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WID mainly publishes articles reporting research results and findings obtained in the field of diabetes and covering a wide range of topics including risk factors for diabetes, diabetes complications, experimental diabetes mellitus, type 1 diabetes mellitus, type 2 diabetes mellitus, gestational diabetes, diabetic angiopathies, diabetic cardiomyopathies, diabetic coma, diabetic ketoacidosis, diabetic nephropathies, diabetic neuropathies, Donohue syndrome, fetal macrosomia, and prediabetic state.

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EDITORIAL

### Recent therapeutic targets for the prevention and management of diabetic complications

Md Shahidul Islam, Lu Cai, Michael Horowitz

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### Abstract

Diabetes and associated complications represent major global public health issues which are associated with impaired quality of life and premature death. Although some diabetic complications have decreased in the developed world, the majority are still prevalent, with an increasing trend in the developing world. Currently used therapies are mainly 'glucocentric', focusing on the optimization of glycemic control to prevent, delay or manage diabetes-associated complications- other common comorbidities, such as dyslipidemia and hypertension are often underestimated. Although a number of novel therapeutic approaches have been reported recently, some of them have not received comparable attention in relation to either further studies or potential clinical implementation. This editorial briefly discusses some recent therapeutic approaches to the prevention and management of diabetes and its associated complications, as well as potential directions for future research and development in this area.

Key Words: Diabetic complications; Oxidative stress; Phytochemicals; Zinc; Silent information regulator 1; FOXO; Micro RNA

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**Core Tip:** 'Glucocentric' approaches are currently being used for the management of diabetes and its associated complications. This articles highlighted some recent therapeutic approaches for the management of diabetes and its associated complications such as the management of oxidative stress by using antioxidative phytochemicals, molecular cell signaling pathways *via* Silent information regulator 1 and FOXOs and micro RNAs.

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### INTRODUCTION

Diabetes and associated complications are major global causes of premature mortality. A minimum of 50% of people who have type 2 diabetes face premature death from diabetes-associated cardiovascular diseases and some 10% from renal failure with a total of 3.8 million deaths per annum[1]. According to a recent review, the rates of myocardial infarction, stroke and limb amputation have decreased among people with diabetes with a concomitant decline in mortality. The majority of these data are, however, sourced from high income countries, when other diabetic complications such as nephropathy, retinopathy and cancers are well represented[2]. It has been reported that the prevalence of diabetic complications are much higher in low and middle-income countries with a range of 12%-16% for microvascular and 2%-6% for macrovascular complications[3].

Diabetes-associated microvascular complications occur frequently in individuals with diabetes, both their prevalence and severity are inversely proportional to the efficacy of management of hyperglycemia. At least 50% of diabetic patients have one or more diabetic complications in their lifetime and many have multiple complications. Microvascular complications such as, diabetic nephropathy, retinopathy, neuropathy and diabetic foot disease represent a major causes of morbidity, impaired quality of life and mortality and are more common than macrovascular complications, such as diabetic cardiomyopathy and peripheral vascular diseases<sup>[4]</sup>. While improved management of hyperglycemia represents a major approach to prevent or delay diabetic complications, currently available therapies are not consistent in maintaining optimum glycemic control, their efficacy in glucose lowering exhibits a substantial interindividual variation and their long-term use is associated with adverse effects<sup>[5]</sup>. Additional major challenges with currently available therapies include, but not limited to, optimizing the dose to control the blood glucose, blood pressure and lipids as well as self-management of diabetes and lifestyle[6]. Hence, there is a need for newer or alternative therapies not only for better glycemic control, but also for the management of blood pressure and blood lipids with an ultimate goal for the prevention of diabetes associated micro- and macro-vascular complications. The outcomes of recent, large cardiovascular prevention trials, initially mandated for regulatory purposes, have provided major insights into the need for a broader, rather than simply 'glucocentric', approach to therapy of type 2 diabetes. Both GLP-1 receptor agonist and SGLT-2 inhibitors are now used widely with the recognition that their beneficial effects include cardiovascular and renal protection[7]. A number of novel therapeutic approaches are currently being evaluated with the potential to improve the prevention and management of diabetic complications, some of which are highlighted below.

Oxidative stress is a major culprit for the induction of diabetic complications[8], since it causes endothelial dysfunction both in small and large vessels, not only by increasing the production of oxidative free radicals and advanced glycation end products, but also by the concomitant reduction of physiological antioxidative status. Over expression of the antioxidative enzyme, superoxide dismutase (SOD) in transgenic diabetic mice has been shown to prevent diabetic micro and macrovascular complications[9]. Accordingly, over expression of antioxidative enzymes, such SOD and catalase, may represent a therapeutic approach to the reduction of diabetic complications, however, the level of over expression also needs to be optimized in order to avoid additional complications[10].

Polyphenols, flavonoids, phenolic acids and zinc have recently been shown to have potent beneficial effects in relation to hyperglycemia, diabetes and its associated complications. Curcumin, the major bioactive compound of turmeric and its analogues, has anti-inflammatory, antioxidant, anti-tumor and epigenetics modulatory effects with potential efficacy against diabetic complications[11,12]. Depletion of zinc in diabetes increases oxidative stress while zinc supplementation has been shown to have a hypoglycemic, antioxidant effect and alleviates some diabetes-associated complications[13]. Resveratrol, a key bioactive compound derived from red grapes, has been shown to have number of benefits including on glycemic control and the management of diabetic complications[14]. Furthermore, nanotechnology or nano-formulations of polyphenols, flavonoids and phenolic acids has the potential to enhance solubility, and intestinal absorption, as well as bioavailability and, therapeutic efficacy in diabetes and its associated complications[15].

Silent information regulator 1 (SIRT1), a member of the sirtuins family when the sirtuins are NAD<sup>+</sup> dependent histone deacetylase. Apart from activating LKB1 mediated AMPK followed by PGC $\alpha$ , PPAR $\alpha$ , eNOS pathways and inhibiting mTOR and NOX or NADPH oxidase pathways[16]; SIRT1, has been reported to regulate the activity of other proteins, such as forkhead box protein of class O or FOXO, which regulates oxidative stress resistance, insulin signaling and metabolism along with its other activities as a transcription factor[17]. Of the many FOXOs, FOXO1 is widely expressed in muscle, liver and pancreas and protects pancreatic  $\beta$ -cells from oxidative stress by increasing the expression of antioxidant genes[18]. On the other hand, there is evidence that FOXO3 can prevent atherosclerosis *via* inhibiting smooth

muscle cell proliferation and activation[19]. FOXO1 and FOXO3 are also involved in many other mechanisms of relevance to glucose metabolism, as well as diabetic complications. Hence, SIRT1 and FOXO1 and FOXO3 may also represent therapeutic targets for the management of diabetic complications.

Like many other molecular pathways, epigenetic factors, including histone modifications, DNA methylations and noncoding RNAs play a major role in the pathogenesis of diabetes and its complications<sup>[20]</sup>. Among many non-coding RNAs, some micro-RNAs have been shown to have a pivotal role in the management of diabetes and diabetic complications, particularly in relation to the diagnosis and prognosis of prevalent microvascular complications e.g. diabetic neuropathy. A number of microRNAs are involved in the signaling pathways of diabetic complications, which can be targeted for the early diagnosis and development of therapeutics for diabetic microvascular complications, particularly for diabetic neuropathy and diabetic foot disease[21].

### CONCLUSION

In conclusion, although many other therapeutic targets are being investigated for the improved management of diabetes and its associated complications, the approach of reducing oxidative stress or increasing antioxidant status using antioxidant phytochemicals or bioactive compounds and mineral such as zinc; molecular metabolic pathways such as SIRT1 and FOXOs and micro RNAs represent important and novel approaches to the diagnosis, prevention and improved management of diabetic complications. We look forward to the outcomes of these ongoing studies, which will be facilitated by an effective collaboration between basic scientists, clinicians and pharma and, hopefully, their prompt translation to clinical practice.

### FOOTNOTES

Author contributions: Islam MS conceptualized and drafted the initial manuscript and revised after receiving reviewers' comments; Cai L and Horowitz M revised and edited the original and revised manuscript before submission.

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MINIREVIEWS

### MicroRNA-155 mediates endogenous angiotensin II type 1 receptor regulation: implications for innovative type 2 diabetes mellitus management

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<b>P-Reviewer:</b> Cai L. United States:	Abstract		
Cen LS, China; Moreno-Gómez-	Type 2 diabetes mellitus (T2DM) is a lifelong condition and a threat to human		
Toledano R, Spain; Papadopoulos	health. Thorough understanding of its pathogenesis is acutely needed in order to		
VP, Greece	devise innovative, preventative, and potentially curative pharmacological		
Received: April 14, 2023	interventions. MicroKNAs (miKNA), are small, non-coding, one-stranded KNA		
Peer-review started: April 14, 2023	translational repression MiR-155 is an ancient evolutionarily well-conserved		
<b>First decision:</b> June 13, 2023	miRNA, with distinct expression profiles and multifunctionality, and a target		
<b>Revised:</b> June 18, 2023	repertoire of over 241 genes involved in numerous physiological and pathological		
Accepted: July 13, 2023	processes including hematopoietic lineage differentiation, immunity, inflam-		
Article in press: July 13, 2023	mation, viral infections, cancer, cardiovascular conditions, and particularly		
Published online: September 15	diabetes mellitus. MiR-155 Levels are progressively reduced in aging, obesity,		
2023	sarcopenia, and T2DM. Thus, the loss of coordinated repression of multiple miR-		
	155 targets acting as negative regulators, such as <i>C</i> / <i>EBP</i> $\beta$ , <i>HDAC</i> 4, and <i>SOCS</i> 1		
	impacts insulin signaling, deteriorating glucose homeostasis, and causing insulin		
	resistance (IR). Moreover, deranged regulation of the renin angiotensin aldo-		
	sterone system (KAAS) through loss of Angiotensin II Type 1 receptor downregu-		
	lation, and negated repression of ETS-1, results in unopposed detrimental		

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Angiotensin II effects, further promoting IR. Finally, loss of BACH1 and SOCS1 repression abolishes cytoprotective, anti-oxidant, anti-apoptotic, and anti-inflammatory cellular pathways, and promotes  $\beta$ -cell loss. In contrast to RAAS inhibitor treatments that further decrease already reduced miR-155 Levels, strategies to increase an ailing miR-155 production in T2DM, *e.g.*, the use of metformin, mineralocorticoid receptor blockers (spironolactone, eplerenone, finerenone), and verapamil, alone or in various combinations, represent current treatment options. In the future, direct tissue delivery of miRNA analogs is likely.

**Key Words:** Angiotensin II; Angiotensin II type 1 receptor; Arginase 2; L-type calcium channel; Mineralocorticoid receptor; MiRNA-155; Renin-angiotensin aldosterone system; Type 1/2 diabetes mellitus; Verapamil

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**Core Tip:** MicroRNAs (miRNA) are small, non-coding, one-stranded RNA molecules that can target and silence over 60% of human genes thereby effectively regulating huge genetic networks. MiRNAs are abundantly found in every human cell and their production is tightly controlled. They play critical roles in regulating almost every cellular pathway, numerous human diseases, and have been linked to the development of diabetes mellitus (DM) and the regulation of blood pressure. In this minireview, we comment on crucial miR-155 effects in type 2 DM (T2DM). Deeper mechanistical understanding of this miRNA's permeating action may lead to innovative therapeutic approaches in T2DM.

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### INTRODUCTION

Diabetes mellitus (DM), until recently considered a lifelong and irreversible condition, is a devastating burden to over half a billion people worldwide[1]. The overwhelming majority of diabetic patients-over 90%-suffer from Type 2 DM (T2DM) caused by an intricate interaction between lifestyle and genetics that through insulin resistance (IR) lead to metabolic syndrome, pre-diabetes, failure of insulin-secreting pancreatic  $\beta$ -cells and ultimately overt disease[2,3]. Uncontrolled T2DM eventually progresses to a myriad of severe health complications [among which cardiovascular disease (CVD), chronic renal failure, and hypertension (HT)], and to an early death[1]. The syndemic of coronavirus disease 2019 and T2DM has affirmed the latter's lethal effect[4]. Ominous future predictions estimate the number of DM-afflicted individuals to be over 800 million by 2045, up from the current 500 million[1]. Increased understanding of the T2DM pathogenesis is, therefore, acutely needed in order to devise innovative, preventative and potentially curative pharmacological interventions[5].

The pathophysiological role of the renin angiotensin aldosterone system (RAAS) and its major effector, Angiotensin II (Ang II) through the Ang II Type 1 receptor (AT1R), in the development of IR in T2DM have long been recognized[6]. Furthermore, convincing evidence exists advocating the use of RAAS inhibition, ACE inhibitors (ACEi) or AT1 receptor antagonists/blockers (ARB), in patients with T2DM, not only for proteinuria and HT, but also as a means to improve IR and glucose homeostasis[6].

MicroRNAs (miRNAs or miRs) are small (21-25 nucleotides), non-coding RNAs, able to translationally repress and downregulate gene expression[7]. Present abundantly in all human cells, miRNAs are endogenously biosynthesized through a strictly regulated process that will ultimately result in a mature miRNA, with a 2-8 nucleotide long seed sequence in its 5'untranslated region (UTR), that will bind to a target messenger RNA (mRNA). If the miRNA seed sequence binds perfectly to the corresponding 3'UTR of a specific mRNA, the latter will be recruited to be degraded by an RNA silencing complex. If the binding is incomplete, mRNA translational machinery will be blocked, thereby inhibiting protein translational efficiency, and repressing (silencing) gene expression[7]. As a specific miRNA can target multiple mRNA molecules, and equally, a single mRNA molecule can bind to multiple miRNAs, the host can modulate response feedback, through regulatory gene networks, in a concerted effort to control diverse aspects of cellular processes[7]. In this minireview, we present additional miRNA- modulated pathways that can modulate AT1R and Ang II effects that are of importance for the pathogenesis of IR, T2DM, and the development of cardiovascular and renal diabetic complications.

MiR-155 is of particular interest as it is intricately involved both in the pathogenesis of DM and the regulation of AT1R and Ang II effects (Figure 1)[6,8-12]. First identified in 1997, miR-155 is a highly conserved and ancient miRNA primarily expressed in the thymus and spleen. It exhibits unique expression profiles and multifunctionality but is minimally detected under normal physiological conditions[13]. With a target repertoire of over 241 genes, miR-155 plays critical roles in various physiological and pathological processes, such as hematopoietic cell line differentiation, inflammation, immunity (especially viral and parasitic infections), cancer, cardiovascular conditions, and notably, DM (Table 1)[5,8,12-25].

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#### Table 1 Direct gene targets of microRNA-155 relevant to type 2 diabetes mellitus

Gene symbol	Full gene name	Action
AGTR1	Angiotensin II type 1 receptor gene	Repressed translation downregulates gene expression mediating endogenous AT1R antagonism[9,10,21]. Human <i>in-vitro</i> and <i>in-vivo</i> studies
ARG2	Arginase-2	Repressed translation prevents L-arginine depletion, supports dendritic cell maturation, and negates lung pathologies[22,23]. Human and mouse <i>in-vitro</i> and <i>in-vivo</i> studies
BACH1	BTB and CNC homology 1, basic leucine zipper transcription factor 1	Translational repression of <i>BACH1</i> leads to potent anti-inflammatory, cytoprotective, antioxidant programs through Heme Oxygenase-1[12]. Review of human <i>in-vitro</i> and <i>in-vivo</i> studies
С/ЕВРβ	CCAAT/enhancer-binding protein $\beta$	Repression downregulates <i>Pyruvate Kinase</i> 4 ( <i>PDK</i> 4) gene expression and negatively regulates Pyruvate kinase complex (PDC) activity, thereby improving glucose utilization [16]. Mouse <i>in-vitro</i> and human <i>in-vivo</i> studies
ETS-1	E26 Transformation-specific Sequence-1	Translational repression averts Ang II effects involving gene regulation of vascular remodeling, angiogenesis, and inflammation[9,10,24]. Review of human <i>in-vitro</i> and <i>in-vivo</i> studies. Mouse <i>in-vitro</i> and <i>in-vivo</i> studies
HDAC4	Histone deacetylase 4	Its repression increases GLUT4 and enhances glucose uptake in insulin-sensitive tissues, <i>i.e.</i> , skeletal muscle [16]. Mouse <i>in-vitro</i> and human <i>in-vivo</i> studies
CACNA1C (Cav1.2)	L-type calcium channel subunit, LTCC	As a subunit of the L-type calcium channel, this pro-constrictive gene contributes to influx of calcium in vascular smooth muscle cells and reactive oxygen species production, thereby mediating the important components of vascular aging: Vasoconstriction and vascular oxidative stress[21]. Human <i>in-vitro</i> and <i>in-vivo</i> studies
SOCS1	Suppressor of cytokine signaling 1	Repression prevents the degradation of IRS-1 (Insulin Receptor Substrate-1) protein that mediates the effect of insulin in muscle, liver, and adipose tissue. Supports the JAK2/Y343/STAT5 pathway through which the protective effects of EPO against ischemic injury are mediated[16,25]. Human <i>in-vivo</i> study. Mouse <i>in-vitro</i> and <i>in-vivo</i> study

AT1R: Angiotensin II Type 1 receptor; Ang II: Angiotensin II; LTCC: L-type calcium channel; EPO: Erythropoietin; ROS: Reactive oxygen species; JAK2: Janus kinase 2; STAT5: Signal transducer and activator of transcription 5.

In T2DM, miR-155 Levels in plasma, peripheral blood cells, platelets, and urine are significantly and consistently decreased, with surprising congruence between different ethnicities[8]. Ranging from obesity to IR to diabetic complications in T2DM, miR-155 Levels are progressively reduced[8,14,15,17]. MiR-155's underlying molecular mechanism in enhancing insulin signaling, improving glucose homeostasis, and alleviating IR in T2DM, occurs partly through the coordinated repression of multiple negative regulators, such as *CCAAT/enhancer-binding protein*  $\beta$  (*C/EBP* $\beta$ ), *Histone Deacetylase* 4 (HDAC4), and *Suppressor of cytokine signaling* 1 (SOCS1) (Table 1)[16]. MiR-155-mediated *C/EBP* $\beta$  repression downregulates *Pyruvate Kinase* 4 (PDK4) gene expression and negatively regulates Pyruvate kinase complex activity, thereby improving glucose utilization[16]. HDAC4 repression increases GLUT4 and enhances glucose uptake in insulinsensitive tissues, *i.e.*, skeletal muscle, while *SOCS1* repression prevents the degradation of Insulin Receptor Substrate-1 (IRS-1) protein that mediates the effect of insulin in muscle, liver, and adipose tissue (Figure 1 and Table 1)[16].

Aging, obesity, sarcopenia, chronic RAAS activation, and IR, invariably predate the development of T2DM[26]. Shared miRNA signatures have been reported, highlighting the central role of miR-155 in the common pathogenesis of those conditions (Figure 1)[8,14,26]. One particularly important observation is the activation of the classical RAAS axis arm that involves Ang II/AT1R signaling in aging skeletal muscle and white adipose tissue (WAT), both fundamentally involved in T2DM pathogenesis[26,27]. In WAT, a chronically activated RAAS axis increases lipogenesis and reduces lipolysis, while in the aging skeletal musculature RAAS hyperactivity promotes protein degradation, and sarcopenia, altogether ultimately leading to oxidative stress, inflammation, fat accumulation, muscle atrophy, and IR[26,27]. In addition, RAAS's protective arm, involving Ang 1-7/AT2R/MasR signaling, is inhibited at the same time, further augmenting an unfavorable AT1R/AT2R imbalance[26,27]. MiR-155, acting as a master regulator, is the key player in chronic RAAS/Ang II/AT1R activation, thereby, intricately associated with the development of IR[6]. Through its repression of the AGTR1 (the gene that codes for the AT1R) miR-155 regulates the homeostasis of the AT1R receptor, its membrane presence, and thus the biological activity of Ang II (Table 1)[9,10,28-30]. Moreover, its regulation of the E26 Transformation-specific Sequence-1 (ETS-1) averts several detrimental vascular Ang II effects involving gene regulation of inflammation, proliferation, remodeling, fibrosis, and angiogenesis (Table 1)[9,13,24]. Furthermore, its repressive effects on Arginase-2 (ARG2) prevent the depletion of l-arginine, the obligate substrate of endothelial nitric oxide (NO) synthase (eNOS), improving substrate availability and further increasing NO-production and NO-bioavailability that further support NO-dependent cardio- and renoprotection in T2DM (Table 1)[13,23]. From the sum of these actions, it is thus evident that the reported loss of miR-155 in T2DM has profound effects leading to persistent RAAS hyperactivity through chronic Ang II stimulation of the AT1R, thereby exerting its detrimental, pro-oxidant, pro-fibrotic, proliferative, and pro- inflammatory actions (Figure 1 and Table 1). Additional miR-155 effects through repressive actions on BTB and CNC homology 1, basic leucine zipper transcription factor 1 (BACH1) and SOCS1, synergistically enhance cytoprotective, anti-oxidant, antiapoptotic, and anti-inflammatory cellular pathways and promote a protective cellular milieu, which is subsequently lost

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**Figure 1 Schematic depiction of coordinated repression of multiple miR-155 targets relevant for T2DM.** Translational repression of *AGTR1*, *ARG2*, *CACNA1C*, and *ETS-1* reshapes RAAS towards cardio-, vasculo-, and renoprotective phenotypes. *BACH1* and *SOCS1* repression promotes cytoprotective phenotypes and preserves β-cell function. *C/EBPβ*, *HDAC4*, and *SOCS1* repression improves glucose homeostasis, enhances insulin signaling, and reverses insulin resistance. Aging, obesity, sarcopenia, AT1R 1169C SNP, and ACEi/ARB treatment negatively impact miR-155 levels and/or function while MR antagonists, metformin, GLP-1 agonists, and verapamil exert beneficial effects. Red arrows or lines represent downregulation, lower Level, inhibition, repression. Green arrows or lines represent increased Level or stimulatory/beneficial action. ACEi: Angiotensin-converting enzyme inhibitors; *AGTR1*: Angiotensin II type 1 receptor gene; Ang II: Angiotensin II; ARB: Angiotensin II type 1 receptor blockers; *ARG2*: Arginase 2; AT1/2R: Angiotensin II type 1/2 receptor; *BACH1*: BTB and CNC homology 1, basic leucine zipper transcription factor 1; *CACNA1C (Cav1.2)*: L-type calcium channel subunit; *C/EBPβ*: CCAAT/enhancer-binding protein β; eNOS: Endothelial nitric oxide synthetase; EPO: Erythropoietin; *ETS-1*: E26 Transformation-specific Sequence-1; GLP-1: Glucagon-like peptide 1; GLUT4: Glucose transporter type 4; HO-1: Heme oxygenase 1; *HDAC4*: Histone Deacetylase 4; IRS-1: Insulin receptor substrate-1; LTCC: L-type Calcium Channel; MasR: Mas Receptor; MicroRNA-155: MiR-155; MR: Mineralocorticoid receptor; NO: Nitric oxide; RAAS: Renin-Angiotensin Aldosterone System; ROS: Reactive oxygen species; *SOCS1*: Suppressor of cytokine signaling 1; SNP: Single nucleotide polymorphism; T2DM: Type 2 Diabetes Mellitus.

following miR-155 downregulation (Table 1)[12,13]. Genetic variants that perturb miR-155's action (such as in carriers of AT1R + 1166C-allele) or that increase its synthesis (such as in trisomy 21 and the rs767649 polymorphism of miR-155) biochemically and molecularly demonstrate this central significance of miR in a plethora of DM-associated pathological conditions[11,18,31-33]. Moreover, clinical data in obese individuals demonstrate that miR-155 Levels correlate with improved insulin sensitivity post-bariatric surgery and are critical in mediating the effects of endurance exercise[34,35].

