Chorionic Villi Expression of Two Vascular Endothelial Growth Factor Proteins in Normal Human Pregnancy

Jayasri Basu PhD*, Samantha Gonzalez-Ramos MD, Vanitha Banajjar MD, Aruna Mishra MD, Magdy Mikhail MD
BronxCare Health System, Department of Obstetrics & Gynecology, Bronx, New York

*Corresponding Author: Jayasri Basu PhD, Director-Graduate Medical Education, Department of OB/GYN, BronxCare Health System, Clinical Assistant Professor of Obstetrics & Gynecology and Reproductive Science Icahn School of Medicine of Mount Sinai, 1276 Fulton Avenue, Room 233, Bronx, NY 10456, New York;
Tel: 7189018476; Email: jbasu@bronxcare.org

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Abstract

Introduction: Alternate splicing of vascular endothelial growth factor (VEGF)-A gene yields a number of proteins of varying amino acid lengths ranging from 121 to 206 amino acids. Recently, other isoforms of VEGF have been recognized that form from alternate splicing of the C-terminal region of exon 8 of VEGF gene. These sister isoforms are known as VEGF\textsubscript{xxx}, where xxx denotes the number of amino acids present in the protein. During human pregnancy, VEGF\textsubscript{165} and its receptors are recognized as master regulator of placental angiogenesis; and both VEGF\textsubscript{165} and VEGF\textsubscript{165b} proteins are secreted by cytotrophoblasts. The present study compares the temporal expression of both isoforms of VEGF in chorionic villi tissues, throughout gestation, in normal human pregnancy.

Methods: Placentas were obtained from elective termination of pregnancy or term delivery. Tissues collected were dissected in saline to identify chorionic villi without associated decidua. VEGF protein expressions were determined by ELISA using monoclonal antibody to human VEGF\textsubscript{165} and VEGF\textsubscript{165b} proteins as capture antibody, DY293B and DY3045, respectively (R&D Systems, Minneapolis, MN). Non-parametric test considered p<0.05 as significant.

Results: Both isoforms of VEGF were detected in 166 chorionic villi samples analyzed. The expression patterns of the two proteins differed markedly throughout gestation. While VEGF\textsubscript{165} showed a minor dip in the second trimester, VEGF\textsubscript{165b} showed a peak during that time. The two isoforms were positively and significantly correlated in second and third trimesters of pregnancy.

Conclusions: The data reveal that both isoforms of VEGF play key roles in regulating the angiogenic balance throughout gestation in normal human pregnancy. We hypothesize that VEGF\textsubscript{165b} protein expression in placental tissues could be a physiological phenomenon to restrain overexpression of VEGF\textsubscript{165} during placentation development, which if left unchecked, could lead to pregnancy-related complications as pregnancy advanced.

Keywords: ELISA, Throughout Gestation, Uncomplicated Human Pregnancy, VEGF\textsubscript{165}, VEGF\textsubscript{165b}

Introduction

Placental angiogenesis plays a pivotal role in instituting a fetomaternal circulation and in establishing the placental villous tree that contributes to the development of the placenta throughout human pregnancy. Of the many angiogenic factors that were investigated e.g., five members of VEGF family, four members of the angiopoietin family, and one member of the ephrin family; vascular endothelial growth factor (VEGF) was recognized as the one specific for blood vessel formation [1–3]. Gene knockout studies have provided convincing evidence for a central role of VEGF in fetal and placental angiogenesis. Targeted homozygous null mutations of VEGF receptors in mice demonstrated failure in hematopoiesis, formation of blood islands and blood vessels, resulting in embryonic death by day 8 of pregnancy [4]. Carmeliet et al. further demonstrated that loss of a single VEGF allele in a mouse model led to gross developmental deformities in vessel formation that resulted in embryonic death between day 11 and 12 of mouse pregnancy. The authors concluded that not only fetal and placental angiogenesis were dependent on VEGF, but a threshold level of VEGF had to be achieved for normal vascular development to occur [5].

