Expression of Human Leukocyte Antigen-G in Placental Tissues from Pregnant Women with Preeclampsia

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Abstract: Objective: To study the gene and protein expressions of human leukocyte antigen-G (HLA-G) in the placental tissues from healthy women in late pregnancy and pregnant women with preeclampsia (PE), and to explore the relationship between HLA-G and PE. Methods: Immunohistochemistry, Western blot and RT-PCR were used to detect the expressions of HLA-G protein and mRNA in the maternal placenta of 60 pregnant women with PE (including 30 cases of mild PE and 30 cases of severe PE) and 30 healthy full-term pregnant women (control group) respectively. Results: HLA-G had its expression mainly on the surface of trophoblast cells, and its response range and reactivity of its immunologic tissue were further reduced and weakened as the patient's condition was aggravated. The protein expression of placental HLA-G in each PE group was significantly lower than control group, with the difference statistically significant (all P<0.05). Compared with mild PE group, the protein expression level of HLA-G was lower in severe PE group (P<0.05). The Spearman's correlation analysis for HLA-G and PE severity showed that the correlation coefficient was -0.892 (rs = -0.892) (P<0.05), indicating that the expression of HLA-G protein was significantly negatively correlated to the severity of PE. The mRNA expression level of placental HLA-G of each PE group was significantly lower than that of control group, and the difference was statistically significant (all P<0.05). But the mRNA expression level of HLA-G was significantly lower in severe PE group than in mild PE group (P<0.05). Conclusion: The decreased expression of HLA-G in placenta may be involved in the pathogenesis of PE, and can also reflect the severity of this disease.

Keywords: HLA-G, Placenta, Preeclampsia, Pregnancy

1. Introduction

Preeclampsia (PE) is a pregnancy-specific multi-system disease that threatens the health of both mothers and babies with unknown causes. It is a relatively severe type in the classification of pregnancy-induced hypertension (PIH), one of the common gestational diseases as well as one of the main causes for maternal mortality. Too shallow invasion of trophoblast cells to endometrium and the remodeling disorders of uterine spiral arteries are its main pathological features [1, 2]. It may be involved in maternal, placental and fetal factors, including abnormal trophoblast invasion, immune dysfunction, endothelial cells injury, genetic and nutritional factors [3, 4]. But there is no one single factor that can explain the pathogenesis of PE’s incidence.

Studies have found that human leukocyte antigen-G (HLA-G) plays an important role in maintaining maternal-fetal immune tolerance and regulating trophoblast infiltration function, and its abnormal expression participates in the incidence of hypertensive disorders in pregnancy [5]. The original transcripts of HLA-G can produce a number of isomer mRNAs through alternative splicing to respectively encode different protein subtypes, of which HLA-G1, G2, G3 and G4 are membrane-bound proteins, while HLA-G5, G6 and G7 are soluble proteins [6].

This study aims to detect the expression, distribution and differences of HLA-G gene and protein between the placenta tissues from pregnant women with PE and healthy women in late pregnancy so as to explore the relationship between HLA-G and the pathogenesis of PE.
2. Methods

2.1. Clinical Data

Thirty patients with mild PE, 30 patients with severe PE and 30 healthy pregnant women who were admitted and delivered in the Obstetrical Department of our hospital from Jan. 2012 to Mar. 2013 were selected, all of which were single pregnancies. They were divided into mild PE group, severe PE group and control group respectively. The ages of mild and severe PE groups and control group were 20-32 (26.7 ± 5.2) years old, 21-33 (26.9 ± 5.8) years old and 20-33 (27.2 ± 5.6) years old respectively. Gestational weeks were (36.7 ± 2.2), (37.2 ± 1.8) and (37.5 ± 1.6) respectively. There were no statistically significant differences in the age, gestational week and times of gestation and delivery among the three groups by balancing test (all P>0.05). The diagnostic criteria for PE were referred to the relevant literatures, previous hypertension, diabetes, heart disease, immune system disorders, liver and kidney diseases and the history of other chronic diseases excluded [7].

2.2. Sample Treatment

For all cases, 3 to 4 pieces of tissue, about 1 cm³ for each, in the central part of maternal surface of placenta attached to the root of umbilical cord were taken by sterile scissor after the delivery of placenta, and the regions of hemorrhage, necrosis and calcification were avoided. One piece was rinsed with sterile normal saline (containing 0.5% DEPC water) to remove the blood on it, and placed into freezing tube processed and sterilized by DEPC water. The other pieces were rapidly placed into liquid nitrogen after being rinsed with normal saline, and transferred to -70°C refrigerator after overnight for standby use. Another piece of fresh placenta of about 1 cm³ was taken and fixed in 3% paraformaldehyde for immunohistochemical detection.