While miR-155 is consistently reduced in serum and tissues in T2DM, it is reported to be upregulated in Type 1 DM (T1DM), highlighting T1DM's autoimmune pathogenesis and miR's crucial and differential role in autoimmunity and innate and adaptive immunity[8,36]. However, even if robustly elevated in newly diagnosed T1DM, miR-155 strikingly diminishes within 5 years of diagnosis[32].

AT1R substrate modulation (ACEi) and/or receptor inhibition (ARBs) may improve glucose homeostasis[6]. However, strategies to increase an ailing miR-155 production in T2DM could prove to be a more appropriate course of action (Figure 1). Metformin with ACEi/ARB improves HbA1c goals[6]. Metformin and the newer Glucagon Like Peptide 1 (GLP-1) analogs have been shown to repress SOCS1 and 3 and increase IRS-1[37]. Metformin mediates miR-155 increases that repress SOCS1 and reduce NF-κB (nuclear factor κB), thereby disrupting NF-κB-mediated high-fat induced inflammatory effects in T2DM[38,39]. The clinical effects of GLP-1 analogs on miR-155 in humans are, to date, unknown, and additional research is needed, but miR-155 has been shown to promote GLP-1 production in the murine pancreas[40]. Moreover, in the resistance vessels of aging humans, elevated expression of mineralocorticoid receptor (MR) is accompanied by a decrease in miR-155 Levels and an upregulation of miR-155 targets such as the CACNA1C (Cav1.2) gene [a subunit of the L-type calcium channel (LTCC)], and the AGTR1 gene. These alterations in gene expression play a role in promoting vasoconstriction and oxidative stress in aging mice (Table 1)[21]. MR inhibition reverses and reinstates the significantly low basal serum miR-155 Levels in the aging blood vessels and blocks two interactive steps involving LTCC and AGTR1 that underlie the pathogenesis of HT[13]. A correlation between improved blood pressure response to therapy with MR antagonists and changes in miR-155 Levels in older individuals has been reported [21]. Moreover, the use of MR-antagonists (spironolactone, eplerenone, finerenone) has shown renal and cardiovascular benefits in T2DM[41-43]. LTCC blockade per se, through verapamil alone, or in combination with MR antagonists/metformin, will offer additional therapeutic options in T2DM[44]. Besides improved blood pressure regulation and cardio-renal protection, verapamil demonstrates additional benefits while avoiding many of the common adverse effects associated with ACEi/ ARB[45]. Verapamil's mode of action is of particular interest in diabetes[46]. Apart from being present in cardiomyocytes,

LTCCs are also present in pancreatic  $\beta$ -cells and participate in insulin homeostasis[47]. In the heart and the pancreas, effective pharmacological LTCC blockage can inhibit the expression of pro-apoptotic thioredoxin-interacting protein, a significant contributor to pancreatic β-cell dysfunction and a key gene regulated in response to hyperglycemia, thereby promoting the survival and proper functioning of  $\beta$ -cells and improving glucose homeostasis[46,48]. Verapamil has, thus, the potential not only to enhance  $\beta$ -cell survival and function, but also improve and even prevent overt diabetes of both types[48,49]. In a recent study, verapamil combined with metformin, significantly improved glycemic control in T2DM [49]. Finally, a drawback in the use of monotherapy as ACEi/ARBs (in conditions that already are associated with low miR-155 Levels) is that they significantly further decrease already reduced miR-155 Levels[50,51]. RAAS inhibition could, thus, theoretically deprive T2DM patients of additional miR-155-engendered favorable immunological and cytoprotective effects and potentially explain ACEi's modest and ARBs' non-existent effects in preventing CVD or improving glycemic indices in DM and HT (Figure 1)[13,50-53].

### CONCLUSION

The data presented above strongly support the role of miR-155 as a major player in the pathogenesis of T2DM and complications, by triggering IR and  $\beta$ -cell loss as well as through RAAS modulatory effects (Figure 1)[5,8]. Large multicenter trials are required to establish this role of miRNA as a reliable biomarker and potential therapeutic target in DM. Then, as increased mechanistic knowledge regarding miR-155 becomes available, novel miRNA-modulating approaches with miR-155 as a target are likely in T2DM. Even though these therapeutic modalities are still in their infancy and might yet be far from the clinic, research must address this knowledge gap in order to devise how to effectively deliver specific, synthetic miRNA mimics (T2DM, aging, obesity, sarcopenia) or inhibitors-antagomiRs (T1DM, cancer), to a specific tissue, in the diabetic patient, as miR-155 actions are tissue-sensitive [54]. In addition, a better understanding is needed on how several miRNAs work synergistically on the same mRNA targets and how miRNA networks function. As disease-specific miRNA expression pattern is ubiquitous in all related tissues, it can prove challenging in a complex disease like DM to accomplish precise delivery to certain tissues/organs and avoid adverse offtarget effects in others[5].

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### **FOOTNOTES**

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MINIREVIEWS

### Hypothesis that alpha-amylase evokes regulatory mechanisms originating in the pancreas, gut and circulation, which govern glucose/insulin homeostasis

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Received: April 24, 2023	Abstract			
Peer-review started: April 24, 2023 First decision: June 14, 2023	The anti-incretin theory involving the abolishment of diabetes type (DT) II by some of methods used in bariatric surgery, first appeared during the early years			
<b>Revised:</b> June 28, 2023	of the XXI century and considers the existence of anti-incretin substances.			
Accepted: August 2, 2023	However, to date no exogenous or endogenous anti-incretins have been found.			
Article in press: August 2, 2023	Our concept of the acini-islet-acinar axis assumes that insulin intra-pancreatically			
Published online: September 15, 2023	stimulates alpha-amylase synthesis ("halo phenomenon") and in turn, alpha- amylase reciprocally inhibits insulin production, thus making alpha-amylase a			



candidate for being an anti-incretin. Additionally, gut as well as plasma alphaamylase, of pancreatic and other origins, inhibits the appearance of dietary glucose in the blood, lowering the glucose peak after iv or oral glucose loading. This effect of alpha-amylase can be interpreted as an insulin down regulatory mechanism, possibly limiting the depletion of pancreatic beta cells and preventing their failure. Clinical observations agree with the above statements, where patients with high blood alpha-amylase concentrations are seldom obese and seldom develop DT2. Obese-DT2, as well as DT1 patients, usually develop exocrine pancreatic insufficiency (EPI) and vice versa. Ultimately, DT2 patients develop DT1, when the pancreatic beta cells are exhausted and insulin production ceases. Studies on biliopancreatic diversion (BPD) and on BPD with duodenal switch, a type of bariatric surgery, as well as studies on EPI pigs, allow us to observe and investigate the above-mentioned phenomena of intra-pancreatic interactions.

**Key Words:** Pancreas; Alpha-amylase; Acini-islet-acinar axis; Hyperglycaemia; Bariatrics; Insulin; Incretins; Glucosedependent insulinotropic polypeptide; Glucagon-like peptide-1; Exocrine pancreatic insufficiency; Pancreatic enzyme therapy; Diabetes

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**Core Tip:** The concept of the acini-islet-acinar axis postulates that insulin stimulates amylase synthesis and amylase in turn inhibits insulin production. This regulation is of particular significance with regards to postoperative glucose regulation after bariatric bypass surgery. The interaction of salivary amylase at the alimentary limb and pancreatic amylase, exclusively at the common channel, results in an immediate metabolic effect that leads to a significant reduction in incretin-induced hyperinsulinaemia and a marked improvement in glucose action on insulin production. As a result, both pancreatic beta cells and acinar cells are effectively protected against depletion.

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### INTRODUCTION

Studies on the dependency of pancreatic enzyme synthesis on insulin release started 60 years ago and resulted in the discovery of the so-called "halo phenomenon" [1-5]. For several years we have explored the 'reverse arm' of regulation; namely the dependency of insulin release on pancreatic enzyme secretion and the role of the interaction between these parameters on glucose metabolism [6]. Our studies indicate the local and peripheral dependency of insulin secretion on pancreatic alpha-amylase. Alpha-amylase, beyond its digestive functions, actively participates in glucose metabolism, both *via* inhibition of glucose absorption from the gut[7,8] and by lowering insulin release [9].

It should be mentioned that other pancreatic-like enzymes, such as proteinases and lipases of microbial origin, were tested with respect to their effect on insulin secretion and glucose utilization. We have shown that lipase does not affect the afore mentioned parameters, while proteinase decreases the insulin sensitivity index and stimulates insulin release[6].

The above-described alpha-amylase-dependent inhibition of glucose absorption, through its storage or metabolism in the gut tissues could reduce insulin release, making the secretion adequate to glucose levels entering the circulation. Thus, there appears to be some sort of intra-pancreatic down regulation effect of alpha-amylase on insulin secretion and/ or synthesis. Insulin stimulates the secretion of alpha-amylase and other pancreatic enzymes, while alpha-amylase inhibits insulin release[10]. In fact, chronic hyperinsulinemia could lead not only to insulin resistance, but also to the destruction of pancreatic acinar cells, with the further development of exocrine pancreatic insufficiency (EPI)[11].

The above observations were confirmed in several *in vitro* and *in vivo* studies in our lab[9,10,12-14]. Both the alphaamylase-insulin reciprocal acini-islet-acinar (AIA) local interactions, as well as the peripheral interactions originating in the gut and blood, which attenuate and downregulate insulin secretion, belong to the postulated new type of regulations where alpha-amylase exhibits a hormonal-like action. In 1976, Arnesjö and Lundquist[15] demonstrated a decrease in insulin release and the improvement of glucose tolerance in rats with long-term CCK-PZ stimulated pancreatic enzymes secretion[15]. We noted the stronger suppression of insulin release after pancreatin administration in the meal glucose tolerance test (44%), when compared to the intravenous glucose tolerance test (31%)[13]. In the same year, Lindqvist *et al* [16] observed a decrease in levels of active glucagon-like peptide-1 (GLP-1) after Roux-and-Y gastric bypass (RYGB) surgery in pigs[16], and in early 2018 our group showed a consequential increase in plasma amylase activity in the EPI porcine model[14]. We recognized that the effects of amylase on the regulation of insulin secretion are para-hormonal and considering all our observations, we thought it would be suitable to locate the anti-incretin action of amylase in the same place where incretins *e.g.*, GLP-1 usually act.

Reciprocal AIA regulations are likely species- and individual-dependent, thus they are in some way immanent. We recognize that dietary factors *e.g.*, carbohydrates (glucose), acting from the duodenum, stimulate insulin and alphaamylase secretion in a species-specific manner and in individuals' adequate amount in islands and acini surrounding them. Previous studies have shown that an iv infusion of glucose does not stimulate the secretion of pancreatic alphaamylase[17]. Considering the observed counteraction of alpha-amylase to glucose-dependent insulinotropic polypeptide (GIP)-1 we used the term anti-incretin to describe metabolic effects of alpha-amylase. This "anti-incretin" hypothesis needs to be confirmed by future studies, with specific focus on enteral-insular regulatory mechanisms. In fact,

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comparative studies on the exclusive effect of pancreatic alpha-amylase on insulin secretion, stimulated by oral or intravenous glucose administration, have not yet been carried out. The effect of alpha-amylase on the secretion of incretins (GLP-1 and GIP) has also not yet been assessed to date.

Incretin-dependent regulation of insulin secretion is coupled with the intake of "quick" energy in the form of carbohydrates and thus could be recognized as temporal. These regulatory mechanisms need to be abruptly inhibited after the digestion of simple, 'easily-digested' carbohydrates in order to prevent the overproduction of insulin by beta cells and thus beta cell exhaustion. The overlooked alpha-amylase dependent downregulation of insulin levels protects both pancreatic beta and acinar cells from metabolic failure.

### BILIOPANCREATIC DIVERSION SURGERY REVEALS THE INHIBITORY EFFECT OF ALPHA-AMYLASE ON INSULIN SECRETION

The clinical efficacy of bariatric surgery has encouraged scientific investigation of the gut as a major endocrine organ. Manipulation of gastrointestinal anatomy, through bariatric surgery, has profound effects. Even though these procedures were designed to restrict food intake and increase nutrient malabsorption, evidence suggests that the contribution of the above-mentioned factors to weight loss is minimal. Instead, these interventions reduce body weight gain by decreasing hunger, increasing satiation during a meal, changing food preferences and energy expenditure. GLP-1 analogues, currently used as a treatment for diabetes type 2 (DT2) or slimming injections, are now recognized as alternatives to bariatric surgery in terms of their weight loss and anti-diabetic effects. However, not much is known about their safety with regards to beta cell function in the long run, since the primary target of GLP-1 analogues is the stimulation of insulin release<sup>[18]</sup>.

It is worth noticing that RYGB and sleeve gastrectomy (SG)[19-21] (Figure 1) have very similar outcomes both in shortterm excessive weight loss and long term total weight loss, as well as in type II diabetes resolution and both procedures are superior when compared to the adjustable gastric banding[22-24]. At the same time, the greatest diabetes remission was observed for patients undergoing biliopancreatic diversion with duodenal switch (BPD-DS) (Figure 2[25,26]) (95.1% resolved), followed by gastric bypass (80.3%), gastroplasty (79.7%), and then laparoscopic adjustable gastric banding (56.7%) The same pattern was observed even for an excessive weight loss and total body weight loss in the long term perspective[27-29]. However, despite better outcomes in terms of obesity and obesity-related comorbidities, BPD-DS counts only for ca 2% of bariatric surgeries performed worldwide due to increased risk of complications and development of malnutrition[30].

We describe the possible pathways of alpha-amylase-dependent AIA regulation of insulin secretion and their integration with food-related "dietary" incretin regulation of insulin secretion, which are usually eliminated or ameliorated by successful bariatric surgery, using the example of BPD or BPD-DS surgery (Figure 3). Breaking contact between the duodenal mucosa and the digesta coming from the stomach through BPD/BPD-DS surgery, in a broad sense interrupts the enteral-insular axis and thus the resulting decrease in insulin secretion could be due to other mechanisms-it is obvious. However, alpha-amylase still flows into the duodenum as a result of stimulation by gastro-pancreatic reflexes and by the attenuated (yet still existing) reflexes which originate in the alimentary limb (stimulated by digesta). Thus, the inhibitory effect of alpha-amylase on insulin secretion can result in the enhanced elimination of incretins.

It seems that BPD/BPD-DS bariatric surgery followed by immediate and long-term remission of DT2[22-24,27-30] has allowed us, for the first time, to observe and reveal these alpha-amylase-dependent inhibitory effects on insulin secretion and glucose metabolism. BPD/BPD-DS surgery prominently highlights the role of pancreatic alpha-amylase in the down regulation of insulin secretion, not only by means of elimination of the duodenal incretin effects on insulin release.

As shown in Figures 3, the remission of DT2 is related to the elimination of the action of incretins from the duodenum in BPD-SD patients. In these patients, insulin production is mainly dependent on blood glucose levels and its local (in pancreas) down regulation by alpha-amylase and in the diverted duodenum, as was proven in a pig model[13].

Attenuation of incretin release, due to the absence of contact between the chyme and the gastro-duodenal mucosa in the biliopancreatic limb after BPD-SD surgery, results in lower insulin release. At the same time, the passage of food to the alimentary limb is responsible for the slight stimulation of pancreatic enzyme secretion. Since alpha-amylase from the pancreatic juice which is being secreted into the biliopancreatic limb does not interact with its' starch substrate, the other functions of alpha-amylase, over and above that of starch digestion, e.g., inhibitory effect on insulin secretion, can be highlighted. Whole or partially digested alpha-amylase attenuates glucose absorption and improves the metabolism and/ or storage of glucose in the gut tissues [13]. One should bear in mind that alpha-amylase, as well as the other pancreatic enzymes being digested in the BP limb, could serve as a source of different types of biologically active peptides which could affect insulin secretion, as has been proven for alpha-amylase-derived peptides[13]. At the same time, alphaamylase from the acini surrounding the pancreatic islets can additionally slow down insulin release, which can be recognized as the inhibitory action of alpha-amylase on insulin release. These events are most probably responsible for the elimination of DT2, following BPD/BPD-DS bariatric surgery.

To highlight the efficacy of classical RYGB bariatric surgery, in terms of DT2 remission, one should consider that the residual secretion of stomach/gastric juice is probably the most important factor which could influence the overall success of bariatric metabolic surgery. Lack of stomach juice entering the biliopancreatic limb-which is what happens in patients after BPD/BPD-DS surgery-eliminates the release of most incretins (GIP, GLP-1), thus insulin secretion is stimulated to a much lower extent. The above is not true for other types of bariatric surgery where small amounts of gastric juice still enter the duodenum, thus stimulating incretin release. The role of alpha-amylase in AIA regulations[10, 13] has been experimentally proven both in the pig model of BPD[13], with complete isolation of the stomach from the



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Figure 1 Main types of bariatric surgery, from left to right. A: Sleeve gastrectomy; B: Roux-en-Y gastric bypass; C: One-anastomosis gastric bypass. Citation: IFSO. Sleeve Gastrectomy. [cited 20 December 2022]. Available from: https://www.ifso.com/sleeve-gastrectomy/[19]; IFSO. Roux-en-Y Gastric Bypass. [cited 20 December 2022]. Available from: https://www.ifso.com/roux-en-y-gastric-bypass/[20]; IFSO. One Anastomosis Gastric Bypass. [cited 20 December 2022]. Available from: https://www.ifso.com/alternative-intestinal-procedures/[21]. Copyright© Dr. Levent Efe, courtesy of IFSO, 2022.



Figure 2 Forms of biliopancreatic diversions, from left to right. A: Biliopancreatic diversion (BPD); B: BPD with duodenal switch. Citation: IFSO. Biliopancreatic Diversion (BPD). [cited 20 December 2022]. Available from: https://www.ifso.com/bilio-pancreatic-diversion/[25]; IFSO. Biliopancreatic Diversion with Duodenal Switch. [cited 20 December 2022]. Available from: https://www.ifso.com/bilio-pancreatic-diversion-with-duodenal-switch/[26]. Copyright© Dr. Levent Efe, courtesy of IFSO, 2022.

biliopancreatic limb, and in *in vitro* experiments, where alpha-amylase directly inhibited insulin release from an insulinoma cell line, BRIN-11[9].

Importantly, the pharmacological duodenal inhibition of the actions of incretins, through the use of GLY-200 polymer [31], which mimics bariatric (metabolic) surgery, is being studied in a clinical trial. Mechanical duodenal exclusion, realized *via* endoscopy, is now being tested in clinics and has produced very good results to date with respect to the remission of DT2[32]. Both strategies assume the exclusion of incretin action and the mechanical prevention of contact between the duodenal mucosa and digesta from the stomach (duodenum isolation), thus allowing AIA pancreatic reflexes, including the anti-incretin action of alpha-amylase, to be realized to the full extent.

Pancreatic alpha-amylase, which is synthesized in the acinar cells and then appears in the pancreatic juice, also leaks into the periinsular interstitial fluid, downregulating insulin secretion. Additionally, digesta entering the alimentary arm (proximal jejunum) stimulates reflexes responsible for insulin and enzyme secretion to a lower extent, since in the proximal jejunum the number of incretins and other gastrointestinal hormone receptors is markedly decreased, when compared to the duodenum.

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Figure 3 A schematic view of acini-islet-acinar- and alpha-amylase-dependent inhibitory pathway involved in the regulation of glucose metabolism before and after Biliopancreatic diversion surgery. A: Before biliopancreatic diversion surgery; B: After biliopancreatic diversion surgery. Incretin-dependent, quick stimulatory pathways (black) and acini-islet-acinar-dependent intrapancreatic inhibitory pathway and downregulating alpha-amylase dependent pathways, originating in the duodenum, of insulin secretion (orange and green respectively). AIA: Acini-islet-acinar.

