Changes of expression levels of serum cystatin C and soluble vascular endothelial growth factor receptor 1 in the treatment of patients with glomerulus nephritis

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Abstract. Expression levels and changes of serum cystatin C (C’s C) and soluble vascular endothelial growth factor receptor 1 (sVEGFR1) in treatment of patients with glomerulus nephritis (GN) were investigated. The medical records of 88 patients with GN who were diagnosed in Weifang People’s Hospital from March 2014 to June 2017 were collected, and their medical records were considered as a study group. The study group was divided into secondary glomerulonephritis (SGN) group (52 cases) and primary glomerulonephritis (PGN) group (36 cases). Physical examination data of 50 healthy volunteers who were examined in the same hospital during the same period were considered as a control group. The correlation between expression of serum C’s C and expression of sVEGFR1 of patients with GN was compared. Expression levels of serum C’s C and sVEGFR1 of patients before treatment in the study group were higher than those in the control group (P<0.05). With the extension of the treatment cycle, C’s C and sVEGFR1 expression levels in PGN and SGN groups reduced gradually (P<0.05). With the extension of the treatment cycle, the renal function indexes of the study group patients showed a downward trend (P<0.05). Expression of C’s C was positively correlated with urea nitrogen and creatinine (P<0.05). In conclusion, C’s C and sVEGFR1 can be used as indicators for monitoring the condition of patients with GN. It is worthwhile to promote C’s C and sVEGFR1 in clinical practice.

Introduction

Glomerulus nephritis (GN) is an autoimmune system disease with complex pathogenesis, diverse pathological classifications and long disease course (1). It can be induced by various causes and can be divided into primary glomerulonephritis (PGN) and secondary glomerulonephritis (SGN) (2). The pathogenic process of GN is divided into several stages: first edema, hematuria or proteinuria can be seen in the body; then, renal function weakens; eventually, patients suffer renal failure (3). At present, the drugs used to treat GN in clinic are few, and the efficacy of some drugs is poor, causing adverse reactions. Moreover, the treatment is expensive (4). Comprehensive preventive measures should be taken in the treatment of GN to control deterioration of renal function, alleviate the symptoms of patients, reduce the incidence of serious complications and improve patients’ life quality (5).

Serum cystatin C (C’s C) is a non-glycosylated small protein, with 120 amino acid residues, and is a member of cysteine protease inhibitors (6). The production rate of C’s C is relatively constant in body fluids and nucleated cells in tissues of the body. C’s C metabolizes and regulates peptides and proteins in cells (7). In the blood circulation of the body, C’s C is filtered by glomerular filtration membrane and is reabsorbed by proximal convoluted tubules. The glomerular filtration function directly determines the level of C’s C in the serum (8,9).

Vascular endothelial growth factor (VEGF), a dynamic source of division and growth of endothelial cells, facilitates angiogenesis. VEGF can combine with soluble vascular endothelial growth factor receptor 1 (sVEGFR1) to prevent VEGF membrane receptors from mediating and taking a biological effect. sVEGFR1 and VEGF membrane receptors are competitive, when they are combined with VEGF, the biological activity of sVEGFR1 and VEGF will be regulated negatively (10,11). Studies have shown that when sVEGFR1 combines with VEGF in the blood circulation, the function of VEGF will be regulated. Increasing the level of sVEGFR1 prevents VEGF from facilitating angiogenesis and injuring endothelial cells, which is closely related to kidney diseases (12).
Studies on the expression of serum C's C and sVEGFR1 in patients with GN and predictive function and evaluation function of serum C's C and sVEGFR1 in diseases are scarce. Therefore, expression levels of serum C's C and sVEGFR1 in the treatment of patients with GN were detected and the monitoring function and significance of serum C's C and sVEGFR1 in GN were investigated in this study, in order to provide specific study data and theoretical basis for clinical treatment of GN.

