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***Retrospective Study***

**Correlation between glycated hemoglobin A1c, urinary microalbumin, urinary creatinine, β2 microglobulin, retinol binding protein and diabetic retinopathy**

Song JJ *et al*. Correlation analysis of DR

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**Abstract**

BACKGROUND

Retinopathy is the most common microvascular disease of type 2 diabetes, and seriously threatens the life, health and quality of life of patients. It is worth noting that the development of diabetic retinopathy (DR) can be hidden, with few symptoms. Therefore, the preliminary screening of diabetic patients should identify DR as soon as possible, delay disease progression, and play a vital role in its diagnosis and treatment.

AIM

To investigate the correlation between glycated hemoglobin A1c (HbA1c), urinary microalbumin (U-mALB), urinary creatinine (U-CR), mALB/U-CR ratio, β2 microglobulin (β2MG), retinol binding protein (RBP) and DR.

METHODS

A total of 180 patients with type 2 diabetes mellitus attending the Second People’s Hospital of Hefei from January 2022 to August 2022 were retrospectively enrolled by ophthalmologists. Based on whether they had combined retinopathy and its degree, 68 patients with diabetes mellitus without retinopathy (NDR) were assigned to the NDR group, 54 patients with non-proliferative DR (NPDR) to the NPDR group, and 58 patients with proliferative DR to the PDR group. General data, and HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR results were collected from the patients and compared among the groups. Pearson's correlation method was used to analyze the correlation between HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR indices, and multiple linear regression was applied to identify the risk factors for DR. Receiver operator characteristic (ROC) curves were also drawn.

RESULTS

The differences in age, gender, systolic and diastolic blood pressure between the groups were not statistically significantly (*P* > 0.05), but the difference in disease duration was statistically significant (*P* < 0.05). The differences in fasting blood glucose, high-density lipoprotein cholesterol, low-density lipoprotein cholesterol, total cholesterol, and triglyceride between the groups were not statistically significant (*P* > 0.05). HbA1c in the PDR group was higher than that in the NPDR and NDR groups (*P* < 0.05). The levels of mALB, β2MG, RBP, mALB/U-CR and U-CR in the PDR group were higher than those in the NPDR and NDR groups (*P* < 0.05). Multiple linear regression analysis showed that disease duration, HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR were risk factors for the development of DR. The ROC curve showed that the area under the curve (AUC) for the combination of indices (HbA1c + mALB + mALB/U-CR + U-CR + β2MG + RBP) was 0.958, with a sensitivity of 94.83% and specificity of 96.72%, which was higher than the AUC for single index prediction (*P* < 0.05).

CONCLUSION

HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP can reflect the development of DR and are risk factors affecting PDR, and the combination of these six indices has predictive value for PDR.

**Key Words:** Diabetic retinopathy; β2 microglobulin; Retinol-binding protein; Urinary microalbumin; Urinary creatinine

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**Core Tip:** Diabetes retinopathy (DR) is a common complication of diabetes, which can eventually lead to blindness in diabetic patients and seriously affect the quality of life of patients. The identification of risk factors for DR is significant for early intervention. Here we retrospectively analyzed 180 patients with type 2 diabetes mellitus to examine the correlation between glycated hemoglobin A1c, microalbumin (mALB), mALB/urinary creatinine (U-CR), U-CR, ββ2 microglobulin, retinol binding protein and DR in diabetic patients in order to provide a scientific basis and guidance for clinical application.