### PANCREATIC ENZYME THERAPY AMELIORATES EPI-DEPENDENT HYPERINSULINEMIA

EPI is frequently diagnosed in both DT1 and DT2 patients[11]. Our own studies on streptozotocin (STZ)-induced diabetes [9] and on EPI[12] pig models have clearly shown that both iv administration of alpha-amylase of microbial origin (Amano, Japan) and the oral administration of pig pancreatic enzymes (enteric coated pancrelipase-Abbot Healthcare Products Ltd, Southampton, United Kingdom), lowers insulin and glucose levels during an iv-glucose tolerance test (IvGTT). However, responses to oral glucose loading are very weak compared to that observed following intravenous loading of a similar amount of glucose. Moreover, studies on healthy pigs[6] have shown that different pancreatic-like enzymes of microbial origin (Sigma-Aldrich), after oral administration, affect insulin release differently; alpha-amylase lowers insulin release, while protease enhances insulin release during an IvGTT and lipase has no effect on insulin

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release. The above-mentioned studies on EPI and STZ pigs have allowed us to explore the potency of the alpha-amylaseinduced, long-lasting AIA regulations of glucose homeostasis, revealing at the same time the protective/downregulating effects of enteral and blood alpha-amylase on the survival of pancreatic beta cells.

Enteral or parenteral alpha-amylase, of microbial or porcine origin, lowers blood levels of insulin and glucose during IvGTT and oral glucose tolerance tests. Undoubtfully, in this scenario, high levels of alpha-amylase in the gut and blood should be treated as a factor which protects pancreatic beta cells from metabolic failure. Nowadays more and more reports about the negative correlation between high plasma amylase activity and the development of metabolic diseases have become available in the literature [33,34].

The "anti-incretin" theory of metabolic surgery mode of action was presented by Rubino et al[35-37] in the beginning of century. The authors considered the gut as a primary organ involved in the regulation of metabolism and describe the balance of "anti-incretins" as a crucial factor which regulates glucose metabolism and could be impaired in metabolic disorders. In their comments to the article by Lindqvist et al[38] describing an increase in b-cell mass after RYGB surgery, Rubino and Amiel[37] point out that proliferation is not always beneficial and in fact, could be due to impaired "incretin/ anti-incretin" balance[37]. The same research group demonstrated an improvement in insulin sensitivity in diabetic rats after duodenal-jejunal bypass surgery, without changes in incretin and insulin secretion[39] and revealed the anti-incretin effect of orally administered glucose[40].

However, no endogenic "anti-incretin" factor have been recognized until now what is conditio sine qua non to complete anti-incretin hypotheses. At the same time, the number of reports confirming the alternative, anti-incretin-like role of alpha-amylase, has been increasing over the last few years. For example, the active role of alpha-amylase of duodenal origin in the regulation of enterocytes' proliferation was shown by Date *et al*[41,42]; there are reports about antiproliferative effects of alpha-amylase in cancer cell lines[43,44]; alpha-amylase is currently even being considered as one of the main energy regulators in the brain [45]. Although almost all tissues express their own amylase and the extra-digestive roles of amylase have become more and more obvious, the possible role of amylase in the regulation of glucose metabolism is still ignored and overlooked.

The overarching question is how are alpha-amylase signals from the gut lumen and from the blood transferred to pancreatic beta and acinar cells? Due to the nature of the portal vascular system in the pancreas[46,47], local peri-islet insulin levels are very high compared to systemic levels. Our supraphysiological doses of iv infused alpha-amylase were much higher than physiological alpha-amylase levels and thus could possibly have increased the alpha-amylase concentration in the peri-islet circulation to a level which inhibited insulin release[12]. The inhibitory action of alpha-amylase on insulin from the gut lumen requires the existence of a gut-pancreas, alpha-amylase-dependent reflex, which ends in the peri-islet interstitial tissue. The more physiological mechanism, however, would be the direct action of alpha-amylase or alpha-amylase-derived peptides on gut glucose absorption [9,13] or on its storage and metabolism in the gut.

To confirm this intriguing postulation, it is necessary to perform in vivo studies on direct infusion of alpha-amylase alone and/or as a mixture with other pancreatic enzymes, to the artery suppling the pancreas, during iv glucose loading. The expected inhibition of insulin release during such an experiment will finally confirm the role of alpha-amylase in the protection of beta cells from the overproduction of insulin.

### CONCLUSION

In summary, we suggest that bariatric BPD/BPD-DS surgery (which is actually seldom performed due to the development of severe EPI syndrome) highlights the never before described alpha-amylase-induced, anti-incretin-like regulation of glucose metabolism, which protects the pancreatic beta cells from exhaustion and subsequent failure. BPD/ BPD-DS surgery eliminates the incretin-dependent vicious cycle driven from the duodenum in obese diabetic patients. Food stimulates the synthesis of insulin and alpha-amylase in parallel. Alpha-amylase synthesis is extraordinarily stimulated by insulin-halo phenomenon. In turn, alpha-amylase, through its ability to inhibit glucose absorption and direct downregulation of insulin release[9] limits hyperglycemia and hyperinsulinemia to a certain degree.

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### FOOTNOTES

Author contributions: Pierzynowski SG conceived and wrote the manuscript; Stier C and Pierzynowska K wrote the manuscript and prepared the figures.

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ORIGINAL ARTICLE

### Genipin relieves diabetic retinopathy by down-regulation of advanced glycation end products via the mitochondrial metabolism related signaling pathway

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### Abstract

### BACKGROUND

Glycation is an important step in aging and oxidative stress, which can lead to endothelial dysfunction and cause severe damage to the eyes or kidneys of diabetics. Inhibition of the formation of advanced glycation end products (AGEs) and their cell toxicity can be a useful therapeutic strategy in the prevention of diabetic retinopathy (DR). Gardenia jasminoides Ellis (GJE) fruit is a selective inhibitor of AGEs. Genipin is an active compound of GJE fruit, which can be employed to treat diabetes.

### AIM

To confirm the effect of genipin, a vital component of GJE fruit, in preventing human retinal microvascular endothelial cells (hRMECs) from AGEs damage in DR, to investigate the effect of genipin in the down-regulation of AGEs expression, and to explore the role of the CHGA/UCP2/glucose transporter 1 (GLUT1) signal pathway in this process.

### **METHODS**

In vitro, cell viability was tested to determine the effects of different doses of glucose and genipin in hRMECs. Cell Counting Kit-8 (CCK-8), colony formation assay, flow cytometry, immunofluorescence, wound healing assay, transwell assay, and tube-forming assay were used to detect the effect of genipin on hRMECs cultured in high glucose conditions. In vivo, streptozotocin (STZ) induced mice were used, and genipin was administered by intraocular injection (IOI). To explore the effect and mechanism of genipin in diabetic-induced retinal



dysfunction, reactive oxygen species (ROS), mitochondrial membrane potential (MMP), and 2-[N-(7-nitrobenz-2oxa-1,3-diazol-4-yl) amino]-2-deoxy-d-glucose (2-NBDG) assays were performed to explore energy metabolism and oxidative stress damage in high glucose-induced hRMECs and STZ mouse retinas. Immunofluorescence and Western blot were used to investigate the expression of inflammatory cytokines [vascular endothelial growth factor (VEGF), SCG3, tumor necrosis factor-alpha (TNF- $\alpha$ ), interleukin (IL)-1 $\beta$ , IL-18, and nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing 3 (NLRP3)]. The protein expression of the receptor of AGEs (RAGE) and the mitochondria-related signal molecules CHGA, GLUT1, and UCP2 in high glucose-induced hRMECs and STZ mouse retinas were measured and compared with the genipin-treated group.

### RESULTS

The results of CCK-8 and colony formation assay showed that genipin promoted cell viability in high glucose (30 mmol/L D-Glucose)-induced hRMECs, especially at a 0.4 µmol/L dose for 7 d. Flow cytometry results showed that high glucose can increase apoptosis rate by 30%, and genipin alleviated cell apoptosis in AGEs-induced hRMECs. A high glucose environment promoted ATP, ROS, MMP, and 2-NBDG levels, while genipin inhibited these phenotypic abnormalities in AGEs-induced hRMECs. Furthermore, genipin remarkably reduced the levels of the pro-inflammatory cytokines TNF- $\alpha$ , IL-1 $\beta$ , IL-18, and NLRP3 and impeded the expression of VEGF and SCG3 in AGEs-damaged hRMECs. These results showed that genipin can reverse high glucose induced damage with regard to cell proliferation and apoptosis in vitro, while reducing energy metabolism, oxidative stress, and inflammatory injury caused by high glucose. In addition, ROS levels and glucose uptake levels were higher in the retina from the untreated eye than in the genipin-treated eye of STZ mice. The expression of inflammatory cytokines and pathway protein in the untreated eye compared with the genipin-treated eye was significantly increased, as measured by Western blot. These results showed that IOI of genipin reduced the expression of CHGA, UCP2, and GLUT1, maintained the retinal structure, and decreased ROS, glucose uptake, and inflammation levels in vivo. In addition, we found that SCG3 expression might have a higher sensitivity in DR than VEGF as a diagnostic marker at the protein level.

### **CONCLUSION**

Our study suggested that genipin ameliorates AGEs-induced hRMECs proliferation, apoptosis, energy metabolism, oxidative stress, and inflammatory injury, partially via the CHGA/UCP2/GLUT1 pathway. Control of advanced glycation by IOI of genipin may represent a strategy to prevent severe retinopathy and vision loss.

Key Words: Genipin; Human retinal microvascular endothelial cells; Angiogenesis; Vascularization; Secretogranin III; Diabetic retinopathy

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**Core Tip:** The formation of advanced glycation end products (AGEs) has been widely validated in pathological changes of diabetic retinopathy (DR). A new vital compound in Gardenia jasminoides Ellis fruit, genipin, can be used to treat DR and decrease AGEs. Genipin ameliorated AGEs-induced human retinal microvascular endothelial cell proliferation, apoptosis, energy metabolism, oxidative stress, and inflammatory injury, partially via the CHGA/UCP2/glucose transporter 1 pathway. Control of AGEs by intraocular injection of genipin may represent a strategy to prevent severe retinopathy and vision loss. Here, we confirmed the effectiveness of genipin to treat DR both in vivo and in vitro, and explored its related molecular mechanism.

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### INTRODUCTION

The number of patients with diabetes mellitus (DM) has increased from less than 110 million in 1980 to approximately 420 million in 2015 worldwide and is expected to reach 642 million in 2040[1]. The estimated overall prevalence of DM is 10.9% among Chinese citizens based on national-wide surveillance<sup>[2]</sup>. Diabetic retinopathy (DR) is the most common microvascular complication of DM and is a vital reason for blindness in citizens aged over 55 years[3]. Some new treatments, such as intravitreal vascular endothelial growth factor (VEGF) inhibitors or steroid hormones, have been introduced for DR[4]. However, up to 50% of the patients failed to respond to such agents. Despite being an inherently destructive procedure, laser photocoagulation remains the mainstay therapy for people with proliferative DR (PDR). This



necessitates other effective methods for DR treatment. In the pathogenesis of DR, the retinal circulation is damaged by microvascular lesions. If this change is not controlled in the early stages, continual progression leads to retinal detachment and vision loss<sup>[5]</sup>. The primary pathogenesis of DR involves angiogenesis, chronic inflammation, and oxidative stress. The retinal capillary endothelium is composed of endothelial cells, pericytes, and the basement membrane. Neovascularization of the retinal surface is the primary stage of PDR[6]. It is also the most important factor in retinal detachment. Endothelial dysfunction leads to angiogenesis, which involves the proliferation, migration, and formation of tubes and is central to the development of vascular complications[7]. Angiogenesis requires energy support such as adenosine triphosphate (ATP) or glucose. Damage to human retinal microvascular endothelial cells (hRMECs) in DR arises from metabolic abnormalities in glucose metabolism, which are principally caused by advanced glycation end products (AGEs).

Glycation is an important step in aging and oxidative stress<sup>[8]</sup>. Persistently elevated glucose concentrations lead to rapid and intensive glycation reactions. Both acute and chronic hyperglycemia can enhance AGEs production<sup>[9]</sup>, which results in endothelial dysfunction and causes severe damage to the eyes or kidneys of diabetics. Inhibition of AGEs and its cell toxicity can be a useful strategy for the prevention of DR[10]. According to previous studies, reducing AGEs receptor/ligand interaction or breaking established AGEs crosslinks prevent AGEs formation[11].

Mitochondrial dysfunction and oxidative stress are largely involved in aging, cancer, age-related neurodegenerative disorder, and metabolic syndrome[12]. Mitochondrial dysfunction may predispose to the development of DM with the accompanying risk of developing DR or may contribute directly to diabetic metabolic dysregulation and thereby increase the risk of late diabetic complications including retinopathy[13,14]. The relation between mitochondrial dysfunction and diabetic eye complications has two elements: (1) Mitochondrial diseases may predispose to the development of DM and such type of diabetes may be accompanied with an increased risk of developing DR; and (2) metabolic dysregulation in DM may increase the risk of development of DR through a disturbance in mitochondrial function[13,14]. A hypothesis has recently been proposed that the metabolic dysfunction in diabetic patients induces the synthesis of a number of reactive oxygen species (ROS) that are normally eliminated in the mitochondria[13]. In addition, enhanced ROS may contribute to the induction of autophagy in the retina. Another mechanism of mitochondrial dysfunction leading to DR is the formation of free radicals[14-17]. These reactive compounds may contribute to a switch in the metabolism, e.g., by changing the activity of glyceraldehyde-3-phosphate dehydrogenase and the polyol pathway and activating protein kinase C, and consequent development of DR[13].

Genipin (Figure 1A), one of the principal bioactive components extracted from the Gardenia jasminoides Ellis (GJE, Chinese herbal name "Zhizi"), has multiple bioactivities, such as anti-inflammation, antitumor, antidepression, and the protection of hippocampal neurons from the toxicity of Alzheimer's amyloid- $\beta$ [18]. GJE, as a selective inhibitor of AGEs, prevents the development of diabetic vascular complications in experimental animal models; however, its possible role and molecular mechanism in the relief of DM symptoms are unknown<sup>[19]</sup>. To explore these, we searched the traditional Chinese medicine systems pharmacology database and analysis platform (TCMSP) (https://old.tcmsp-e.com/tcmsp.php) using the keyword "Zhizi". Genipin was the third most active constituent of Zhizi (Table 1). Other active constituents of Zhizi include quercetin and lutein. Recent studies demonstrated that quercetin inhibited the overexpression of TLR4 and NF-KB p65, and reduced the expression of VEGF and soluble intercellular adhesion molecule-1, thus exerting therapeutic effects in DR[20,21]. Many basic and clinical studies have reported that lutein has anti-oxidative and anti-inflammatory properties in the eye, suggesting its beneficial effects in protection and alleviation of ocular diseases such as age-related macular degeneration (AMD), DR, retinopathy of prematurity, myopia, and cataract[22,23]. However, some publications report that genipin, but not quercetin or lutein, can influence AGEs. In the present study, we administered streptozotocin (STZ)-treated mice by intraocular injection (IOI) of genipin to investigate whether genipin could attenuate the development of DR in experimental models through AGEs inhibition[24].

### MATERIALS AND METHODS

#### Reagents

Genipin (99%, Figure 1) was purchased from Selleck (Shanghai, China). D-glucose (PB180418) and the complete endothelial cell medium of hRMECs (CM-H130) were purchased from Procell (Wuhan, Hunan Province, China). Trypsinethylenediamine tetraacetic acid (EDTA) solution (S310JV), Dulbecco's modified Eagle's medium (L170KJ), minimum essential medium (L510KJ), penicillin-streptomycin solution (S110JV), and phosphate buffer solution (B320KJ) were purchased from BasalMedia (Shanghai, China). Cell Counting Kit-8 (BS350A) and 4% paraformaldehyde fix solution (BL539A) were purchased from Biosharp (Hefei, Anhui Province, China). Endothelial cell medium (1001), endothelial cell growth supplement (1052), fetal bovine serum (0025), and penicillin and streptomycin solution (0503) were purchased from ScienCell (Shanghai, China).

### Animals and ethical statement

All animal procedures were conducted in accordance with the China Animal Welfare Legislation. We purchased 4-wkold C57/BL6 male or female mice (3-8 g, n = 60) from the Animal Ethics Committee of Chongqing Medical University (Yuzhong, Chongqing, China). They were housed in a light-temperature automatically controlled room with free access to food and water. One female mouse and four male mice were housed in a single cage. All animal experimental protocols were approved by the Institutional Animal Care and Use Committee of Chongqing Medical University (2022-K45). All experiments were performed in accordance with the Guidelines and Regulations for the Care and Use of Laboratory Animals by the Chongqing Medical University (Yuzhong, Chongqing, China) and the Association for



Table 1 Active components of Gardenia jasminoides Ellis							
MolID	Active component	OB (%)	DL	BBB			
MOL009038	GBGB	45.58	0.83	-5.43			
MOL001652	1H-2,6-dioxacyclopent(cd)inden-1-one, 4-((acetyloxy)methyl)-5-(beta-D-glucopyranosyloxy)-2a,4a,5,7b-tetrahydro-, (2aS-(2aalpha,5alpha,7balpha))-	26.43	0.71	-2.00			
MOL001648	Genipin	26.06	0.10	-0.98			
MOL000098	Quercetin	46.43	0.28	-0.77			
MOL000422	Kaempferol	41.88	0.24	-0.55			
MOL007245	3-Methylkempferol	60.16	0.26	-0.49			
MOL001406	Crocetin	35.30	0.26	-0.83			
MOL000551	Hederagenol	22.42	0.74	-0.51			
MOL003515	(3S,4S,4aR,6aR,6bS,8aS,12aS,14aR,14bR)-3-hydroxy-4,6a,6b,11,11,14b-hexamethyl-1,2,3,4a,5,6,7,8,9,10,12,12a,14,14a-tetradecahydropicene-4,8a-dicarboxylic acid	27.21	0.72	-0.68			
MOL013377	Lutein	22.59	0.55	-0.99			
MOL009548	Desacetyl asperulosidic acid_qt	32.49	0.10	-1.85			
MOL004560	SHANZHISIDE_qt	117.77	0.10	-1.44			
MOL001667	Deacetyl asperuloside acid_qt	62.46	0.11	-1.62			
MOL007148	Shanzhiside methyl ester_qt	109.77	0.12	-3.18			
MOL004555	GARDENOSIDE_qt	52.77	0.12	-0.91			
MOL007994	Ilexoside A_qt	22.43	0.74	-0.33			

Oral bioavailability ≥ 20%, drug like ≥ 0.1, and brain-blood barrier ≤ -0.30. OB: Oral bioavailability; DL: Drug like; BBB: Brain-blood barrier.

Research in Vision and Ophthalmology Statement for the Use of Animals in Ophthalmic and Vision Research.

### STZ-induced diabetic mice and intraperitoneal injections

STZ-induced hyperglycemia is a widely used DM model. At the beginning of the experiment, the mice were fed a highsugar and high-fat diet (rodent diet with 60% calories from fat; XTHF60, XieTong Scientific Diets, Wuhan, China) for 70 d (20-35 g, n = 60). Using 1 mL syringes and 25 G needles, intraperitoneal injection (IPI) of freshly prepared STZ solution at 40 mg/kg (1.0 mL/100 g of citrate buffer, pH 4.5) was performed. This step was repeated on days 2 and 3 after fasting [25]. We used similar forage and feeding management practices. Fasting blood glucose (FBG) levels were measured with a Precision Plus blood glucose meter 14 d after STZ IPI. Mice with FBG levels > 11.1 mmol/L were considered diabetic.

### Drug treatment

In the genipin-treated group, IOI of genipin [genipin dissolved in penicillin-streptomycin solution (PBS)] at a dose of 10 mmol/L was administered to the right eye (OD) (0.4 µL/eye) of mice after 3 mo of STZ administration. The same volume of PBS was administered to the left eye (OS) of the same mouse. Mice were maintained on a high-sugar and high-fat diet with free access to water under standard conditions and were assessed once every 2 wk and 1 mo for their body weight and FBG, respectively. All animals were killed at the end of the third time of IOI administration. Both eyes were collected and frozen at -80 °C or fixed in 4% paraformaldehyde fix solution for further experiments.

### Hematoxylin and eosin staining

The eyes of the mice were fixed overnight with 4% paraformaldehyde fix solution and embedded in paraffin. Coronal sections (5 mm thick) were dewaxed and stained with hematoxylin and eosin stain (H&E). The wound areas of each sample were observed under a Leica DM2000 microscope (Leica, Wetzlar, Germany).

### Cell culture

hRMECs (BNCC 358978) were purchased from BeNa Culture Collection (Beijing, China). The cells were plated in T<sub>25</sub> flasks and incubated in a humidified incubator with 5% CO<sub>2</sub> at 37 °C.

### Cell counting kit-8 assay

hRMEC pellet was resuspended in 100 µL of endothelial cell medium (ECM) without FBS, incubated in 96-well plates for 1 d, and treated with various concentrations of D-glucose. Cell viability was assessed using the Cell Counting Kit-8 (CCK-8). We added CCK-8 (10%) with a culture medium of 110 µL was added to each well and incubated for 2.5 h (37 °C, 5%





Figure 1 Optimization of dose of genipin and high glucose in human retinal microvascular endothelial cells. A: Chemical structure of genipin; B and C: Cell viability of human retinal microvascular endothelial cells (hRMECs) treated with different concentrations of glucose and genipin; D: Cell viability of hRMECs treated with 30 mmol/L glucose and 0.4 µmol/L genipin for different durations; E-I: RAGE, SCG3, and vascular endothelial growth factor protein expression in cells treated with genipin for different durations. VEGF: Vascular endothelial growth factor.

 $CO_2$ ). Absorbance at 450 nm and 600 nm was measured using a microplate reader (Thermo Fisher). The cell inhibition rate (I%) was calculated as  $(A_{control} - A_{treated})/A_{control} \times 100\%$ .

### Drug preparation

hRMEC pellet was resuspended in 100 mL of ECM supplemented with 30 mmol/L of D-glucose (HG), incubated in 96well plates for 1 d, and then treated with various concentrations of genipin. After 7 d of treatment, cell viability was assessed using CCK-8 assay.

