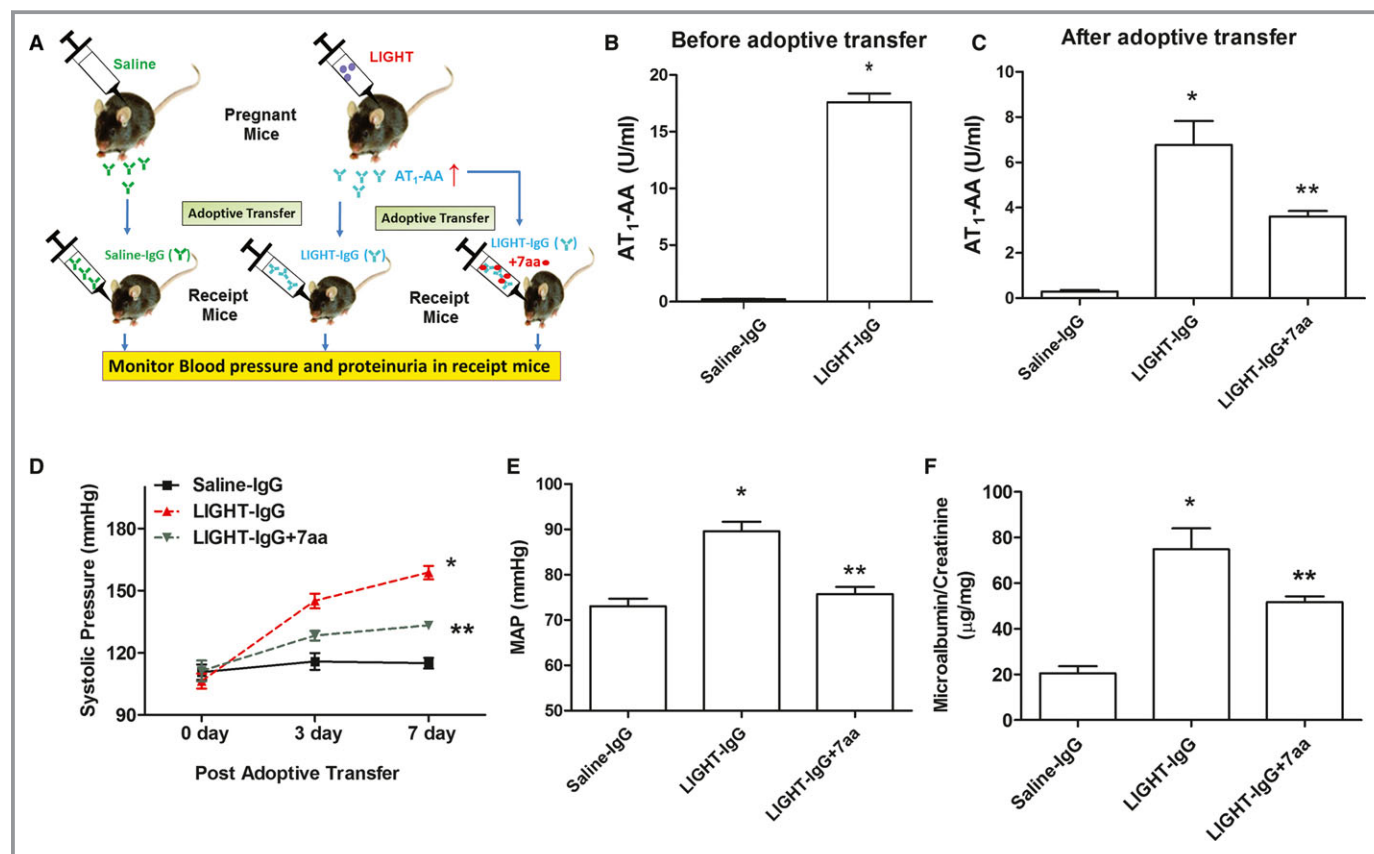
maternal PE features and AT<sub>1</sub>-AA production, we co-injected LIGHT with PBS or cystamine, a competitive inhibitor of TGase,<sup>41,50</sup> into pregnant mice on GD 13.5 and 14.5. To continuously inhibit TGase activity, cystamine-injected mice were also provided drinking water containing 0.9 g/L cystamine. First, we confirmed that cystamine treatment significantly inhibited the increase in plasma TGase activity in LIGHT-injected pregnant mice (Figure 2E). Subsequently, we found that cystamine treatment significantly attenuated LIGHT-induced PE features including hypertension and proteinuria (Figure 2A and 2B). Finally, we used both the cell-based ELISA and the bacterial peptide display assay to quantify the effect of cystamine treatment on LIGHT-induced

