

# VEGF levels in plasma in relation to metabolic control, inflammation, and microvascular complications in type-2 diabetes

## A cohort study

Qin Zhang, MMed, Wen Fang, MD, Li Ma, MD, Zhao-Di Wang, MD, Yun-Mei Yang, MD, Yuan-Qiang Lu, MD, PhD\*

### Abstract

The vascular endothelial growth factor (VEGF) level in human circulation may reflect the severity of endothelial dysfunction in patients with diabetes mellitus, which leads to diabetic microvascular complications.

We determined plasma VEGF levels as well as metabolic control and inflammatory factors in 26 healthy subjects and 52 type-2 diabetes mellitus (T2DM) patients with or without diabetic microvascular complications. Pearson correlation coefficient was used to evaluate the associations among those indices.

The results showed that VEGF levels in plasma were positively correlated with fasting blood glucose level, glycosylated hemoglobin (HbA1c) level, type 1 helper T cell (Th1) percentage, and Th1/Th2 ratio, while they were negatively correlated with regulatory T cell percentage. Multiple linear regression analysis showed that HbA1c and Th1/Th2 ratio were the independent predictors of VEGF levels in T2DM patients.

Thus, in T2DM patients with poor glycemic control as well as an elevated Th1/Th2 cell ratio, more VEGF might be released.

**Abbreviations:** BMI = body mass index, CRP = C-reactive protein, DN = diabetic nephropathy, DPN = diabetic peripheral neuropathy, DR = diabetic retinopathy, FBG = fasting blood glucose, HbA1c = glycosylated hemoglobin, Hcy = homocysteine, SD = standard deviation, T2DM = type-2 diabetes mellitus, Th = helper T, Treg = regulatory T, VEGF = vascular endothelial growth factor, WHR = waist-to-hip ratio.

**Keywords:** diabetic microvascular complication, T lymphocyte, type-2 diabetes mellitus, vascular endothelial growth factor

Individuals with type-2 diabetes mellitus (T2DM) are often at high risk for microvascular complications, including diabetic retinopathy (DR), diabetic nephropathy (DN), and diabetic peripheral neuropathy (DPN) due to diabetic microvascular dysfunction.<sup>[1,2]</sup> Angiogenesis, as an essential biological process, involves the progression of diabetic microvascular complications. Meanwhile, vascular endothelial growth factor (VEGF) is the most potent proangiogenic growth factor that increases vascular permeability *in vivo* and activates endothelial cells *in vitro*.<sup>[3]</sup>

Taken together, VEGF may play an important role in diabetic endothelial dysfunction, which leads to diabetic microvascular complications. Previous studies have found that VEGF is involved in the pathogenesis of diabetic complications.<sup>[4-7]</sup> Plasma VEGF levels were reported to be higher in diabetic patients than in healthy control individuals, and a correlation of plasma VEGF levels with proliferative DR and DN has also been noticed.<sup>[8,9]</sup> The synthesis and secretion of VEGF are affected by several factors, including gender, hypoxia, hyperglycemia, smoking, blood lipids, inflammatory reaction, and activated stress axes.<sup>[10]</sup> However, there are few research studies that illustrate the detailed mechanism of the association between VEGF and diabetic microangiopathy, particularly in T2DM.<sup>[11]</sup> To better understand it, T2DM patients with or without diabetic microvascular complications and healthy volunteers were selected, and their VEGF plasma levels as well as other clinical parameters were assessed in this study. We investigated the relationship between plasma VEGF levels and parameters of metabolic control, inflammation, and the presence of diabetic microvascular complications.

Editor: Gaurav Malhotra.

This work was supported in part by grants from the National Key Clinical Specialist Construction Program of China (geriatric medicine) and the Foundation of Key Discipline Construction of Zhejiang Province for Traditional Chinese Medicine (grant number 2012-XK-A20).

The authors have no conflicts of interest to disclose.

Department of Emergency and Geriatrics Medicine, The First Affiliated Hospital, School of Medicine, Zhejiang University, Hangzhou, Zhejiang, China.

\* Correspondence: Yuan-Qiang Lu, The First Affiliated Hospital, School of Medicine, Zhejiang University, Hangzhou, Zhejiang, China (e-mail: luyuanqiang@zju.edu.cn).