The VEGF A gene has eight exons. Alternate splicing of VEGF mRNA accounts for five isoforms of VEGF proteins: VEGF\textsubscript{121}, VEGF\textsubscript{145}, VEGF\textsubscript{165}, VEGF\textsubscript{186} and VEGF\textsubscript{206}, of which VEGF\textsubscript{165} isoform is most abundant in vivo [6]. In the past decade, the complexity of VEGF\textsubscript{165} biology further intensified when it was found that alternate splicing of exon 8 at the C-terminal region of VEGF gene could yield yet other isoforms of VEGF. While the number of amino acids in the sister isoforms remained the same, there were however alternate open reading frames of six amino acids at the C-terminal region of the sister isoforms. The sister isoforms were identified as VEGF\textsubscript{xxx},xxx referring to the number of amino acids present in the protein [7]. The splicing event modifies the C-terminal region of the VEGF protein from CDKPRR (VEGF\textsubscript{xxx}) to SLTRKD (VEGF\textsubscript{xxx}). The replacement of the six amino acids alters the tertiary structure of the newly formed protein, as the disulfide bond that was previously formed between cysteine 160 at the C-terminal region with cysteine 146 of exon 7, could no longer be formed. The terminal two arginine molecules (RR)
being replaced with lysine and aspartic acid (KD) further modified the overall charge of the protein; and replacement of proline (P) residue with arginine (R) additionally transformed the structure of the C-terminal domain [7].

In a previous study, placental expressions of VEGF\textsubscript{165} and VEGF\textsubscript{165,b} were examined in which the comparison was made between uncomplicated and complicated pregnancies at term [8]. In an earlier study, we have examined placental expression of VEGF\textsubscript{165} throughout gestation in normal human pregnancy [9]. The two growth factors of VEGF are secreted by cytotrophoblasts; and are implicated in placental angiogenesis in human pregnancy. We, therefore, undertook the present study to simultaneously investigate the expressions of both VEGF\textsubscript{165} and VEGF\textsubscript{165,b} in placental tissues, throughout gestation, in women with uncomplicated pregnancy. Understanding the simultaneous temporal changes in these two growth factors of VEGF as placenta develops, we consider, would be extremely valuable.

**Materials and Methods**

The investigative protocol for the study was approved by the Human Subject Ethics Committee of the BronxCare Health System, New York, protocol # 10101304. Discarded placental tissue samples were collected within approximately 30 minutes of the procedures from normal pregnant women who underwent elective termination of pregnancy at 6 weeks to 23 weeks and 6 days; and from normal women who delivered uncomplicated singleton pregnancies at term. Placentas from missed abortion or from pregnancies complicated with diabetes, hypertension, chronic renal disease, and chronic peripheral vascular disease or with major fetal anomalies were excluded. The approved protocol allowed the collection of the following clinical information regarding the women from whom placental tissues were obtained. These included: maternal age, parity, race/ethnicity (self-reported), gestational age (as determined by ultrasound or by initial date of the last menstrual period), reason for pregnancy termination (whether elective, for maternal medical reasons or for fetal indications), and medicine(s) administered to induce termination of pregnancy. Placentas from pregnancies 7–23 weeks and 6 days were collected from the elective termination group, and those from 37 to 42 weeks of gestation were collected from term delivery group. Placentas delivered below 37 weeks of gestation were not included because these placentas are considered as preterm.

Soon after collection, the placental tissues were processed to obtain choriocarcinoma villi samples by a method described earlier [10]. Briefly, portions of each placenta were first thoroughly washed in cold saline to remove maternal blood and were then dissected in saline to collect free floating choriocarcinoma villi that were not anchored to the basal plate nor were emerging from the chorionic plate surface vessels. Pieces of these choriocarcinoma villi samples were placed in separate tubes, each tube bearing the same ID# designated for that placenta. Tissues and clinical information were de-identified before exiting the delivery suites. The sample collection tubes were then transported to the laboratory on ice and stored at -80°C until assay.