### Flow cytometry

Apoptosis was detected using an Annexin V-FITC Apoptosis Detection Kit (Beyotime) according to the manufacturer's instructions (CytoFLEX, United States). Cells ( $2 \times 10^6$ ) were seeded in 6-well plates and incubated with a high concentration of glucose and genipin for 1 wk. They were collected after digestion with trypsin-EDTA (Canada). After washing with PBS twice, the supernatant was discarded. The pellet was resuspended using ice-cold 70% ethanol (30 min at 4 °C) and incubated at 37 °C for 30 min. Next, 500 µL of 1 × binding buffer was added, and the cells were stained with 5 µL of annexin V-FITC and 5 µL of propidine iodide in the dark at 4 °C for 20 min, followed by flow cytometry analysis.

### Western blot analysis

Cells were lysed with ice-cold radioimmunoprecipitation assay (RIPA) lysis buffer (Beyotime) and centrifuged at 12000 r/ min for 15 min at 4 °C, and the supernatant was collected. Protein concentrations in the supernatant were measured using the BCA protein assay kit (ThermoFisher, Shanghai, China). Proteins (40  $\mu$ g) were resolved by SDS-PAGE and transferred onto polyvinylidene difluoride membranes. The membranes were blocked with 5% BSA in TBST (1 × Tris buffered saline, 0.1% TWEEN 20) for 2 h at room temperature. The blocked membranes were immunoblotted with rabbit anti-UCP2



(A4178, ABclonal, Wuhan, Hubei Province, China), rabbit anti-glucose transporter 1 (GLUT1)/SLC2A1 (A11208, ABclonal), rabbit anti-CHGA (A1668, ABclonal), rabbit anti-AGER (A13264, ABclonal), rabbit anti-SCG3 (A7799, ABclonal), and rabbit anti-VEGF (A12303, ABclonal) antibodies overnight at 4 °C. The membranes were then washed and probed with secondary antibody (horseradish peroxidase-labeled goat anti-rabbit IgG; A0208, Beyotime) for 2 h at room temperature. Chemiluminescence detection was performed using a SuperSignal West Atto chemiluminescence detection kit (Thermo Fisher, Shanghai, China) according to the manufacturer's instructions.

### Immunofluorescence

hRMECs were seeded in 24-well plates with a cover glass (REF.10212424C, Shitai, Jiangsu, China) at a final cell density of  $5.0 \times 10^4$  cells/mL. In the genipin group, 4 mmol/L genipin was added. When the cell confluence reached approximately 50%, the cells were transferred to the FBS-free and antibiotic-free medium and cultured for 1 d. They were washed with PBS three times. The cells were fixed in 4% paraformaldehyde fix solution (P0099, Beyotime) for 30 min and blocked for 0.5 h using goat serum (C0265, Beyotime). Then, the membranes were washed and immunoblotted with primary antibodies overnight at 4 °C, followed by probing with Alexa Fluor 488-labeled Goat Anti-Rabbit IgG (H + L) (A0423, Beyotime) as secondary antibody for 2 h at room temperature. 2-(4-amidinophenyl)-6-indolecarbamidine dihydrochloride staining solution (C1005, Beyotime) was used for counterstaining. Antifade mounting medium (P0126, Beyotime) was used to reduce the luminescence delay. All pictures were obtained using a Leica fluorescence microscope (DMR, Deerfield, IL, United States).

### Colony formation assay

hRMECs were seeded into 6-well plates, and the number of cells was adjusted to approximately 2000 cells/well. After allowing cells to attach the plates, ECM with 20% FBS was added. After 2 wk, the cells were fixed in 4% paraformal-dehyde fix solution for 0.5 h and stained with crystal violet (Beyotime Institute of Biotechnology, Shanghai, China) for 15 min. Each well with > 50 cells was photographed and observed under a phase-contrast microscope.

### Wound healing assay

hRMECs were seeded in 6-well plates at a final cell density of  $2.0 \times 10^5$  cells/mL. After treatment for 5 d, the confluent cell monolayer was scraped with a sterile 200 µL tip to create a scratch across the center of the circle when the hRMECs confluence reached > 95%. The wound areas of each sample were observed under a microscope and evaluated using ImageJ software (Media Cybernetics, Silver Springs, MD, United States) at 0 h, 12 h, 24 h, 48 h, and 72 h.

### Transwell assay

We used 8  $\mu$ m sized Transwell BD Matrigel chambers (Costar 3470, Corning, NY, United States) to measure cell migration. hRMECs were seeded into the upper chamber with 100  $\mu$ L of serum-free ECM, and the number of cells was adjusted to approximately 2000 cells/well. The lower chamber contained 600  $\mu$ L ECM with 20% of FBS. After 30 h of culturing, the chambers were removed and fixed with 4% paraformaldehyde fix solution for 0.5 h. The cells were stained with crystal violet for 15 min. Medical Cotton Stickers (Sanhe, Sichuan Province, China) were used to remove non-migratory cells from the upper chamber. Photographs were captured by using a microscope. Five fields with evenly distributed cells were examined.

### Tube formation assay

To analyze the influence of high glucose and genipin on the tube-forming activity of hRMECs, we plated 50  $\mu$ L Matrigel (BioCoat Matrigel 356234, Corning, NY, United States) onto a cold 96-well plate. hRMECs were added into each well at a cell density of 4.0 × 10<sup>5</sup>. After 6 h, images were captured under a microscope.

### ATP concentration determination

hRMECs were plated onto 6-well plates and grown until they reached confluence. The cells were starved in an endothelial basal medium (EBM) without serum for 1 d. They were collected after transfection and washed twice with PBS. The cells were collected into 1.5 mL Eppendorf tubes, and the ATP concentration was measured using an ATP fluorometric assay kit (S0026, Beyotime).

### ROS and mitochondrial membrane potential assay

hRMECs were plated onto 6-well plates and grown until they reached approximately 60% confluence. The cells were starved in an EBM medium without serum for 1 d. The ROS Assay Kit (S0033, Beyotime) and Apoptosis Detection Kit with Mito-Tracker Red CMXRos and annexin V-FITC (C1071S, Beyotime) were used to measure the ROS and mitochondrial membrane potential (MMP) changes.

### Glucose uptake assay

hRMECs were plated onto 6-well plates and grown until they reached confluence. The cells were starved in an EBM medium without serum for 1 d. D-glucose analog (100 mmol/L; 2-NBDG) assay (HY-116215, MCE, Shanghai, China) was performed to measure the D-glucose transported into the hRMECs. Photographs were captured under a fluorescence microscope.

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### Cytokine analysis

Pro-inflammatory cytokines [tumor necrosis factor-alpha (TNF- $\alpha$ ), interleukin (IL)-1 $\beta$ , IL-18, and NLRP3] and other hallmark proteins of DR (VEGF and SCG3) were assayed by Western blot and immunofluorescence as described previously.

### Statistical analysis

Statistical analyses were performed using IBM SPSS Statistics 25 (IBM Corp., Armonk, NY, United States) and GraphPad Prism 8 (GraphPad Software, LLC). All experiments were repeated at least three times independently. All data are presented as the mean  $\pm$  SEM. Independent sample *t*-test was used for comparing variables with a normal distribution. One-way analysis of variance and multiple comparisons for trends were used for comparing continuous variables. The chi-square test (or Fisher's exact test, when appropriate) was used to analyze dichotomous variables. Pearson's correlation coefficient was used to determine the relationship between variables. A *P* value less than 0.05 was considered statistically significant.

### RESULTS

### Effects of genipin on hRMECs viability and proliferation

First, we investigated the effects of high glucose and genipin on hRMEC viability and cytotoxicity using CCK-8 assay. Figure 1B depicts that 30 mmol/L D-glucose could decrease approximately 30% of the cell viability by approximately 30%, and cell viability decreased by < 30% until high glucose reached 90 mmol/L. Thus, 30 mmol/L of D-glucose was used for subsequent experiments. After treatment with different doses of genipin (0.1 µmol/L to 1.0 µmol/L) in a 30 mmol/L glucose medium, we detected cell proliferation by CCK-8 assay. The administration of 0.4 µmol/L genipin exerted a notable effect on hRMECs proliferation (Figure 1C). However, genipin at 0.8 µmol/L remarkably inhibited cell proliferation (P < 0.001). The optimal effective dose of high glucose was 30 mmol/L to 50 mmol/L and that of genipin was approximately 0.3-0.8 µmol/L; these doses did not exert any cytotoxic effects on the cells.

Next, we explored the optimal action time of genipin by CCK-8 assay (Figure 1D). Compared with normal glucose medium without genipin, treatment with 0.4  $\mu$ M genipin for 4 d exerted a notable effect on high glucose-induced HRMECs (*P* < 0.001) (Figure 1D). In hRMECs cultured in ECM containing 30 mmol/L glucose, we observed the highest cell viability after 7 d of treatment with 0.4  $\mu$ mol/L genipin (Figure 1D). We verified these results using Western blot (Figure 1E-H). Based on these results, treatment with 30 mmol/L high glucose and 0.4  $\mu$ mol/L genipin for 7 d was used for the subsequent experiments.

### Genipin protects hRMECs from high-glucose damage with regard to cell proliferation, apoptosis, angiogenesis, energy metabolism, oxidative stress, and inflammatory injury in vitro

Action of glucose and genipin on hRMECs proliferation, apoptosis, and angiogenesis: CCK-8 assay demonstrated that genipin, particularly 0.4 µM genipin for 7 d, increased the viability of hRMECs in high-glucose ECM. The colony formation results supported this finding (Figure 2A). Results of flow cytometry (Figure 2B) suggested that the number of apoptotic hRMECs increased after the stimulation with 30 mmol/L glucose (61.05%), compared with normal medium stimulation (30.30%). However, the cell apoptosis ratio decreased after genipin treatment, and particularly, early cell apoptosis decreased from 33.98% to 11.07%.

To explore the effects of genipin on cell migration, we performed scratch wound healing and transwell migration assays. High glucose decelerated the closure of hRMECs, and genipin could restore this change (Figure 2C). Furthermore, high glucose levels increased the number of migrated cells, and genipin restored this change (Figure 2D).

Then, we analyzed the tube-forming capacity of hRMECs in high glucose, with or without genipin. High glucose significantly increased the number of newly formed tubes, while genipin reversed this change (Figure 2E).

Action of glucose and genipin on intracellular ATP levels and MMP: Hyperglycemia in patients with DM is associated with abnormally elevated cellular glucose levels. ATP, as the primary energy storage molecule, is central to cell survival, proliferation, and migration. Increased cellular glucose levels alter glucose metabolism and influence intracellular ATP levels by inhibiting complexes I and II[26]. Subsequently, we measured the intracellular ATP levels and observed that high glucose increased the ATP level by 60%, and genipin could reduce this level by approximately 20% (Figure 3A).

Next, we explored the mitochondrial changes in DR. MMP increased after high glucose stimulation (Figure 3B and C), indicating that DR produced more ATP, which promotes the formation of mitochondria[27]. Simultaneously, these mitochondrial changes may lead to increased ROS accumulation.

Action of glucose and genipin on intracellular oxidative stress levels: To confirm the oxidative stress damage caused by high glucose levels, we measured intracellular ROS production. High glucose increased the fluorescence significantly; however, this increase was alleviated by genipin (Figure 3B-D).

Action of glucose and genipin on intracellular glucose metabolism levels: NBDG is a fluorescence-labeled 2-deoxy-glucose analog that works as a tracer for evaluating cellular glucose metabolism[28]. High glucose increased the fluorescence, and genipin might reverse the effects.

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Figure 2 Genipin protects from advanced glycation-induced human retinal microvascular endothelial cell proliferation, apoptosis, and angiogenesis. A and B: Genipin protects human retinal microvascular endothelial cell (hRMECs) from high glucose-induced damage with regard to cell proliferation as revealed by colony formation assay and flow cytometry; C-E: Wounding healing, migration and tube formation of hRMECs treated with 30 mmol/L glucose and 0.4 µmol/L genipin. NG: 5 mmol/L glucose containing ECM-treated hRMECs group; HG: 30 mmol/L glucose containing ECM-treated hRMECs.

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Figure 3 Genipin protects human retinal microvascular endothelial cells from high glucose damage with regard to energy metabolism, oxidative stress, and inflammatory injury in vitro. A: ATP levels among normal glucose, high glucose, and genipin-treated human retinal microvascular endothelial cells (hRMECs); B-D: Reactive oxygen species, mitochondrial membrane potential, and 2-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) amino]-2-deoxy-d-glucose levels in hRMECs treated with high glucose and genipin; E-L: Western blot and immunofluorescence to measure the expression of inflammatory factors expression. IL: Interleukin; TNF: Tumor necrosis factor-alpha; NLRP3: Nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing 3; VEGF: Vascular endothelial growth factor; ROS: Reactive oxygen species; MMP: Mitochondrial membrane potential; 2-NBDG: 2-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) amino]-2deoxy-d-glucose; hRMECs: Human retinal microvascular endothelial cells. NG: 5 mmol/L glucose containing ECM-treated hRMECs group; HG: 30 mmol/L glucose containing ECM-treated hRMECs group; GG: 0.4 µM genipin + 30 mmol/L glucose containing ECM-treated hRMECs.

Action of glucose and genipin on inflammation: To evaluate the effects of genipin on inflammation following HG stimulation, we performed Western blot and Immunofluorescence to detect the protein levels of inflammatory cytokines (TNF-α, IL-1β, IL-18, and NLRP3) and DR biomarkers (VEGF and SCG3) in hRMECs inclubated with high glucose (Figures 3E-L and 5D). High glucose significantly increased the protein expression of TNF- $\alpha$ , IL-1 $\beta$ , IL-18, and NLRP3, and DR biomarkers. Interestingly, genipin down-regulated the expression of these proteins. Therefore, genipin may alleviate the damage caused by inflammation. Next, we will investigate the mechanisms underlying this effect.

### Genipin IOI protects the retina of STZ mice from high-glucose damage in vivo

After continuous STZ IPI for 3 d, the blood glucose levels were > 7.5 mmol/L in 2 wk and > 11.1 mmol/L in 1 mo in C57 mice.

DM-induced neuronal loss was assessed by measuring the retinal thickness at 3 mo after DM induction using retinal sections from untreated eyes (OS) and genipin-injected eyes (Figure 4B and C). At 3 mo after DR induction, H&E staining was performed to assess the structural morphology of the retinal cells. We measured four spots of the retina; the thickness of genipin-treated retina (190.70  $\mu$ m ± 58.32  $\mu$ m) was lesser than that of DR-untreated retina (262.10  $\mu$ m ± 55.52  $\mu$ m, P = 0.045). Representative microphotographs demonstrate a disorder of the entire retina, vacuolization, and dissolution of the

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Figure 4 Genipin reduces retinal oxidant stress, glucose metabolism, and inflammation in streptozotocin-induced mice. A: Blood sugar levels of mice induced with streptozotocin (STZ) for 2 wk and 4 wk; B: Hematoxylin and eosin staining of STZ mouse retina; C: Retinal thickness in different eyes of STZ mice; D: Inflammatory factor expression measured by Western blot; E and F: Reactive oxygen species and 2-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) amino]-2deoxy-d-glucose levels in STZ mice; G and F: RAGE, SCG3, and vascular endothelial growth factor expression in STZ mice. IL: Interleukin; TNF: Tumor necrosis factor-alpha; NLRPa: Nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing 3; VEGF: Vascular endothelial growth factor; ROS: Reactive oxygen species; MMP: Mitochondrial membrane potential; 2-NBDG: 2-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) amino]-2-deoxy-d-glucose; OS: Left eye, without genipintreated; OD: Right eye, genipin-treated; GCL: Ganglion cell layer; INL: Inner nuclear layer; ONL: Outer nuclear layer.

ganglion cell layer in the left eye.

### Genipin reduces retinal oxidant stress, glucose metabolism, and inflammation in STZ mice

We performed ROS and 2-NBDG assays to measure oxidative stress and glucose metabolism in the frozen sections. Genipin-treated cells had significantly increased oxidant stress and glucose metabolism. SCG3 and VEGF expression increased in the untreated group.

In addition, the protein levels of inflammatory cytokines (TNF- $\alpha$ , IL-1 $\beta$ , IL-18, and NLRP3) and DR biomarkers (VEGF and SCG3) were measured by Western blot. Genipin protected the cells from inflammation.

### Mechanism of genipin to relives retinal endothelial damage induced by high glucose

UCP2/GLUT1 pathway is involved in the protective effect of genipin on hRMECs (Figure 5): The decrease in glucose uptake in genipin-treated cells was mediated by the reduction of GLUT1 expression, as demonstrated by Western blot (Figure 5C and D). To investigate the glycation process that occurred after glucose, we explored the AGE-receptor interactions in both cells and tissues, which showed that genipin treatment decreased AGEs.

We used STRING (Version: 11.5, https://cn.string-db.org/cgi/input) to analyze the relationships among "RAGE", "SCG3", "UCP2", "GLUT1", and "VEGF" (Figure 5A). GLUT1 may be influenced by UCP2 expression in oxygen-induced retinopathy (OIR) at both the mRNA and protein levels, whereas elevated circulating CHGA can lead to UCP2 overexpression in the hypertensive phenotype[9]. Next, we performed Western blot and immunofluorescence to verify the expression of these proteins. The protein expression of these three proteins was increased in high-glucose conditions, and genipin could restore such change.

Genpin relives retinal endothelial damage in high-glucose conditions by influencing energy metabolism, oxidative stress, and inflammatory injury: We identified 33 and 4328 DR-related genes using GeneCards (https://www.genecards. org/) and OMIM (https://omim.org/), respectively, with the keywords "DR" or "diabetic retinopathy". Next, we used STRING to analyze the relationships among these genes (Figure 6A); Gene Ontology (GO)/Kyoto Encyclopedia of Genes and Genomes function enrichment analyses revealed that AGE-RAGE plays a vital role in DR (Figure 6B). Restraining AGEs and alleviating toxicity may be a novel way to control DR.



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Figure 5 Mechanism of genipin to relive retinal endothelial damage caused by high glucose. A and B: Schematic summary of the results; C-I: Western blot and immunofluorescence (IF) analysis of expression of proteins related to CHGA/UCP2/GLUT1; J: RAGE, SCG3, and vascular endothelial growth factor expression detected by IF.

We used TCMSP to identify 21 genes possibly targeted by genipin. UniProt (https://www.uniprot.org/) was used to assess and convert these gene names. A Wayne diagram was plotted to display the intersection of the genipin target and DR-related genes (Figure 6C and D). To determine the available phenotypic abnormalities, we used GO function enrichment and pathway enrichment analyses to explore the relationships among genes that were mixed by DR-related and genipin targets. Finally, oxidoreductase activity and arachidonic acid metabolism demonstrated distinct correlations (Figure 6E), indicating that genipin may influence ROS and inflammation as two vital phenotypic abnormalities in DR.

### DISCUSSION

Vision loss in DR principally occurs in either diabetic macular edema (DME) or PDR[29]. DR is classified based on visible ophthalmologic changes and manifestations of retinal neovascularization[30]. Therefore, the early diagnosis and treatment of DR cover the complete range of retinopathies and are complex. Fortunately, regardless of the type of DR, abnormal AGE accumulation has the same basic reason.

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Figure 6 Pathway enrichment analysis. A: Prediction of diabetic retinopathy (DR)-related genes; B: Pathway enrichment analysis of DR-related genes; C and D: Intersection of genipin target and DR-related genes; E: Pathway enrichment analysis of genes which are mixed by DR-related and genipin targets. EPO: Erythropoietin; BP: Biological process-pathways and larger processes made up of the activities of multiple gene products; MF: Molecular Function-molecular activities of gene products; KEGG: Kyoto Encyclopedia of Genes and Genomes.

DR has a mixed pathogenesis. It involves cell cycle regulation, apoptosis signal transduction, oxidative stress response, protein biosynthesis, carbohydrate metabolism, and other important proteins and proteases. The ROS assay demonstrated a high level of oxidative stress response in both cells in the high glucose group and STZ mice without treatment. Carbohydrate metabolism and mitochondrial injury in the high glucose group led to increased ATP levels and cell apoptosis. Glucose uptake was higher in the high glucose group than in the control group, which can be validated by GLUT1 expression. Genipin, an aglycone derived from the iridoid glycoside, is also used as an anti-tumor drug in oral squamous cell carcinoma, hepatocellular carcinoma, and gastric cancer *via* the signal transducer and activator of the transcription-3 pathway and upregulated MMP genes[31,32]; the relationships between genipin and glucose was rarely reported. We explained whether genipin could change the GLUT family in different cell types. This finding may help explain how genipin influences tumors and apoptosis in various diseases. VEGF is a classical biomarker of DR, and it is widely used to treat inflammation[32-35]. NLRP3 inflammasomes are involved in the production and persistence of

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inflammation in diabetic nephropathy [36-38]. We observed that high glucose stimulation could up-regulate IL-1 $\beta$ , IL-18, TNF-α and NLRP3, but not IL-6.

Upon exposure to sugar, proteins, lipids, and nucleic acids are oxidized and rearrange to form stable products, which can further undergo crosslinking, thus yielding AGEs[39]. AGEs are associated with a series of diseases such as Alzheimer's disease, skin senescence, face splash capillary ectasia, and pathologies such as AMD, diabetic keratopathy, and DR. AGEs are reactive intermediates of chronic hyperglycemia with proteins, lipids, and nucleic acids in vivo, leading to the accumulation of pro-inflammatory cytokines and ROS. Furthermore, they damage the vasculature, eye, heart, and kidneys in patients with DM.

Consequently, we focused on glucose metabolism and AGEs production in hRMECs and addressed the hypothesis that high-glucose medium affects glucose uptake and mitochondrial metabolism, leading to intravitreal neovascularization in DR. UCP2, a mitochondrial carrier protein, can increase the cellular sensitivity to glucose and promote cellular glucose uptake by mediating proton leak across the inner mitochondrial membrane<sup>[40]</sup>. Genipin is a standard UCP2-inhibiting agent in various experimental models of ROS generation and mitochondrial activity[41]. Most tumor growth and proliferation processes involve excessive glucose uptake mediated by GLUTs[42]. The retina takes up glucose, principally through GLUT1[43]. Genipin promotes physiological retinal vascular development and improves endothelial function by inhibiting the UCP2/GLUT1 pathway in premature retinopathy.