AT<sub>1</sub>-AA production in pregnant mice. We found that cystamine treatment significantly reduced AT<sub>1</sub>-AA production in the LIGHT-injected pregnant mice based on the cell-based ELISA (Figure 2C) and the bacterial peptide display assay (Figure 2D). Thus, 2 independent assays provide *in vivo* evidence that elevated TGase activity is required for LIGHT-induced AT<sub>1</sub>-AA production in pregnant mice and that the LIGHT-induced autoantibodies recognized the same 7-aa epitope as AT<sub>1</sub>-AA present in women with PE. Altogether, we established the importance of elevated TGase activity in the pathophysiology of PE and AT<sub>1</sub>-AA production *in vivo*.

### IgG from LIGHT-Injected Pregnant Mice Causes Hypertension and Proteinuria Following Transfer into Recipient Mice

To determine whether the AT<sub>1</sub>-AAs produced in LIGHT-injected pregnant mice have the ability to contribute to the

pathophysiology of PE, we conducted adoptive transfer experiments as illustrated in Figure 3A. For this purpose we isolated total IgG from the saline or LIGHT-injected pregnant mice described above and tested the isolated IgG for the presence of AT<sub>1</sub>-AA before and after introduction into recipient mice. We found that IgG from LIGHT-injected pregnant mice (ie, LIGHT-IgG), in contrast to IgG from saline-injected mice (saline-IgG), displayed significant titers of AT<sub>1</sub>-AA (Figure 3B). Next, LIGHT-IgG or saline-IgG was infused into recipient mice for 7 days by osmotic minipump, systolic blood pressure was monitored by tail cuff plethysmography, and urine was collected in metabolic cages on the seventh day for determination of urinary protein. To validate successful antibody transfer, we analyzed AT<sub>1</sub>-AA titers in the sera from infused mice at the end of experiments. We found that AT<sub>1</sub>-AA was readily detected in the animals infused with LIGHT-IgG compared to mice infused with saline-IgG (Figure 3C). A pronounced increase in systolic blood pressure



**Figure 3.** Adoptive transfer experiments show that IgG from LIGHT-injected mice causes hypertension and proteinuria following introduction into recipient mice. A, Illustration of adoptive transfer experiments. B, Prior to infusion AT<sub>1</sub>-AA levels in IgG from saline (n=6) or LIGHT injected mice (n=8) were determined by ELISA. These IgG preparations, termed saline-IgG or LIGHT-IgG ( $\pm$  pre-incubation with 7-aa epitope peptide) were subsequently infused into mice for 7 days. Following IgG infusion the recipient mice were analyzed for (C) plasma AT<sub>1</sub>-AA levels on day 7, (D) systolic blood pressure measured by tail cuff plethysmography on different days, (E) intracarotid mean arterial blood pressure was measured on day 7, and (F) urinary protein on day 7. \* $P$ <0.01 vs saline-IgG injected mice, \*\* $P$ <0.05 vs LIGHT-IgG injected mice. Saline-IgG (n=5), LIGHT-IgG (n=6), LIGHT-IgG+7-aa (n=4); AT<sub>1</sub>-AA indicates angiotensin II receptor 1 agonistic autoantibody; MAP, mean arterial pressure.

was observed following infusion of LIGHT-IgG and no significant change in blood pressure was observed in the animals infused with saline-IgG (Figure 3D). To validate the tail cuff measurements of blood pressure, the intracarotid mean arterial blood pressure was measured in the mice under anesthesia on the final day of saline- or LIGHT-IgG infusion. As shown in Figure 3E, the mean arterial pressure was significantly elevated in the LIGHT-IgG-injected mice in contrast to that of the saline-IgG-injected mice. A significant increase in urinary protein was also observed in mice infused with LIGHT-IgG in contrast to mice infused with saline-IgG (Figure 3F). Thus, the pathologic properties of LIGHT-induced AT<sub>1</sub>-AA were clearly demonstrated by introduction into recipient mice followed by the appearance of hypertension and proteinuria.