Patients and methods

General data. The medical records of 88 patients with GN who were diagnosed in Weifang People's Hospital (Weifang, China) from March 2014 to June 2017 were collected, and their medical records were considered as study group. The study group was divided to SGN group and PGN group. There were 52 patients in SGN group and 36 patients in PGN group. Physical examination data of 50 healthy volunteers who were examined in Weifang People's Hospital during the same period were considered as the control group. Among those patients, there were 66 males and 72 females, with an average age of 45.0±15.7 years and a BMI of 21.0±3.6. There was no significant difference in some data in the patients in two groups, including gender, age and BMI (P>0.05). There were significant differences in hypertension, edema and some other aspects (P<0.05) (Table I).

Inclusion and exclusion criteria. Inclusion criteria: Patients who were diagnosed with GN by histopathology were included; Serum creatinine (SCr) <550 mol/l, and blood urea nitrogen (BUN) <17 mmol/l; urine RBC was increased and showed tubular shape (13).

Exclusion criteria: i) patients with malignant tumors; ii) patients complicated with solid lesions of other organs; iii) patients complicated with blood diseases and infection; iv) patients with cognitive disorder or communication disorder; v) patients with poor compliance.

All the patients and their family members agreed to participate in the experiment, and an informed consent form was signed. This experiment was approved by the Medical Ethics Committee of Weifang People's Hospital.

Reagents and materials. The serum separation centrifuge was purchased from Zhengzhou Zhishi Changyun Technology Co., Ltd. C's C was detected by automatic biochemical analyzer BS-600 (Shenzhen Mindray Biomedical Electronics Co., Ltd.); C's C test kits (Shenzhen ziker Biological Technology Co., Ltd., ZK-H068); sVEGFR1 kits were from R&D Systems (ML-ELISA). The procedures were carried out strictly according to the instructions of the kits.

Serum C's C and sVEGFR1 in serum of the patients in the study group and the control group were compared before treatment. ii) The changes of expression levels of serum C's C of the patients in PGN group and SG group were compared before they were treated, after one cycle of treatment, after three cycles of treatment and after discharged from hospital. iii) The changes of expression levels of sVEGFR1 of the patients in PGN group and SG group were compared before they were treated, after one cycle of treatment, after three cycles of treatment and after discharged from the hospital. iv) The correlation between the expression of C's C and the expression of sVEGFR1 in GN was analyzed by correlation analysis.

Statistical analysis. In this study, SPSS 19.0 (Bizinsight (Beijing) Information Technology Co., Ltd.) was used to statistically analyze the experimental data. The count data were analyzed by χ² test. The measurement data were expressed in the form of mean value ± standard deviation. t-test was used in comparison of data between two groups. Bonferroni test was the post hoc test. Repeated measures analysis of variance was used in comparison of the data between two groups at different time points in the treatment of patients. Pearson analysis was used to carry out correlation analysis of the data. The illustrations of this experiment were drawn by GraphPad Prism 8. P<0.05 indicates statistically significant difference.

Results

Comparison of expression levels of C's C and sVEGFR1 of the patients in the study and the control groups. The expression levels of C's C and sVEGFR1 of the patients in the study group and the subjects in the control group were compared before treatment. The expression level of C's C of the patients in the study group was significantly higher than that in control group. The expression level of sVEGFR1 of the patients in study group was significantly higher than that in the control group. Differences in expression levels of C's C and sVEGFR1 of the patients in the two groups were statistically significant (P<0.05) (Table II).

Comparison of the changes of expression level of C's C of the patients in PGN and SG groups before treatment, one cycle after treatment, three cycles after treatment and after discharge. The expression level of C's C of the patients in PGN group and SG group was compared before treatment, after one cycle of treatment, after three cycles of treatment, and after discharge from hospital. It was found that the expression level of C's C of the patients in PGN group and SG group decreased gradually after one cycle of treatment.
The expression level of C's C of the patients in PGN group and SGN group after one cycle of treatment was lower than that before they were treated (P<0.05). The expression level of C's C of the patients in PGN group and SGN group decreased gradually after three cycles of treatment. The expression level of C's C of the patients in PGN group and SGN group after three cycles of treatment was lower than that after one cycle of treatment (P<0.05). The expression level of C's C of the patients in PGN group and SGN group decreased gradually after discharge from hospital. The expression level of C's C of the patients in PGN group and SGN group after discharged from the hospital was lower than that after three cycles of treatment (P<0.05). There was no significant difference between the expression level of C's C of the patients in PGN group and that in SGN group (P>0.05) (Table III and Fig. 1).