**INTRODUCTION**

Diabetic retinopathy (DR) is an irreversible blindness-causing disease[1]. The prevalence of diabetes in China accounts for 26.2% of the global diabetic population, and the prevalence of DR is approximately 35%-50%[2]. The prevalence of DR in Singapore and the United States is 20.1% and 25.7%, respectively[3]. The disease progresses rapidly and if not diagnosed and treated early, it will seriously affect the visual field and vision. In severe cases, patients may even lose their sight, which causes many inconveniences to their life and work and hinders their normal life. Therefore, early clinical diagnosis is important for the subsequent treatment of DR patients[3]. Currently, the clinical diagnosis of this disease is mainly based on fundus photography and fluorescein angiography, but the application process is complicated and may cause adverse reactions in diabetic patients. In addition, there is a lack of convenient and intuitive biochemical markers providing guidance for the diagnosis of DR[4]. Therefore, it is important to identify relevant biochemical markers to predict DR. Urinary β2 microglobulin (β2MG) has been found to be closely associated with microvascular complications such as diabetic nephropathy. It is known that DR is a microvascular complication, so it is assumed that the pathogenesis of the two diseases is similar and β2MG may be a useful marker for predicting DR[5]. Retinol-binding protein (RBP), a lipid-derived cytokine, has been shown to be closely associated with the development of diabetes mellitus and diabetic vasculopathy[6]. Urinary microalbumin (U-mALB), urinary creatinine (U-CR) and the mALB/U-CR ratio are predictors of diabetic vasculopathy and are risk factors for endothelial cell function and microvascular function[7]. In this study, we aimed to examine the correlation between glycated hemoglobin A1c (HbA1c), β2MG, RBP, mALB, U-CR, mALB/U-CR and DR lesions in patients with DR. The innovation of this study is determination of the predictive value of the combined detection of HbA1c, mALB, mALB/U-CR, U-CR, β2MG, and RBP in DR using real clinical data. The clinical significance is to provide a scientific basis and guidance for the clinical use of the combined detection of HbA1c, mALB, mALB/U-CR, U-CR, β2MG, and RBP to evaluate the risk of DR.

**MATERIALS AND METHODS**

***General data***

A total of 180 type 2 diabetic patients attending the Second People’s Hospital of Hefei from January 2022 to August 2022 were enrolled retrospectively, including 68 patients with diabetes without retinopathy (NDR group), 54 patients with non-proliferative diabetic retinopathy (NPDR group), and 58 with proliferative diabetic retinopathy (PDR group).

***Inclusion criteria***

(1) The study subjects met the diagnostic criteria for type 2 diabetes mellitus[8]; (2) The diagnosis of DR was based on the International Clinical Classification Criteria for Diabetic Retinopathy[9]. NPDR: microaneurysm alone was observed or 4 quadrants with intraretinal hemorrhage and microangioma; or moderate retinal mesangiopathy occurring in more than 2 or more quadrants; PDR: If the retina had new abnormal blood vessels, this was considered PDR. The diagnosis was confirmed by satisfying one or more of the following: neovascularization, vitreous hematopoiesis or anterior retinal hemorrhage; and (3) None of the study subjects had a history of trauma or ocular surgery.

***Exclusion criteria***

(1) Those with combined non-fundus pathology, *e.g.,* cataract, glaucoma; (2) Those with poorly graded fundus visual field images due to blurring of large blood vessels adjacent to the optic disc, and whose diagnosis was more difficult to further confirm on fundus examination; (3) Those with organ disease, such as coronary artery disease, heart failure, diabetic nephropathy, *etc.*; (4) Combined with diabetic complications, such as diabetic gangrene, stroke, or atherosclerosis; and(5) Difficult to cooperate in the completion of the study.

***Methods***

General information of the patients was collected, including age, gender, duration of disease, systolic and diastolic blood pressure. Blood was collected in the morning after a 12-h fast to measure HbA1c, fasting blood glucose (FPG), high-density lipoprotein cholesterol (HDL-C), low-density lipoprotein cholesterol (LDL-C), total cholesterol (TC), and triglyceride (TG) using a glycated hemoglobin analyzer and supporting reagents.

Urinary mALB and U-CR concentrations were measured using a special protein analyzer and the urinary mALB/U-CR ratio was calculated three times. β2MG was measured by the immunoturbidimetric method and RBP was measured using an automatic biochemical analyzer.