Thus, we focused on the effects and molecular mechanisms of genipin on the proliferation, apoptosis, angiogenesis, energy metabolism, oxidative stress, and inflammation of AGEs-induced hRMECs.

Genipin have been used to treat several diseases due to its choleretic, anti-inflammatory, antioxidant, anti-apoptotic, autophagy-inducing, anti-necroptotic, and anti-pyroptotic properties[41,44-47]. Simultaneously, genipin is an aglycone of an iridoid glycoside termed geniposide. Moreover, it is a commonly used biological agent, e.g., as a cross-linker, because it can be better tolerated by cells. Despite its benefits and good biocompatibility, genipin is still unpopular in classical medicine because of its unclear mechanism of action. We analyzed the role of genipin as an important element of Zhizi against AGEs accumulation in DR. First, we evaluated the intracellular concentration of AGEs by assessing the expression level of RAGE. RAGE expression increased significantly with high glucose stimulation, and genipin decreased this increase. High glucose stimulation increased apoptosis, ROS, ATP, mitochondrial formation, glucose metabolism, and inflammation. High glucose could increase the accumulation of AGEs via pathways that can be affected by genipin and finally decrease GLUT1 expression. UCP2, as a binding site for genipin, decreases GLUT1 expression at both the mRNA and protein levels<sup>[40]</sup>. Then, we tried to connect RAGE and DR biomarkers with these pathways. We identified RAGE and DR biomarkers in these pathways. We analyzed the relationships between these five proteins and identified the "CHGA" protein. Elevated CHGA levels result in UCP2 overexpression in hypertension[48]. We evaluated these changes in hRMECs treated with high glucose and STZ mouse retinas and observed increased protein expression of CHGA, UCP2, and GLUT1. Hence, high glucose leads to the addition of AGEs, increases CHGA expression, and promotes UCP2 and GLUT1 expression, which influences cell proliferation, apoptosis, oxides, cellular energy metabolism, and inflammation. Genipin can inhibit this CHGA/UCP2/GLUT1 pathway not only by controlling UCP2 but also through other mechanisms.

Genipin is the third most active constituent of Zhizi (Table 1). The oral bioavailability of genipin is 26.06% and its topological polar surface area is 75.99, which suggests that genipin is preferentially administered by injection than orally. However, genipin is principally used as an oral formulation in previous research, whereas injection formulations have been less investigated. In addition, genipin is an extremely good natural cross-linking agent that contributes to sustainedrelease or delayed-release joint use with other drugs<sup>[49]</sup>. AGEs are a type of cross-linker that combines sugars and proteins and can also reduce this step by linking with proteins by physical methods. Furthermore, genipin is widely used in the clinic to treat corneal or scleral diseases[46,49].

Ocular drug delivery is difficult because of the lacrimal fluid-eye barrier and the retina-blood barrier[50]. Retinal diseases, such as retinal detachment, AMD, and DR, require an intravitreal injection to inhibit the formation of ocular neovascularization. Such common drugs include anti-VEGF medications and hormones. Our previous report describes the use of an AGE inhibitor that can penetrate the vitreous humor and treat retinal disease in STZ mice. We used the most effective dose of genipin in hRMECs for injection (C = m  $\times$  V). The vitreous volume of mice was approximately 10  $\mu$ L, and the maximum injection volume was 1  $\mu$ L. We used 4  $\mu$ mol/eye (5 mmol/L × 0.8  $\mu$ L/eye, 10 mmol/L × 0.4  $\mu$ L/eye) and 8  $\mu$ mol/eye (10 mmol/L × 0.8  $\mu$ L/eye), three different doses for IOI, and the results did not have a measurable difference. Finally, we selected the minimal volume for subsequent experiments.

Eventually, we identified the biomarkers of DR. Previous studies have reported on VEGF, monocyte chemoattractant protein-1, TNF- $\alpha$ , transforming growth factor- $\beta$ , and nicotinamide adenine dinucleotide phosphate oxidase as the markers of inflammation. L-citrulline, indoleacetic acid, chenodeoxycholic acid, and eicosapentaenoic acid have been reported as markers of metabolism[51]. SCG3 is an angiogenic factor restricted to pathological conditions in DR[52]. Jiao et al[53] demonstrated that the concentration of SCG3 in the vitreous increased in 77 patients with DR. LeBlanc et al[54] reported that SCG3 antibodies alleviated retinal vascular leakage in diabetic mice with high efficacy. Furthermore, we assessed SCG3 expression in cells and tissues. Our results implied that SCG3 may be another biomarker of DR, which displayed a similar growing tendency in high glycogen to VEGF expression but with a higher sensitivity.

However, this study had several limitations. First, how the CHGA/UCP2/GLUT1 pathway is regulated by AGEs-RAGE binding requires further research. Some studies have demonstrated that they may form a GLUT1-UCP2 complex attached to the mitochondrial membrane; nonetheless, some details are still unknown. Second, we did not explain whether different doses of AGEs influence the severity of DR. Consequently, we added quantitative exogenous AGEs and observed changes in the cells and tissues. Third, the IOI group comprised limited doses in vivo. Next, we will add more dose groups injected with perfect genipin drug transduction in vivo and evaluate clinical medicine. Finally, we determined that SCG3 may replace VEGF in evaluating DR; however, the function of SCG3 in angiogenesis is unclear.

Our future work will focus on the changes in SCG3 during abnormal physiological processes. Some researchers have mentioned that SCG3 is central to tumors, apoptosis, and other vital processes[55]. Togayachi et al[56] demonstrated that SCG3 is activated by adding N-glycosylation to change short-form SgIII into a long form in small-cell lung carcinoma. This biological process may be related with the GLUT family. To further explore the relevant mechanisms, we used UniProt (https://www.uniprot.org/) to determine the structure of SCG3 and explained the potential target site of glycosylation (Figure 7A, Table 2). The ZDOCK score for RAGE and SCG3 was 1277.337, which means that RAGE can be linked with SCG3 easily. There are totally three positions of RAGE associated with SCG3 (at sites 216, 231, and 359; Figure 7B). PNGase F, used to split SCG3, decreased SCG3 protein expression (Figure 7C).

# CONCLUSION

Overall, apoptosis, angiogenesis, proliferation, ROS production, carbohydrate metabolism, and inflammation can reflect DR damage both in vivo and in vitro. Genipin is useful against AGEs, and its protective role in DR has been confirmed. This action may occur via AGEs-RAGE binding to control the CHGA/UCP2/GLUT1 pathway. SCG3 is a novel and sensitive biomarker for DR.

Table 2 Features for signal, chain, modified residue, and glycosylation in SCG3				
Туре	Position	Description		
Signal	1-19	-		
Chain	20-468	Secretogranin-3		
Modified residue	37	Phosphoserine		
Glycosylation1	216	O-linked (GalNAcan) threonine		
Glycosylation 2	231	O-linked (GalNAcan) threonine		
Glycosylation 3	359	O-linked (GalNAcan) threonine		
Modified residue	362	Phosphoserine		



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Figure 7 SCG3 plays a vital role in diabetic retinopathy via glycosylation. A: Structure of SCG3; B: Molecular docking simulation with RAGE (green, left) and SCG3 (orange, right); C: PNGase F cleavaged Scg<sub>3</sub>. NG: 5 mmol/l glucose containing ECM-treated human retinal microvascular endothelial cells (hRMECs) group; HG: 30 mmol/L glucose containing ECM-treated hRMECs group; GG: 0.4 µM genipin + 30 mmol/L glucose containing ECM-treated hRMECs.

# ARTICLE HIGHLIGHTS

# Research background

Diabetic retinopathy (DR) is a serious and common complication of diabetes. Advanced glycation end products (AGEs) are a group of reversible and poisonous products formed by nonenzymatic glycation of glucose with protein and lipids under hyperglycemia conditions. Both acute and chronic hyperglycemia can enhance AGEs production, which results in endothelial dysfunction and causes severe damage to diabetic retina. Some traditional Chinese herb like Gardenia jasminoides Ellis (GJE) fruit is a selective inhibitor of AGEs.



#### **Research motivation**

To confirm the effect of genipin, a vital component of GJE fruit, in preventing human retinal microvascular endothelial cells (hRMECs) from AGEs damage in DR and explored its mechanism.

#### **Research objectives**

To demonstrate whether genipin could lessen the development of DR in experimental models (4-wk-old C57/BL6 mice) through AGEs inhibition.

#### **Research methods**

Cell Counting Kit-8 (CCK-8) assay, colony formation assay, flow cytometry, immunofluorescence, wound healing assay, transwell assay, and tube-forming assay were used to detect the effect of genipin on hRMECs *in vivo*. Streptozotocin induced mice were used to explore retinal dysfunction with DM *in vitro*. Reactive oxygen species (ROS), mitochondrial membrane potential (MMP), and 2-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) amino]-2-deoxy-d-glucose assays were used to evaluate energy metabolism and oxidative stress damage in high glucose-induced hRMECs and STZ mouse retinas. Immunofluorescence and Western blot were used to investigate the expression of inflammatory cytokines (VEGF, SCG3, TNF- $\alpha$ , IL-1 $\beta$ , IL-1 $\beta$ , and NLRP3).

#### **Research results**

Our study confirmed that genipin ameliorated AGEs-induced hRMECs proliferation, apoptosis, energy metabolism, oxidative stress, and inflammatory injury, partially *via* the CHGA/UCP2/glucose transporter 1 pathway. Control of AGEs by IOI of genipin may represent a strategy to prevent severed retinopathy and vision loss.

#### Research conclusions

Our study suggested that intraocular injection of genipin can ameliorate AGEs to control DR. Control of AGEs and using principal bioactive components extracted from herb by intraocular injection may represent strategies to prevent DR.

#### **Research perspectives**

Our future work will focus on the changes in SCG3 during abnormal physiological processes.

# FOOTNOTES

**Author contributions:** Sun KX and Hu K were involved in design and conduct of the study, and preparation of the manuscript; Sun KX, Chen YY, and Li Z participated in the collection of the data; Zheng SJ, Wan WJ, Ji Y, and Hu K participated in the management of this program; all authors have read and approved the final manuscript.

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ORIGINAL ARTICLE

# **Basic Study** XB130 inhibits healing of diabetic skin ulcers through the PI3K/Akt signalling pathway

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# Abstract

# BACKGROUND

Diabetic skin ulcers, a significant global healthcare burden, are mainly caused by the inhibition of cell proliferation and impaired angiogenesis. XB130 is an adaptor protein that regulates cell proliferation and migration. However, the role of XB130 in the development of diabetic skin ulcers remains unclear.

# AIM

To investigate whether XB130 can regulate the inhibition of proliferation and vascular damage induced by high glucose. Additionally, we aim to determine whether XB130 is involved in the healing process of diabetic skin ulcers, along with its molecular mechanisms.

# **METHODS**

We conducted RNA-sequencing analysis to identify the key genes involved in diabetic skin ulcers. We investigated the effects of XB130 on wound healing using histological analyses. In addition, we used reverse transcription-quantitative polymerase chain reaction, Western blot, terminal deoxynucleotidyl transferasemediated dUTP nick end labeling staining, immunofluorescence, wound healing, and tubule formation experiments to investigate their effects on cellular processes in human umbilical vein endothelial cells (HUVECs) stimulated with high glucose. Finally, we performed functional analysis to elucidate the molecular mechanisms underlying diabetic skin ulcers.

# **RESULTS**



2023

RNA-sequencing analysis showed that the expression of XB130 was up-regulated in the tissues of diabetic skin ulcers. Knockdown of XB130 promoted the healing of skin wounds in mice, leading to an accelerated wound healing process and shortened wound healing time. At the cellular level, knockdown of XB130 alleviated high glucose-induced inhibition of cell proliferation and angiogenic impairment in HUVECs. Inhibition of the PI3K/Akt pathway removed the proliferative effects and endothelial protection mediated by XB130.

#### CONCLUSION

The findings of this study indicated that the expression of XB130 is up-regulated in high glucose-stimulated diabetic skin ulcers and HUVECs. Knockdown of XB130 promotes cell proliferation and angiogenesis *via* the PI3K/Akt signalling pathway, which accelerates the healing of diabetic skin ulcers.

Key Words: XB130; Diabetes mellitus; Diabetic skin ulcers; PI3K/Akt signalling pathway

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**Core Tip:** The role of XB130 in the occurrence and development of diabetic skin ulcers healing is unclear. This study showed that the expression of XB130 was up-regulated in tissues of diabetic skin ulcers and human umbilical vein endothelial cells (HUVECs) stimulated by high glucose. Knockdown of XB130 promote the healing of skin wounds in mice, leading to an accelerated wound healing process and shortened wound healing time and alleviated hyperglycemia-induced cell proliferation inhibition and angiogenic impairment in HUVECs.

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### INTRODUCTION

Diabetes mellitus (DM) is a chronic metabolic disease in which the body does not produce enough insulin or reacts abnormally to the hormone, leading to typically high blood sugar levels. In 2021, more than 37 million Americans have diabetes. Of those, 28.7 million have been diagnosed and 8.5 million are undiagnosed[1]. Impaired wound healing often results in severe skin ulcers[2]; a high percentage (15%-27%) of diabetic foot skin ulcer cases require lower extremity amputation owing to treatment failure[3]. Therefore, developing effective strategies to prevent and treat cutaneous wounds in patients with diabetes is crucial.

Impaired healing of diabetic wounds is characterised by significant deficits in cellular proliferation and migration, and a notable decrease in protein synthesis[4,5]. These impairments result in delayed re-epithelialisation, angiogenesis, and granulation tissue formation[6]. Numerous molecular pathways play crucial roles in cell proliferation and protein synthesis during the wound healing process[7]. Laplante *et al*[8] discovered that mTOR signalling could effectively treat diabetic skin damage by promoting cell proliferation, inhibiting skin cell apoptosis, and accelerating epithelial regeneration.

XB130 is a bridging protein that serves as the mediator of multiple tyrosine kinases, playing an important role in regulating cell proliferation, survival, migration, and invasion[9,10]. As a tumor suppressor in carcinogen-induced skin tumorigenesis, knockout of XB130 significantly up-regulated epidermal tumor cell proliferation[11]. XB130 also found in stomach, oesophagus, and thyroid epithelial cells, and has been studied as a signal transduction protein[12-14]. The expression of XB130 were up-regulated in cholangiocarcinoma, non-small cell-lung cancer, prostate cancer, esophageal squamous cell carcinoma and gastric cancer, regulating XB130 expression may inhibit the development and progression of tumors[12,13,15-17]. However, whether XB130 regulate wound healing in patients with diabetes and the underlying molecular mechanisms have not been reported.

In this study, we first confirmed that the expression of XB130 was up-regulation in tissues of diabetic skin ulcer mice. Knockdown of XB130 can accelerate the healing process and shorten the healing time of skin wound in mice. At the cellular level, down-regulation of XB130 alleviated hyperglycaemia-induced inhibition of cell proliferation and angiogenic impairment in human umbilical vein endothelial cells (HUVECs). Mechanistically, down-regulation of XB130 accelerates healing of diabetic skin ulcers by promoting cell proliferation and angiogenesis through the PI3K/Akt signalling pathway.

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# MATERIALS AND METHODS

# Animals

C57BL/6 mice (20-25 g, 6 wk old) were obtained from Beijing Weitong Lihua Experimental Animal Technology, China. The mice were provided with free access to food and water; the temperature and relative humidity of the room were maintained at 20 °C-24 °C and 40%-60%, respectively. The mice were allowed to acclimatize to their surroundings for 1 wk prior to the experiment.

# Diabetes induction and wound creation

Diabetes was induced in mice by injecting them with streptozotocin (STZ; 40 mg/kg) for 5 consecutive days. The control group received an equivalent dose of citrate buffer solution[18]. Diabetic status was determined in mice if their blood glucose levels exceeded 250 mg/dL following STZ injection. Both diabetic and normal mice were anaesthetised with 2% isoflurane. Full-thickness excisional wounds measuring 8 mm in diameter were created on the dorsal skin of the mice.

# Adenovirus

Adenovirus knockdown of XB130 targeting sequence and its corresponding control were constructed by ABM (Nanjing, China). Mice were given tail vein injections of adenovirus at the concentration of  $5 \times 10^{11}$  genome-equivalents five days before surgery.

# Haematoxylin & eosin and Masson staining

On day 14 post-treatment, the mice were euthanised and their wound tissues were fixed in 4% paraformaldehyde and embedded in paraffin. Haematoxylin & eosin (HE) and Masson staining were performed using kits from Solarbio (China) and Sigma (United States), respectively, following the manufacturer's protocols. Images were captured using a Ci-L microscope (Nikon, Japan) and analysed using the Image J software (version 1.8.0).

# Cell culture and transfection

HUVECs were sourced from Guangzhou Saliai Stemcell Co., Ltd. And cultured using EGM-2 BulletKit (Lonza). Subconfluent cells from passages five to seven were used for the experiments. Before cell culture procedures, stock media were replaced with phenol red-free low-glucose DMEM (Gibco, United States) supplemented with 1% calf serum (Gibco, CA, United States) and left for 12 h. HUVECs were then exposed to EGM-2 supplemented with either normal glucose (NG, 5.5 mmol/L) or high glucose (HG, 33 mmol/L) for 72 h, while using D-mannitol as an osmotic control for the HG condition. The knockdown plasmid of XB130 and its corresponding control plasmid were obtained from Guangdong Ruibo Biotechnology Co., Ltd. Transfection efficiency was confirmed by reverse transcription-quantitative polymerase chain reaction (RT-qPCR) and Western blot.

# Transferase-mediated dUTP nick end labeling assay

Transferase-mediated dUTP nick end labeling (TUNEL) staining was performed according to the instruction of TUNEL apoptosis detection kit (Promega, United States). The cells were seeded onto 24-well plates and incubated at 37 °C for 12 h. After staining, cell apoptosis was counted by photographing. The apoptosis cells were brown considered as the TUNEL-positive cells. Apoptotic rate was evaluated.

# Tubular formation assay

Upon completion of the experimental protocol, HUVECs were stained with calcein (Corning, United States). The stained cells were then replated onto Matrigel-precoated 24-well plates containing growth-factor reduced Matrigel (150 µL/well) and incubated at 37 °C for 12 h. To assess capillary-like tubule formation, a computer-assisted microscope was used to detect tube-like structures that were at least four times longer than their widths.

# Wound healing assay

To evaluate cell migration, a scratch assay was performed as described previously[19]. Following overnight incubation, the cells were seeded onto a 3.5-cm-diameter dish and allowed to form a confluent monolayer. Subsequently, a wound was created by scratching with a 200-mL pipette tip. The wounded cell monolayers were then imaged using a CoolSNAP HQ CCD (Nippon Roper Japan) at 0 h and 24 h post-wounding.

# Immunofluorescence staining

Well-grown HUVECs were fixed with 4.0% paraformaldehyde in phosphate-buffered saline for 10 min and permeabilised with 0.5% Triton for 15 min. Antibodies against PCNA, Ki67, Bcl2, Bax, and Cleaved caspase-3 obtained from Abcam (United States) were then used to incubate the cells overnight, followed by a 1 h incubation with or without Alexa Fluor 488-conjugated anti-mouse IgG secondary antibody (Abcam, United States) at room temperature. Nuclei were labelled with the fluorescent dye DAPI after 5 min of incubation. The cells were then observed under a confocal microscope (Leica, Germany) under each experimental condition.

# RT-qPCR

Total RNA was extracted from skin tissues or HUVECs using TRIzol reagent (Invitrogen, United States) and reversetranscribed into cDNA using a Tiangen Biotechnology (China) kit. RT-qPCR was conducted with β-Actin as an internal



Table 1 Primer sequences				
Gene	Forward (5'→3')	Reverse (5'→3')		
XB130	AAGCAGCAGCTCTGATGAGG	GGTCTGGAAGGCTCTTCTGA		
β-Actin	GGCTGTATTCCCCTCCATCG	CCAGTTGGTAACAATGCCATGT		

#### Table 2 Antibodies used in this study

Gene	Brand	Provenance
XB130	Abcam	Rabbit
Cleaved-caspase-3	Abcam	Rabbit
Bcl2	Abcam	Rabbit
Bax	Abcam	Rabbit
Ki67	Abcam	Rabbit
PCNA	Abcam	Rabbit
АКТ	Abcam	Rabbit
p-AKT	Abcam	Rabbit
p85α	Abcam	Rabbit
p-p85α	Abcam	Rabbit
β-Actin	Sigma	Mouse

control, using 2 × SYBR Green QPCR Master Mix (Shanghai Dongsheng Biotechnology, China). The relative gene expression was determined using the  $2^{-\Delta\Delta Ct}$  method, and the primer sequences are provided in Table 1.

#### Western blot

The protein from cell or tissue was extracted by RIPA and PMSF (Shanghai Life Mode Engineering, Shanghai, China), and the concentration of protein was determined by BCA kit (Shanghai Dongsheng Biotechnology, shanghai, China). The PVDF membrane (0.22 µm, Millipore ISEQ00010, United States) was then incubated with primary antibodies (Table 2) overnight at 4 °C, followed by incubation with HRP-conjugated secondary antibodies (Abcam, United States). Protein bands were detected using Prime Western Blot Detection Reagent (Cytiva, United Kingdom). A ChemiDoc MP imaging system (Tanon 4800, China) was used to detect chemiluminescence, and the ImageJ software was used to analyse the grey values of the bands.

#### Statistical analysis

Experimental data were analysed using GraphPad Prism 9.0, SPSS 24.0, and R software 4.2.0. Statistical tests were chosen based on the distribution and variance homogeneity of the data. For normally distributed and homogeneous variance data, t-tests were used to compare two groups. For multiple group comparisons Dunnett's-T3 test was used based on the distribution and variance. Data are presented as mean  $\pm$  SD, and statistical significance was set at P < 0.05.