Finally, to determine whether LIGHT-IgG induced hypertension and proteinuria via interaction with the 7-aa epitope peptide of AT<sub>1</sub> receptor, the LIGHT-IgG was pre-incubated with the 7-aa epitope peptide prior to introduction into recipient mice. ELISA analysis of IgG from injected mice showed that the titer of LIGHT-IgG pre-incubated with the 7-aa epitope peptide was significantly lower than that of LIGHT-IgG that was not pre-incubated with the 7-aa epitope peptide (Figure 3C). Moreover, the stimulatory effects of LIGHT-IgG on blood pressure and urinary protein were significantly reduced when the LIGHT-IgG was incubated with the 7-aa epitope peptide prior to injection (Figure 3D through 3F). Overall, these data provide functional evidence that AT<sub>1</sub>-AAs produced in LIGHT-injected pregnant mice specifically recognize the 7-aa epitope peptide of AT<sub>1</sub>Rs and adoptively transfer hypertension and proteinuria to recipient mice.

### Increased Plasma Transglutaminase Activity in PE Patients is Positively Correlated with AT<sub>1</sub>-AA Titer, Systolic Blood Pressure and Urinary Protein

The results presented above indicate that increased plasma TGase is required for AT<sub>1</sub>-AA production in pregnant mice. To translate our mouse findings to humans, we measured TGase activity and AT<sub>1</sub>-AA titer in plasma samples from normotensive and PE patients. Similar to our mouse results, we found that plasma TGase activity was significantly elevated in PE patients compared to normotensive pregnant women (Figure 4A). Additionally, AT<sub>1</sub>-AA levels were independently determined by the cell-based whole receptor ELISA (Figure 4B). Thus, we provide evidence that plasma TGase activity and AT<sub>1</sub>-AA titer were significantly elevated in PE patients compared to normotensive individuals. However, because these values were determined at term, and because women with PE usually deliver early, we cannot rule out the possibility that the observed differences are related to differences in gestational age.

To assess the potential significance of plasma TGase activity in the pathophysiology of PE, we examined the relationship of circulating TGase activity with AT<sub>1</sub>-AA titer, systolic blood pressure, and urinary protein in women with PE. We found that plasma TGase activity was positively and significantly correlated with AT<sub>1</sub>-AA titer blood pressure, and urinary protein (Figure 4C through 4E). Altogether, our translational studies with human samples are consistent with our mouse findings and suggest a pathological role for TGase activity in AT<sub>1</sub>-AA production and subsequent disease development.

### Discussion

The presence of AT<sub>1</sub>-AAs in the circulation of PE patients was first reported in 1999.<sup>5</sup> Since then, a large and growing body of evidence has demonstrated that AT<sub>1</sub>-AAs stimulate the G<sub>q</sub>-coupled AT<sub>1</sub> receptor<sup>5,8,9</sup> and contribute to the pathophysiology of PE.<sup>6,7,10,11</sup> Although the pathogenic role of these autoantibodies has been extensively studied, the molecular basis triggering their generation remains a mystery. Research reported here revealed that TGase is required for the production of AT<sub>1</sub>-AA against 7-aa epitope peptide of AT<sub>1</sub>R in a cytokine-induced model of PE in pregnant mice. Our findings are strongly supported by multiple lines of evidence: (1) In vitro studies demonstrated that TG2 directly modifies the 7-aa epitope peptide of the AT<sub>1</sub>R; (2) In vivo data revealed that elevated TGase activity is essential for the production of AT<sub>1</sub>-AA with specificity against the 7-aa epitope peptide and for disease development in PE; and (3) Human translational studies showed that plasma TGase activity is elevated in PE patients and positively correlated with AT<sub>1</sub>-AA levels, hypertension, and proteinuria. Overall, in vitro direct evidence of TG2-mediated modification of 7-aa epitope peptide of AT<sub>1</sub>R and in vivo mouse findings coupled with human translational studies have added significant new insight to the pathogenesis of PE and open up innovative diagnostic and therapeutic possibilities.