Table I. General data of the patients in the study group and the control group [n (%)].

| Factors         | Study group (n=88) | Control group (n=50) | χ²/t  | P-value |
|-----------------|--------------------|----------------------|-------|---------|
| Sex             |                    |                      | 0.105 | 0.746   |
| Male            | 43 (48.86)         | 23 (46.00)           |       |         |
| Female          | 45 (51.14)         | 27 (54.00)           |       |         |
| Age (years)     |                    |                      | 0.001 | 0.975   |
| ≤45             | 46 (52.27)         | 26 (52.00)           |       |         |
| >45             | 42 (47.73)         | 24 (48.00)           |       |         |
| BMI (kg/m²)     |                    |                      | 0.178 | 0.673   |
| ≤21             | 42 (47.73)         | 22 (44.00)           |       |         |
| >21             | 46 (52.27)         | 28 (56.00)           |       |         |
| Hypertension    |                    |                      | 48.391| <0.001  |
| Yes             | 63 (71.59)         | 5 (10.00)            |       |         |
| No              | 25 (28.41)         | 45 (90.00)           |       |         |
| Edema           |                    |                      | 63.971| <0.001  |
| Yes             | 62 (70.45)         | 0                    |       |         |
| No              | 26 (29.55)         | 50 (100.00)          |       |         |
| Hematuria       |                    |                      | 60.311| <0.001  |
| Yes             | 60 (68.18)         | 0                    |       |         |
| No              | 28 (31.82)         | 50 (100.00)          |       |         |
| Proteinuria     |                    |                      | 76.171| <0.001  |
| Yes             | 68 (77.27)         | 0                    |       |         |
| No              | 20 (22.73)         | 50 (100.00)          |       |         |
| Urea nitrogen (mmol/l) | 14.34±1.23 | 5.65±0.83 | 44.500 | <0.001  |
| Creatinine (µmol/l) | 293.43±32.32 | 81.34±11.25 | 44.820 | <0.001  |

Table II. Comparison of expression levels of C's C and sVEGFR1 in the two groups.

| Groups          | Study group (n=88) | Control group (n=50) | t    | P-value |
|-----------------|--------------------|----------------------|------|---------|
| C's C (mg/l)    | 2.43±0.45          | 0.83±0.18            | 24.040 | <0.001  |
| sVEGFR1 (µg/l)  | 30.42±7.42         | 7.34±1.54            | 21.700 | <0.001  |

C's C, serum cystatin C; sVEGFR1, soluble vascular endothelial growth factor receptor 1.

Comparison of the changes of expression level of sVEGFR1 of the patients in PGN and SGN groups before treatment, one cycle after treatment, in three cycles after treatment and after discharge. The expression level of sVEGFR1 of the patients in PGN group and SGN group was compared before they were treated, after a cycle of treatment, after three cycles of treatment, and after they discharged from the hospital. It was found that the expression level of sVEGFR1 of the patients in PGN group and SGN group decreased gradually after a cycle of treatment. The expression level of sVEGFR1 of the patients in PGN group and SGN group after a cycle of treatment was lower than that before they were treated (P<0.05). The expression level of sVEGFR1 of the patients in PGN group and SGN group after three cycles of treatment was lower than that after one cycle of treatment (P<0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and that in SGN group (P>0.05) (Table III and Fig. 1).
Table III. Comparative analysis of the expression level of C’s C of the patients in PGN group and SGN group before and after treatment.