***Observation indicators***

General information: age, gender, duration of disease, systolic and diastolic blood pressure. Clinical indicators: FPG, HDL-C, LDL-C, TC, TG, and HbA1c. Combined indicators: mALB, mALB/U-CR, U-CR, β2MG, and RBP levels.

***Statistical analysis***

GraphPad Prism 9 was used to analyze the study data and for image export. The measurement data were expressed as mean ± SD, and compared by one-way ANOVA for multiple groups of data or for two groups of data. The count data were expressed by *n* (%), and compared using the *χ*2 test. Correlation analysis and risk factor identification were performed using Pearson’s correlation method and multiple linear regression, respectively. A receiver operator characteristic (ROC) curve was plotted to predict the value of PDR. *P* < 0.05 was considered statistically significant.

**RESULTS**

***General information in each group***

The differences in age, gender, systolic and diastolic blood pressure between the three groups were not significant (*P* > 0.05), but the differences in disease duration were significant (*P* < 0.05, Table 1).

***Clinical indicators among the groups***

No significant differences in FPG, HDL-C, LDL-C, TC and TG were observed among the groups (*P* > 0.05); HbA1c in the PDR group was higher than that in the NPDR and NDR groups (*P* < 0.05, Table 2).

***Comparison of mALB, mALB/U-CR, U-CR, β2MG and RBP levels among the groups***

The levels of mALB, β2MG, RBP, mALB/U-CR, and U-CR in the PDR group were higher than those in the NPDR and NDR groups (*P* < 0.05, Table 3).

***Correlation analysis***

By Pearson's correlation analysis, mALB, mALB/U-CR, U-CR, β2MG, and RBP were positively correlated with disease duration and HbA1c, (*P* < 0.05, Figure 1).

***Risk factors for the development of PDR***

With PDR as the dependent variable (yes = 1, no = 0) and the above meaningful results as independent variables all included as original values, multiple linear regression analysis was performed and the results revealed that disease duration, HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR were all risk factors for the development of PDR (Table 4).

***ROC curve analysis of HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP for predicting PDR***

As shown in Table 5 and Figure 2, the ROC curve indicated that the combined diagnostic area under the curve of the indicators was 0.904, with a sensitivity of 92.53% and specificity of 90.65%, which was higher than the prediction of HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP alone (*P* < 0.05).

**DISCUSSION**

DR is a diabetes-induced retinal vascular complication and causes irreversible visual impairment and vision loss[10]. Currently, irreversible visual impairment due to DR accounts for approximately 1.9% worldwide, while visual loss accounts for approximately 2.6% worldwide. However, there are significant reported differences in the prevalence of DR in China and abroad[11]. Some scholars have reported that the prevalence of DR in diabetes is about 34.6% globally, and is 16.4% and 25.9% in the UK and Australia, respectively. The incidence of PDR is approximately 7.0%. In China, the results of the six provinces of the Guangdong Provincial Flow Survey showed that the prevalence of DR in 13473 diabetic patients ranged from 33.28% to 34.88%[12,13]. The above studies suggest that DR is a common and highly prevalent chronic microangiopathy, which endangers public health safety. Therefore, early diagnosis of DR in diabetic patients is essential in clinical settings.