# RESULTS

#### Construction of a diabetic mice model

The toxicity of STZ to pancreatic cells makes it a popular choice for inducing diabetes mellitus in mice and rats[20]. After administering STZ for 5 consecutive days, the model group showed significantly elevated fasting blood glucose levels compared with the control group (Figure 1A). As shown in Figure 1B, the model group exhibited slower wound healing time and rate. Tissue samples from the model and control groups were collected on day 14 for HE and Masson staining. The wounds of the model mice showed severe tissue damage, disorganised tissue granulation, and reduced collagen deposition (Figure 1C and D).

# HG inhibited proliferation and tubule formation in HUVECs

To simulate the HG environment during diabetic wound healing, we incubated HUVECs in an HG medium. Immunofluorescence staining for Bax and Cleaved caspase-3 in HG-treated HUVECs was higher than that in the NG group, whereas staining for PCNA, Ki67, and Bcl2 was lower (Figure 2A). The results of immunofluorescence analysis were confirmed by Western blot analysis (Supplementary Figure 1A and B). Moreover, HG treatment significantly impaired





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Figure 1 Establishment of diabetic model using streptozotocin. A: The blood glucose level of mice, control (n = 10) and model (n = 10); B: Representative skin wound images from control and model mice at day 0 d, 7 d, and 14 d; C and D: HE and Masson staining of skin wound in each group. <sup>a</sup>P < 0.0001, compared to the control group.

HUVEC migration and tube-forming activity (Figure 2B and C). We used the TUNEL assay to assess apoptosis in different groups and revealed an increased number of TUNEL-positive cells in the HG group (Figure 2D).

#### HG led to increased expression of XB130 in vivo and in vitro

To investigate the genes involved in diabetic wound healing, RNA sequencing was performed on the ulcer tissues from control and model mice group. Prior to differential expression analysis, we conducted background correction, normalisation, and gene filtering. Total 1547 differentially expressed genes (DEGs) were yielded (Figure 3A-C). We employed the STRING database for protein-protein interaction (PPI) network analysis to better understand the interactions between DEGs. Using the CytoHubba plug-in in Cytoscape, we identified hub genes based on their degree, betweenness, and closeness centrality, and found XB130 to be the most significant hub gene (Figure 3D).

To confirm the accuracy of the microarray results, we assessed XB130 mRNA and protein levels using RT-qPCR and Western blot. Compared with the control group, the mRNA level of XB130 was significantly increased in the model group (Figure 4A). HG also resulted in increased mRNA expression of XB130 in HUVECs (Figure 4D). Western blot analysis validated the RT-qPCR results (Figure 4B, C, E, and F).

#### Down-regulation of XB130 promoted wound healing in diabetic mice

Next, we established a mouse model knockdown XB130 (XB130-KD) in diabetic mice by injection of adenovirus vectors carrying short hairpin RNA targeting XB130. The XB130-KD group showed a significant decrease in protein expression of XB130 compared with the vector group (Figure 5A and B). Subsequently, we evaluated the wound healing process and found that knocking down XB130 Led to a shorter wound healing time and a higher wound healing rate (Figure 5C). Furthermore, histological analysis revealed that knocking down XB130 resulted in better granulation formation, collagen deposition, and denser alignment compared with the model group (Figure 5D).

#### Down-regulation of XB130 attenuated HG-induced inhibitory effects on proliferation and tubule formation in vitro

We evaluated the potential function of XB130 in HUVECs. After transfection, the expression of XB130 was significantly reduced in the XB130-KD group (Figure 6A and B). Immunofluorescence results showed that XB130 down-expression had a restraining effect on cell apoptosis and promoted cell proliferation in HUVECs (Figure 6C). These findings were further confirmed by Western blot analysis (Supplementary Figure 2A and B). Additionally, the TUNEL assay indicated that knocking down of XB130 reduced HG-induced apoptosis in HUVECs (Supplementary Figure 2C). Furthermore, XB130 down-expression counteracted the negative impact of hyperglycaemia on the migration and tube-forming activity of HUVECs (Figure 6D-E).



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Figure 2 High glucose inhibited proliferation and tubule formation of human umbilical vein endothelial cells. A: Immunofluorescence staining of Bax, cleaved caspase-3, PCNA, Ki67, and Bcl2 in human umbilical vein endothelial cells (HUVECs); B: Wound healing assay in HUVECs, scale bars = 200 µm; C: Capillary-like tubule formation, scale bars = 200 µm; D: Transferase-mediated dUTP nick end labeling assay.

# HG inhibited the PI3K/Akt signalling pathway in vivo and in vivo

To investigate the mechanism underlying delayed wound healing in mice patients with diabetes, we conducted a functional enrichment analysis of the previously obtained hub genes. Gene Ontology (GO) analysis revealed that the hub genes were involved in the of cell proliferation, differentiation, and tube morphogenesis (Figure 7A). Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway analysis indicated that these genes were predominantly associated with the PI3K/Akt, Ras, p53, and NF-kappa B signalling pathways. Notably, the PI3K/Akt signalling pathway had the highest number of differentially expressed genes (Figure 7B).

To validate this finding, we performed Western blot analysis and confirmed that the ratio of p-AKT to AKT was significantly decreased in diabetic mice (Figure 7C and D). Similar to results in vivo, HG levels inherited the PI3K/Akt signalling pathway in HUVECs (Figure 7E and F).

#### Down-regulation of XB130 accelerated wound healing in diabetic mice via the PI3K/Akt signalling pathway

To investigate the effect of XB130 on wound healing via this pathway, diabetic mice with reduced XB130 expression were treated with LY294002, an inhibitor of the PI3K/Akt signalling pathway. Western blot analysis revealed that knocking down XB130 stimulated the Akt phosphorylation, however, the AKT phosphorylation was suppressed by LY294002 treatment (Figure 8A and B). Additionally, wound healing time was significantly longer in the LY294002 group than in the XB130-KD group (Figure 8C). Histological analysis showed that LY294002 treatment decreased collagen deposition in the wound area and angiogenesis in the granulation tissue (Figure 8D).

# Down-regulation of XB130 reversed HG-induced inhibitory effects on proliferation and tubule formation via the PI3K/Akt signalling pathway

Western blot analysis showed that knocking down XB130 stimulated the PI3K/Akt signalling pathway, which was subsequently inhibited by LY294002 in HUVECs (Figure 9A and B). In LY294002-treated HUVECs, the protective effect of knocking down XB130 against HG was lost, resulting in increased apoptosis and decreased proliferation (Figure 9C). Western blot analysis yielded similar results (Supplementary Figure 3A and B). The TUNEL assay further demonstrated that LY294002 reversed the inhibitory effect of knocking down XB130 on HUVEC apoptosis (Supplementary Figure 3C). Moreover, in LY294002-treated HUVECs, the effects of knocking down XB130 on the pro-migration and tube-forming



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Figure 3 Differential expression genes identification and hub genes identification. A: Volcano plot; B: Histogram; C: Heatmap; D: Top 10 hub genes obtained from protein-protein interaction network.



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**Figure 4 High glucose environment increased the expression of XB130.** A: Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analysis of XB130 mRNA expression in mice; B: Western blot analysis of XB130 protein expression in mice; C: Quantification of Western blot results; D: RT-qPCR analysis of XB130 mRNA expression in human umbilical vein endothelial cells (HUVECs); E: Western blot analysis of XB130 protein expression in HUVECs; F: Quantification of Western blot results. °P < 0.05, °P < 0.01, °P < 0.001, compared to the control or NG group.

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**Figure 5 Knock-down of XB130 accelerated wound healing in diabetic mice.** A: Western blot analysis of XB130 protein expression in each group; B: Quantification of Western blot results; C: Representative skin wound images from control and model mice at day 0 d, 7 d, 14 d, control (n = 10), model (n = 10), Vector (n = 10), and XB130-KD (n = 10); D: HE and Masson staining of skin wound in each group. <sup>a</sup>P < 0.001, compared to the Control group; <sup>b</sup>P < 0.01, compared to the Vector group.

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HG + XB130-KD NG HG HG + Vector Bax Cleaved caspase-3 PCNA Ki67 Bcl2

D

A

С





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Figure 6 Knockdown of XB130 attenuated hyperglycemia-induced inhibitory effects on proliferation and tubule formation of human umbilical vein endothelial cells. A: Western blot analysis of XB130 protein expression in human umbilical vein endothelial cells; B: Quantification of Western blot results; C: Immunofluorescence staining of Bax, cleaved caspase-3, PCNA, Ki67, and Bcl2 in each group; D: Wound healing assay in each group, scale bars =  $200 \mu m$ ; E: Capillary-like tubule formation, scale bars =  $200 \mu m$ . <sup>a</sup>P < 0.001, compared to the high glucose group; <sup>b</sup>P < 0.001, compared to the HG + Vector group.



**Figure 7 High glucose inhibited PI3K/Akt signalling pathway** *in vivo* and *in vitro*. A: Barplot of Gene Ontology enrichment analysis; B: Dotplot of Kyoto Encyclopedia of Genes and Genomes pathway analysis; C: Western blot analysis of AKT and p-AKT expression in mice; D: Quantification of Western blot results; E: Western blot analysis of AKT and p-AKT expression in human umbilical vein endothelial cells (HUVECs); F: Quantification of Western blot results. <sup>a</sup>P < 0.01, compared to the control or NG group.

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**Figure 8 Knock-down of XB130 accelerated wound healing in diabetic mice via PI3K/Akt signalling pathway.** A: Western blot analysis of AKT and p-AKT expression in each group; B: Quantification of Western blot results; C: Representative skin wound images from control and model mice at day 0, 7, and 14, model (n = 10), XB130-KD (n = 10), LY294002 (n = 10), XB130-KD + LY294002 (n = 10); D: HE and Masson staining of skin wound in each group. <sup>a</sup>P < 0.01, compared to the control group; <sup>b</sup>P < 0.05, compared to the XB130-KD group.

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Figure 9 Knock-down of XB130 attenuated hyperglycemia-induced inhibitory effects on human umbilical vein endothelial cells via PI3K/Akt signalling pathway. A: Western blot analysis of AKT and p-AKT expression in human umbilical vein endothelial cells; B: quantification of Western blot results; C: Immunofluorescence staining of Bax, cleaved caspase-3, PCNA, Ki67, and Bcl2 in each group; D: Wound healing assay in each group, scale bars = 200  $\mu$ m; E: Capillary-like tubule formation, scale bars = 200  $\mu$ m. <sup>a</sup>P < 0.01, compared to the high glucose group; <sup>b</sup>P < 0.05, compared to the HG + XB130-KD group.

activities against HG impairment were removed (Figure 9D and E).

# DISCUSSION

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Diabetic skin ulcers, resulting from both internal and local pathological changes caused by diabetes, are a serious complication that can lead to amputation, despite therapeutic management[21,22]. The process of wound healing is complex, dynamic, and orderly, and is characterised by cell proliferation and migration[23,24]. In this study, we chose STZ, a well-established method for chemically inducing diabetes, to construct diabetic mice as an experimental model [18]. HUVECs were cultured *in vitro* in a high-glucose medium to mimic diabetic conditions. Our findings demonstrated that a high-glucose environment prolongs wound healing by inhibiting cell proliferation and tubule formation.

Aberrant gene expression is associated with various pathological conditions, including diabetic skin ulcers[25]. However, key driver genes that trigger and exacerbate this condition are not fully understood. We performed RNA sequencing the ulcer tissues from control and model mice group and identified 980 upregulated and 567 downregulated genes. Moreover, by performing PPI network analysis, we identified ten hub genes associated with diabetic skin ulcers (XB130, ITGAM, FCGR3A, PECAM1, CXCR4, CD34, CXCL8, CCL3, VCA11, and CD19). Among them, XB130 had the highest score based on topological algorithms. The expression of XB130 was up-regulated in HG induced diabetic skin ulcers tissue and HUVECs using RT-qPCR and Western blot.

XB130 is vital for regulating signal transduction and affects the cell cycle, proliferation, survival, and migration[14,16, 17]. Although limited research has explored the relationship between XB130 and diabetic wound healing, our study aimed to address this knowledge gap by creating a model of XB130-downexpressed diabetic mice. Our findings indicate that knockdown of XB130 can enhance the healing of diabetic wounds, as evidenced by shorter epithelialisation time and rapid wound contraction. Moreover, histological analysis revealed that knockdown of XB130 not only improved diabetic wound healing, but also significantly promoted angiogenesis. To investigate the mechanism and function of XB130, we establish a knockdown of XB130 with HUVCEs. Our results revealed that the expression of PCNA, Ki67 and Bcl2 were significantly increased, while decreasing the expression of Bax and Cleaved caspase-3 in XB130-KD HUVECs treated with HG. Furthermore, knockdown of XB130 enhanced the migration and tube-forming activities of HUVECs. These findings indicate that knockdown of XB130 promotes diabetic wound healing by enhancing cell proliferation and tubule formation.

To investigate how HG environment affects wound healing, functional analysis of hub genes was performed. GO analysis revealed that the hub genes were enriched in cell proliferation, differentiation, migration, apoptosis, and tube morphogenesis. These results confirmed that HG levels hindered wound healing by affecting these cellular processes. Meanwhile, KEGG pathway analysis showed that these genes were involved in PI3K/Akt, Ras, p53, and NF-kappa B signalling pathways, and in apoptosis. Among these, the PI3K/Akt signalling pathway was the most important, with seven enriched genes. Our findings further prove that HG conditions reduce the ratio of p-AKT/AKT, leading to inhibition of the PI3K/Akt pathway.

The PI3K/AKT pathway plays a pivotal role in regulating cell proliferation and metabolism and is involved in the progression of various diseases[26-29]. When RTKs and GPCRs are activated, PI3K phosphorylates phosphatidylinositol 3,4-diphosphate, resulting in the production of PIP3. PIP3 activates AKT, which in turn phosphorylates and activates the mTORC complexes. In turn, activated mTORC complexes promote cell proliferation by phosphorylating and activating matrix metalloproteinase[30-32]. Our results indicate that knockdown of XB130 increases the levels of phosphorylated AKT. LY294002 and Wortmannin are commonly used inhibitors of the PI3K/Akt pathway. Our findings indicate that LY294002 delays wound healing in diabetic mice. Furthermore, inhibition of the PI3K/Akt signalling pathway reverses the negative effects of XB130 on proliferation and tubule formation in HUVECs.

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# CONCLUSION

This study shows that knockdown of XB130 can prevent high glucose-induced inhibition of proliferation and angiogenic impairment through the PI3K/Akt pathway. Based on our results, decreasing the expression of XB130 could serve as a promising therapeutic approach to accelerate the healing of diabetic skin ulcers.

# **ARTICLE HIGHLIGHTS**

#### Research background

Diabetic skin ulcers are mainly caused by the inhibition of cell proliferation and impaired angiogenesis, a high percentage (15%-27%) of diabetic foot skin ulcer cases require lower extremity amputation owing to treatment failure. XB130 is an adaptor protein that regulates cell proliferation and migration.

#### Research motivation

To explore the role of XB130 in the development of diabetic skin ulcers.

#### Research objectives

To investigate whether XB130 can regulate the inhibition of proliferation and vascular damage induced by high glucose. Additionally, we aim to determine whether XB130 is involved in the healing process of diabetic skin ulcers, along with its molecular mechanisms.

#### Research methods

We conducted RNA-sequencing analysis to identify the key genes. The RT-qPCR, Western blot, TUNEL staining, immunofluorescence, wound healing, and tubule formation experiments were used to investigate their effects on cellular processes in human umbilical vein endothelial cells (HUVECs) stimulated with high glucose. Finally, we performed functional analysis to elucidate the molecular mechanisms underlying diabetic skin ulcers.

#### Research results

RNA-sequencing analysis showed that the expression of XB130 was up-regulated in the tissues of diabetic skin ulcers. Knockdown of XB130 promoted the healing of skin wounds in mice, leading to an accelerated wound healing process and shortened wound healing time. At the cellular level, knockdown of XB130 alleviated high glucose-induced inhibition of cell proliferation and angiogenic impairment in HUVECs. Inhibition of the PI3K/Akt pathway removed the proliferative effects and endothelial protection mediated by XB130.

#### Research conclusions

The expression of XB130 is up-regulated in high glucose-stimulated diabetic skin ulcers and HUVECs. Knockdown of XB130 promotes cell proliferation and angiogenesis via the PI3K/Akt signalling pathway, which accelerates the healing of diabetic skin ulcers.

#### Research perspectives

Decreasing the expression of XB130 could serve as a promising therapeutic approach to accelerate the healing of diabetic skin ulcers.

# FOOTNOTES

Author contributions: Pan WH, Liao WQ, Fang WJ, Lei WZ, and Zhu XL contributed to conceptualization; Zhu XL, Zeng ZX, and Jiang WW contributed to data curation; Zhu XL, Jiang WW, Zeng ZX, Hu DY, Chen TY, Chen TC, Liao WQ, Lei WZ, Fang WJ, and Pan WH contributed to investigation; Zhu XL, Zeng ZX, Jiang WW, Fang WJ, and Pan WH contributed to methodology; Pan WH contributed to project administration; Pan WH, Fang WJ, and Lei WZ contributed to supervision; Zeng ZX, Hu DY, and Zhu XL contributed to visualization; Zhu XL contributed to writing-original draft preparation; Liao WQ, Fang WJ, Pan WH, and Zhu XL contributed to writingreview and editing; Zhu XL, Hu DY, and Zeng ZX contribute equally to this paper.

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ORIGINAL ARTICLE

# Effects of paricalcitol combined with hemodiafiltration on bonemetabolism-related indexes in patients with diabetic nephropathy and chronic renal failure

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# Abstract

# BACKGROUND

Diabetic nephropathy (DN) is frequently seen in the development of diabetes mellitus, and its pathogenic factors are complicated. Its current treatment is controversial, and there is a lack of a relevant efficacy prediction model.

#### AIM

To determine the effects of paricalcitol combined with hemodiafiltration on bonemetabolism-related indexes in patients with DN and chronic renal failure (CRF), and to construct an efficacy prediction model.

# **METHODS**

We retrospectively analyzed 422 patients with DN and CRF treated in Cangzhou Central Hospital between May 2020 and May 2022. We selected 94 patients who met the inclusion and exclusion criteria. Patients were assigned to a dialysis group (n = 45) and a joint group (n = 49) in relation to the rapeutic regimen. The clinical efficacy of the two groups was compared after treatment. The changes in laboratory indexes after treatment were evaluated, and the two groups were compared for the incidence of adverse reactions. The predictive value of laboratory indexes on the clinical efficacy on patients was analyzed.

# RESULTS

The dialysis group showed a notably worse improvement in clinical efficacy than the joint group (P = 0.017). After treatment, the joint group showed notably lower serum levels of serum creatinine, uric acid (UA) and blood urea nitrogen (BUN) than the dialysis group (P < 0.05). After treatment, the joint group had lower serum levels of phosphorus, procollagen type I amino-terminal propeptide (PINP) and intact parathyroid hormone than the dialysis group, but a higher calcium level (P < 0.001). Both groups had a similar incidence of adverse reactions (P > 0.001).



0.05). According to least absolute shrinkage and selection operator regression analysis, UA, BUN, phosphorus and PINP were related to treatment efficacy. According to further comparison, the non-improvement group had higher risk scores than the improvement group (P < 0.0001), and the area under the curve of the risk score in efficacy prediction was 0.945.

#### **CONCLUSION**

For treatment of CRF and DN, combined paricalcitol and hemodiafiltration can deliver higher clinical efficacy and improve the bone metabolism of patients, with good safety.

Key Words: Paricalcitol; Hemodiafiltration; Diabetic nephropathy; Chronic renal failure; Serum calcium; Serum phosphorus; Intact Paricalcitol hormone

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**Core Tip:** This study confirmed that paricalcitol combined with hemodiafiltration can effectively improve the condition of patients with diabetic nephropathy (DN) and chronic renal failure (CRF) and alleviate calcium-phosphorus metabolism disorder. This study has also successfully constructed a predictive model. It provides a new reference for evaluating the efficacy of treatment of combined DN and CRF.

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# INTRODUCTION

Improvement of socioeconomic and living standards has resulted in an increase in the global prevalence of diabetes mellitus (DM), bringing an increasing incidence of diabetic nephropathy (DN)[1]. DN is one of the most frequent causes of chronic renal failure (CRF) worldwide, and 30%-40% of DM patients develop DN[2]. The pathogenesis of DN is complex, and is bound up with abnormal balance in the body, including hemodynamics, metabolic disorder, inflammation and fibrosis[3]. The imbalance between renal injury and renal protection factors is the primary cause of disease progression[4]. DN has a heavy burden on the global economy and a high mortality risk in DM patients.

DN is a most common complication in DM patients and a common cause of CRF[5]. Approximately 30%-40% of the patients with DM will develop DN, leading to CRF[6,7]. When DN progresses to the advanced stage, the patients may need dialysis or kidney transplantation to help maintain renal function and quality of life[8]. Hemodiafiltration is an effective method of treatment for CRF, which can prolong the survival of patients[9]. However, patients receiving hemodialysis often have abnormal serum calcium and phosphorus, and adverse reactions such as joint calcification, bone pain, arrhythmia, pruritus and platelet insufficiency[10].

Paricalcitol is a vitamin D analog that is used to treat secondary hyperparicalcitolism triggered by renal insufficiency [11]. Paricalcitol can promote intestinal absorption of calcium and inhibit secretion and differentiation of Paricalcitol cells by binding to vitamin D receptor, thus reducing the synthesis and secretion of Paricalcitol hormone (PTH)[12]. Paricalcitol can also alleviate kidney damage triggered by inflammation and apoptosis, thus helping to protect kidney function<sup>[13]</sup>.

However, there is controversy about the impact of paricalcitol combined with hemodiafiltration on serum calcium, phosphorus and intact PTH (iPTH) in patients with both DN and CRF. Accordingly, this study aimed to provide a basis for clinical treatment of comorbid DN and CRF.