| Groups              | Before patients were treated | In one cycle after patients were treated | In three cycles after patients were treated | After patients were discharged | F-value | P-value |
|---------------------|------------------------------|-----------------------------------------|-------------------------------------------|---------------------------------|---------|---------|
| SGN group (n=52)    | 2.42±0.42                    | 2.11±0.34<sup>a</sup>                  | 1.62±0.30<sup>a,b</sup>                  | 0.45±0.22<sup>c</sup>          | 0.842   | <0.001  |
| PGN group (n=36)    | 2.43±0.45                    | 2.12±0.42<sup>a</sup>                  | 1.64±0.35<sup>a,b</sup>                  | 0.42±0.24<sup>c</sup>          | 0.812   | <0.001  |
| t-test              | 0.107                        | 0.123                                   | 0.287                                     | 0.775                           | 0.546   |         |
| P-value             | 0.915                        | 0.902                                   | 0.775                                     | 0.546                           |         |         |

<sup>a</sup>P<0.05, compared with before treatment in the same group; <sup>b</sup>P<0.05 compared with one cycle of treatment in the same group; <sup>c</sup>P<0.05 compared with three cycles of treatment in the same group. C’s C, serum cystatin C; PGN, primary glomerulonephritis; SGN, secondary glomerulonephritis.

Table IV. Comparative analysis of the expression level of sVEGFR1 of the patients in PGN group and SGN group before and after treatment.

| Groups              | Before the patients were treated | In one cycle after the patients were treated | In three cycles after the patients were treated | After the patients were discharged | F-value | P-value |
|---------------------|----------------------------------|---------------------------------------------|-----------------------------------------------|---------------------------------|---------|---------|
| SGN group (n=52)    | 30.51±7.13                      | 26.43±6.46<sup>a</sup>                     | 21.42±5.25<sup>a,b</sup>                     | 17.34±5.13<sup>c</sup>          | 0.409   | <0.001  |
| PGN group (n=36)    | 30.42±7.42                      | 26.32±6.42<sup>a</sup>                     | 21.53±5.63<sup>a,b</sup>                     | 17.45±5.26<sup>c</sup>          | 0.387   | <0.001  |
| t-test              | 0.057                           | 0.079                                      | 0.094                                       | 0.926                           | 0.922   |         |
| P-value             | 0.955                           | 0.937                                      | 0.926                                       | 0.922                           |         |         |

<sup>a</sup>P<0.05, compared with before treatment in the same group; <sup>b</sup>P<0.05 compared with a cycle of treatment in the same group; <sup>c</sup>P<0.05 compared with three cycles of treatment in the same group. sVEGFR1, soluble vascular endothelial growth factor receptor 1; PGN, primary glomerulonephritis; SGN, secondary glomerulonephritis.

Figure 1. Comparison of the expression level of C’s C of the patients in PGN group and SGN group before treatment, in one cycle after treatment, in three cycles after treatment and after they were discharged from hospital. The result of immunoturbidimetry showed that the expression level of C’s C of the patients in PGN group and SGN group decreased gradually in one cycle after treatment. The expression level of C’s C of the patients in PGN group and SGN group one cycle after treatment was lower than that before treatment (P<0.05). Expression level of C’s C of the patients in PGN group and SGN group decreased gradually in three cycles after treatment. Expression level of C’s C of the patients in PGN group and SGN group in three cycles after treatment was lower than that in one cycle after treatment (P<0.05). Expression level of C’s C of the patients in PGN group and SGN group decreased gradually after they were discharged from hospital. There was no significant difference between the expression level of C’s C of the patients in PGN group and SGN group in three cycles after they were treated (P≥0.05). There was no significant difference between the expression level of C’s C of the patients in PGN group and SGN group in three cycles after they were treated (P>0.05). P<0.05, C’s C, serum cystatin C; PGN, primary glomerulonephritis; SGN, secondary glomerulonephritis.