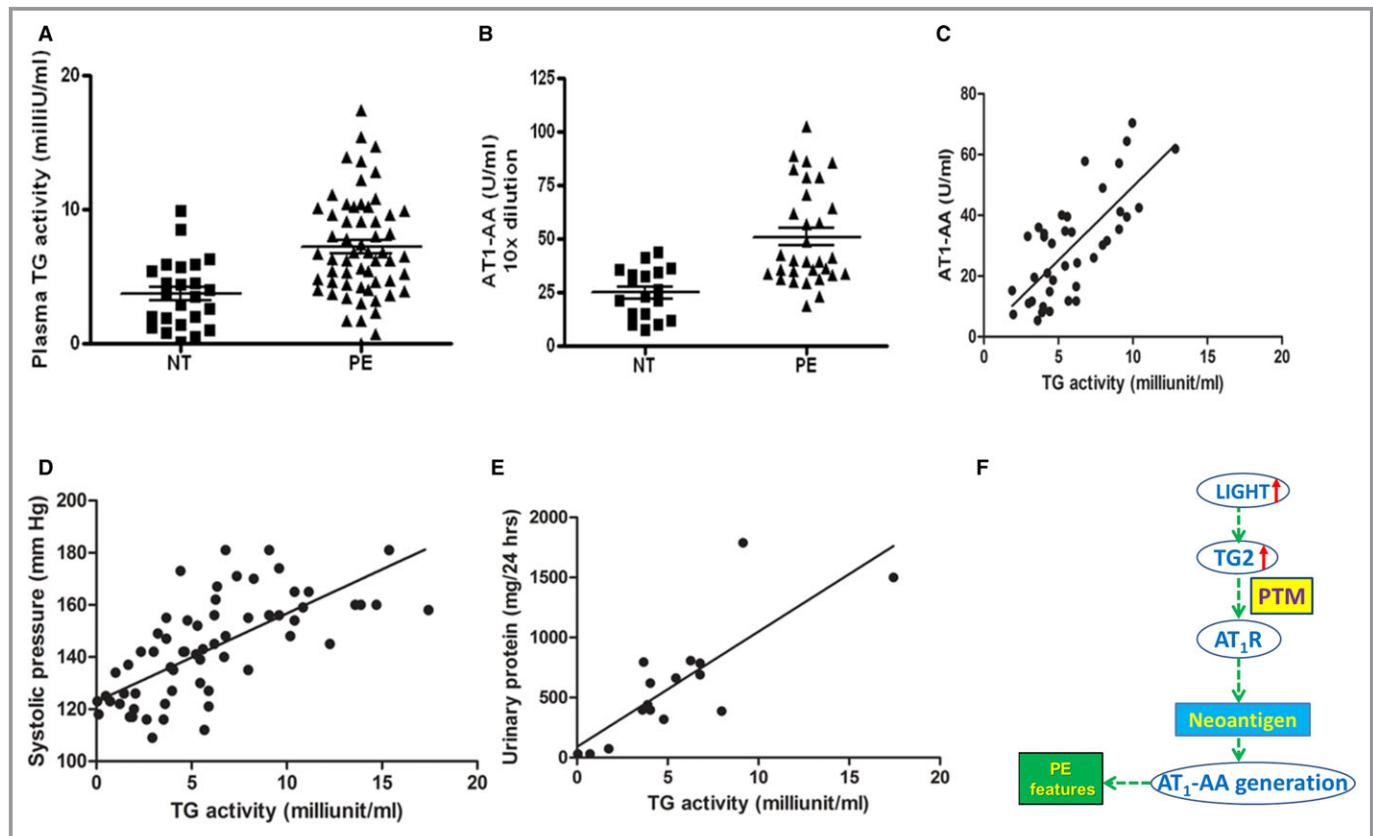
Our results show that most women with PE have elevated AT<sub>1</sub>-AA titers and increased plasma TGase activity compared to those with normotensive pregnancies. The fact that not all women with PE harbor AT<sub>1</sub>-AA or have elevated TGase activity is consistent with the mounting evidence that PE is a heterogeneous disease. It is also noteworthy that a small portion of pregnant women who were not diagnosed with PE had AT<sub>1</sub>-AA titers and plasma TGase activities that overlapped with the lower end of the distribution for women diagnosed with PE. We have previously reported similar observations using a functional bioassay for AT<sub>1</sub>-AA.<sup>11</sup> Because our studies only determined AT<sub>1</sub>-AA titers and plasma TGase activities at term, we do not know how long these values were elevated. Thus, it is



possible that these values were only recently elevated in these women, in comparison to those observed in women with PE, and had not been elevated for a sufficient length time to cause features of PE. This possibility underscores the need for a prospective clinical study measuring these parameters throughout pregnancy to determine when they first appear and what temporal relationship this has with the onset of PE symptoms.