Copyright © 2018 the Author(s). Published by Wolters Kluwer Health, Inc.

This is an open access article distributed under the Creative Commons Attribution License 4.0 (CCBY), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Medicine (2018) 97:15(e0415)

Received: 6 June 2017 / Received in final form: 14 March 2018 / Accepted: 21 March 2018

<http://dx.doi.org/10.1097/MD.00000000000010415>

## 1. Methods

### 1.1. Subjects

The study included 26 healthy volunteers as a control group (category 1, n = 26) and 52 patients diagnosed with T2DM under the care of the First Affiliated Hospital, School of Medicine, Zhejiang University, China between October 2015 and Decem-

ber 2016. The T2DM patients were divided into 2 categories: T2DM without microvascular complications (category 2,  $n=26$ ) and T2DM with microvascular complications (category 3,  $n=26$ ). Twenty-six individuals were selected in category 3, including 1 case of DR, 7 cases of DN, 11 cases of DPN, 3 cases of DR combined with DPN, 2 cases of DR combined with DN, and 2 cases of DPN combined with DN. To minimize the statistical error caused by small samples, we put all T2DM patients with various diabetes microvascular complications into category 3.

To qualify for the study, patients had to satisfy the following criteria: a level of glycosylated hemoglobin (HbA1c) of  $\geq 6.5\%$  and fasting blood glucose (FBG)  $\geq 7.0$  mmol/L, or with a glucose tolerance test, 2 h after the oral dose, a plasma glucose level  $\geq 11.1$  mmol/L. Patients meeting any of the above criteria can be categorized as T2DM. The control category selected healthy participants without any type of diabetes, as well as without hypertension, hyperlipidemia, and other metabolic syndromes. DR was diagnosed according to the Clinical Guidelines of Diabetic Retinopathy in China (2014).<sup>[12]</sup> DN was diagnosed when albuminuria  $> 300$  mg/24h, or glomerular filtration rate  $< 60$  mL/min lasting for 3 months. DPN was diagnosed with the following symptoms: abnormal temperature sense, foot sense loss detected by nylon yarn, decreased vibration perception, disappeared ankle reflexes, and nerve conduction tests showing reduced functioning of the peripheral nerves. Patients matched 2 or more of the above clinical peripheral neuropathy and electrophysiological symptoms can be defined as DPN. T2DM patients who had serious medical comorbidities, such as macroangiopathy, acute inflammation, autoimmune diseases, endocrine diseases, malignancies, hematological diseases, unstable angina or myocardial infarction, end-stage cardiac insufficiency, cerebral infarction, and pulmonary or hepatic diseases, were excluded from this study.

This study was approved by the Medical Ethical Committee of First Affiliated Hospital, School of Medicine, Zhejiang University, and all participants gave written informed consent. The methods in this study were performed in accordance with the relevant guidelines and regulations.

### 1.2. Measurements of plasma VEGF, metabolic and inflammatory parameters

From each patient or healthy volunteer, the 10 mL of venous blood was collected from an elbow vein between 6:30 and 7:30 AM in a fasting state. Blood samples were kept in 3 types of tubes. Six milliliters of blood was collected in a tube containing ethylenediaminetetraacetic acid to determine the plasma VEGF level and lymphocyte cell counts in peripheral blood. The blood sample was centrifuged for 10 min at 400g to obtain plasma. The plasma VEGF concentration was determined using human VEGF-A platinum enzyme-linked immunosorbent assay kit (eBioscience, Vienna, Austria). The remaining peripheral blood mononuclear cells were collected after sequentially adding RBC Lysis Buffer and Ficoll-Paque PLUS (Sigma-Aldrich, St Louis, MO) to remove the red blood cells and granulocytes for further lymphocyte cell analysis. A flow cytometer (FC500, FACSCalibur; Beckman-Coulter, Brea, CA) was used to detect regulatory T cells (Treg: CD4+CD25+FoxP3+), type 1 helper T cells (Th1: CD4+T-bet+), and type 2 helper T cells (Th2: CD4+GATA3+), and the ratio of Th1/Th2 cells was calculated. Th1 and Th2 antibodies used for flow cytometry were purchased from BD Biosciences (Franklin Lakes, NJ); Tregs antibodies were purchased from eBioscience, Inc. (San Diego, CA). In addition,

2 mL of blood was drawn into a tube containing sodium fluoride to determine the FBG and HbA1c. Another 2 mL of blood was collected in a serum tube, and the serum C-reactive protein (CRP) and homocysteine (Hcy) levels were then analyzed.

