Expression of nephrin in the human placenta and fetal membranes

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Abstract. Nephrin is the signature molecule in the podocyte of the glomerulus that forms the renal slit diaphragm, the main functional unit of the glomerulus. The present study focused on the expression of nephrin in the human placenta, which may also have a role in filtration and the maintenance of homeostasis in the kidneys. A total of nine placentas from normal healthy pregnant females at full term were investigated. Reverse transcription-quantitative polymerase chain reaction, western blotting and immunofluorescence were performed. The expression of nephrin mRNA was relatively increased in the chorion compared with that in the villi and the amnion. The nephrin gene was detected in the villous cytotrophoblast cells and the endothelium of the intravillous vessels. It was also present in the chorionic and amniotic membranous lining, with its distribution being particularly dense in the amnion. The identification of nephrin in the human placenta, particularly at the maternal-fetal interface, provides a novel insight into the molecular basis of the selective permeability of the placental barrier, which requires further elucidation.

Introduction

The podocyte is the key component of the glomerular slit diaphragm. The podocyte and its associated proteins form the diaphragm and these cells have an intimate interaction with capillary endothelial cells, which exhibit an important role in selective filtration acting as a barrier. Previously, several studies have identified podocyte-associated proteins as the main factors in nephropathy demonstrating glomerular basement membrane alteration, such as in diabetic nephropathy (1,2). Notably, the blood-brain barrier of the central nervous system, another representative homeostatic barrier system in the human body, contains astrocytes, which are similar to podocytes in structure and function (3,4).

Nephrin is a podocyte-associated protein, a constituent of the slit diaphragm of the glomerulus, which appears to have a critical role in maintaining glomerular filtration. Previously, inactivation of the nephrin gene in the mouse embryo demonstrated a detrimental loss of the ultrafiltration function of the kidneys (5).

Similar to the kidney, the placenta is a major organ of the fetus, which filtrates and exchanges fetal and maternal blood, enabling the placental cotyledon and the glomerulus to undertake homeostatic functions. Notably, the expression of the nephrin gene has been reported in the fetal membranes and placenta of pregnant Sprague-Dawley rats (6). This led us to hypothesize that nephrin may be involved in regulating homeostasis at the maternal-fetal interface. However, thus far, evidence of nephrin expression in the human placenta is lacking. Therefore, the objective of the present study was to evaluate the presence and localization of nephrin, a signature molecule of podocytes (7), in the human placenta.

Materials and methods

Participants. Females that were pregnant with a single fetus, who delivered a normal infant between 37 and 40 weeks of gestation by Cesarean section at Severance Hospital, Yonsei University College of Medicine (Seoul, Republic of Korea), between January 2011 and July 2012 were enrolled in the present study. Patients with a history of active labor prior to the surgery, multiple pregnancies, prior or current diagnosis of any medical illness (diabetes, gestational diabetes, hypertension, thyroid disease or infectious disease), placenta previa, fetal anomaly, oligohydramnios, hydramnios, fetal aneuploidy, preterm labor...
or premature rupture of the membranes during the present pregnancy were excluded from the present study. The present study was approved by the Institutional Review Board of Yonsei University College of Medicine and all patients consented to be involved in the present study.

Sample collection. When the placenta was delivered, the amnion and chorion were dissected under aseptic conditions. Dissected membranes were sampled at a size of 1.0x1.0x1.0 cm\(^3\). Villous tissue measuring 1.0x1.0x1.0 cm\(^3\) was sampled at the site near the cord insertion. Following the tissue sampling, the amnion, chorion and villous tissue were frozen in liquid nitrogen and stored for further analysis.

Quantification of mature mRNA levels using reverse transcription-quantitative polymerase chain reaction (RT-qPCR). RNA fractions were initially isolated from the tissue samples using an RNA extraction kit (Intron Biotechnology, Inc., Gyeonggi, Korea). RNA concentrations were measured using absorbance at 260/280 nm with a SuperScript First-Strand RT kit (Invitrogen Life Technologies, Carlsbad, CA, USA). Total RNA (1 µg) was converted to cDNA using oligo-dT as the primer and reverse transcriptase (Invitrogen Life Technologies). The reaction mixture (20 µl) was incubated at 65˚C for 5 min, 50˚C for 50 min and 85˚C for 5 min. RT-qPCR was performed with Applied Biosystems 7500 real-time PCR system (Applied Biosystems Life Technologies, Foster City, CA, USA) using the Taqman® Gene Expression Assays kit (Taqman Life Technologies, Foster City, CA, USA). A total of 20 µl of the RT-qPCR reaction mixture contained 2 µl RT-qPCR products, 10 µl Taqman® 2X universal PCR master mix and 1 µl 20X Taqman® Gene Expression Assay mix (RT-qPCR primers). Nuclease-free water was used to adjust the final volume to 20 µl. The reactions were incubated in a 96-well optical plate at 95˚C for 10 min, followed by 40 cycles of 95˚C for 15 sec and 60˚C for 60 sec. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as an endogenous reference gene for normalizing the results. The relative abundance of each mRNA was calculated using the comparative cycle threshold \(2^{-\Delta\Delta Ct}\) method. Individual samples were assayed in triplicate, with three independent biological replicates. The primer sequences for the endogenous control (GAPDH) and nephrin are shown in Table I.

