Evaluating the Serum Transforming Growth Factor-Beta 1 Level in Chronic Kidney Disease Caused by Glomerulonephritis

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Abstract

Background: The transforming growth factor-beta 1 (TGF-β1) has been demonstrated as one of the main factors in the progression of fibrosis and sclerosis glomerular damages. Glomerulonephritis is one common cause of chronic kidney disease (CKD) with the promotion of inflammatory renal damage containing fibrosis and sclerosis glomerular.

Objectives: This study aimed to evaluate the TGF-β1 level in CKD patients and compare it with the healthy control group.

Methods: This cross-sectional case-control study was carried out on 212 subjects admitted to the Nghe An Friendship General Hospital in Vietnam from March 2018 to February 2020. The case group included 152 patients diagnosed with CKD caused by glomerulonephritis, and the control group included 60 healthy individuals. The TGF-β1 was determined in serum by ELISA method.

Results: The serum TGF-β1 concentration of the healthy control group and CKD group was 13.45 ± 7.17 and 32.35 ± 11.74, respectively. The CKD group had a significantly higher TGF-β1 level than the control group (P < 0.05). The CKD group with the eGFR ≥ 60 mL/min/1.73 m² group had a higher TGF-β1 level than the eGFR < 60 mL/min/1.73 m² group, and the TGF-β1 level increased from stage 1 to stage 5 (P < 0.001). The TGF-β1 had a medium correlation to urea, creatinine, and hs-CRP.

Conclusions: The concentration of TGF-β1 in the CKD group was higher than the control group so that it increased early from the first stage of the disease.

Keywords: Inflammatory, Fibrosis, Sclerosis, Glomerular

1. Background

The transforming growth factor-beta 1 (TGF-β1) is essential for normal development, repair of the organization, and maintenance of the organ function. However, an increase in the concentration of TGF-β1 causes pathological changes in chronic kidney disease (CKD). In human glomerulonephritis, such as focal segmental glomerulosclerosis (FSGS), IgA glomerulonephritis, sickle-shaped glomerulonephritis, lupus glomerulonephritis, and diabetic nephropathy, TGF-β1 plays an important role in contributing glomerular sclerosis (1-4). The main feature of these pathologies is the excessive accumulation of extracellular material and the increased effect of TGF-β1 in the glomerulus and interstitial nephropathy. In the glomerulus, TGF-β1 contributes mainly to the transformation of the glomerular membrane, fibrosis, and glomerular sclerosis. In the renal tubules, TGF-β1 is involved both directly and indirectly in tubular degeneration (5). Several studies have demonstrated the increase of TGF-β1 in CKD patients (6, 7), leading to a new therapy in the prevention and treatment of CKD (8). This study aimed to evaluate the serum level of TGF-β1 in CKD patients and the related changes during other stages of the disease.

2. Objectives

This study aimed to evaluate the TGF-β1 level in CKD patients and compare it with the healthy control group.

3. Methods

This cross-sectional case-control study was carried out on 212 subjects admitted to the Nghe An Friendship General Hospital in Vietnam from March 2018 to February 2020. The case group included 152 patients diagnosed with CKD caused by glomerulonephritis, and the control group
included 60 healthy individuals. The case group was divided into two small groups based on eGFR value (the CKD group with the eGRP ≥ 60 mL/min/1.73 m² and the CKD group with eGRP < 60 mL/min/1.73 m²).

The inclusion criteria were: age ≥ 18; having one of the following symptoms: (1) markers of kidney damage such as albuminuria (ACR ≥ 30 mg/g), or urine sediment abnormalities, or electrolyte and others abnormalities in tubular disorders, or abnormalities detected in histology, or structural abnormalities detected in imaging, or edema, or hypertension; and (2) decreased estimated glomerular filtration rate (eGFR) < 60 mL/min/1.73 m².

Exclusion criteria were: (1) pregnancy; (2) transplantation; (3) diabetes mellitus (DM); (4) hypertension (HTN); (5) CKD caused by HTN or DM; (6) acute period of chronic kidney failure; (7) acute kidney failure; (8) patients who had heart diseases before CKD; (9) patients who had a fever during sampling phase; and (10) patients using statin, corticoid, and the drugs which increased creatinine clearance in healthy controls compared to CKD with eGFR ≥ 60 mL/min/1.73 m² group and the CKD with eGFR < 60 mL/min/1.73 m² group. The plasma urea and creatinine were significantly higher in healthy controls compared to CKD with eGFR ≥ 60 mL/min/1.73 m² group and the CKD with eGFR < 60 mL/min/1.73 m² group. The plasma TGF-β1 levels had a negative correlation with plasma urea, creatinine, and hs-CRP.