Clinically, FBG and HbA1c are the most widely used parameters for glycemic control. Hcy is the marker of amino acid metabolism disorder and has been demonstrated to be an independent risk factor for cardiovascular and cerebrovascular diseases.<sup>[13,14]</sup> The CRP, Tregs, and Th1/Th2 ratios are the common clinical indicators for immune and inflammatory responses.

### 1.3. Statistical analysis

All statistical analyses were performed using SPSS version 18.0 software (SPSS Inc., Chicago, IL). The categorical variables are expressed as numbers or percentages, and the continuous variables that were close to a normal distribution are presented as the mean  $\pm$  standard deviation. The chi-squared test was applied for comparisons of categorical variables. For continuous variables, the 1-way analysis of variance with the least significant difference *t* test was applied for comparisons differences among the 3 categories. Pearson correlation coefficients were calculated between the VEGF level and clinical indicators of metabolic control, and markers of inflammation. Forward multiple linear regression analysis was performed with VEGF levels as the dependent variable and those determinants that correlated in the univariate analysis with  $P < .05$  as independent variables. For multiple comparisons,  $P < .05$  was considered significant.

## 2. Results

### 2.1. General clinical data

Data were obtained from 78 participants (41 males and 37 females). The demographic and clinical parameters of 3 categories were recorded: gender, age, diabetes duration, body mass index (BMI), waist-to-hip ratio (WHR), comorbidity, and smoking habits (Table 1). No differences in sex ratio, age, or rate of smokers were found among the 3 categories (all  $P > .05$ ). The diabetes duration in category 3 was longer than that in category 2 ( $P < .05$ ). The WHRs were higher in the T2DM categories, but no difference was found between categories 2 and 3 ( $P > .05$ ). BMI was elevated significantly in category 3 compared with category 1 ( $P < .05$ ). The percentages of hypertensive patients among the 3 categories were significantly different ( $P < .001$ ), but no difference was found between categories 2 and 3 ( $P > .05$ ).

### 2.2. Metabolic and inflammatory parameters

The metabolic and inflammatory parameters of the 3 categories are summarized in Table 2. The indicators of glycemic control (FBG and HbA1c) in the T2DM categories were significantly higher than those in category 1 (all  $P < .05$ ), but no difference was found between the 2 T2DM categories (all  $P > .05$ ). In addition, the Hcy concentration was elevated significantly in category 3 compared with that in the other 2 categories (both  $P < .05$ ). As markers of inflammation, Th1 percentages and Th1/Th2 ratio were significantly higher, as well as CD4+CD25+FoxP3+Treg percentages being lower in the 2 T2DM categories than in category 1 (all  $P < .05$ ). In particular, the Th1 percentages and Th1/Th2 ratio in category 3 were even higher than in category 2 (all  $P < .05$ ). There were no marked differences in Th2 percentages among all

**Table 1****Demographic and clinical parameters according to category.**

Parameters	Type 2 DM			F or $\chi^2$	P
	Healthy control subjects	Without microvascular complications	With microvascular complications		
	Category 1	Category 2	Category 3		
N (males)	26 (13)	26 (12)	26 (16)	0.726	.696
Age, y	57.6±7.1	57.8±13.3	60.9±9.2	0.084	.432
Diabetes duration, y	0	5.6±4.0*	10.3±6.8*†	33.144	<.001
BMI, kg/m <sup>2</sup>	22.79±2.92	24.32±3.06	25.22±3.26*	4.417	.020
WHR	0.84±0.07	0.93±0.06*	0.94±0.07*	19.018	<.001
No. of smokers (%)	4 (15.38)	6 (23.08)	7 (26.92)	1.053	.591
No. of hypertensive patients (%)	0 (0)	10 (38.46)	18 (69.23)	27.189	<.001

BMI=body mass index, DM=diabetes mellitus, WHR=waist-to-hip ratio.

\* P &lt; .05 compared with category 1.

