

# Evaluation Gene Expression of Kinase Insert Domain (KDR) as Prediction of Preeclampsia with Novel Biomarkers

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### ABSTRACT

Preeclampsia (PE) is the complexity the case of pregnancy with an indication of the presence of defective placentas, and inadequate blood supply to the placenta defect and endothelial dysfunction .the aim of study. The variation between endothelial growth factor angiogenesis (VEGF), placental growth factor (PIGF) and their receptors kinase domain that contains the receptor (KDR) and tyrosine kinase 1 (Flt-1) plays a valuable role in the pathophysiology of PE .We conducted a cross-sectional study of 50 women with Preeclampsia: 25 women with mild PE, and 25 patients with severe PE. While 24 pregnant women work normally with age as a control group. placental mRNA gene expression of KDR was measured using real time PCR (RT/PCR),The mean of KDR expression was statistically significantly higher in the control group compared to any of the PE groups (p < 0.0005) and there was no statistically significant difference between sever PE and mild PE groups. The Information from different groups that reinforce KDR theory, useful in identifying an opportunity for ladies with Preeclampsia, to accurately diagnose Preeclampsia and anticipate progression of difficulties.

#### Keywords

Preeclampsia, KDR, endothelial dysfunction, placental ischemia.

#### 1. INTRODUCTION

Preeclampsia is a pregnancy related illness that occurs in 5-8% of pregnancy and is an important cause of disease and maternal mortality and fetus. The description of the disease symptoms of mothers that may occur from pregnancy for 20 weeks, high blood pressure> 140/90 mm Hg and proteinuria (> 300 mg / 24 hours) is also characterized by the Royal College of Obstetricians and Gynecologists [1]. The exact cause of Preeclampsia confused, but it is known that this infection incited by placentas factor assumes that oxidative stress may participate equally in dentistry [2]. It is recognized that the primary component of PE hinders the placenta due to lack of infiltration trophoblast in the arteries winding motherly [3]. Lead expansion is full of

the arteries of the placenta to the maternal lack of blood supply to the developing fetus [4]. It produces power ischemia and placental hypoxia [5].

A possible critical system of disease (PE) is a mismatch between the placenta proteins that generate the placenta and the antiangiogenic proteins. Proteins VGFC and PLGF are linked to the formation of blood guidelines for the growth of blood vessels choroid is expected to reduce maternal lining during pregnancy [6]. To regulate the vascular changes chorionic early, family can VEGF for vascular growth factors to do so. The intervention in improving VEGF systematically through the link receptors VEGFR-1, which are expressed in vascular endothelial cells in the trophoblast cells of the choroid in individual cells and large cells and KDR, which is surprising receptors that are expressed mainly in venous endothelial cells waiting for the exclusive development of new blood vessels [7].The current study aims to study the expression of KDR mRNA in the placenta of the breed vessels to link the degree of disruption of blood vessels with PE degree and gain further evidence of her job in Preeclampsia.

## 2. SUBJECTS AND METHODS

## 2.1 Subjects

Fifty patients determined to have PE were assembled from the Obstetrics and Gynecology division, Mansoura University, Egypt . Their mean age (years)  $\pm$  SD were 28.15 $\pm$ 5.25) during the period from July 2015 and May 2017. Preeclamptic women were partitioned into 25 women analyzed as mild preeclampsia (MPE), diagnosed as blood pressure  $\geq$ 140/90 mmHg and 25 women were analyzed as severe preeclampsia (SPE) ,diagnosed as blood pressure  $\geq$ 160/110 mmHg.The control group included 24 normotensive pregnant women . Exclusion criteria included pregnancy twins, women who have a background characterized by diabetes, liver or kidney disease or a continuous rise in the previous blood pressure, which has no end integration criteria: Normotensive pregnant women had blood pressure less than 120/80 mm Hg and there was no history of high blood pressure or proteinuria. Every pregnant woman referred to the gestational age of 20 to 40 weeks. Control group were taken from women who entered Mansoura University Hospital, Egypt, Obstetric department, Cohort study of mothers within two weeks of each case .Ethical approval: A composed educated assent was taken before test withdrawal. The examination was acted as per the moral guidelines set down in Mansoura Faculty of Medicine.