One of the commonly accepted causes for autoantibody production is PTM of proteins to generate neoantigens.<sup>21–26</sup> Celiac disease is the most common autoimmune disease caused by TG2-mediated PTM. Specifically, in celiac disease TG2 mediates PTM of glutamine-rich gluten peptides, resulting in the generation of neoantigens that subsequently trigger autoantibody production.<sup>28–30</sup> We have recently found that TGase activity is increased in the placentas and plasma of PE patients.<sup>40</sup> Moreover, TGase mediates PTM of the AT<sub>1</sub>R in the placentas from PE patients.<sup>40</sup> However, whether TGase-

mediated modification of AT<sub>1</sub>R contributes to the production of autoantibodies in PE has not been determined. Our mutational analysis of exogenous human AT<sub>1</sub>R produced in Chinese Hamster Ovary (CHO) cells determined that intracellular TG2 modifies AT<sub>1</sub>R on the cytoplasmic tail glutamine residue (Q315) and thereby increases receptor stability by inhibiting ubiquitin-mediated degradation.<sup>40</sup> We found that 1 of the 5 glutamine residues (Q187) present in AT<sub>1</sub>R is present at the end of the 7-aa epitope peptide (AFHYESQ) in the second extracellular loop of the receptor. This finding raises a possibility that TG2 functions extracellularly to mediate PTM of glutamine residue Q187 of the 7-aa epitope peptide on the second extracellular loop of AT<sub>1</sub>R. Indeed, TG2 is also known to function extracellularly in the microparticles shed from the cell membrane.<sup>51,52</sup> In contrast to celiac disease where TG2-mediated PTM of a short dietary peptide (gliadin) is easily detected,<sup>28</sup> it has been challenging to purify sufficient quantities of AT<sub>1</sub>R from placentas of women with PE



**Figure 4.** Increased plasma TGase activity in PE patients is positively correlated with AT<sub>1</sub>-AA titer, systolic blood pressure, and urinary protein. A, Plasma TGase activity in normotensive pregnant women (NT, n=24) and PE patients (PE, n=55) ( $P<0.05$ ). B, Cell-based whole receptor ELISA assay of circulating AT<sub>1</sub>-AA (NT, n=18; PE, n=31,  $P<0.05$ ). C through E, The correlation of plasma TGase activity to AT<sub>1</sub>-AA (ELISA) (correlation coefficient  $r>0.5$ ,  $P<0.01$ , n=39), systolic blood pressure (correlation coefficient  $r>0.5$ ,  $P<0.01$ , n=61), and urinary protein (correlation coefficient  $r>0.5$ ,  $P<0.01$ , n=16). F, Working model-essential role of TGase in the production of AT<sub>1</sub>-AA in PE. Our data support a model in which inflammatory cytokines such as LIGHT induce TGase, which modifies AT<sub>1</sub> receptors. TGase modified AT<sub>1</sub> receptors serve as neoantigens that stimulate the autoimmune production of AT<sub>1</sub>-AA. Thus, our findings identify novel mechanisms for autoantibody production and reveal innovative therapeutic possibilities for PE. AT<sub>1</sub>-AA indicates angiotensin II receptor 1 agonistic autoantibody; AT<sub>1</sub>R, AT<sub>1</sub> angiotensin receptor; NT, normotensive; PE, preeclampsia; PTM, posttranslational modification; TGase, transglutaminase.

to determine whether glutamine Q187 is modified endogenously. However, here we overcame this hurdle by conducting in vitro functional assays to determine whether TG2 can specifically modify the 7-aa epitope peptide of the AT<sub>1</sub>R. With the antibody specifically developed against the 7-aa epitope peptide,<sup>32</sup> Western blot analysis following denaturing gel electrophoresis and mass spectral analysis of the protein immunoprecipitated with the antibody show that the 7-aa epitope peptide was modified and covalently cross-linked to TG2 (Figure 1). These in vitro findings support the possibility that TG2-mediated PTM of AT<sub>1</sub>R may contribute to autoantigen origination of AT<sub>1</sub>-AA and account for its epitope specificity.