† P &lt; .05 compared with category 2.

categories ( $P > .05$ ). The CRP and plasma VEGF levels in the 2 T2DM categories were obviously higher than those in category 1; meanwhile, CRP and VEGF levels in category 3 were also higher than those in category 2 (all  $P < .05$ ).

### 2.3. Relationships between VEGF level and its putative determinants

Pearson correlation coefficients were calculated to analyze the associations between plasma VEGF level and clinical indicators of metabolic control or inflammation in healthy subjects and T2DM patients. Figure 1 demonstrated that VEGF levels were correlated positively with FBG level ( $r=0.483$ ,  $P < .001$ ), HbA1c level ( $r=0.531$ ,  $P < .001$ ), Th1 percentage ( $r=0.366$ ,  $P=.001$ ), and Th1/Th2 ratio ( $r=0.373$ ,  $P=.001$ ) in the T2DM patients and controls. By contrast, a negative correlation between the VEGF level and CD4+CD25+FoxP3+Treg percentage was also found. Multiple linear regression analysis indicated that only the HbA1c level and Th1/Th2 ratio were independent predictors of VEGF levels in T2DM. The regression equation of VEGF levels was  $VEGF = -5.275 + 4.099 \times \text{HbA1c} + 6.471 \times \text{Th1/Th2 ratio}$  ( $r^2 = 0.332$ ;  $P < .001$ ). The residual sum of the squares indicated that 33.2% of variation in VEGF levels was explained by the HbA1c level and Th1/Th2 ratio.

### 3. Discussion

In this study of people with and without T2DM, we found that VEGF levels in plasma were positively correlated with glycemic control indicators (FBG and HbA1c), and inflammatory parameters (Th1, Th1/Th2 ratio), while they were negatively correlated with Treg percentage; further multiple linear regression analysis revealed that HbA1c and Th1/Th2 ratio were the independent predictors of VEGF levels in plasma of T2DM patients. This indicates a close association among hyperglycemia, inflammation, and VEGF in T2DM patients.

The number of diabetes cases worldwide is approximately 366 million according to the estimation by the International Diabetes Federation, of which T2DM constitutes approximately 90% to 95%.<sup>[15]</sup> Up to 80% of mortality with diabetes is directly associated with vascular diseases affecting micro- or macro-circulation.<sup>[16]</sup> Angiogenesis, about the formation and differentiation of blood vessels, is an essential biological process existing in embryogenesis and in the development of major diseases, such as cancer, inflammation, and diabetes.<sup>[17]</sup> The fundamental regulator most widely known to be involved in angiogenesis is VEGF. VEGF is also associated with tumor progression and poor outcomes in various human cancers.<sup>[18,19]</sup> The activated platelets and leukocytes are the main sources of VEGF in blood.<sup>[20]</sup> In cultured endothelial cells, VEGF has been proven to be induced