Western blotting. Nephrin expression was examined using western blotting analysis of the tissue homogenate samples. Following isolating each protein using the cell lysis buffer (Cell Signaling Technology, Inc., Danvers, MA, USA), 30 µg of total protein loading was used to run 8% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and then transferred to polyvinylidene difluoride membranes (Millipore, Eten-Leur, The Netherlands). The blot membranes were blocked with 5% skimmed milk in Tris-buffered saline-Tween 20 (containing 10 mM Tris, pH 8.0, 150 mM NaCl, 0.0025% Tween-20) at room temperature for 1 h. Subsequently, the membrane was incubated overnight at 4˚C with specific antibodies against nephrin (polyclonal rabbit anti-human, 1:1,000; #2265; ProSci, Portway, CA, USA) and β-actin (monoclonal mouse anti-human, 1:2,000; A5441; Sigma-Aldrich, St. Louis, MO, USA). The membrane was then washed six times with TBS-T for 5 min. Thereafter, the membrane was incubated with anti-rabbit antibody (IgG, 1:4,000; ab6802; Abcam, Cambridge, MA, USA) and anti-mouse antibody conjugated with horseradish peroxidase (IgG, 1:5,000; ab6808; Abcam) for 1 h at room temperature. The membrane was washed eight times with TBS-T for 5 min at room temperature. The protein was visualized on the ImageQuant LAS-4000 imager using enhanced chemiluminescence (GE Healthcare, Arlington Heights, IL, USA). Relative densities for nephrin expression were normalized using tubulin expression for each sample.

Immunofluorescence (IF). Cells were fixed with acetone for 15 min and washed three times with PBS-T (PBS/0.1% Tween-20

### Table I. Sequence of primers for GAPDH and the human nephrin gene.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer sequence (5’-3’)</th>
<th>Size (bp)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GAPDH</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sense</td>
<td>AGGCCAACCGCGAGAAGATGACC</td>
<td>320</td>
</tr>
<tr>
<td>Antisense</td>
<td>GAAGTCCAGGGCGACGTAGCAC</td>
<td></td>
</tr>
<tr>
<td>Human nephrin</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sense</td>
<td>CCAACATCGTTTTCACTTGG</td>
<td>349</td>
</tr>
<tr>
<td>Antisense</td>
<td>GGGTGGTACGACATCCACAC</td>
<td></td>
</tr>
</tbody>
</table>

### Table II. Clinical characteristics of pregnancies included in the present study.

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>Variable</th>
</tr>
</thead>
<tbody>
<tr>
<td>Maternal age (years)</td>
<td>34 (30-42)</td>
</tr>
<tr>
<td>Primigravida (n)</td>
<td>2</td>
</tr>
<tr>
<td>Nulliparity (n)</td>
<td>4</td>
</tr>
<tr>
<td>Gestational days</td>
<td>38±3 (37±3-39±1)</td>
</tr>
<tr>
<td>Cesarean section indication (n)</td>
<td></td>
</tr>
<tr>
<td>Prior cesarean section</td>
<td>6</td>
</tr>
<tr>
<td>Prior myomectomy</td>
<td>1</td>
</tr>
<tr>
<td>Advanced maternal age</td>
<td>1</td>
</tr>
<tr>
<td>Cephalopelvic disproportion</td>
<td>2</td>
</tr>
</tbody>
</table>

Data are expressed as median and range.
Following blocking with 5% goat serum for 1 h, the cells were incubated with anti-nephrin antibody (polyclonal rabbit anti-human, 1:1,000; #2265; ProSci) overnight at 4°C. Negative controls consisted of cells incubated with rabbit IgG (1:200; cat no. ab27478; Abcam) at the same concentration as the antibody against nephrin. The cells were washed three times with PBS-T and incubated with goat polyclonal secondary antibody to rabbit IgG (1:200, DyLight® 594; cat no. ab96901; Abcam) for 1 h. The cells were washed three times in PBS-T and mounted with a fluorescent mounting medium (Vector Laboratories, Inc., Burlingame, CA, USA). Fluorescent microscopy was performed using the LSM700 microscope (Carl Zeiss; Oberkochen, Germany).