Chorionic villi VEGF\textsubscript{165} and VEGF\textsubscript{165,b} protein expressions were simultaneously determined by enzyme-linked immunoassay (ELISA) methods. The capture antibodies for the ELISA kits were monoclonal antibodies to human proteins VEGF\textsubscript{165} (DY293B) and VEGF\textsubscript{165,b} (DY3045), respectively (R&D Systems, Minneapolis, MN). Chorionic villi samples were homogenized in Reagent Diluent 2 (R&D Systems, Minneapolis, MN), and supernatants following centrifugation of the homogenate at 13000 rpm for 2 minutes were used to determine the free forms of the isoforms based on the manufacturer’s protocols. Simultaneous analysis of both VEGF isoforms meant that we carried out both VEGF\textsubscript{165} and VEGF\textsubscript{165,b} proteins assays using chorionic villi tissues samples isolated from the same placentas, the assays were not carried out using the same aliquots. A Tecan infinite 200 Pro microplate reader (Tecan Systems Inc., San Jose, CA) set at 450 nm with wavelength correction set at 540 nm was used to measure absorbance. The sensitivity of VEGF\textsubscript{165} ELISA was 31.3 pg/ml and that of VEGF\textsubscript{165,b} ELISA was 62.5 pg/ml. Intra-assay and inter-assay variations for both assay kits were between 7–10%.

**Statistical Analyses**

The statistical software package SPSS, version 25 (IBM Corporation, Armonk, NY) was used for statistical analyses. First, the normality of the data was statistically analyzed which showed that the data was not normally distributed. Hence, non-parametric statistics were applied. The data was grouped by trimesters and the following non-parametric tests were performed: 1) Kruskal Wallis test (an alternative test to One-Way ANOVA) was used to explore the differences in VEGF isoform expressions among the trimester groups. (It is important to note that KW test does not provide post-hoc data the way one-way ANOVA does). 2) The non-parametric Mann Whitney U test was applied to compare two trimester groups at a time against each other. 3) Spearman Rank correlation coefficient test was applied to summarize the strength and direction of a relationship between the variables as well as between the variables and gestational age in days. A p<0.05 was considered statistically significant.

**Results**

The demographic characteristics of women from whom placental samples were obtained showed comparable maternal age among the three trimester groups (28.2 ± 6.5, 28.2 ± 6.1 and 30.0 ± 6.8 years, for first, second and third trimester groups, respectively). Race/ethnicity was self-reported, and the distribution pattern was 30% Black, 60% Hispanic, 2% Caucasian and 8% of other ethnic origin. When 166 placentas were grouped by trimester the data showed that 71 placentas collected from elective termination of pregnancy were in the first trimester, with an average gestational age of 8 weeks; 33 placentas collected also from elective termination of pregnancy were in the second trimester, with an average gestational age of 16 weeks; and 62 placentas collected from term delivery were from third trimester, with an average gestational age of 39 weeks and 2 days.

In this study, non-parametric statistics were applied because the data were skewed. Kruskal Wallis test results revealed that VEGF\textsubscript{165} protein expression was not statistically different among the three trimester groups, the expression of VEGF\textsubscript{165,b} protein, however, was significantly different (p=0.0001). In (Table 1) the 25th, 50th and 75th percentile values of the two VEGF isoforms are shown. The expression
patterns of the two proteins are shown graphically in (Figure 1), which depicts two entirely different expression patterns for the two isoforms of VEGF throughout gestation in normal human pregnancy. While VEGF\textsubscript{165} protein showed a minor dip in the second trimester of pregnancy, the expression patterns of VEGF\textsubscript{165b} showed a significant peak during that time. Pairwise comparison of trimester groups using Mann Whitney U test further revealed that the expression pattern of VEGF\textsubscript{165} was not significantly different, but VEGF\textsubscript{165b} protein expression in each trimester was significantly different from the other (p=0.0001).