In recent years, studies have found that persistent poor glycemic control was a risk factor for the development and progression of DR, disrupting polyol metabolic pathways, contributing to the release of protein kinase C in large amounts and stimulating the onset of oxidative stress, inflammatory cell infiltration and other metabolic imbalances[14]. The above cascade of reactions further affects endothelial cells and microcirculatory function, leading to abnormal retinal microvascular biology and hemodynamics, and the development of DR. It has been found that persistent poor glycemic control is associated with alterations in mALB and U-CR, which are stimulated by oxidative stress and inflammation, and persistent high expression of mALB and U-CR[15]. The mALB/U-CR ratio is a novel index that is more accurate and reliable than traditional 24 h urine protein quantification, and is a valid marker for qualitative or quantitative prediction of proteinuric changes in the clinic[16]. DR severity has been reported to be positively correlated with decreased renal function and is independent of renal pathology[17]. An 8-year follow-up study reported that patients with DR with upregulated expression of mALB/U-CR had a progressively reduced glomerular filtration rate[18]. In the current results, mALB, mALB/U-CR, U-CR, β2MG, and RBP levels were found to be consistently increased as DR progressed from NDR, NPDR, to the PDR stage. It is hypothesized that mALB, mALB/U-CR, U-CR, β2MG, and RBP upregulated expression in DR patients is closely associated with progressive loss of renal function in diabetic patients.

Urinary β2MG was also expressed at high levels with the progressive of DR, which is a recognized early predictor of diabetic nephropathy in the clinic with high sensitivity and specificity[19]. This is consistent with previous studies by Cheng *et al*[20] and others, although altered β2MG levels have been associated with systemic lupus erythematous nephritis and globular nephropathy. However, the present study combined urinary mALB, mALB/U-CR, U-CR, and RBP to positively verify the association between DR occurrence and altered renal function. RBP is a low molecular mass vitamin A transporter protein, synthesized by the liver, expressed in large amounts in urine, blood, and cerebrospinal fluid, and reaches the blood *via* retinol in the liver[21]. It has been found that free RBP can normally be filtered by the glomerulus in healthy populations[22]. Lu *et al*[23] reported that urinary RBP correlated significantly with changes in renal function as the disease progressed in patients with diabetic nephropathy, elevating the rate of thylakoid cell proliferation, basement membrane synthesis and impaired glomerular filtration in patients with diabetic nephropathy, with subsequent upregulation of urinary RBP. Our study showed that mALB was involved in the regulation of renal function.

In addition, the results showed that mALB, mALB/U-CR, U-CR, β2MG and RBP were related to disease duration and HbA1c (*P* < 0.05); and disease duration, HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR were risk factors for the development of PDR. This indicates that the progression of diabetic microangiopathy is related to duration of the disease and the degree of abnormal glucose metabolism. It was found that persistent elevation of HbA1c accelerates damage to structural proteins in the glomerular basement membrane, causing disruption of polyol pathways, oxidative stress onset, and inflammatory infiltration involved in microvascular injury[24]. With the onset and progression of DR, disease duration and HbA1c levels increased abnormally, suggesting that persistent disease duration and abnormal HbA1c expression are involved in the development of diabetic microangiopathy, consistent with the findings of Casadei *et al*[25] and others. mALB, mALB/U-CR, U-CR, β2MG, RBP, disease duration and HbA1c were positively correlated in DR patients suggesting a synergistic role in promoting disease progression. The physiological characteristics of the glomerular and retinal vasculature, both of which are microcirculatory systems, suggest that persistent disease progression and elevated HbA1c levels induce disruption of the body's metabolic homeostasis and activation of oxidative stress, leading to damage to the vascular endothelium and the release of large amounts of inflammatory cytokines, inducing damage to the blood-retinal barrier and the glomerular filtration membrane barrier. In a state of persistently high glucose levels, oxides in vascular endothelial cells cannot be excreted, activating multiple signaling pathways and accelerating the impairment of vascular endothelial function, which may manifest as diabetic nephropathy if the abnormality is only in the kidney, or as DR if it occurs in the retina. Therefore, further studies found that the combination of HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP levels is predictive of the occurrence of PDR and can be used as a biochemical marker of DR. However, this study is a single center small sample study, and the results require further verification by follow-up multicenter and large sample studies.

**CONCLUSION**

HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP levels were up-regulated in DR patients, and their levels were closely related to disease duration, HbA1c and severity, all of which are risk factors for the development of PDR and can be used as markers to screen for DR progression. In the future, multi-center or propensity matching methods will be adopted to exclude the interference of multiple factors and provide new directions for clinical targeted therapy.