2.3. Detection of HLA-G Protein Expression in Placental Tissues by Immunohistochemistry Method (SP Method)

The placental tissue pieces fixed in 3% paraformaldehyde mentioned above were conducted conventional dehydration, paraffin-embedding and section at 5 μm. The slides were cleaned and coated with APES to prevent section flaking, 3 slices in each piece for immunohistochemical staining using the SP method. The main steps were as follows: The slices were incubated with mouse anti-HLA-G monoclonal antibody (1: 200, U.S. Sigma) overnight at room temperature and then biotinylated rabbit anti-mouse IgG (1: 100, Amresco) at room temperature for 6 h. DAB/H2O2 was used for color development and hematoxylin for mild counterstaining. The primary antibody was replaced with antibody diluent as the negative control, and the reaction was negative. Five high-power fields (~200) were randomly selected under optical microscope, observed and photographed. The absorbance values of HLA-G immune reaction products on each slice were measured using Image-Pro Plus Version 6.0 color image analysis system and made semi-quantitative statistical analysis.

2.4. Detection of HLA-G Protein Expression in Placental Tissues by Western Blot

Placental tissue (500 mg) was added 5 ml of cell lysis buffer, homogenized, and centrifuged at 3000 r/min and 4°C for 10 min to take the supernatant which was separated by electrophoresis for protein samples. The proteins were transferred to NC membrane by constant current at 275 mA for 1 h for Western blot. The samples were incubated with mouse anti-HLA-G monoclonal antibody (1: 2000, U.S. Sigma) at 4°C overnight, with mouse anti-β-actin monoclonal antibody (1: 1000, USA sigma) as internal reference, and horse radish peroxidase (HRP)-labeled rabbit anti-mouse IgG (1: 1000, Shanghai RICKY Bio-tech Co., Ltd.) at room temperature for 6 h. NC membrane was made color development with DAB/H2O2 and the results were obtained by scanner. The bio-electrophoresis image analysis system (OLYMPUS, Japan) was used to analyze the absorbance values of bands. The target protein level was expressed by the ratio of the average absorbance values of the target band and the internal reference β-actin, and then made analysis.

2.5. Detection of HLA-G mRNA Expression in Placental Tissues by RT-PCR

Placental tissue (500 mg) was extracted total RNA by Trizol method (Trizol kit was purchased from Invitrogen, USA). HLA-G mRNA level was detected by RT-PCR. Primers were designed and synthesized by Shanghai Jianglai Bio-tech Co., Ltd. β-Actin was amplified as an internal reference. The annealing temperatures for the sequences of all target gene primers, base positions, lengths of amplified fragments and PCR amplification were shown in Table 1.

<table>
<thead>
<tr>
<th>Primer sequence</th>
<th>Length</th>
<th>Cycle</th>
</tr>
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<tbody>
<tr>
<td>HLA-G</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Upstream primer 5′-ACAGGCACACAATAACCCCGGC-3′</td>
<td>30</td>
<td></td>
</tr>
<tr>
<td>Downstream primer 5′-CTCCTAGTGGCCTTGATCTTC-3′</td>
<td>497 bp</td>
<td></td>
</tr>
<tr>
<td>β-Actin</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Upstream primer 5′-ATGAGTACATCTGTCACTCGG-3′</td>
<td>30</td>
<td></td>
</tr>
<tr>
<td>Downstream primer 5′-TGAGTACATTGGTACACCCGGGA-3′</td>
<td>341 bp</td>
<td></td>
</tr>
</tbody>
</table>

PCR amplification products were conducted 3.0% agarose gel electrophoresis, and gel imaging system (Labnet) was used for analysis of absorbance values of bands. The β-actin mRNA level was expressed by the ratio of the mean absorbance values of the target gene and β-actin, and then made semi-quantitative analysis.

2.6. Statistical Analysis

The measurement data were expressed as x±s and analyzed by SPSS 15.0. Means of groups were compared by F test. Two groups were compared by LSD method. Correlation was subjected to Spearman's rank correlation analysis. P<0.05 was
considered statistically significant.

3. Results

3.1. Immunoreactive Expressions of HLA-G in Placental Tissues

HLA-G protein had expression in the placental tissues of both control group and PE groups. The positive products of HLA-G immunoreaction were brown granular or punctate, expressed mainly on the surface of trophoblast cells. Its response range and reactivity of its immunologic tissue were further reduced and weakened as the patient's condition was aggravated (Figure 1).