<table>
<thead>
<tr>
<th>Groups</th>
<th>N</th>
<th>VEGF\textsubscript{165} (pg/100 mg tissue)</th>
<th>VEGF\textsubscript{165b} (pg/100 mg tissue)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>25\textsuperscript{th} Percentile</td>
<td>50\textsuperscript{th} Percentile</td>
</tr>
<tr>
<td>1\textsuperscript{st} Trimester</td>
<td>71</td>
<td>68.80</td>
<td>87.45</td>
</tr>
<tr>
<td>2\textsuperscript{nd} Trimester</td>
<td>33</td>
<td>50.80</td>
<td>68.40</td>
</tr>
<tr>
<td>3\textsuperscript{rd} Trimester</td>
<td>62</td>
<td>77.25</td>
<td>110.95</td>
</tr>
</tbody>
</table>

VEGF\textsubscript{165}: Vascular Endothelial Growth Factor\textsubscript{165}; VEGF\textsubscript{165b}: Vascular Endothelial Growth Factor\textsubscript{165b}; GA: gestational age in days. The first trimester placental chorionic villi samples were from 70/7-120/7 weeks gestation, the average GA was 81/7 weeks; second trimester were from 121/7 to 239/7 weeks, the average GA was 160/7 weeks; and the third trimester term were from 370/7 to 414/7 weeks of gestation, the average GA 392/7 weeks. Homogenized human placental chorionic villi samples were analyzed using assay kit from R&D Systems, Minneapolis, MN; to determine VEGF\textsubscript{165} and VEGF\textsubscript{165b} protein expressions. Data were not normally distributed hence non-parametric statistics were used and 25\textsuperscript{th}, 50\textsuperscript{th} and 75\textsuperscript{th} percentile values of VEGF\textsubscript{165} and VEGF\textsubscript{165b} are shown.

Figure 1. GA: gestational age.
The first trimester placental chorionic villi samples were from 70/7-120/7 weeks gestation, the average GA was 81/7 weeks; second trimester were from 121/7 to 239/7 weeks, the average GA was 160/7 weeks; and the third trimester term were from 370/7 to 414/7 weeks of gestation, the average GA 392/7 weeks.

Correlation between the two proteins of VEGF as well as between the isoforms of VEGF and gestational age were examined in this study. Spearman’s correlation test results revealed that expression of VEGF\textsubscript{165b} protein was significantly and positively correlated with gestational age in days in the first trimester of normal pregnancy (r= +0.299, p=0.011, (Table 2). The table also depicts the test results of Spearman’s correlation between the two isoforms. The data reveal a significant positive correlation in the second (r=+0.376, p=0.031) and third trimester (r=+0.271, p=0.033), of normal pregnancy.
Table 2. Correlation between VEGF$_{165}$, VEGF$_{165b}$ and Gestational Age in Days throughout gestation

<table>
<thead>
<tr>
<th>Trimester (N)</th>
<th>VEGF$_{165}$</th>
<th>VEGF$_{165b}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>First (N=71)</td>
<td>Correlation 1.000</td>
<td>0.070</td>
</tr>
<tr>
<td></td>
<td>Coefficient 0.029</td>
<td>0.299*</td>
</tr>
<tr>
<td></td>
<td>Sig. (2-tailed) 0.159</td>
<td>0.011</td>
</tr>
<tr>
<td></td>
<td>GA Correlation 1.000</td>
<td>0.376*</td>
</tr>
<tr>
<td></td>
<td>Coefficient 0.013</td>
<td>0.031</td>
</tr>
<tr>
<td></td>
<td>Sig. (2-tailed) 0.409</td>
<td>0.944</td>
</tr>
<tr>
<td>Second (N=33)</td>
<td>Correlation 0.944</td>
<td>0.271*</td>
</tr>
<tr>
<td></td>
<td>Coefficient 0.916</td>
<td>0.033</td>
</tr>
<tr>
<td></td>
<td>Sig. (2-tailed) 0.833</td>
<td>0.385</td>
</tr>
<tr>
<td>Third (N=62)</td>
<td>Correlation 0.027</td>
<td>-0.112</td>
</tr>
<tr>
<td></td>
<td>Coefficient 0.013</td>
<td>0.385</td>
</tr>
<tr>
<td></td>
<td>Sig. (2-tailed) 0.833</td>
<td>0.385</td>
</tr>
</tbody>
</table>