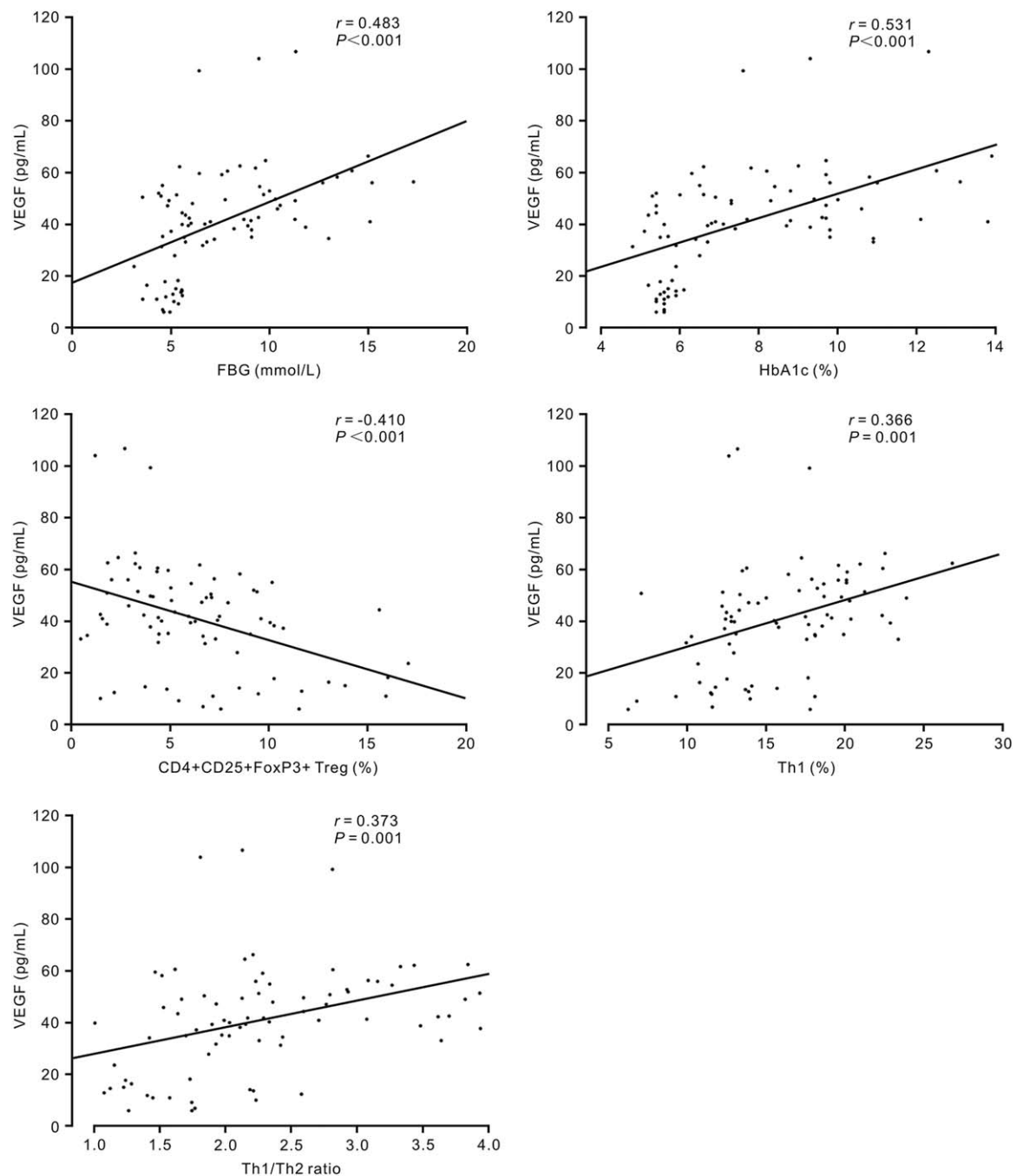
**Table 2****Metabolic and inflammatory parameters according to category.**

Parameters	Type 2 DM			F	P
	Healthy control subjects	Without microvascular complications	With microvascular complications		
	Category 1 (n=26)	Category 2 (n=26)	Category 3 (n=26)		
FBG, mmol/L	4.87±0.68	8.23±3.05*	9.51±3.21†	22.272	<.001
HbA1c, %	5.55±0.27	8.49±2.28*	9.03±2.09*	28.223	<.001
Hcy, $\mu\text{mol/L}$	13.78±7.62	15.39±3.54	18.88±4.58*†	5.795	.005
CD4+CD25+FoxP3+ Treg, %	2.52±1.30	1.81±1.10*	1.41±0.64*	7.361	.001
Th1, %	12.92±3.33	16.20±3.93*	18.52±3.56*†	15.715	<.001
Th2, %	7.85±2.88	7.51±1.66	6.97±1.59	1.152	.322
Th1/Th2 ratio	1.79±0.58	2.21±0.59*	2.78±0.75*†	15.210	<.001
CRP, mg/L	1.98±1.35	3.49±2.57*	5.42±2.82*†	14.190	<.001
VEGF, pg/mL	21.70±14.75	45.38±15.45*	55.48±16.13*†	32.787	<.001

CRP = C-reactive protein, DM = diabetes mellitus, FBG = fasting blood glucose, HbA1c = glycated hemoglobin, Hcy = homocysteine, Th = helper T, Treg = regulatory T, VEGF = vascular endothelial growth factor.

\* P &lt; .05 compared with category 1.

† P &lt; .05 compared with category 2.