Figure 2. Comparison of the expression level of sVEGFR1 of the patients in PGN group and SGN group before treatment, in one cycle after treatment, in three cycles after treatment and after they were discharged from hospital. The result of ELISA showed that the expression level of sVEGFR1 of the patients in PGN group and SGN group decreased gradually in one cycle after treatment. Expression level of sVEGFR1 of the patients in PGN group and SGN group one cycle after treatment was lower than that before treatment (P<0.05). Expression level of sVEGFR1 of the patients in PGN group and SGN group decreased gradually in three cycles after treatment. Expression level of sVEGFR1 of the patients in PGN group and SGN group in three cycles after treatment was lower than that in one cycle after they were treated (P<0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and SGN group in three cycles after they were treated (P>0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and SGN group after discharge was lower than that in three cycles after they were treated (P<0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and SGN group after discharge was lower than that in three cycles after they were treated (P>0.05). P<0.05, sVEGFR1, soluble vascular endothelial growth factor receptor 1; PGN, primary glomerulonephritis; SGN, secondary glomerulonephritis.
three cycles of treatment was lower than that after one cycle of treatment (P<0.05). The expression level of sVEGFR1 of the patients in PGN group and SGN group decreased gradually after discharge from the hospital. The expression level of sVEGFR1 of the patients in PGN group and SGN group after discharge from hospital was lower than that after three cycles of treatment (P<0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and that in SGN group (P>0.05) (Table IV and Fig. 2).

Changes in renal function in different periods of treatment.

The results showed that with the increase of the treatment cycle, the renal function indexes of patients in the study group showed a downward trend (P<0.05). There was no significant difference between the expression level of sVEGFR1 of the patients in PGN group and that in SGN group (P>0.05) (Table IV and Fig. 2).

Correlation analysis of expression of C’s C and sVEGFR1 in GN and the renal function indexes. According to Spearman’s correlation analysis, the expression of C’s C and sVEGFR1 was positively correlated in GN (r=0.740, P<0.001). Due to the positive correlation between the expression of C’s C and urea nitrogen, creatinine, urea nitrogen, creatinine were analyzed. Spearman’s correlation results showed that C’s C expression was positively correlated with urea nitrogen (r=0.797, P<0.001), and C’s C expression was positively correlated with creatinine (r=0.841, P<0.001) (Fig. 3).

Discussion

At present, it is believed that GN can lead to end-stage renal diseases and its pathological changes are complex. For many GN diseases, cellular immunity plays a decisive role, and humoral immunity is also clearly expressed during the pathogenic process of GN diseases (14). In addition, the disorder of blood coagulation mechanism and the change of blood rheology have an effect on the occurrence and progression of glomerular diseases (15). In clinic, the pathogenesis and treatment of GN are known, the drugs used to treat GN are immunosuppressive agents and glucocorticoid, but the efficacy of many glomerular diseases is still unsatisfactory (16,17). In order to evaluate the changes of patients’ condition in the occurrence and progression of GN disease and improve clinical symptoms of the patients, and the cure rate, the expression of C’s C and sVEGFR1 in GN was investigated. C’s C is an independent endogenous marker, only filtered and metabolized by kidneys. C’s C is important in the early diagnosis of renal injury and is highly recognized for evaluating glomerular function in clinic (18,19). sVEGFR1 is produced in endothelial cells, monocytes and placenta. As a soluble receptor of VEGF, it negatively regulates the activity of VEGF (20). When sVEGFR1 is combined with VEGF, the signal transduction acts on angiogenesis, when this process is disrupted, a series of pathological conditions will occur, such as the deterioration of renal function, which is caused by injury of vascular endothelial cell function (21).