VEGF$_{165}$ : Vascular Endothelial Growth Factor$_{165}$, VEGF$_{165b}$: Vascular Endothelial Growth Factor$_{165b}$; GA: gestational age in days. Data were not normally distributed hence Spearman’s correlation was applied. Significant positive correlation was seen between VEGF$_{165b}$ and GA in the first trimester of pregnancy. The two VEGF proteins were significantly and positively correlated in the second and third trimester of normal pregnancy.

Conclusion

In this study, chorionic villi samples were isolated from each placenta, and VEGF$_{165}$ and VEGF$_{165b}$ protein expressions were determined by ELISA that used monoclonal antibody to human VEGF$_{165}$ or VEGF$_{165b}$ protein as the capture antibody in each case. VEGF protein expression and that of its receptors are closely regulated to times of vasculogenesis and angiogenesis [11]. In this study, both isoforms of VEGF proteins were identified in all 166 chorionic villi samples that were analyzed suggesting that both proteins of VEGF may exert regulatory roles in pregnancy-linked angiogenesis in normal human pregnancy.

The angiogenic potential of VEGF$_{165}$ has been confirmed by several bioassays that measured VEGF receptor expression during embryogenesis and tissue repair [11], capillary growth in vivo in developing chick chorioallantoic membrane [12], and a temporal and spatial correlation between VEGF and ocular angiogenesis in a primate model [13]. These studies affirmed VEGF signaling pathways to be the master regulator of angiogenesis [11–14]. The pro-angiogenic potential of VEGF$_{165}$ results from six amino acids at its C-terminus e.g., CDKPRR [7, 15]. Involvement of VEGF$_{165}$ in human pregnancy, particularly in placental development has also been recognized. Reports on the placental expressions of VEGF$_{165}$ protein state that the protein increases in first 10 weeks of normal pregnancy, but as pregnancy advances, VEGF and its receptor VEGFR-2 concentrations decline. In these reports, VEGF was suggested to increase vascular permeability, vasodilation and angiogenesis [16–18]. It is also suggested to be involved in the formation and maintenance of the trophoblastic plugs that block the spiral arteries in the first trimester of normal pregnancy [19]. Figure 1 depicts that VEGF$_{165}$ protein expression is somewhat lower in the second trimester of normal pregnancy. This data is consistent with the finding of an immunohistochemical study that reported VEGF$_{165}$ antigen staining to be weaker in mid-gestational placental tissues, compared to the intensity of staining for the protein in the first or third trimester placental samples [20].