### 3. Samples

The plasma of all subjects were collected to determine the concentration of urea and creatinine in the AU system (Beckman Coulter). The serum was used to qualify TGF-β1 in serum. First, the subjects of this study were concentrated by double-antibody sandwich enzyme-linked immunosorbent assay (ELISA) with the reagent of DRG manufacturer (American, ELA1864) in the Evolis Twin Plus automation machine (American). BMI was calculated as the ratio of the weight to the height (kg/m²).

### 3.1. Samples

The plasma of all subjects were collected to determine the concentration of urea and creatinine in the AU system (Beckman Coulter). The serum was used to qualify TGF-β1. The EDTA blood was collected to determine red blood cells, hematocrit, and hemoglobin. The clinical characteristics such as height, weight, body mass index (BMI), and waist circumference were also collected. The eGFR was calculated using the CKD-EPI creatinine equation 2009, recommended by the National Kidney Foundation (NKF).

In the equation, the unit of eGFR (estimated glomerular filtration rate) was mL/min/1.73 m²; the unit of Scr (standardized serum creatinine) was mg/dL; α = 0.7 (females) or 0.9 (males); β = -0.329 (females) or -0.411 (males); min, the minimum of Scr/κ; or 1; max, the maximum of Scr/κ; or 1; age, years.

TGF-β1 was concentrated by double-antibody sandwich enzyme-linked immunosorbent assay (ELISA) with the reagent of DRG manufacturer (American, ELA1864) in the Evolis Twin Plus automation machine (American). BMI was calculated as the ratio of the weight to the height (kg/m²).

### 3.2. Statistical Analysis

The data were analyzed by Excel 2013 and SPSS software version 18.0. The non-normal quantitative variables were presented as median and quartiles (25th and 75th). The quantitative variables with normal distribution were presented as mean ± standard deviation, maximum, minimum. The difference between the means of the groups was calculated by the ANOVA test. The results were considered to be significant at P < 0.05.

### 4. Results

#### 4.1. Clinical Characteristic of the Subjects

As Table 1 shows, there were no significant differences between the healthy control group, CKD with eGFR ≥ 60 mL/min/1.73 m² group, and CKD with eGFR < 60 mL/min/1.73 m² group in age, gender, height, weight, waist circumference, and BMI. The RBC, hematocrit, and hemoglobin were significantly lower in healthy controls compared to CKD with eGFR ≥ 60 mL/min/1.73 m² group and the CKD with eGFR < 60 mL/min/1.73 m² group. The plasma TGF-β1 level increased from stage 1 to 5 of CKD (P < 0.001) (Table 2). In CKD group, the eGFR < 60 mL/min/1.73 m² group had higher TGF-β1 level than the eGFR ≥ 60 mL/min/1.73 m² group (P < 0.001) (Table 3). The TGF-β1 level increased from stage 1 to 5 of CKD (P < 0.001) (Table 4).

#### 4.2. The TGF-β1 Levels

The TGF-β1 levels in the CKD group were higher than the control (P < 0.001) (Table 2). In CKD group, the eGFR < 60 mL/min/1.73 m² had higher TGF-β1 level than the eGFR ≥ 60 mL/min/1.73 m² group (P < 0.001) (Table 3). The TGF-β1 level increased from stage 1 to 5 of CKD (P < 0.001) (Table 4).

#### 4.3. The Correlation Between TGF-β1 Levels and Other Characteristics

While the TGF-β1 levels had a negative correlation with eGFR, hematocrit, red blood cell, and hemoglobin, it had a positive correlation with plasma urea, creatinine, and hs-CRP.

### 5. Discussion

The TGF-β1 has been considered as one of the main cytokines in the pathogenesis of renal inflammation and fibrosis (9). Some pathogenic evidence showed that the inflammation progressed silently and almost unrecovered in the CKD even during the early stage (10). TGF-β1 has been found to be increasing in CKD patients than in healthy individuals. Our results were similar to the reports of some other studies (11-15). Three main mechanisms increase the TGF-β1 in serum. First, the subjects of this study were CKD patients caused by glomerulonephritis, and the patients with CKD caused by other chronic diseases such as
\[
eGFR = 141 \times \min \left( \frac{SCr}{K}, 1 \right) \alpha \times \max \left( \frac{SCr}{K}, 1 \right) - 1.209 \times 0.993 \text{Age} \times 1.018 \text{ (if female)} \times 1.159 \text{ (if black)}
\]