In this study, the expression levels of C’s C and sVEGFR1 of the patients in the study group and the control group were compared. It was found that the expression levels of C’s C and sVEGFR1 of the patients in the study group were significantly higher than those in the control group. The results indicated that the expression levels of C’s C and sVEGFR1 were diagnostic markers in the occurrence and progression of glomerular.
studies on sVEGFR1 and the correlation between C’s C and expression was an important indicator to evaluate the filtration rate of glomeruli. Endogenous creatinine clearance and expression of C’s C were related to the pathogenic process of GN. At present, studies (28) demonstrated that endogenous creatinine clearance and expression of C’s C and sVEGFR1 were associated with the progression of GN. Some studies showed that the expression of C’s C and sVEGFR1 was positively correlated in GN. This result suggests that C’s C and sVEGFR1 play important roles in monitoring the condition of GN patients during treatment.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions

XX analyzed and interpreted the patient general data. LQ performed ELISA. LY was responsible for analysis of the results. XX wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The study was approved by the Ethics Committee of Weifang People's Hospital (Weifang, China). Patients who participated in this research had complete clinical data. Signed informed consents were obtained from the patients and/or the guardians.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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nephropathy. Some studies reported that the occurrence and progression of GN diseases are related to cellular immunity, serum free radicals, and coagulation mechanism, and that the occurrence and development of GN diseases are closely related to cellular immunity (22). C’s C is a cysteine protease inhibitor that can be used to detect the filtration rate of glomeruli in kidney injury. The practical method to detect C’s C is immunoassay (23). The signal transduction of sVEGFR1 improves hematopoietic function, proliferative function, differentiation function and immune function of monocytes and macrophages of hematopoietic stem cells (HSC) (24). The changes of cellular immunity exist in the expression of C’s C and sVEGFR1, which indirectly confirms the conclusion of this study. To investigate the specific changes of the expression levels of C’s C and sVEGFR1 in GN, the changes of expression levels of C’s C and sVEGFR1 were investigated before and after treatment of patients. The results showed that expression levels of serum C’s C and sVEGFR1 of the patients in PGN group and SGN group decreased gradually after a cycle of treatment. Expression levels of serum C’s C and sVEGFR1 of the patients in PGN group and SGN group after a cycle of treatment were lower than those before treatment (P<0.05). Expression levels of serum C’s C and sVEGFR1 of the patients in PGN group and SGN group after three cycles of treatment. Expression levels of serum C’s C and sVEGFR1 of the patients in PGN group and SGN group after three cycles of treatment were lower than those after a cycle of treatment (P<0.05). Expression levels of serum C’s C and sVEGFR1 of the patients in PGN group and SGN group after discharged from hospital were lower than those after three cycles of treatment (P<0.05). Previous studies have shown that the pathogenesis of secondary glomerular diseases and primary glomerular diseases is completely different, while their clinical manifestations and pathological changes are similar (25). Previous studies have demonstrated that C’s C can be expressed effectively in the occurrence and progression of kidney diseases. With the deterioration of pathological conditions, the filtration rate of glomeruli decreases and the expression level of C’s C increases (26). When the glomerular filtration barrier is injured and the repair of endothelial cells is blocked, the biological effects of VEGF will be inhibited and respond to the expression of sVEGFR1 negatively (27). In this study, the changes of expression of C’s C and sVEGFR1 were investigated before and after treatment. The result showed that the expression levels of C’s C and sVEGFR1 decreased with the continuous treatment of the patients. This result indicates that the expression of C’s C and sVEGFR1 can be used as an indicator for monitoring the condition of patients with GN when they are treated. The correlation between expression of C’s C and expression of sVEGFR1 was analyzed. The result showed that the expression of C’s C and sVEGFR1 was positively correlated in GN. This result suggests that C’s C and sVEGFR1 are associated with the progression of GN. Some studies (28) demonstrated that endogenous creatinine clearance was an important indicator to evaluate the filtration rate of glomeruli. Endogenous creatinine clearance and expression of C’s C are related to the pathogenic process of GN. At present, studies on sVEGFR1 and the correlation between C’s C and sVEGFR1 are few, so the specific mechanism of C’s C and sVEGFR1 needs further investigation.

In conclusion, C’s C and sVEGFR1 are highly expressed in serum of GN patients. As patients are treated, the expression levels of C’s C and sVEGFR1 decrease. C’s C and sVEGFR1 are few, so the specific mechanism of C’s C and sVEGFR1 needs further investigation.
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