# MATERIALS AND METHODS

#### Patient characteristics

We retrospectively analyzed 422 patients with both DN and CRF treated in Cangzhou Central Hospital between May 2020 and May 2022. The patients were screened according to the following criteria. Inclusion criteria: patients who met the diagnostic criteria of DN[14] and were not allergic to the drugs used in this study. Exclusion criteria: patients with primary nephropathy or secondary DM; patients comorbid severe dysfunction of the heart, liver or brain; patients with proteinuria due to other reasons; and patients who had received angiotensin-converting enzyme inhibitors or diuretics before admission. According to the above standards, 94 patients with DN and CRF were selected. The patients were assigned to a dialysis group (n = 45) and a joint group (n = 49) according to therapeutic regimen. This study was carried out with approval of the Medical Ethics Committee of Cangzhou Central Hospital.



# Therapeutic regimen

Both groups received symptomatic treatment, including correction of fluid and electrolyte imbalance, anti-allergy, antihemolysis and correction of acid poisoning. The dialysis group was treated by hemodiafiltration through an Fx80 dialyzer (Fresenius, Germany), with blood flow of 200-300 mL/min, and dialysis time of 4 h (3 times/wk). Twelve sessions were taken as a course of dialysis treatment, and two courses of treatment were conducted. The joint group was treated with paricalcitol while being treated with hemodiafiltration. Paricalcitol was injected at a dose of  $0.04-0.1 \, \mu g/kg$  (2.8-7.0  $\mu g$ ) each time, and the dose was adjusted according to the serum iPTH level. The frequency of administration was no more than once every other day, and it was administered for eight continuous weeks. During treatment, serum calcium, phosphorus and iPTH were detected every 2 wk, and the dose was adjusted according to the test results.

# Clinical data collection

The clinical data were collected through the medical record system of our hospital and included: age, gender, body mass index, course of renal decompensation, history of hypertension, history of hyperlipidemia, and history of smoking and alcoholism. Laboratory indexes included renal-function-related indexes [serum creatinine (SCr), uric acid (UA) and urea nitrogen (BUN)], and bone-metabolism-related indexes [serum phosphorus, serum calcium, procollagen type I Nterminal propeptide (PINP) and iiPTH]. Improvement of clinical efficacy in patients after treatment and the incidence of adverse reactions during treatment were evaluated.

# Evaluation criteria of clinical efficacy

Markedly effective: After treatment, the related symptoms of the patients subsided, and the levels of renal-functionrelated indexes such as SCr, UA and BUN decreased by over 30%. Effective: After treatment, the related symptoms were relieved, and the levels of SCr, UA and BUN decreased by 5%-30%. Ineffective: The related symptoms and condition of the patients were not notably alleviated, and there was even some aggravation of disease, with the indexes such as SCr, UA and BUN decreased by < 5%, or even increased. Total effective rate = (number of cases with markedly effectively treatment + number of cases with effective treatment]/total number of cases × 100%.

# Outcome measures

Primary outcome measures: The clinical efficacy on the two groups was compared. The changes in laboratory indexes in patients after treatment were evaluated. Secondary outcome measures: The clinical data of the two groups were compared. The adverse reactions of the two groups were also compared. The predictive value of laboratory indexes on the clinical efficacy on patients was analyzed.

# Statistical analysis

This study used R language 4.1.1 software (R Foundation for Statistical Computing, Vienna, Austria) for data reduction and data analysis, and established a model. The predictors of non-zero coefficient were screened by least absolute shrinkage and selection operator (LASSO) regression, and the nomogram was drawn by R (R3.5.3) software package and rms package. The consistency index (C-index) was calculated by rms package, and its clinical value was verified by the receiver operating characteristic curve. We used Graph Pad Prism 8.0 (La Jolla, CA, USA) for visualization of data. P < 0.05 implies a notable difference.

# RESULTS

# Comparison of clinical data

According to inter-group comparison of clinical data, the dialysis and joint groups were similar (P > 0.05, Table 1).

# Evaluation of clinical efficacy

The dialysis group showed a significantly lower improvement in clinical efficacy than the joint group (P = 0.017, Table 2).

# Changes in renal-function-related indexes

Before treatment, SCr, UA and BUN levels were similar in the dialysis and joint groups (P > 0.05). After treatment, SCr, UA and BUN levels in both groups decreased significantly (P < 0.05), with greater decreases in the joint group than in the dialysis group (P < 0.05, Figure 1).

# Changes in bone-metabolism-related indexes

Before treatment, the levels of phosphorus, calcium, PINP and iPTH did not differ significantly between the dialysis and joint groups (P > 0.05). After treatment, phosphorus, PINP and iPTH in both groups decreased significantly (P < 0.0001), while calcium increased significantly. After treatment, the joint group had significantly lower serum levels of phosphorus, PINP and iPTH than the dialysis group, and a significantly higher Ca level (P < 0.001, Figure 2).

# Occurrence of adverse reactions

The incidence of adverse reactions was similar in the joint and dialysis groups (P > 0.05, Table 3).



Table 1 Clinical data				
Factors	Dialysis group ( <i>n</i> = 45)	Joint group ( <i>n</i> = 49)	χ² value	P value
Age (yr)			2.042	0.153
> 60	25	20		
≤ 60	20	29		
Gender			0.641	0.423
Male	22	28		
Female	23	21		
BMI (kg/m <sup>2</sup> )			0.300	0.583
> 25	15	19		
≤ 25	30	30		
Course of renal decompensation (yr)			0.526	0.468
>2	29	35		
≤2	16	14		
History of hypertension			0.512	0.474
Yes	12	10		
No	33	39		
History of hyperlipidemia			0.566	0.451
Yes	8	6		
No	37	43		
History of smoking			0.641	0.423
Yes	22	28		
No	23	21		
History of alcoholism			0.222	0.637
Yes	6	5		
No	39	44		

BMI: Body mass index.

Table 2 Efficacy evaluation, n (%)						
Group	Markedly effective	Effective	Ineffective	Total effective rate		
Dialysis group ( $n = 45$ )	25 (55.56)	8 (17.78)	12 (26.66)	33 (73.34)		
Joint group ( $n = 49$ )	33 (67.35)	12 (24.48)	4 (8.16)	45 (91.84)		
$\chi^2$ value	5.686					
<i>P</i> value				0.017		

#### Value of laboratory indexes in predicting efficacy

The value of laboratory indexes for predicting treatment efficacy was determined using LASSO regression. Patients with markedly effective treatment and those with effective treatment were assigned to an improvement group (n = 78), while patients with ineffective treatment were assigned to a non-improvement group (n = 16). Through LASSO regression analysis, UA, BUN, phosphorus and PINP were related to treatment efficacy (Figure 3). A predictive equation was constructed according to the risk coefficient: = -0.0056463379 × UA + 0.1334320595 × BUN + 0.9356101817 × P + - 0.0002297355 × PINP. The non-improvement group had higher risk scores than the improvement group (Figure 4A, P < 0.0001), and the area under the curve of risk score in forecasting efficacy was 0.945 (Figure 4B).

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Table 3 Adverse reactions, n (%)						
Group	Nausea and vomiting	Loss of appetite	Phlebitis	Gastrointestinal reaction	Rash	Total incidence rate
Dialysis group ( $n = 45$ )	2 (4.44)	2 (4.44)	1 (2.22)	2 (4.44)	1 (2.22)	8 (17.78)
Joint group ( $n = 49$ )	1 (2.04)	1 (2.04)	2 (4.08)	1 (2.04)	1 (2.04)	6 (12.24)
$\chi^2$ value						0.566
<i>P</i> value						0.451



Figure 1 Changes in renal-function-related indexes in the patients before and after treatment. A: Comparison of uric acid changes in the two groups before and after treatment; B: Comparison of serum creatinine changes in the two groups before and after treatment; C: Comparison of blood urea nitrogen changes in the two groups before and after treatment. \*P < 0.05, \*P < 0.01 \*P < 0.001, 4P < 0.0001. SCr: Serum creatinine; UA: Uric acid; BUN: Blood urea nitrogen.



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Figure 2 Changes in bone-metabolism-related indexes in patients before and after treatment. A: Comparison of phosphorus changes before and after treatment: B: Comparison of calcium changes before and after treatment: C: Comparison of procollagen type I amino-terminal propeptide changes before and after treatment; D: Comparison of intact Paricalcitol hormone changes before and after treatment. \*P < 0.05, \*P < 0.01, \*P < 0.001, \*P < 0.0001. PINP: Procollagen type I amino-terminal propeptide; iPTH: intact Paricalcitol hormone.

#### DISCUSSION

DM is characterized by high incidence, low control rate and various complications worldwide[15]. DN is a common complication of DM, especially in patients with a disease course > 10 years [16]. The long-term presence of hyperglycemia triggers activation of the polyol pathway and protein kinase C pathway, resulting in a series of pathophysiological changes, such as oxidative stress, disorder of renal glucose metabolism, inflammatory reaction, abnormal metabolism, and abnormal hemodynamics, and finally inducing CRF[17]. The kidneys of patients with CRF shrink and lose the ability to maintain normal renal function, which results in serious consequences such as imbalance of acid and potassium, retention of metabolites, and fluid and electrolyte imbalance, which increases the risk of death[18].

Hemodiafiltration is a type of renal replacement therapy, which is primarily used for blood purification in patients with renal failure<sup>[19]</sup>. This method removes metabolites, toxins, and excess liquids and electrolytes from the blood through a series of filtration and dialysis processes. In hemodiafiltration treatment, the patient's blood is filtered and dialyzed by special filters and dialyzers, and the removed waste and excess liquid are excreted through the urine<sup>[20]</sup>. Hemodiafiltration can ameliorate the symptoms and quality of life of patients with kidney disease, but it also needs close monitoring and adjustment of treatment regimens to avoid complications[21]. According to prior research[22], patients



Figure 3 Least absolute shrinkage and selection operator-based screening of predictors. A, B: The coefficient distribution of the least absolute shrinkage and selection operator repression analysis and the calculation of adjustment parameters (lambda) based on partial likelihood deviation of 10 times cross-validation.



Figure 4 Risk score in patients with different efficacy. A: Risk score in the non-improvement group and improvement group; B: Receiver operating characteristic curve of risk score in predicting efficacy. <sup>d</sup>P < 0.0001.

often have calcium-phosphorus metabolism disorder during dialysis, which compromises its efficacy. Continuous hemodialysis can reduce urinary toxins, but it cannot replace normal renal metabolism and endocrine function. In this study, the clinical efficacy and renal function improvement of the joint group were significantly higher than those in the dialysis group after treatment. The results suggest that paricalcitol combined with hemodiafiltration can deliver significantly greater efficacy in patients with comorbid DN and CRF, and improve renal function.

Due to the loss of renal function, dialysis patients often have abnormal bone metabolism, including hyperphosphatemia, hypocalcemia, and hyperparicalcitolism. These abnormalities increase the risk of osteoporosis and fracture[23]. In hemodialysis therapy, the use of auxiliary drugs can improve this situation and reduce the occurrence of adverse reactions[24]. Therefore, in the bone metabolism management of dialysis patients, it is necessary to comprehensively consider factors including the clinical situation of patients and serum bone-metabolism-related indexes. In this study, the joint group showed significantly lower levels of phosphorus, PINP and iPTH than the dialysis group, and a significantly higher calcium level. The two groups showed no significant difference in the incidence of adverse reactions. The results indicate that paricalcitol combined with hemodiafiltration can control serum phosphorus and calcium levels, lower iPTH level, and improve bone-metabolism-related indexes, without increasing safety risks.

Predicting the clinical efficacy on patients is important in the treatment process[25]. For example, predictive models can help healthcare providers make more informed decisions and recommend specific treatments for individual patients, which can bring better results and more personalized care[26]. Predictive models can also help providers identify high-risk patients in certain situations, so that they can carry out early intervention to prevent or mitigate the development of these situations. In this study, we established a risk model to predict the efficacy based on laboratory indexes. In this study, UA, BUN, phosphorus and PINP were related to the efficacy in patients. The risk score of each patient was calculated. The non-improvement group had significantly higher risk scores than the improvement group. According to receiver operating characteristic curve-based analysis, the area under the curve of risk score for predicting efficacy was 0.945.

Our study confirmed that paricalcitol combined with hemodiafiltration improved the condition of patients with comorbid DN and CRF and corrected calcium-phosphorus metabolism disorder. We also successfully constructed a predictive model. However, the study still had some limitations. First, we did not collect data about long-term prognosis, and we only evaluated short-term efficacy, so whether there is any influence on long-term efficacy after treatment needs further study. Second, we need to verify whether the predictive model has high generality or not needs verification by more data. Finally, in such a single-center study, the lack of research samples may have led to bias in the analysis. We hope to carry out more studies in the future to improve the research conclusions.



# CONCLUSION

In the treatment of comorbid DN and CRF, the combined use of paricalcitol and hemodiafiltration delivered greater clinical efficacy and improved the bone metabolism of patients, with good safety.

# ARTICLE HIGHLIGHTS

#### Research background

Diabetic nephropathy (DN) is one of the common complications of diabetes, mainly manifested as glomerular damage. As it progresses, DN may lead to chronic renal failure (CRF), which seriously affects quality of life and life expectancy.

#### Research motivation

Hemodialysis filtration is an effective method for treating CRF, but patients receiving hemodialysis often experience abnormalities in blood calcium and phosphorus. Paroxycarbinol can promote intestinal calcium absorption and inhibit the secretion and differentiation of Paricalcitol cells by binding to vitamin D receptors. However, it is still unclear whether periostenol has an effect on calcium phosphate metabolism disorder in patients with CRF due to DN.

#### Research objectives

Supplement the blank of paracalcitol combined with hemodiafiltration in the treatment of CRF, and increases the clinical treatment plan for disorder of calcium and phosphate metabolism during hemodialysis and filtration.

#### Research methods

We retrospectively analyzed and observed the effect of paricalcitol combined with hemodiafiltration on calcium phosphate metabolism disorder in patients with CRF due to DN. For the first time, a risk model for predicting efficacy was established using a least absolute shrinkage and selection operator (LASSO) regression model.

#### Research results

We found that the combination of paricalcitol and hemodialysis filtration significantly improved the metabolic disorder of calcium phosphate metabolism disorder in patients, and improved efficacy. Using the LASSO model, we established a risk score to predict efficacy, which provides a new reference for clinical treatment and efficacy prediction.

#### Research conclusions

Paricalcitol can improve calcium phosphate metabolism disorder in hemodialysis patients, and the risk model established by LASSO regression model effectively predicts clinical efficacy.

#### Research perspectives

As a retrospective study, we cannot collect more samples and observe the prognosis of patients. We hope to conduct randomized controlled trials in future studies to observe the long-term prognosis.

# FOOTNOTES

Author contributions: Ma XY designed and performed the research and wrote the paper; Sun FY designed the research and supervised the report; Sheng YP and Yang XM designed the research and contributed to the analysis; Zhang HR provided clinical advice.

Institutional review board statement: The study was reviewed and approved by the Cangzhou Central Hospital.

Informed consent statement: All participants have signed an informed consent form.

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

Data sharing statement: Clinical data for this study can be obtained from the corresponding author.

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# Early neonatal complications in pregnant women with gestational diabetes mellitus and the effects of glycemic control on neonatal infection

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# Abstract

#### BACKGROUND

Gestational diabetes mellitus (GDM) has become increasingly prevalent globally. Glycemic control in pregnant women with GDM has a critical role in neonatal complications.

#### AIM

To analyze the early neonatal complications in GDM, and examine the effect of blood glucose control level on neonatal infection.

#### **METHODS**

The clinical data of 236 pregnant women with GDM and 240 healthy pregnant women and newborns during from March 2020 to December 2021 the same period were retrospectively analyzed, and the early complications in newborns in the two groups were compared. The patients were divided into the conforming glycemic control group (CGC group) and the non-conforming glycemic control group (NCGC group) based on whether glycemic control in the pregnant women with GDM conformed to standards. Baseline data, immune function, infectionrelated markers, and infection rates in neonates were compared between the two groups.

#### **RESULTS**

The incidence of neonatal complications in the 236 neonates in the GDM group was significantly higher than that in the control group (P < 0.05). Pregnant women with GDM in the NCGC group (n = 178) had significantly higher fasting plasma glucose, 2 h postprandial blood glucose and glycated hemoglobin A1<sub>c</sub> levels than those in the CGC group (n = 58) (P < 0.05). There were no differences in baseline data between the two groups (P > 0.05). Additionally, the NCGC group had significantly decreased peripheral blood CD3<sup>+</sup>, CD4<sup>+</sup>, CD8<sup>+</sup> T cell ratios, CD4/CD8



ratios and immunoglobulin G in neonates compared with the CGC group (P < 0.05), while white blood cells, serum procalcitonin and C-reactive protein levels increased significantly. The neonatal infection rate was also significantly increased in the NCGC group (P < 0.05).

#### **CONCLUSION**

The risk of neonatal complications increased in pregnant women with GDM. Poor glycemic control decreased neonatal immune function, and increased the incidence of neonatal infections.

Key Words: Gestational diabetes mellitus; Early neonatal complications; Glycemic control; Neonatal infection

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Core Tip: Gestational diabetes mellitus (GDM) is an important complication that affects pregnancy outcome. Pregnant women with GDM and long-term abnormal glucose metabolism are closely associated with the risk of adverse maternal and neonatal outcomes. Some studies suggest that the immune function of newborns may be significantly affected by GDM, and that the effect of glycemic control is related to pregnancy outcomes and neonatal prognosis. In this study, we confirmed that the risk of neonatal complications increased in pregnant women with GDM, and poor glycemic control leads to impairment of fetal immune system and ultimately increases the risk of neonatal infections.

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# INTRODUCTION

Gestational diabetes mellitus (GDM) is a glucose metabolism disorder during pregnancy. Studies have indicated that GDM tends to cause diabetes mellitus and cardiovascular disease in neonates after delivery by pregnant women with GDM[1]. Recently, studies reported that GDM incidence has shown a significant increasing trend worldwide[2] and has become a global health concern. Previous studies [3,4] have shown that GDM is considered to be one of the major risk factors for maternal and child complications in the perinatal phase and the incidence of cesarean section, polyhydramnios, premature birth, fetal deformities, neonatal infection, hyperbilirubinemia, fetal macrosomia, and neonatal hypoglycemia is higher in pregnant women with GDM who have abnormal glucose metabolism for a long period. Rational diet control, exercise, and glucose-lowering treatments have achieved good results in terms of glycemic control in pregnant women with GDM. However, 30% of pregnant women with GDM were reported to be affected by multiple factors, including irregular diet, lack of exercise, hormone secretion disorder, etc[5]. In addition, glycemic control was poor. This ultimately affected maternal and child health, and increased the risk of neonatal complications. A study found that the immune function of neonates may be significantly affected by GDM, and the effectiveness of glycemic control is strongly associated with adverse pregnancy outcomes and neonatal complications[6]. Early detection of abnormal glucose metabolism and achieving good glycemic control during pregnancy can effectively prevent adverse maternal and child outcomes in the perinatal stage of GDM patients<sup>[7]</sup>. However, there is still unclear whether glycemic control that does not conform to the standards in pregnant women with GDM decreases immune function in neonates and increases the incidence of neonatal infections.

In this study, we investigated the differences in neonatal complications between pregnant women with GDM and healthy controls to examine the effects of GDM on neonatal prognosis. We then divided the pregnant women with GDM into two groups based on whether glycemic control conformed to standards. Subsequently, we compared the blood glucose levels in pregnant women with GDM, immune function, infection-related marker levels, and the incidence of infection in neonates of the pregnant women with GDM in these two groups. The purpose of this study was to analyze the effects of glycemic control in GDM pregnant women on neonatal immune function and infection.

# MATERIALS AND METHODS

#### Patient characteristics

The newborns delivered by 236 pregnant women with GDM in Taizhou People's Hospital of Jiangsu Province from March 2020 to December 2021 were retrospectively included in the GDM group. The neonates of 240 healthy pregnant women during the same period were selected as the control group. The patients were divided into the conforming glycemic control group (CGC group) and the non-conforming glycemic control group (NCGC group) based on whether glycemic control in pregnant women with GDM conformed to standards.



Inclusion criteria: (1) Natural singleton pregnancy; (2) age 20-40 years; and (3) for the GDM group, blood glucose measurement at week 24-28 of pregnancy conformed to the diagnostic criteria for GDM formulated by the American Diabetes Association in 2013[8]: 75 g oral glucose tolerance test result showed fasting plasma glucose (FPG)  $\geq$  5.1 mmol/ L, blood glucose 1 h after test  $\geq$  10.0 mmol/L or blood glucose 2 h after test  $\geq$  8.5 mmol/L; the GDM group received diet and/or glucose-lowering treatment.

Exclusion criteria: (1) Comorbid hypertension, anemia, polyhydramnios, and other underlying diseases or pregnancy complications; (2) past history of adverse pregnancy outcomes; (3) gestational age at delivery < 28 wk; and (4) presence of heart, brain, lung, liver, and other organ diseases.

The screening process is shown in Figure 1. The study was reviewed and approved by the Taizhou People's Hospital of Jiangsu Province Institutional Review Board.

#### Treatment methods

Dietary control and/or insulin treatment was carried out in all pregnant women with GDM, which was specified as follows: total daily caloric intake was calculated based on 130 J/kg145 J/kg, and the proportions of carbohydrates, proteins, and fats were 55%-65%, 20%-25%, and 15%-25%, respectively. A routine diet complied with the principle of eating less in more meals and 4-6 meals were consumed each day. Insulin treatment of 0.6 U/kg-0.8 U/kg was administered every day, blood glucose level was closely monitored, and insulin dose was promptly adjusted in pregnant women with GDM with abnormal blood glucose after comprehensive dietary intervention. Follow-up was carried out every 2 wk in the form of hospital visits. Glycemic control criteria [9] were FPG  $\leq$  5.6 mmol/L, 2 h postprandial blood glucose (P2h-PG)  $\leq 6.7$  mmol/L, and glycated hemoglobin (HbA1<sub>c</sub>) < 6%. All three criteria must be met to conform to glycemic control standards. Otherwise, the patient was considered not to conform to glycemic control standards.