**Figure 1.** Relationship between VEGF levels in plasma and its putative determinants in healthy subjects and T2DM patients. FBG = fasting blood glucose, HbA1c = glycated hemoglobin, T2DM = type-2 diabetes mellitus, Th = helper T, Treg = regulatory T, VEGF = vascular endothelial growth factor.

by the elevated levels of glucose and advanced glycation end products.<sup>[11]</sup> It is interesting that the correlations of VEGF with blood glucose concentrations and immune levels are both found in our cohort study, suggesting an interaction may exist among hyperglycemia, inflammation, and VEGF.

Chronic hyperglycemia has been reported to stimulate the synthesis and secretion of VEGF-A. It triggers a chain reaction that contributes to VEGF-A accumulation and then leads to DM microvascular complications.<sup>[21]</sup> The major physiological stimulus for VEGF production is cellular hypoxia. Hyperglycemia can act as toxin to the endothelium through increasing oxidative stress. The high concentration of blood glucose increases the

production of vasoconstrictor substances, particularly endothelin-1.<sup>[22]</sup> Hyperglycemia-induced pathological mechanism affects the expression of VEGF and its receptors VEGFR1 and VEGFR2. The elevated circulating VEGF-A levels are already found in adult T1DM patients with DN, and in T1DM prepubertal and pubertal children.<sup>[8,23]</sup> Similarly, urinary VEGF-A was elevated in T2DM patients and the diabetic mice model holding no correlation with their albuminuria.<sup>[24,25]</sup> VEGF-A polymorphisms are associated with DR and DN as well.<sup>[21]</sup> The characteristic parameter to evaluate glucose control in the blood is the level of HbA1c. Mahdy et al<sup>[7]</sup> measured the serum VEGF level in T2DM patients before glycemic control and at 4 months follow-up and observed

a significant decrease in the serum level of VEGF in patients with glycemic control.<sup>[7]</sup> These previous research studies are in line with the results from our study, which show that a significant correlation exists between the concentrations of VEGF and glycemic control.

Hcy has been reported to inhibit endothelial cell proliferation and induce endothelial dysfunction as well as endothelial cell apoptosis.<sup>[26–28]</sup> The serum Hcy concentration of T2DM with microvascular complications group was elevated significantly, compared with the 2 other groups in our research. These results suggest that there may be more serious endothelial damage and metabolic disorders in the T2DM patients with microvascular complications.

In our study, CRP, Th1 percentage, and Th1/Th2 ratio in the T2DM patients with microvascular complications were significantly higher than those in the control and T2DM without microvascular complications. This indicated that persistent inflammatory activity was involved in the progression of microvascular complications in diabetes. Inflammatory reaction can lead to an increase in vascular permeability, endothelial cell apoptosis, and chronic inflammation. Actually, CRP as common clinical indicator of inflammatory status could upregulate the VEGF-A expression by activating hypoxia inducible factor-1 $\alpha$  in adipose-derived stem cells.<sup>[29,30]</sup> Furthermore, disruption in immune homeostasis with a shift toward a Th2-dominant or chronic inflammatory state by tumor-derived VEGF has been reported previously.<sup>[31]</sup> By contrast, we noticed that the Th1/Th2 immune imbalance with a shift to a Th1-dominant was associated with plasma VEGF accumulation in T2DM. The Th1/Th2 ratio has switched directions in diabetes possibly due to the diverse immune-related cytokines activated in these diseases and then triggered the proliferation of different helpers T cells. Treg is an indicator of immune response, which has the potent immunosuppressive function to maintain immune homeostasis.<sup>[32,33]</sup> VEGF is proved to be a promoter of Treg activation in antitumor immunity. Conversely, a negative correlation between the plasma VEGF level and Treg percentage in T2DM was noticed in our study, although further analysis demonstrated that the Treg concentration was not an independent predictor of VEGF levels in T2DM. The above results showed that the inflammation levels of T2DM patients were higher than those of the healthy control group. Particularly in T2DM patients with microvascular complications, there may exist a more serious immune dysfunction.

In next multivariate analysis, only HbA1c and Th1/Th2 ratios were found to be independent determinants of the VEGF plasma level. The independent correlation between VEGF and HbA1c has been reported only in T1DM.<sup>[11]</sup> Combined with our data, this observation suggests that in both T1DM and T2DM, poor glycemic control possibly leads to more VEGF released. We can also speculate that the possibility of higher release of VEGF in patients with poor glycemic control and persistent inflammation activity could be explored as a potential contributor to endothelial dysfunction in diabetic patients.