**ARTICLE HIGHLIGHTS**

***Research background***

Diabetic retinopathy (DR) is a common complication of diabetes, which can eventually lead to blindness and seriously affect the quality of life of diabetic patients. Therefore, identification of the risk factors of DR is significant for early intervention.

***Research motivation***

This study explored the risk factors for DR and their predictive effect on retinopathy.

***Research objectives***

This study aimed to investigate the correlation between glycated hemoglobin A1c (HbA1c), urinary microalbumin (U-mALB), urinary creatinine (U-CR), mALB/U-CR ratio, β2 microglobulin (β2MG), retinol binding protein (RBP) and DR.

***Research methods***

Based on real population data, a retrospective study was carried out.

***Research results***

Duration of disease, HbA1c, mALB, β2MG, RBP, mALB/U-CR and U-CR were found to be risk factors for the development of DR. The area under the curve of the combined indices (HbA1c + mALB + mALB/U-CR + U-CR + β2MG + RBP) was 0.958.

***Research conclusions***

The combination of HbA1c, mALB, mALB/U-CR, U-CR, β2MG and RBP has predictive value for proliferative DR.

***Research perspectives***

Large multicenter studies are needed to further verify these results.

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**Footnotes**

**Institutional review board statement:** This study was approved by the Medical Ethics Committee of the Second People’s Hospital of Hefei (No. 2023014).

**Informed consent statement:** This study only used anonymous data in the system, and did not require informed consent according to institutional policy.

**Conflict-of-interest statement:** The authors declare no conflicts of interest for this article.

**Data sharing statement:** According to institutional policy, the third party has no access to obtain the data.

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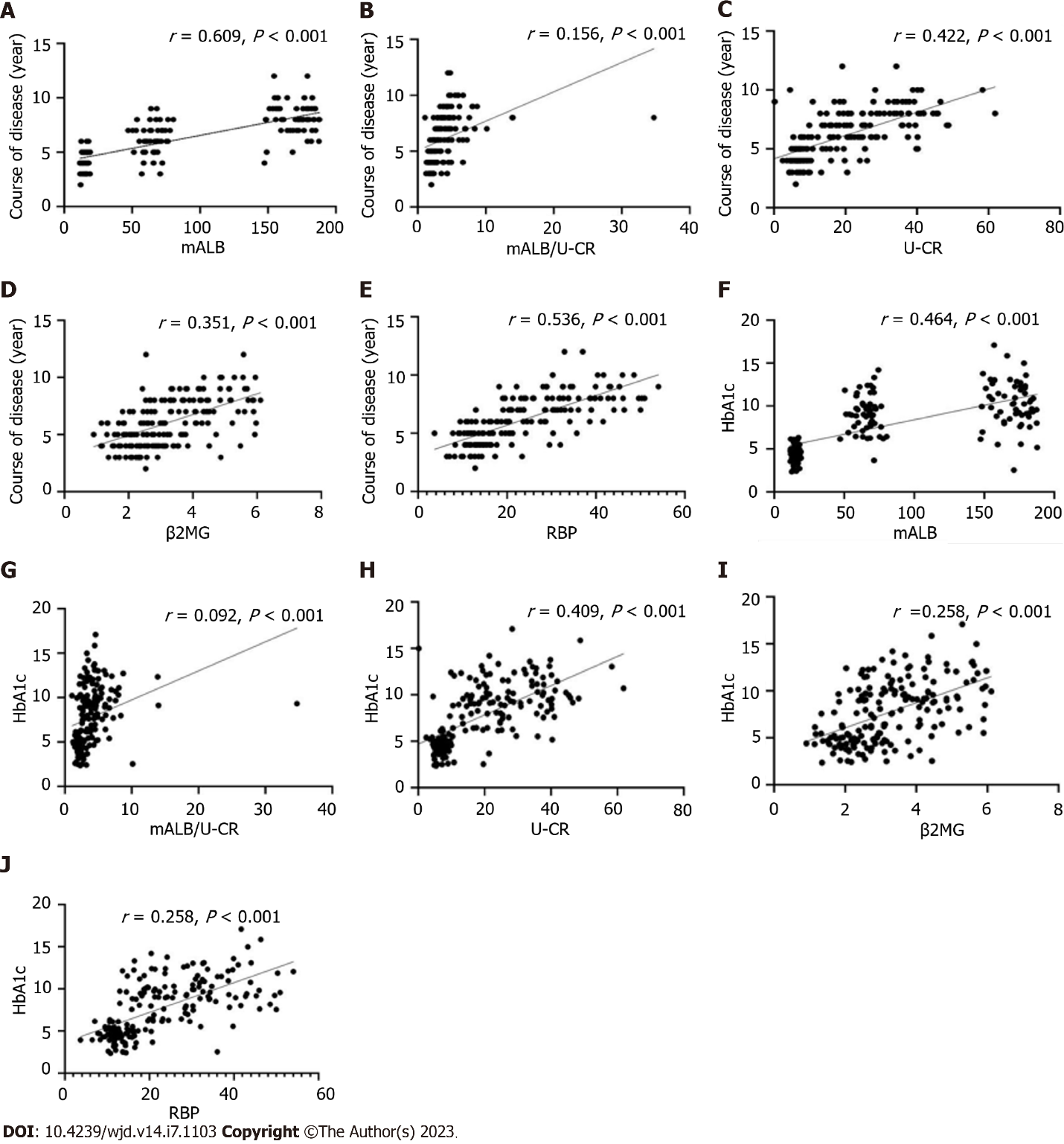
Grade C (Good): C