In vitro experiments are unable to determine whether TGase modification of AT<sub>1</sub>R promotes AT<sub>1</sub>-AA production. Thus, we extended our in vitro studies to in vivo to address the importance of TGase in AT<sub>1</sub>-AA production in PE. PE is associated with an increased maternal inflammatory response.<sup>33</sup> A pathogenic role for the inflammatory response is provided by data showing that infusion of inflammatory cytokines (TNF- $\alpha$ , IL-6, and IL-17)<sup>15–17</sup> results in key features of PE in pregnant rats. Supporting this finding, our studies presented here demonstrated that infusion of LIGHT,<sup>32</sup> a TNF superfamily member,<sup>53,54</sup> into pregnant mice results in PE features and AT<sub>1</sub>-AA production. Using an adoptive transfer experimental strategy, we demonstrated for the first time that AT<sub>1</sub>-AAs produced in LIGHT-infused pregnant mice are sufficient to regenerate PE features including hypertension and proteinuria in recipient mice, indicating the pathogenic role of LIGHT-induced autoantibody. More importantly, we further demonstrated that pre-incubation of autoantibody purified from LIGHT-infused pregnant mice with the 7-aa epitope peptide significantly attenuated PE features in recipient mice. Thus, our adoptive transfer experiments provide compelling evidence that LIGHT-induced AT<sub>1</sub>-AA has a pathogenic role and is directed to the same 7-aa epitope as the AT<sub>1</sub>-AAs present in women with PE.

Successful establishment of an animal model with production of AT<sub>1</sub>-AA specific for the 7-aa epitope peptide of AT<sub>1</sub>R provides us with an important investigative tool to test the importance of TGase in AT<sub>1</sub>-AA production in vivo. Consistent with AT<sub>1</sub>-AA production triggered in LIGHT-infused pregnant mice, we discovered that plasma TGase activity was also induced by LIGHT. To test the importance of elevated TGase in AT<sub>1</sub>-AA production and disease development, we treated LIGHT-infused pregnant mice with cystamine, a well-accepted and commonly used TGase inhibitor.<sup>32–38,45</sup> We found that inhibition of TGase activity significantly reduced LIGHT-induced AT<sub>1</sub>-AA production and maternal disease development in pregnant mice. While it is clear that cystamine inhibits TGase activity, the possibility of off-target effects cannot be ruled out at this time. Nevertheless, the fact that cystamine

blocks LIGHT-induced features of PE, including AT<sub>1</sub>-AA production, is a very important finding with significant mechanistic and therapeutic implications. Taken together, our findings suggest that elevated TGase activity is essential for LIGHT-induced pathophysiology of PE including hypertension, proteinuria, and AT<sub>1</sub>-AA production.

Our preclinical studies with mouse models suggest that elevated TGase may play an important role in the pathophysiology of PE and the production of AT<sub>1</sub>-AA. To extend these findings to humans, we conducted translational studies and found that plasma TGase activity was significantly elevated in patients with PE. A cell-based ELISA was used to determine the presence of antibody that binds to intact AT<sub>1</sub> receptors<sup>55</sup>. We observed a strong positive correlation between plasma TGase activity and circulating AT<sub>1</sub>-AA titers, blood pressure, and urinary protein in women with PE. This strong correlation is consistent with a crucial role for TGase in autoantibody production, a possibility supported by the ability of cystamine to block LIGHT-induced AT<sub>1</sub>-AA production in pregnant mice.

PE is a complicated pregnancy disorder featuring an increase in maternal inflammatory cytokines.<sup>3,32–38</sup> Rodent models of PE based on infusion of inflammatory cytokines (TNF- $\alpha$ , IL-6, IL-17<sup>15–17</sup>) are characterized by the presence of AT<sub>1</sub>-AA. It is of special note that the promoter region of the TG2 gene contains gene regulatory elements for the binding of transcription factor NF- $\kappa$ B,<sup>39</sup> which functions in response to inflammatory cytokines such as TNF, or members of the TNF superfamily (e.g., LIGHT), as well as regulatory elements that respond to IL-6.<sup>39</sup> These cytokines (TNF, LIGHT, and IL-6) are increased in most, but not all, women with PE and are likely to contribute to the increased TG2 production<sup>39,56,57</sup> observed in most PE pregnancies.<sup>40</sup> Consistent with this expectation, we have shown here that circulating levels of TGase activity are significantly increased in women with PE and in LIGHT-injected pregnant mice. We have previously shown that increased TG2 modifies AT<sub>1</sub>R by ubiquitination-preventing isopeptide modification in PE placental syncytiotrophoblasts,<sup>40</sup> where the expression of TG2 and AT<sub>1</sub>R is the highest in the placenta. We believe that the TG2-mediated AT<sub>1</sub> receptor modification and stabilization antagonizes AT<sub>1</sub>R downregulation in PE and may thereby contribute to disease development by providing increased abundance of AT<sub>1</sub>R for activation and downstream signaling and for presentation to the immune system, resulting in autoimmunity and autoantibody production.