The isoform VEGF$_{165b}$ has not been investigated in as much detail as VEGF$_{165}$. VEGF$_{165b}$ protein is expressed in normal tissues of lung, pancreas, colon, and brain [7, 21, 22]. Reports reveal that VEGF$_{165b}$ constitute 50% or more of total VEGF expressed in most non-angiogenic tissue [23]. Investigators have alleged that VEGF$_{165b}$ is not pro-angiogenic in vivo [23] and it actively inhibits VEGF$_{165}$ mediated endothelial cell proliferation and migration in vitro [4, 7, 21–23]. VEGF$_{165b}$ is further reported to inhibit physiological angiogenesis in developing mammary tissue, in transgenic animals, overexpressing VEGF$_{165b}$ [24]. Moreover, VEGF$_{165b}$ is reported to inhibit angiogenesis in vivo in six other different tumor models [21–25]. The results of the present study on chorionic villi VEGF$_{165b}$ protein expression throughout gestation in normal human pregnancy are similar to the results we have reported earlier [9]. Our studies underscore the importance of VEGF$_{165b}$ antigen in normal human pregnancy. Identification of VEGF$_{165b}$ protein in this study, in all 166 chorionic villi samples that were collected throughout all trimesters of human pregnancy validates this notion. This temporal variation in VEGF$_{165b}$ protein in human pregnancy perhaps reflects that VEGF$_{165b}$ protein expression is more stringently controlled at each phase of human gestation, and that the isoform may play a more active role in placental development. The spatial activation of VEGF$_{165b}$ protein at various phases of human gestation as seen in the study emphasizes that placental angiogenesis could be more dependent on VEGF$_{165b}$ isoform.

VEGFR-2 is the predominant signaling receptor for VEGF-mediated angiogenesis [26]. Phosphorylation of VEGFR-2 on tyrosine residue at 1052 position of the molecule generates a highly dynamic and complex signaling system that triggers angiogenic response [4, 21, 26]. The binding affinities of VEGFR-2 are identical between VEGF$_{165}$ and VEGF$_{165b}$ isoforms. However, VEGF$_{165b}$ is less efficient than VEGF$_{165}$ in inducing phosphorylation on Y1052 residue [27]. The lower ability of VEGF$_{165b}$ to induce VEGFR2 phosphorylation on Y1052 is due to its incapacity to induce optimal rotation of the intracellular domain of the receptor molecule that is required for phosphorylation to occur [27]. Reduced Y1052 phosphorylation induces rapid inactivation of the kinase domains of the receptors resulting in weak activation of the signaling pathways [23]. Recent studies have shown that VEGF$_{165b}$ can stimulate VEGFR-2, ERK1/2 and Akt phosphorylation in endothelial cells, however, the induced phosphorylation is weaker than that promoted by VEGF$_{165}$ [21, 28]. VEGF$_{165b}$ is currently not considered as anti-angiogenic but as a weakly angiogenic form of VEGF [28].

Establishment of functional fetal and placental circulations is some of the earliest strategic events during embryonic/placental development [29, 30]. The significant positive correlation seen between chorionic villi VEGF$_{165b}$ protein expression and gestational age in first trimester of pregnancy in this study (Table 2) suggests that VEGF$_{165b}$

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may contribute to the development of villous vascular network in early pregnancy. An increase in expression of plasma VEGF$_{bb}$ protein in first trimester of normal pregnancy has been reported by other investigators, and the authors suggest that failure in the upregulation of VEGF$_{bb}$ protein in the plasma in the first trimester of pregnancy was a predictive marker of preeclampsia [31]. Simultaneous increase of both isoforms of VEGF in human placental tissues in the third trimester of pregnancy has also been reported by other investigators [8]. The large increase in transplacental exchange which supports the exponential increase in fetal growth and uterine blood flow during the last half of gestation is suggested to depend primarily on the dramatic growth of placental vascular beds [32]. Vascular density of the placental cotyledons remains relatively constant throughout mid-gestation and increases dramatically during the last third of gestation in association with dramatic fetal growth [32, 33]. In third trimester of normal pregnancy, both fetal development and demands are at its peak; and our data show a positive correlation between VEGF$_{165}$ and VEGF$_{bb}$ during this time. Hence, it may be suggested that angiogenic modification of placental vasculature that allows maximum blood to flow through may depend on the synergistic action of both isoforms of VEGF.