Grade D (Fair): 0

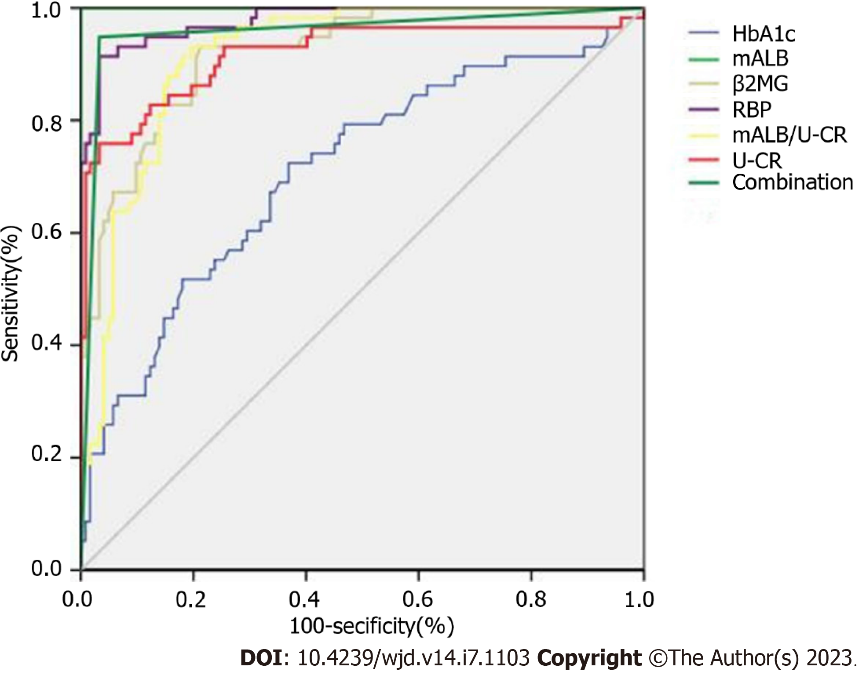
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**Figure Legends**

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**Figure 1 Correlation analysis.** A-E: The relationship between microalbumin (mALB), mALB/urinary creatinine (U-CR), U-CR, β2 microglobulin (β2MG), retinol binding protein (RBP) and course of disease; F-J: The relationship between mALB, mALB/U-CR, U-CR, β2MG, RBP and glycated hemoglobin A1c. U-mALB: Urinary microalbumin; U-CR: Urinary creatinine; β2MG: β2 microglobulin; RBP: Retinol binding protein; HbA1c: Glycated hemoglobin A1c.