Table 1. Characteristics of the Subjects *

| Variables          | Healthy Control Group | Case Group | P Value * |
|--------------------|-----------------------|------------|----------|
| No. (%)            | 60 (100)              | 79 (100)   | 73 (100) |
| Age (years)        | 47.07 ± 17.44         | 49.03 ± 13.40 | 51.26 ± 16.32 | P_all > 0.05 |
| Gender             |                       |            |          |
| Male               | 30 (50)               | 38 (48.1) | 41 (56.1) | P_all > 0.05 |
| Female             | 30 (50)               | 22 (31.9) | 22 (43.9) |          |
| Height (m)         | 1.61 ± 0.08           | 1.62 ± 0.08 | 1.59 ± 0.08 | P_all > 0.05 |
| Weight (kg)        | 57.22 ± 8.54          | 56.28 ± 8.58 | 51.55 ± 9.61 | P_all > 0.05 |
| Waist (cm)         | 75.37 ± 6.47          | 77.35 ± 7.09 | 77.50 ± 7.10 | P_all > 0.05 |
| BMI (kg/m^2)       | 21.88 ± 2.33          | 21.54 ± 2.74 | 20.50 ± 2.75 | P_all > 0.05 |
| Red blood cell (T/L)| 4.76 ± 0.44           | 4.78 ± 0.76 | 3.60 ± 1.01 | P1 > 0.05; P2 < 0.05; P3 < 0.05 |
| Hematocrit (%)     | 40.57 ± 4.08          | 40.39 ± 7.09 | 29.84 ± 9.21 | P1 > 0.05; P2 < 0.05; P3 < 0.05 |
| Hemoglobin (g/L)   | 131.85 ± 14.26        | 138.88 ± 25.79 | 102.72 ± 31.49 | P1 > 0.05; P2 < 0.05; P3 < 0.05 |
| Urea (mmol/L)      | 7.58 ± 1.50           | 7.13 ± 7.99 | 18.51 ± 11.76 | P_all > 0.05 |
| Creatinine (μmol/L)| 84.60 ± 17.06         | 83.20 ± 22.27 | 362.90 ± 324.95 | P_all < 0.05 |
| eGFR (mL/min/1.73m²)| 11.74 ± 7.17          | 12.45 ± 7.17 | 32.15 ± 11.74 | < 0.001 |

* Values are expressed as mean ± SD unless otherwise indicated.

Table 2. The Concentration of TGF-β1 in Control and Case Groups

| TGF-β1 (ng/mL) | Healthy Control Group (n = 60) | Case Group (n = 152) | P Value |
|----------------|-------------------------------|----------------------|---------|
| Χ ± SD         | 13.45 ± 7.17                  | 23.86 ± 8.28         | 37.88 ± 10.29 | P1 < 0.0001; P2 < 0.0001; P3 < 0.0001 |
| Median (25th; 75th) | 12.44 (8.28; 17.59) | 22.08 (20.06; 27.79) | 36.13 (29.50; 45.89) |

DM and HTN were excluded. With glomerulonephritis, the renal injuries often occurred with primary renal diseases due to some inflammation agents and the progression of the autoinflammation, so that inflammation were considered to be the main cause of CKD (16). Many inflammation cytokines, such as TGF-β1, had a motivating role in the progression of the disease. The TGF-β1 increase is associated with the increase of indoxyl sulfate in CKD patients, which is derived from protein in food. Some of the tryptophan from dietary protein is converted to indole by the tryptophanase of intestinal bacteria such as E. Coli indole, which is absorbed into the bloodstream from the intestines.
Table 4. The TGF-β1 Levels in Five Stages of CKD

| Variables | Control Group; n = 60 | CKD Stage | P |
|-----------|-----------------------|-----------|---|
|          | X ± SD                |           | n = 30 | n = 31 | n = 31 | n = 31 | n = 31 |
| Urea      | 13.45 ± 7.27          | 22.07 ± 8.59 | 25.65 ± 7.69 | 35.17 ± 11.41 | 37.87 ± 9.20 | 40.52 ± 9.80 | P = 0.001 |
| Creatinine| 0.49 ± 0.24           | 0.42 ± 0.11 | -0.59 | -0.54 | -0.45 | -0.50 | -0.49 |
| eGFR      | 0.35 ± 0.19           | 0.27 ± 0.14 | -0.36 | -0.28 | -0.20 | -0.17 | -0.18 |
| hsCRP     | 9.80 ± 4.79           | 11.41 ± 5.97 | -0.98 | -0.92 | -0.88 | -0.85 | -0.88 |
| RBC       | 27.30 ± 10.20         | 37.87 ± 11.41 | -0.92 | -0.88 | -0.85 | -0.85 | -0.85 |
| Hct       | 0.34 ± 0.08           | 0.37 ± 0.09 | -0.31 | -0.24 | -0.21 | -0.19 | -0.19 |
| Hem       | 0.28 ± 0.08           | 0.31 ± 0.09 | -0.25 | -0.18 | -0.15 | -0.13 | -0.13 |

Footnotes

Authors’ Contribution: All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

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