The limitations of this study are that the population of T2DM with microvascular complications patients is not large enough to be subdivided based on different types of complications. A large population can minimize statistical variance. Next step, we will verify if the elevated VEGF levels could be reversed after controlling the diabetic states of T2DM patients.

In conclusion, our study showed a close association among hyperglycemia, inflammation, and VEGF, which link with microvascular diseases in T2DM patients. The elevated circulat-

ing levels of Th1 and VEGF may contribute to the pathogenesis of T2DM microangiopathy.

## Author contributions

**Conceptualization:** Yuan-Qiang Lu.

**Formal analysis:** Qin Zhang, Li Ma.

**Funding acquisition:** Yun-Mei Yang, Yuan-Qiang Lu.

**Investigation:** Qin Zhang, Zhao-Di Wang, Yun-Mei Yang.

**Methodology:** Qin Zhang, Zhao-Di Wang, Yun-Mei Yang.

**Project administration:** Yuan-Qiang Lu.

**Software:** Li Ma.

**Supervision:** Yuan-Qiang Lu.

**Writing – original draft:** Wen Fang, Yuan-Qiang Lu.

**Writing – review & editing:** Qin Zhang, Wen Fang, Yuan-Qiang Lu.

## References

- World Health Organization. Global Status Report on Noncommunicable Diseases; 2014. Available at: <http://www.who.int/nmh/>.
- DeFronzo RA, Ferrannini E, Groop L, et al. Type 2 diabetes mellitus. *Nat Rev Dis Primers* 2015;1:15019. doi: 10.1038/nrdp.2015.19.
- Asselbergs FW, de Boer RA, Diercks GF, et al. Vascular endothelial growth factor: the link between cardiovascular risk factors and microalbuminuria? *Int J Cardiol* 2004;93:211–5.
- Aiello LP, Wong JS. Role of vascular endothelial growth factor in diabetic vascular complications. *Kidney Int Suppl* 2000;77:S13–9.
- Watanabe T. Is vascular endothelial cell growth factor (VEGF) involved in the pathogenesis of diabetic nephropathy? *Nephrology (Carlton)* 2007;12(suppl 3):S27.
- Xie XJ, Yang YM, Jiang JK, et al. Association between the vascular endothelial growth factor single nucleotide polymorphisms and diabetic retinopathy risk: a meta-analysis. *J Diabetes* 2017;9:738–53.
- Mahdy RA, Nada WM, Hadhoud KM, et al. The role of vascular endothelial growth factor in the progression of diabetic vascular complications. *Eye* 2010;24:1576–84.
- Hovind P, Tarnow L, Oestergaard PB, et al. Elevated vascular endothelial growth factor in type 1 diabetic patients with diabetic nephropathy. *Kidney Int Suppl* 2000;75:S56–61.
- Ma Y, Zhang Y, Zhao T, et al. Vascular endothelial growth factor in plasma and vitreous fluid of patients with proliferative diabetic retinopathy patients after intravitreal injection of bevacizumab. *Am J Ophthalmol* 2012;153:307–13.
- Guo L, Jiang F, Tang YT, et al. The association of serum vascular endothelial growth factor and ferritin in diabetic microvascular disease. *Diabetes Technol Ther* 2014;16:224–34.
- Schlingemann RO, Van Noorden CJ, Diekman MJ, et al. VEGF levels in plasma in relation to platelet activation, glycemic control, and microvascular complications in type 1 diabetes. *Diabetes Care* 2013;36:1629–34.
- ETDRS Research Group. Classification of diabetic retinopathy from fluorescein angiograms. Early treatment diabetic retinopathy study report number 11. *Ophthalmology* 1991;98:807–22.
- Jung JM, Kwon DY, Han C, et al. Increased carotid intima-media thickness and plasma homocysteine levels predict cardiovascular and all-cause death: a population-based cohort study. *Eur Neurol* 2013;70:1–5.
- Hooshmand B, Polvikoski T, Kivipelto M, et al. Plasma homocysteine, Alzheimer and cerebrovascular pathology: a population-based autopsy study. *Brain* 2013;136:2707–16.
- Whiting DR, Guariguata L, Weil C, et al. IDF diabetes atlas: global estimates of the prevalence of diabetes for 2011 and 2030. *Diabetes Res Clin Pract* 2011;94:311–21.
- Silvestre JS, Levy BI. Molecular basis of angiopathy in diabetes mellitus. *Circ Res* 2006;98:4–6.
- Shibuya M. Structure and function of VEGF/VEGF-receptor system involved in angiogenesis. *Cell Struct Funct* 2001;26:25–35.
- Salven P, Teerenhovi L, Joensuu H. A high pretreatment serum vascular endothelial growth factor concentration is associated with poor outcome in non-Hodgkin's lymphoma. *Blood* 1997;90:3167–72.
- Brychtova S, Bezdekova M, Brychta T, et al. The role of vascular endothelial growth factors and their receptors in malignant melanomas. *Neoplasma* 2008;55:273–9.