**Statistical analysis.** Statistical analysis was performed using the SPSS software package, version 18.0 (IBM Inc., Armonk, NY, USA). The mRNA expression level of the nephrin gene was expressed as the mean ± standard deviation. Descriptive statistics were used for the baseline characteristics and the linear mixed model for comparing the mRNA expression of the nephrin gene in the villi, chorion and amnion. P<0.05 was considered to indicate a statistically significant difference.

**Results**

In all, nine healthy pregnant females were enrolled in the present study. The basal characteristics of the patients are shown in Table II. Due to the small sample size, data was expressed as the median and the range. One participant had a history of a previous cesarean section and concurrent myomectomy.

**mRNA and protein expression in the placenta.** The expression of nephrin mRNA and protein in the uteroplacental unit were confirmed using RT-qPCR and western blotting,
The gene expression was relatively higher in the chorion when compared with the villus and amnion (mean, $2^{\Delta\Delta Ct} = 3.79\pm 4.49$ vs. $0.50\pm 0.45$ and $2.12\pm 2.41$); however, the difference was not statistically significant ($P=0.37$).

**Immunolocalization of nephrin.** Marked positive staining for nephrin was observed in the villi, chorion and amnion in the IF studies. Nephrin expression was clearly localized to the syncytiotrophoblast and endothelial cells of the arteries and veins of the chorionic plate, where the labeling was pronounced at the apical membrane of the syncytiotrophoblast. In the fetal membranes, intense immunoreactivity was localized to the cuboidal amniotic epithelium, particularly at the apical membrane and in the stromal cells (Fig. 2).
Discussion

The main role of the placenta is the separation of maternal and fetal circulation, while also facilitating the transport of nutrients and substances from the mother to the fetus. As the placenta has extensive contact between the mother and the fetus in structures, such as the villus, amnion-chorion and the chorion-decidua, it is hypothesized that a filtering, barrier-like structure exists within the placenta at such interfaces. As hypothesized, evidence of the presence of nephrin in cells at the maternal-fetal interface was identified and immunoreactivity to nephrin was most prominent at the apical membrane of the syncytiotrophoblasts and amniotic epithelium, and in the stromal cells of the chorion. The present findings suggest the existence of a barrier-like structure formed by nephrin in the placenta similar to that of the glomerulus.

Although the present study did not investigate the function of nephrin in the placenta, it is likely that in the placenta, the protein may have a role similar to that observed in other organs (8-13). Its presence in astrocytes, which interact with the capillaries in the blood-brain barrier of the central nervous system suggested it exhibits a similar role as in the glomerular slit membrane (13). The role for nephrin as a barrier system has also been suggested in the testis due to the observation of nephrin in the Sertoli cells colocalized with zona occludens-1 along the basement membrane of the seminiferous tubule in the testis (9). However, a previous study demonstrated that nephrin is expressed in the radial glial cells, which are involved in the directional migration of neurons and development of glial cell lineages (14), and that it binds with glutamate receptors and scaffolding molecules of the primary neuronal cells (5). Furthermore, its active role in the vesicle and actin interaction involved in insulin release was described in the pancreatic β islet cells (11,12). These findings suggest that nephrin not only functions as a barrier, but also exhibits functional roles, such as in cell maturation and development, cell-to-cell interaction and signaling (15-19).

It has been hypothesized that nephrin, expressed in the villi and the membranes holds such functions as well as other organs. Nephrin may be involved in the seclusion of fetal materials from the maternal circulation and thereby provide an immunologically aergic environment, which may be crucial to the exchange of nutrients between the mother and the fetus or it may have a role in placental development. If so, impairment of placental nephrin may result in an adverse pregnancy outcome, such as preeclampsia, restricted fetal growth or abnormal placental growth. As serum from a patient with preeclampsia has been revealed to damage the podocyte and shed nephrin via endothelin-mediated endothelin-1 (20), theoretically, nephrin in the syncytiotrophoblasts or the chorioamniotic membranes in close contact with the maternal serum may also be affected or disrupted via a similar mechanism. This may partly contribute to the pathophysiological mechanism of adverse fetal outcomes.

To the best of our knowledge, the present study is the first to provide evidence of the presence of nephrin in the human placenta and placental membranes. The identification of podocyte-associated nephrin expression in the placenta, particularly at the maternal-fetal interface, provides novel insight into the molecular basis of the selective permeability of the placental barrier, which requires further elucidation. The present findings highlight the requirement for further investigation into the placental expression of podocyte-associated proteins in pregnancies with complications and their specific roles in the human placenta.

Acknowledgements

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References