### 2.2 Methods

## 2.2.1. Urine Collection:

All pregnant women were asked for ten millilitres of urine for protein detection by dipstick in at least two random urine specimens obtained at least 4 h apart was required. proteinuria means protein  $\geq 3$  gr in a 24 hr (medi-test combi).

### 2.2.2. RNAextraction from placental tissue

Fresh samples of 74 placentas humanity within 30 min after placental delivery, After a wash of the samples with normal saline were performed, the amniotic layers and the maternal decidua were removed, then the samples were snap frozen in liquid nitrogen and stored at -70C until RNA isolation by using miRNeasy mini kit (Qiagen, cat no. 217004, Germany). RNA was quantified by spectrophotometry [8]. we measured the expression of KDR gene in placenta by used TaqMan real-time PCR ((provided by Thermo Scientific, USA, cat No. # K1641).

# 2.2.3. Produce DNA from RNA by Reverse Transcription

For each sample, was used 1µgof total RNA Adjust by reverse transcription using the Maxima® First Strand cDNAsynthess kit provided by Thermo Scientific, USA, cat. No.K1641. Following the manufacturer's instructions, Reverse transcription and the quantitative PCR were performed as previously described[9,10].

Genetic prefixes own purchased from Invitrogen by Thermo Fisher Scientific. It was selected primer sets to amplify the gene PCR after three of the gene sequence database NCBI [11].,It included the following assay targeting specific mRNAs in the study: KDR: forward 5-GTAGGACTGCTCAGTTCAAACAT-3, reverse 5- ACAGTTACTACACCCGTAAGG-3.

## 2.2.4. Quantitative PCR Analysis.

PCR was performed in real-time serial to estimate the quantification of genes VEGF and Flt1 and KDR using SensiFAST SYBR® Lo-ROX (purchased from Bioline, London, United Kingdom, Catalog Number: BIO-94005) following the manufacturer's instructions[12] .Analysis was performed fusion curve after amplification to confirm the product's privacy and exclude the presence Altmhedy- dimmers. Genetic expression was performed using the relative delta delta cycle threshold (DDCT)by using method  $2-\Delta\Delta ct[13-14]$ .

## 3. STATISTICAL ANALYSIS

Data analysis was performed by Statistical package for social science software (SPSS) version 25. Data are presented as mean  $\pm$  SD and compared by One-Way ANOVA and they are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Significance was considered at P<0.05 [31]. Data are presented as frequency (percentage) and compared by Chi-Square test (Monte Carlo significance).\*\*Data are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Pairwise comparisons are presented as capital letters (similar letters = no significant difference, while different letters = significant difference).

## 4. **RESULTS**

This study involved 74 subjects divided into 3 groups: The first is control group: 24 pregnantwomen with no current or previous PE. Their mean age (years)  $\pm$ SD= 27.8 $\pm$ 4.7 years. The second is mPE group: 25 pregnant ladies fulfilling the criteria of mPE. Their mean age was found to be 27.5 $\pm$ 4.3 years. The third is sPE group: 25 pregnant women fulfilling the criteria of sPE. Their mean age  $\pm$  SD = 28.8 $\pm$ 6.2 years. There was statistically significant difference in the level of protein in urine with trace and + in mild cases, but ++ and +++ in severe cases (P<0.0005)(Table 1).

|            | Group      |            |                   |         |
|------------|------------|------------|-------------------|---------|
| Variable   | Mild PE    | Severe PE  | Statistic         | P value |
|            | (n=25)     | (n=25)     |                   |         |
| Protein in |            |            |                   |         |
| urine:     | 1 (4%) a   | 0 (0%) a   |                   |         |
| Nil        | 13 (52%) a | 0 (0%) b   | 2                 | **<0.00 |
| Trace      | 11 (44%) a | 0 (0%) b   | $\chi^2 = 50.000$ | 05      |
| +          | 0 (0%) a   | 12 (48%) b |                   |         |
| ++         | 0 (0%) a   | 13 (52%) b |                   |         |
| +++        |            |            |                   |         |

Table(1):Comparison of proteinuria and previous Preeclampsia between mild and severe Preeclampsia.