TG2 is also known to be present in microparticles shed from cell membranes.<sup>51,52</sup> We show here that TG2 is able to modify and cross-link the 7-aa epitope peptide in vitro. Thus, similar to celiac disease,<sup>28–30</sup> where TG2 modifies a dietary peptide gliadin, it is possible that in PE the circulating autoantigen of AT<sub>1</sub>-AA is created by elevated TG2 modification of the AT<sub>1</sub>-AA epitope from the second extracellular loop of

the receptor in the microparticles shed from placental syncytiotrophoblasts.<sup>58</sup> In this way, the TG2-generated autoantigen may be presented to the immune system in the circulation where we identified an increased level of TGase activity in PE. From this perspective, our results are in good agreement with earlier results<sup>15–17</sup> showing that TNF, IL-6, and IL-17 can stimulate AT<sub>1</sub>-AA production in pregnant rats. An especially important finding reported in our study is that cytokine-mediated induction of AT<sub>1</sub>-AA production is inhibited by cystamine, an inhibitor of TGase. Our findings implicate TGase-mediated modification of AT<sub>1</sub> receptors as a key factor contributing to cytokine-induced autoantibody production. The increased AT<sub>1</sub>-AA will in turn activate AT<sub>1</sub> receptors, leading to increased TG2 production and AT<sub>1</sub> receptor modification.<sup>32</sup> Taken together, we believe that AT<sub>1</sub>-AA, AT<sub>1</sub> receptor activation, and TG2-mediated receptor modification form a vicious cycle driving disease development.

In conclusion, nothing was known about the role of TGase in AT<sub>1</sub>-AA production and disease development prior to our studies. Therefore, our findings about elevated TGase activity in AT<sub>1</sub>-AA production are novel. First, a strong correlation of plasma TGase activity with AT<sub>1</sub>-AA titers and disease features in humans is highly significant for diagnostic purpose. Our data also support a new working model (Figure 4F) in which a heightened inflammatory response characteristic of PE contributes to the induction of TGase, which modifies AT<sub>1</sub>Rs. The TGase-modified AT<sub>1</sub>Rs serve as neoantigens that stimulate the autoimmune production of AT<sub>1</sub>-AA. These studies suggest that PE may be an autoimmune condition in which cytokine- and/or placental hypoxia-induced production of TG2 results in increased AT<sub>1</sub> receptor modification leading to AT<sub>1</sub>-AA production and thereby contributing to other disease features. The novel but compelling concept of PE as an autoimmune condition offers new insight regarding the origin and management of PE.<sup>59</sup> Finally, our findings that elevated TGase underlies autoantigen origination is likely not limited to PE and may also apply to the generation of agonistic autoantibodies in other autoimmune-related cardiovascular diseases. Examples include several forms of hypertension associated with autoantibodies that activate  $\alpha$ 1-adrenergic receptors,<sup>60–62</sup> idiopathic dilated cardiomyopathy associated with  $\beta$ 1-adrenergic receptor activating autoantibodies,<sup>63,64</sup> idiopathic dilated cardiomyopathy associated with autoantibodies that activate the muscarinic M2 receptor,<sup>65,66</sup> and scleroderma associated with autoantibodies that activate the endothelin type A receptor.<sup>67</sup> Thus, our discovery of the contribution of TGase in autoantibody production in PE may provide a clue about why self-reacting antibodies are generated in other autoimmune-related cardiac diseases that are also associated with hypoxia and increased inflammatory cytokines. Taken together, our current studies highlight TG2 as an

attractive therapeutic target in PE and potentially other cardiovascular and hypertensive disorders.

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## Disclosures

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