Our present findings suggest a notable difference between tumor and placental angiogenesis. It is widely reported that in human tumors, up-regulation of VEGF$_{165}$ protein occurs with a proportional drop in VEGF$_{bb}$ levels; indicating that in tumor angiogenesis the balance between the two isoforms of VEGF is lost [34]. However, the findings of the present study suggest that gestational age-specific expression of both VEGF isoforms is necessary for optimal placental angiogenesis and growth to results in a successful viable outcome.

That hypoxic environment favors VEGF$_{165}$-induced angiogenesis and tumorigenic growth is well known [35]. In an earlier study we have demonstrated that placental environment switches from a hypoxic to a normoxic environment beyond the first trimester of normal human pregnancy [36]. We hypothesize that perhaps during this transitional period from first to second trimester of normal pregnancy, placental oxidative environment per se regulate the alternate splicing of exon 8 of the VEGF gene. In hypoxic condition proximal splicing of exon 8 may be favored, which up-regulates the expression of VEGF$_{165}$ protein. In a normoxic state, distal splicing of exon 8 may be favored, whereby the expression of VEGF$_{bb}$ protein gets up-regulated. That switch in the partial pressure of oxygen at the end of first trimester of normal pregnancy can modify the expression pattern of two proteins has been reported earlier, between placental derived growth factor and VEGF [37]. It is feasible that this switch in the splicesome at the end of the first trimester may be the contributing factor that may have resulted in the peak VEGF$_{165}$ protein expression seen in the second trimester of this study. It is likely that this dependence of placental growth on VEGF$_{bb}$ beyond the first trimester could be a physiological phenomenon to restrain any overexpression of VEGF$_{165}$ which if left unchecked, could perhaps lead to pregnancy-related complications as pregnancy advanced.

The limitation of our study is that we have not confirmed our data by western blot, polymerase chain reaction or by immunohistochemical methods. We acknowledge that cells respond to extracellular stimuli through a series of signaling cascades. When a receptor is activated, varieties of proteins are recruited to interact with each other to generate a cascade of sequential steps that result in a biological effect [38]. Signaling molecules are common to several pathways and often form a complex intracellular network [38]. Yet, in this study we have not examined other isoforms of VEGF, their receptors or other factors that may have also been involved in placental angiogenesis. We believe that addition of all these factors would have made the study more complex and any conclusions drawn from the study would have been extremely difficult to interpret. The strength of our study is our relatively larger sample size in each of the trimester groups. Simultaneous expressions of these two important VEGF proteins in human chorionic villi tissue throughout gestation in normal human pregnancy have not been reported before. In this study, stringent ELISA methods using monoclonal antibodies to human VEGF$_{165}$ and VEGF$_{bb}$ human proteins were used as capture antibodies. The manufacturer claims that DY293B capture antibody that was used does not cross react with placental growth factor (PIGF), VEGF-B$_{165}$, VEGF-B$_{165}$, VEGF-C, VEGF-D, recombinant (rh) human VEGF R1(Flt-1)/Fc Chimera or with rhVEGF R2(KDR)/Fc Chimeras. Similarly, the monoclonal capture antibody DY3045 does not cross react with rhhuman VEGF$_{165}$, VEGF$_{bb}$, VEGF$_{165}$, rhVEGF R3/Flc, VEGF R1/Fc or rhVEGF R2/Fc Chimeras. Furthermore, the detection methods used were sensitive, allowing the detection of VEGF$_{165}$ as low as 31.2 and VEGF$_{bb}$ as low as 62.5 pg/ml, respectively. The study underscores the importance of VEGF$_{bb}$ in placental angiogenesis in normal human pregnancy. We speculate that placental oxidative environment might influence the C-terminal VEGF angiogenic switch, whereby different VEGF variants could be formed from a single VEGF gene; that play key roles in regulating angiogenic balance during normal human pregnancy. It may be suggested that gestational age-specific expression of both VEGF$_{165}$ and VEGF$_{bb}$ isoforms may be necessary for a successful viable outcome.

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Declaration of Interest

The authors declare no conflicts of interest with respect to the research, authorship and/or publication of this article.

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