\*\*Data are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Pairwise comparisons are presented as capital letters (similar letters = no significant difference, while different letters = significant difference). \*\*Data are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Pairwise comparisons are presented as capital letters (similar letters = no significant difference, while different letters = significant letters = significant difference).

KDR is the major mediator of migration, proliferation and differentiation of endothelial cells. Activation of KDR results in an expansion in both mitogenic cell signaling and migratory activity of endothelial cells[15]. The present study showed that, KDR expression was statistically significantly higher in control group as compared to either of the two PE groups (p<0.0005) and there is no statistically significantly difference between sPE and mPE (Table 2). Our studywas consistent with numerous studies which indicated that the KDR expression is low in pregnancies comlicated by PE than control[16-19, 24]. Other studies, have detailed additional levels of maternal plasma from sKDR during PE or early pregnancy while reducing intrauterine growth [20-22].

George et al., stated that the expression of KDR is low in PE women result from the decline in access to free VEGF to stimulate the synthesis of KDR and move away from the cell surfaces of endothelial cell surfaces, as seen in other models [23]. PE is characterized by the destroying of the motherly vessels that call for the failure of the placenta, which in turn are associated with high speed in the mother's blood flow to the chorionic villi, however, changes in the composition of vessels within the placenta in any case questionable [25]. It is expected to improve the increase in vessels, which were seen in the villi separated from the preeclamptic placenta, to facilitate the transfer of oxygen supplementation between the mother and the fetus [15].

| Variabl -<br>e | Group         |               |               |           |          |
|----------------|---------------|---------------|---------------|-----------|----------|
|                | Control       | Mild PE       | Severe PE     | Statistic | P value  |
| C              | (n=24)        | (n=25)        | (n=25)        |           |          |
| KDR            | $1.28\pm0.13$ | $0.61\pm0.23$ | $0.61\pm0.24$ | F =       | *<0.0005 |
|                | А             | В             | В             | 86.265    |          |

Table (2): Comparison of the KDR between the three study groups

Data are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Pairwise comparisons are presented as capital letters (similar letters = no significant difference, while different letters = significant difference). \*\*Data are presented as median (IQR) and compared by Kruskal-Wallis (K-W) test. Pairwise comparisons are presented as capital letters (There was no significant differences between means have the same letter, while different letters = significant difference).

Vessels exceed the development of the growth medium villi mature, blood and turn into circles and wrapped structure, which spread from the average surface villi mature, to form a new terminal villi [26].However, it has been assumed that a greater number of vessels in the villi, may reduce the diffusion of oxygen in the placenta, due to diffuse examination [15]. This decrease in oxygen conductivity is manifested by an increase in the placental hypoxia factor in PE [27]. These lines make sense to increase the placenta formation in the placenta before sagging, most likely not distinguishing the fetus. It is also plausible that increased vascular formation is not sufficient to overcome both the reduced oxygen conduction caused by the placenta infarction and the expected decrease in oxygen delivery while reshaping the maternal spiral arteries that provide the overlapping space [28]. In a few investigations there is no qualification in the expression of KDR mRNA between PE and controls [28-30].

The biomarker KDR with other angiogenic factors which may have ability to investigation and diagnosis of severe-early onset PE but couldn't be utilized alone for intervention, but in conjunction with other angiogenic factors like soluble endoglin, Doppler sonography and other clinical and biochemical biomarkers they are progressively important for predicting.

## 5. CONCLUSION

Overall, though, we found a significant decrease in KDR mRNA expression in the placenta for two preeclampsia groups compared to the Normotensive women's group, Further studies are needed to understand the molecular defect in preeclampsia and the role of KDR biomarker for diagnosis, prognosis and treatment of this obstetric condition.

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