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**Figure 2 Receiver operator characteristic curve analysis of glycated hemoglobin A1c, microalbumin, microalbumin/urinary creatinine, urinary creatinine, β2 microglobulin, and retinol binding protein for predicting proliferative diabetic retinopathy.** U-mALB: Urinary microalbumin; U-CR: Urinary creatinine; β2MG: β2 microglobulin; RBP: Retinol binding protein; HbA1c: Glycated hemoglobin A1c.

**Table 1 General information of the three groups (mean ± SD)**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Group** | **Age (yr)** | **Sex (M/F)** | **Duration of illness (yr)** | **Systolic blood pressure (mmHg)** | **Diastolic blood pressure (mmHg)** |
| NDR (*n*=68) | 57.71 ± 7.18 | 37/31 | 4.21 ± 0.81 | 117.47 ± 19.38 | 76.05 ± 9.48 |
| NPDR (*n*=54) | 58.00 ± 8.93 | 29/25 | 6.22 ± 1.26 | 118.32 ± 16.02 | 75.34 ± 11.91 |
| PDR (*n*=58) | 56.59 ± 7.12 | 31/37 | 8.12 ± 1.47 | 111.33 ± 18.09 | 75.69 ± 7.96 |
| *F/χ*2value | 0.534 | 0.013 | 169.133 | 2.606 | 0.178 |
| *P* value | 0.587 | 0.994 | < 0.001 | 0.078 | 0.836 |

NDR: Diabetes mellitus without retinopathy; NPDR: Non-proliferative diabeticretinopathy; PDR: Proliferative diabeticretinopathy.

**Table 2 Clinical indicators among the groups (mean ± SD)**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Group** | **HbA1c (%)** | **FPG (mmol/L)** | **TC (mmol/L)** | **TG (mmol/L)** | **LDL-C (mmol/L)** | **HDL-C (mmol/L)** |
| NDR (*n*=68) | 8.01 ± 1.86 | 8.60 ± 1.96 | 4.86 ± 0.98 | 1.68 ± 0.21 | 2.61 ± 0.42 | 1.15 ± 0.22 |
| NPDR (*n*=54) | 9.14 ± 2.12 | 8.55 ± 1.94 | 4.42 ± 0.75 | 1.69 ± 0.27 | 2.62 ± 0.41 | 1.24 ± 0.20 |
| PDR (*n*=58) | 10.28 ± 2.66 | 8.92 ± 2.16 | 4.55 ± 0.84 | 1.77 ± 0.29 | 2.74 ± 0.54 | 1.22 ± 0.27 |
| *F* value | 15.385 | 0.572 | 0.319 | 2.216 | 1.476 | 1.073 |
| *P* value | < 0.001 | 0.565 | 0.726 | 0.112 | 0.231 | 0.344 |

NDR: Diabetes mellitus without retinopathy; NPDR: Non-proliferative diabeticretinopathy; PDR: Proliferative diabeticretinopathy; HbA1c: Glycated hemoglobin A1c; FBG: Fasting blood glucose; HDL-C: High-density lipoprotein cholesterol; LDL-C: Low-density lipoprotein cholesterol; TC: Total cholesterol; TG: Triglyceride.