The results of semi-quantitative analysis showed that compared with control group, the positive staining of placental HLA-G immunoreaction in PE groups became shallow, and the average absorbance values were reduced, therefore, the protein expression level of placental HLA-G in each PE group was decreased, with the difference statistically significant (all P<0.05). Compared with mild PE group, the positive staining of placental HLA-G immunoreaction in severe PE group was more shallow, so the protein expression level of placental HLA-G was lower in severe PE group than in mild PE group, with the difference statistically significant (P<0.05) (Table 2).

![Figure 1. Immunoreactive expressions of HLA-G in placental tissues (×400). A: Control group; B: mild PE group; C: severe PE group.](image)

### Table 2. Absorbances of HLA-G immunoreaction products.

<table>
<thead>
<tr>
<th>Group</th>
<th>n</th>
<th>HLA-G expression</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>30</td>
<td>0.85±0.12</td>
</tr>
<tr>
<td>Mild PE</td>
<td>30</td>
<td>0.57±0.09*</td>
</tr>
<tr>
<td>Severe PE</td>
<td>30</td>
<td>0.26±0.08*#</td>
</tr>
</tbody>
</table>

Compared with the control group, *P<0.05; compared with mild PE group, #P<0.05.

3.2. HLA-G protein Expression in Placental Tissues Detected by Western Blot

The molecular weights of HLA-G and β-actin proteins were 97 kD and 56 kD respectively. The absorbance values of bands in each group were corrected with β-actin as internal reference and then made analysis. The results showed that the protein expression level of placental HLA-G in each PE group was significantly lower than control group, with the difference statistically significant (Figure 2), with the difference statistically significant (all P<0.05). Compared with mild PE group, the protein expression level of HLA-G was lower in severe PE group (P<0.05) (Figure 2). The Spearman's correlation analysis for HLA-G and PE severity showed that the correlation coefficient was -0.892 (rs = -0.892) (P<0.05), indicating that the expression of HLA-G protein was significantly negatively correlated to the severity of PE. In other words, HLA-G protein expression level can be further decreased as the patient’s condition was aggravated.

![Figure 2. HLA-G protein expression in placental tissues detected by Western blot. 1: Control group; 2: mild PE group; 3: severe PE group. Compared with the control group, *P<0.05; compared with mild PE group, #P<0.05.](image)

3.3. HLA-G mRNA in Placental Tissues Detected by RT-PCR

![Figure 3. HLA-G mRNA in placental tissues detected by RT-PCR. M: Marker; 1: control group; 2: mild PE group; 3: severe PE group. Compared with the control group, *P<0.05; compared with mild PE group, #P<0.05.](image)
The lengths of amplified fragments of HLA-G and β-actin mRNAs were 497 bp and 341 bp respectively, with clear bands. The absorbance values of bands in each group were corrected with β-actin as internal reference and then made analysis. The results showed that the mRNA expression level of placental HLA-G in each PE group was significantly lower than that of control group, and the difference was statistically significant (all P<0.05). But the mRNA expression level of HLA-G was significantly lower in severe PE group than in mild PE group (P<0.05) (Figure 3).

4. Discussion

HLA-G belongs to a non-classical HLA- I type (HLA- I b) gene family, which is located on human chromosome 6 short arm 6p21.3, with encoding membrane binding heavy chain (HC). The whole HC consists of the extracellular region (containing α1, α2 and α3 domains), transmembrane region and cytoplasmic region. HC is bound with β2 microglobulin into a functional protein by non-covalent bond. HLA-G transcript is composed of eight exons, seven introns and two untranslated regions, of which exon 1 encodes the signal peptide, exons 2-4 extracellular α1-α3 domains and exons 6-8 cytoplasmic region [8]. Its original transcript can generate seven kinds of isomer mRNA through alternative splicing, which can encode seven kinds of protein molecule: the four membrane-bound molecules are HLA-G1 (containing 3 extracellular functional domains), HLA-G2 (lacking α2 domain), HLA-G3 (lacking α2 and α3 domains), HLA-G4 (lacking α3 domains) respectively, and the three soluble molecules are HLA-G5, HLA-G6 and HLA-G7 respectively. Both the intron 4 contained in HLA-G5 and HLA-G6 transcripts and the intron 2 contained in HLA-G7 mature mRNA have terminating codons, resulting in the transmembrane and cytoplasmic regions unable to be translated, so as to turn into a soluble form [9]. HLA-G has the features of low-degree polymorphism and limitation of distribution, mainly expressed in extravillous trophoblast [10]. Studies have found that HLA-G molecule can inhibit the activity of maternal NK cells and protect trophoblast cells from their killing, and it can also induce the generation of Th2-type cytokines and promote Th1/Th2-type equilibrium to migrate towards the Th2-type direction conducive to pregnancy, taking trophic effects on trophoblast cells [11]. In recent years, it has been found that its non-full-length molecules, i.e. HLA-G2, HLA-G3 and HLA-G4 can also inhibit the cytoplastic effects of NK cells and antigen-specific CTL. In individuals with homozygous deletion of HLA-G1, these subtypes play a leading role in inducing maternal-fetal immune tolerance [12]. Therefore, most scholars believe that HLA-G is an immune-tolerance molecular, which can maintain normal pregnancy through the induction of immune tolerance of maternal-fetal interface, while its abnormal expression may lead to pathological pregnancy [13].