- [20] Gunsilius E, Petzer A, Stockhammer G, et al. Thrombocytes are the major source for soluble vascular endothelial growth factor in peripheral blood. *Oncology* 2000;58:169–74.
- [21] Tufro A, Veron D. VEGF and podocytes in diabetic nephropathy. *Semin Nephrol* 2012;32:385–93.
- [22] Ruzkowska-Ciastek B, Sokup A, Socha MW, et al. A preliminary evaluation of VEGF-A, VEGFR1 and VEGFR2 in patients with well-controlled type 2 diabetes mellitus. *J Zhejiang Univ Sci B* 2014;15:575–81.
- [23] Chiarelli F, Spagnoli A, Basciani F, et al. Vascular endothelial growth factor (VEGF) in children, adolescents and young adults with type 1 diabetes mellitus: relation to glycaemic control and microvascular complications. *Diabet Med* 2000;17:650–6.
- [24] Veron D, Bertuccio CA, Marlier A, et al. Podocyte vascular endothelial growth factor (VEGF(1)(6)(4)) overexpression causes severe nodular glomerulosclerosis in a mouse model of type 1 diabetes. *Diabetologia* 2011;54:1227–41.
- [25] Kim NH, Kim KB, Kim DL, et al. Plasma and urinary vascular endothelial growth factor and diabetic nephropathy in type 2 diabetes mellitus. *Diabet Med* 2004;21:545–51.
- [26] Nagai Y, Tasaki H, Takatsu H, et al. Homocysteine inhibits angiogenesis in vitro and in vivo. *Biochem Biophys Res Commun* 2001;281:726–31.
- [27] Lee SJ, Kim KM, Namkoong S, et al. Nitric oxide inhibition of homocysteine-induced human endothelial cell apoptosis by down-regulation of p53-dependent Noxa expression through the formation of S-nitrosohomocysteine. *J Biol Chem* 2005;280:5781–8.
- [28] Oosterbaan AM, Steegers EA, Ursem NT. The effects of homocysteine and folic acid on angiogenesis and VEGF expression during chicken vascular development. *Microvasc Res* 2012;83:98–104.
- [29] Chen J, Gu Z, Wu M, et al. C-reactive protein can upregulate VEGF expression to promote ADSC-induced angiogenesis by activating HIF-1alpha via CD64/PI3k/Akt and MAPK/ERK signaling pathways. *Stem Cell Res Ther* 2016;7:114.
- [30] Tian Y, Li JL, Hao L, et al. Association of cytokines, high sensitive C-reactive protein, VEGF and beta-defensin-1 gene polymorphisms and their protein expressions with chronic periodontitis in the Chinese population. *Int J Biol Markers* 2013;28:100–7.
- [31] Agostino NM, Saraceni C, Kincaid H, et al. A prospective evaluation of the role of vascular endothelial growth factor (VEGF) and the immune system in stage III/IV melanoma. *Springerplus* 2015;4:186.
- [32] Zhang Q, Lu YQ, Jiang JK, et al. Early changes of CD4(+)CD25(+) Foxp3(+) regulatory T cells and Th1/Th2, Tc1/Tc2 profiles in the peripheral blood of rats with controlled hemorrhagic shock and no fluid resuscitation. *Chin Med J (Engl)* 2012;125:2163–7.
- [33] Lu YQ, Gu LH, Zhang Q, et al. Hypertonic saline resuscitation contributes to early accumulation of circulating myeloid-derived suppressor cells in a rat model of hemorrhagic shock. *Chin Med J (Engl)* 2013;126:1317–22.