**Table 3 Comparison of microalbumin, microalbumin/urinary creatinine, urinary creatinine, β2 microglobulin and retinol binding protein levels in each group (mean ± SD)**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Group** | **mALB (mg/L)** | **mALB/U-CR (mg/mmoL)** | **U-CR (μmol/L)** | **β2MG (mg/L)** | **RBP (μg/L)** |
| NDR (*n*=68) | 15.04 ± 1.94 | 2.19 ± 0.86 | 6.86 ± 1.67 | 2.28 ± 0.66 | 12.29 ± 2.82 |
| NPDR (*n*=54) | 65.69 ± 7.30 | 3.29 ± 1.26 | 19.97 ± 5.81 | 3.13 ± 0.84 | 21.58 ± 4.83 |
| PDR (*n*=58) | 170.29 ± 11.63 | 5.09 ± 1.02 | 33.35 ± 11.45 | 4.53 ± 0.97 | 36.78 ± 7.84 |
| *F* value | 147.103 | 121.668 | 206.027 | 117.619 | 69.460 |
| *P* value | < 0.001 | < 0.001 | < 0.001 | < 0.001 | < 0.001 |

NDR: Diabetes mellitus without retinopathy; NPDR: Non-proliferative diabeticretinopathy; PDR: Proliferative diabeticretinopathy; U-mALB: Urinary microalbumin; U-CR: Urinary creatinine; β2MG: β2 microglobulin; RBP: Retinol binding protein.

**Table 4 Multiple linear regression analysis of risk factors associated with the development of proliferative diabetic retinopathy**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Independent variable** | **B value** | **SE** | **β value** | ***t* value** | ***P* value** |
| Course of disease | 1.203 | 0.293 | 0.220 | 4.106 | < 0.001 |
| HbA1c | 0.942 | 0.192 | 0.755 | 4.906 | < 0.001 |
| mALB | 0.874 | 0.128 | 0.256 | 6.828 | < 0.001 |
| mALB/U-CR | 0.743 | 0.284 | 0.525 | 6.959 | < 0.001 |
| U-CR | 0.842 | 0.121 | 0.254 | 6.959 | < 0.001 |
| β2MG | 1.048 | 0.123 | 0.157 | 8.520 | < 0.001 |
| RBP | 1.262 | 0.184 | 0.215 | 3.271 | < 0.001 |

HbA1c: Glycated hemoglobin A1c; U-mALB: Urinary microalbumin; U-CR: Urinary creatinine; β2MG: β2 microglobulin; RBP: Retinol binding protein.

**Table 5 Receiver operator characteristic curve analysis of glycated hemoglobin A1c, microalbumin, microalbumin /urinary creatinine, urinary creatinine, β2 microglobulin, retinol binding protein for predicting proliferative diabetic retinopathy**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Item** | **Cut-off** | **Standard error** | **AUC** | **95%CI** | **Sensitivity (%)** | **Specificity (%)** |
| mALB | 56.84 mg/L | 0.040 | 0.641 | 0.530-0.688 | 68. 82 | 71.24 |
| mALB/U-CR | 2.45 mg/mmoL | 0.046 | 0.726 | 0.728-0.876 | 70.38 | 73.85 |
| U-CR | 25.96 μmol/L | 0.004 | 0.757 | 0.508-0.722 | 72.49 | 75.58 |
| β2MG | 3.18 mg/L | 0.027 | 0.748 | 0.637-0.882 | 76.84 | 79.84 |
| RBP | 26.58 μg/L | 0.036 | 0.807 | 0.637-0.882 | 82.48 | 79.38 |
| HbA1c | 9.05% | 0.043 | 0.710 | 0.638-0.775 | 72.41 | 63.11 |
| Combination | - | 0.017 | 0.958 | 0.917-0.982 | 94.83 | 96.72 |

AUC: Area under the curve; U-mALB: Urinary microalbumin; U-CR: Urinary creatinine; β2MG: β2 microglobulin; RBP: Retinol binding protein; HbA1c: Glycated hemoglobin A1c; Combination: Glycated hemoglobin A1c + microalbumin + microalbumin/urinary creatinine + urinary creatinine + β2 microglobulin + retinol binding protein.



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