Some studies have found that compared with normal placental tissue, the protein expression of placental HLA-G was significantly decreased in patients with severe PE [14]. Thus, it can be speculated that due to the defect of HLA-G protein expression, trophoblast cells have a reduced ability of inhibiting the attack of maternal decidual NK cells and some cytotoxic cytokines, so that extravillous trophoblast cannot be effectively implanted into the maternal spiral arteries, which can result in vascular casting blockage and placental hypoperfusion, causing a series of adverse consequences, and ultimately developing to hypertensive disorders in pregnancy [15].

In recent years, more and more studies have shown that trophoblast infiltration is the key for the incidence and progress of PE. The implantation of fertilized egg of normal pregnancy into uterine cavity requires twice physiological intrusions of trophoblast cells to destroy the muscle elastic fibers, smooth muscle fibers and nerve tissues on vessel wall, so that the spiral artery appears progressive dilation, and loses its elasticity and its reactivity to vasoconstrictor substances, and then it succumbs to physiological expansion. This is spiral arterial recasting, which can be found in the decidual layer and part of the superficial muscle segments [16, 17]. But extravillous trophoblast cells have superficial infiltration in PE, and uterine spiral arteries, muscular spiral arteries in particular, are imperfectly rebuilt in the decidua of uteroplacental bed and superficial muscle segment. The lumen diameters of most spiral arteries are only half of those in healthy pregnant women in the same period, and in addition, the arterial muscular layer is still thick, vascular endothelial cells have not yet been replaced with trophoblast cells, so there are rare endovascular trophoblast cells in the spiral arteries in the muscular layer [18, 19]. Zhao et al. studied the placental bed biopsies from normal pregnant women and those with PE, and found that HLA-G had sustained high expression in placental cells in the former, but on the contrary, HLA-G had low expression in the latter [20]. Thus, it can be inferred that the primary reason for the incidence of PE lies in that placental cells in PE may experience abnormal differentiation at the molecular level, with reduced trophoblast infiltration, resulting in poor spiral arterial recasting and shallow placental implantation.

In this study, immunohistochemistry, Western blot and RT-PCR were used to study the differences in the expression of placental HLA-G in healthy pregnant women and those with PE from transcription to translation in both qualitative and quantitative way. The experimental results show that HLA-G protein had expression in the placental tissues of both control group and PE groups, mainly on the surface of trophoblast cells. However, the expression levels of HLA-G protein in PE placentas were significantly lower than that of normal group, and as the condition of PE was aggravated, their levels were further decreased markedly. The RT-PCR results show that the expression levels of HLA-G mRNA transcription were consistent with those of HLA-G protein in all the three groups. Therefore, it can be inferred that HLA-G has decreased impression in placental tissues from pregnant women with PE. On the one hand, it may reduce the binding of HLA-G with the extracellular matrix ligand fibronectin, affect the normal transition of trophoblast cells from adhesion
phenotype to invasion phenotype, and inhibit the ability of trophoblast invasion, while on the other hand, HLA-G decline may affect the activation of its downstream signaling molecules, so as to cause the abnormality of multiple signal transduction pathways of trophoblast cells, affect their growth, proliferation, differentiation, apoptosis and invasion, etc., eventually leading to decreased trophoblast invasion, which is closely related to the pathogenesis of PE.

5. Conclusion

In summary, low HLA-G expression plays a major role in the incidence of hypertensive disorders in pregnancy. Its expression level monitoring is conducive to the early diagnosis of this disease, and regulating its gene expression may also become a new target for the treatment of hypertensive disorders of pregnancy, so as to provide a new idea for its pathogenesis, diagnosis and treatment.

References