#### **Observation markers**

Baseline data and blood glucose in pregnant women and immune function, infectionrelated markers, and infection rate in neonates were observed and compared. (1) Blood glucose: This was measured 24 h before delivery. Venous whole blood was collected from pregnant women, and a low-speed centrifuge was used to extract serum samples at 3000 r/min for 10 min. A Mindray glucose assay kit (glucose oxidase assay) was used to measure FPG and P2h-PG in serum samples. Heparin anticoagulant tubes were used to collect venous whole blood from pregnant women. Ion exchange chromatography and gradient elution were used to measure  $HbA1_c$  after hemolysis of blood samples using hemolysin; (2) immune function: 5 mL of umbilical vein blood was collected from neonates after delivery. Blood samples were mixed with allophycocyanin (APC)/cyanine dye 7 (Cy7) fluorescently labeled mouse anti-human CD3 antibody, phycoerythrin/Cy7 mouse anti-human CD4 antibody, and APC/Cy7 mouse anti-human CD8, and incubated at room temperature for 15 min. BD FACS™ lysis solution (BD Inc., USA) was added and incubated in the dark for 15 min. A BD FACSCanto II flow cytometer (BD Inc., USA) was used to measure the proportions of CD3+ T cells, CD4+T cells, and CD8+ T cells with different fluorescent labels in peripheral blood. The CD4/CD8 ratio was then calculated; (3) the levels of immunoglobulin G (IgG), IgA, and IgM in the peripheral blood of newborns in both groups were measured by immunoturbidimetry; (4) a Roche Cobas 8000 fully automatic biochemical analyzer was used to measure the white blood cell (WBC) count in the umbilical vein blood in neonates; (5) enzyme-linked immunosorbent assay (ELISA) was used to measure procalcitonin (PCT) and C-reactive protein (CRP) levels. ELISA kits were purchased from Beyotime Biotechnology Co., Ltd; and (6) neonatal infections were observed and recorded, including upper respiratory tract infection, lower respiratory tract infection, skin infection, intestinal infection, and sepsis.

#### Statistical analysis

Statistical data were processed using SPSS 20.0 software, and quantitative data were tested for normal distribution. Quantitative data with normal distribution were expressed as mean ± SD. An independent sample *t*-test was used for inter-group comparisons. Quantitative data with abnormal distribution are represented by median (quartile), and a nonparametric test was used for inter-group comparisons. Qualitative data were expressed as % and the  $\chi^2$  test was used. A P value < 0.05 was considered statistically significant.

#### RESULTS

#### Comparison of baseline data

In the GDM group, the age of pregnant women ranged from 21-39 years and the mean age was  $29.98 \pm 4.65$  years; body mass index (BMI) was 19.7-38.1 kg/m<sup>2</sup> and mean BMI was (29.75  $\pm$  2.68) kg/m<sup>2</sup>. There were 143 primipara and 93 multipara women. In the control group, the age ranged from 22-40 years and the mean age was  $30.26 \pm 4.74$  years; BMI was 20.3–37.9 kg/m<sup>2</sup> and mean BMI was  $30.01 \pm 3.12$  kg/m<sup>2</sup>; there were 159 primipara and 181 multipara women. There were no differences in age, BMI, parity, and gravidity between the two groups and the groups were comparable (P >0.05). as shown in Table 1.

#### Comparison of early neonatal complications

The incidence of premature births, fetal macrosomia, hypoglycemia, hypocalcemia, and hyperbilirubinemia in neonates delivered by pregnant women with GDM was significantly higher than that in neonates delivered by women in the



Table 1 Comparison of baseline data in pregnant women with gestational diabetes mellitus between conforming glycemic control and non-conforming glycemic control groups

Groups	Cases	Age (yr)	BMI (kg/m²)	Type of pregnant woman, n (%)	
				Primipara	Multipara
CGC group	178	$30.05 \pm 4.46$	29.55 ± 2.82	109 (61.24)	69 (38.76)
NCGC group	58	29.47 ± 3.75	$30.08 \pm 2.57$	34 (56.90)	25 (43.10)
$t/\chi^2$ value		0.893	1.270	0.344	
<i>P</i> value		0.373	0.206	0.558	

BMI: Body mass index; CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control.





Figure 1 Screening, randomization and analysis of populations. OGTT: Oral glucose tolerance test; FPG: Fasting plasma glucose; P1h-PG: 1 h postprandial blood glucose; GDM: Gestational diabetes mellitus.

control group. These differences were statistically significant (P < 0.05) as shown in Table 2.

#### Comparison of blood glucose markers

Pregnant women with GDM were divided into the CGC group and the NCGC group based on whether their glycemic control conformed to standards. Blood glucose markers in the NCGC group, such as FPG, P2h-PG, and HbA1<sub>c</sub> levels, were significantly higher (P < 0.05) compared with pregnant women with GDM in the CGC group. These results are shown in Table 3.

#### Comparison of peripheral blood T cell subsets

Results of the flow cytometry analysis of peripheral blood T cell subsets were compared between the two groups of neonates. The ratio of peripheral blood CD3<sup>+</sup>T cells, CD4<sup>+</sup>T cells, and CD8<sup>+</sup>T cells, and the CD4/CD8 ratio in neonates in the NCGC group were all significantly lower than those in the CGC group (P < 0.05). These results showed that immune function was significantly decreased in neonates from the NCGC group. These results are shown in Table 4.

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Table 2 Comparison of early neonatal complications between the gestational diabetes mellitus and control groups, n (%)						
Complications			χ² value	<i>P</i> value		
Premature birth	45 (19.07)	12 (5.00)	22.341	< 0.001		
Fetal macrosomia	57 (24.15)	23 (9.58)	18.064	< 0.001		
Hypoglycemia	42 (17.80)	7 (2.92)	28.53	< 0.001		
Hypocalcemia	22 (9.32)	5 (2.08)	11.653	< 0.001		
Hyperbilirubinemia	29 (12.29)	13 (5.42)	6.984	0.008		
Polycythemia	38 (16.10)	22 (9.17)	5.195	0.023		
Hyaline membrane disease	13 (5.51)	2 (0.83)	8.522	0.004		
Fetal distress	34 (14.41)	6 (2.50)	21.917	< 0.001		
Congenital malformation	11 (4.66)	1 (0.42)	8.723	0.003		
Neonatal asphyxia	21 (8.90)	6 (2.50)	9.104	0.003		
Neonatal infection	35 (14.83)	11 (4.58)	14.312	< 0.001		

GDM: Gestational diabetes mellitus.

Table 3 Comparison of blood glucose markers in pregnant women with gestational diabetes mellitus between conforming glycemic control and non-conforming glycemic control groups

Groups	Cases	FPG (mmol/L)	P2h-PG (mmol/L)	HbA1 <sub>c</sub> (%)
CGC group	178	$4.68\pm0.60$	$5.51 \pm 0.85$	$5.11 \pm 0.45$
NCGC group	58	$5.96 \pm 0.68$	$7.14 \pm 1.04$	$6.38 \pm 0.74$
$t/\chi^2$ value		13.645	11.979	15.691
<i>P</i> value		< 0.001	< 0.001	< 0.001

CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control; FPG: Fasting plasma glucose; P2h-PG: 2 h postprandial blood glucose; HbA1<sub>C</sub>: Hemoglobin.

Table 4 Comparison of CD3 <sup>+</sup> , CD4 <sup>+</sup> , CD8 <sup>+</sup> and CD4 <sup>+</sup> /CD8 <sup>+</sup> ratio between the two groups of neonates (mean ± SD, <i>n</i> )						
Groups	Cases	CD3 <sup>+</sup> (%)	CD4 <sup>+</sup> (%)	CD8⁺ (%)	CD4/CD8	
CGC group	178	$52.01 \pm 10.78$	39.21 ± 7.80	$25.69 \pm 5.47$	$1.61 \pm 0.54$	
NCGC group	58	$45.25 \pm 7.33$	$22.46 \pm 5.48$	$19.42 \pm 2.95$	$1.17 \pm 0.33$	
<i>t</i> value		4.449	15.170	8.335	5.854	
<i>P</i> value		< 0.001	< 0.001	< 0.001	< 0.001	

CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control.

#### Comparison of immunoglobulin levels

The levels of IgG, IgM, and IgA, were compared between neonates in the two groups. The results showed that the level of IgG in the CGC group was significantly higher than that in the NCGC group, with a statistically significant difference (P < 0.05). There were no significant differences in the levels of IgM and IgA in peripheral blood between the two groups of newborns (P > 0.05), as shown in Table 5.

#### Comparison of infection-related inflammatory markers

Infection-related inflammatory markers, including WBC, serum PCT and CRP levels, were compared between neonates in the two groups. WBC, PCT, and CRP levels in neonates in the NCGC group were significantly higher than those in the CGC group, and these differences were statistically significant (P < 0.05), as shown in Table 6.

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Table 5 Comparison of immunoglobulin G, immunoglobulin M and immunoglobulin A levels between the two groups of neonates (mean ± SD, <i>n</i> )						
Groups	Cases	lgG (g/L)	IgM (g/L)	lgA (g/L)		
CGC group	178	9.78 ± 1.38	0.181 ± 0.043	0.31 ± 0.08		
NCGC group	58	$7.21 \pm 1.32$	$0.173 \pm 0.040$	$0.29\pm0.07$		
<i>t</i> value		12.447	1.251	1.688		
<i>P</i> value		< 0.001	0.212	0.093		

IG: Immunoglobulin; CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control.

Table 6 Comparison of white blood cell, procalcitonin and C-reactive protein levels between the two groups of neonates (mean ± SD, <i>n</i> )							
Groups	Cases	WBC (×10 <sup>9</sup> /L)	PCT (µg/L)	CRP (mg/L)			
CGC group	178	$15.56 \pm 5.47$	$0.43 \pm 0.12$	7.22 ± 2.07			
NCGC group	58	$25.80 \pm 8.61$	$0.81 \pm 0.24$	12.38 ± 3.22			
<i>t</i> value		10.618	15.920	14.212			
<i>P</i> value		< 0.001	< 0.001	< 0.001			

WBC: white blood cell; PCT: procalcitonin; CRP: C-reactive protein; CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control.

#### Comparison of neonatal infections

The incidence of upper respiratory tract infection, lower respiratory tract infection, and skin infection in neonates in the NCGC group was 31.03%, which was significantly higher than neonates in the CGC group (9.35%). These differences were statistically significant (P < 0.05) and are shown in Table 7.

#### DISCUSSION

Glucose metabolism disorders and long-term blood glucose abnormalities in pregnant women with GDM are associated with decreased pancreatic islet function and insulin resistance[10], and GDM has a major impact on adverse maternal and child outcomes in the perinatal phase compared to the normal diabetic population. Currently, there are 210 million neonates affected by GDM globally, as shown in the 2017 International Diabetes Federation report[11]. Furthermore, GDM has become a major global public health concern. The results of this study show that the incidence of premature births, fetal macrosomia, hypoglycemia, hypocalcemia, hyperbilirubinemia, and infection in neonates delivered by pregnant women with GDM was significantly higher than in neonates delivered by healthy pregnant women. These results demonstrate that the health of neonates is severely affected by blood glucose abnormalities in the mother during pregnancy. The main harm caused by GDM is an increase in maternal and child adverse outcomes and mortality rate during the perinatal phase, resulting in fetal distress, developmental abnormalities, and increases the risk of hypoglycemia, deformities, and infection in neonates[12]. Therefore, it can be seen from these results that stringent glycemic control in pregnant women with GDM is an essential measure to prevent neonatal complications in the perinatal stage. Capobianco et al[13] also reported the GDM is an important complication that affects maternal and pregnancy outcomes, and good glycemic control can decrease the risk of pregnancy complications and the cesarean section rate, increase the rate of natural vaginal delivery, and has positive effects in decreasing premature births, fetal macrosomia, hypoglycemia, asphyxiation, and infection.

A recent study[14] found that interleukin (IL)-6 and tumor necrosis factor (TNF)- $\alpha$  levels were significantly increased in the umbilical vein blood from neonates delivered by women with GDM. The neonates also showed varying degrees of immune dysfunction, suggesting that the risk of infection is higher in neonates of GDM patients. Our study results showed that compared with healthy pregnant women, the incidence of neonatal infection was significantly higher in pregnant women with GDM. Blood glucose markers and neonatal infection rates were significantly increased in pregnant women with GDM in the NCGC group compared to those in the CGC group. This suggested that glycemic control has significant effects in decreasing neonatal infection caused by blood glucose abnormalities in GDM patients. Zarrin *et al* [15] found that the condition of GDM patients worsened as gestational age increased, and glucose metabolism abnormalities during pregnancy may directly affect maternal and fetal immune function. Maternal immune defects and fetal T lymphocyte developmental abnormalities will affect neonatal immune function. Neonatal immune dysfunction is an independent risk factor for infection[16]. T lymphocytes are the most important cell population in the immune system, of which the CD3+ subset represents mature T lymphocytes and immune function[17]. CD4+ T cells mainly regulate

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Table 7 Comparison of neonatal infections between the two groups, n (%)								
Groups	Cases	Upper respiratory tract infection	Lower respiratory tract infection	Skin infection	Intestinal infection	Sepsis	Total	
CGC group	178	10 (5.62)	4 (2.25)	1 (0.56)	2 (1.12)	0 (0.00)	17 (9.55)	
NCGC group	58	7 (12.07)	3 (5.17)	2 (3.45)	5 (8.62)	1 (1.72)	18 (31.03)	
$\chi^2$ value							15.985	
P value							< 0.001	

CGC: Conforming glycemic control; NCGC: Non-conforming glycemic control.

humoral immunity while CD8+ T cells are inhibitory/cytotoxic T lymphocytes that mainly regulate cellular immunity and play a critical role in regulating CD3+ and CD4+ functions [18,19]. The CD4+/CD8+ ratio can normally be used to reflect the equilibrium between humoral/cellular immunity. This cell ratio is an important marker for evaluating immune function and a low CD4+/CD8+ ratio usually means that the body is in an immunosuppressed state[20]. Kugler et al[21] confirmed that neonatal immune function defects were related to the inheritance of abnormal T lymphocyte development and maternal immune function defects, and were independent risk factors for neonatal infection. We compared the differences in peripheral blood T cell subsets in this study between neonates in the two groups. The results showed that peripheral blood CD3<sup>+</sup>T cells, CD4<sup>+</sup>T cells, and CD8<sup>+</sup>T cells, and the CD4/CD8 ratio of neonates in the NCGC group were all significantly lower than those in the CGC group. These results demonstrated that blood glucose abnormalities in pregnant women with GDM may affect peripheral blood T cell subsets in neonates, resulting in decreased immune function and immune regulation disorders, thus reducing infection resistance.

Immunoglobulins are an important class of immune effector molecules, including IgG, IgA, IgM, and so on. IgG in the peripheral blood of newborns is mainly from the mother, accounting for about 75% of the total serum immunoglobulin content, and plays an important role in preventing infection[22]. This study found that IgG in neonates from women with GDM in the NCGC group significantly decreased compared to that in the CGC group, suggesting that poor blood glycemic control in GDM pregnant women can lead to a decline in neonatal immune function. It is speculated that for patients with GDM, abnormal glucose metabolism itself is an inflammatory reaction, which hinders the production of IgG, thereby reducing the amount of IgG entering the fetus *via* the placenta<sup>[23]</sup>. However, IgA and IgM cannot pass through the placental barrier, resulting in extremely low levels in the peripheral blood of newborns, leading to insignificant changes in levels.

WBC are the most commonly used marker for early diagnosis and treatment of neonatal infections, and PCT and CRP are important serum markers for diagnosing neonatal infection[24]. This study compared these markers in pregnant women with GDM and neonates from the two groups. The results revealed that WBC, serum PCT and CRP levels in neonates in the NCGC group were significantly greater than those in the CGC group, suggesting that GDM patients who did not meet glycemic control standards have neonates with elevated inflammatory markers and an increased risk of infection. This may be because blood glucose abnormalities during pregnancy can promote the transcription of placental CRP, IL-6, and PCT, which are important mechanisms that directly affect the fetal immune system[25]. Li *et al*[26] found that a hyperglycemic environment activated placental HIF-1a and TLR4/MyD88/NF-kB pathways in pregnant women with GDM, induced IL-6 and IL-8 secretions, promoted placental inflammation and autophagy to disrupt placental homeostasis and cell renewal, and increased the risk of infection in neonates delivered by pregnant women with GDM. From these findings combined with the results from the present study, we believe that poor glycemic control in pregnant women with GDM can result in long-term blood glucose abnormalities, which may stimulate inflammatory responses in the fetal placenta, thereby affecting the fetal immune system and ultimately increase the risk of neonatal infections.

The innovation of this study is that the relationship between the blood glucose control level in pregnant women with GDM and neonatal immune function was analyzed, which opens up a new direction for predicting neonatal infectious pathology. However, a larger multicenter clinical study is needed in the future to validate the results of this study as this was a single center study with a limited sample size due to the strict screening conditions. In addition, immunoglobulin, as an antibody related to the immune response in vivo, can be used as an early diagnostic indicator of infection, and IgG antibodies can also enhance the anti-infection ability of newborns and prevent related infectious diseases. However, this study lacks the assessment of neonatal peripheral blood immunoglobulin level to verify neonatal immune function. Finally, the neonates included in this study were only followed up for a short time. The influence of blood glucose control level in GDM patients on the long-term immune function of neonates still requires further research.

#### CONCLUSION

In summary, we speculate that glucose metabolism disorders and long-term blood glucose abnormalities in GDM patients with poor glycemic control may be considered a type of inflammatory response which affects the T lymphocyte subsets in neonates, resulting in immune dysfunction, and ultimately decreasing immune function and increasing the risk of infection.



## **ARTICLE HIGHLIGHTS**

#### Research background

Gestational diabetes mellitus (GDM) is related to obesity in pregnant women, older age in pregnant women, excessive nutrition during pregnancy, lack of exercise, genetic history of familial type 2 diabetes, excessive sugar consumption and other factors. GDM often causes obstetric complications, which seriously threaten the life and health of pregnant women and newborns. Blood sugar control measures have a considerable impact on pregnancy outcome and newborn status in patients with GDM.

#### Research motivation

The long-term abnormal glucose metabolism in GDM pregnant women affects the immune function of newborns, and it is unclear whether poor glucose control in GDM pregnant women increases the risk of neonatal infectious diseases.

#### Research objectives

The purpose of this study was to determine the correlation between GDM pregnant women and neonatal complications, and to analyze the impact of blood glucose control on the risk of neonatal infectious diseases.

#### Research methods

The clinical data of 236 pregnant women with GDM and 240 healthy pregnant women and newborns were retrospectively analyzed to compare early neonatal complications in the two groups of pregnant women. The 236 pregnant women with GDM were divided into two groups based on whether their blood sugar control reached the standard. The baseline data, neonatal immune function, infection related indicators, and neonatal infection rate in the two groups of pregnant women with GDM were compared.

#### **Research results**

The incidence of neonatal complications in GDM pregnant women was significantly higher than that in normal pregnant women. Compared with GDM pregnant women who achieved glycemic control, the proportion of CD3+, CD4+, and CD8+T cells in peripheral blood and the ratio of CD4/CD8 cells in newborns from mothers who did not achieve glycemic control significantly decreased, while the white blood cell count, serum procalcitonin, and C-reactive protein levels significantly increased, and the neonatal infection rate significantly increased.

#### Research conclusions

The risk of neonatal complications is increased in pregnant women with GDM, and poor glycemic control leads to impairment of the fetal immune system and ultimately increases the risk of neonatal infections.

#### Research perspectives

The effect of blood glucose control is related to pregnancy outcome and neonatal prognosis.

## FOOTNOTES

Author contributions: Wang BB conceived the study; Wang BB and Xue M performed the data collection and extraction and analyzed the data; Wang BB and Xue M interpreted and reviewed the data and drafts; Xue M reviewed the final draft.

Institutional review board statement: The study was reviewed and approved by the Taizhou People's Hospital of Jiangsu Province Institutional Review Board.

Informed consent statement: The analysis used anonymous clinical data that were obtained after each patient agreed to treatment by written consent. According to institutional policy, this study was exempt from the informed consent process.

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

Data sharing statement: statement: The dataset is available from the corresponding author at xuemei009260@163.com.

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**Retrospective Study** 

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ORIGINAL ARTICLE

## Risk factors of concurrent urinary sepsis in patients with diabetes mellitus comorbid with upper urinary tract calculi

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## Abstract

## BACKGROUND

Urinary sepsis is frequently seen in patients with diabetes mellitus (DM) complicated with upper urinary tract calculi (UUTCs). Currently, the known risk factors of urinary sepsis are not uniform.

#### AIM

To analyze the risk factors of concurrent urinary sepsis in patients with DM complicated with UUTCs by logistic regression.

## **METHODS**

We retrospectively analyzed 384 patients with DM complicated with UUTCs treated in People's Hospital of Jincheng between February 2018 and May 2022. The patients were screened according to the inclusion and exclusion criteria, and 204 patients were enrolled. The patients were assigned to an occurrence group (n= 78) and a nonoccurrence group (n = 126). Logistic regression was adopted to analyze the risk factors for urinary sepsis, and a risk prediction model was established.

## RESULTS

Gender, age, history of lumbago and abdominal pain, operation time, urine leukocytes (U-LEU) and urine glucose (U-GLU) were independent risk factors for patients with concurrent urinary sepsis (P < 0.05). Risk score =  $0.794 \times$  gender + 0.941 × age + 0.901 × history of lumbago and abdominal pain - 1.071 × operation time + 1.972 × U-LEU + 1.541 × U-GLU. The occurrence group had notably higher risk scores than the nonoccurrence group (P < 0.0001). The area under the curve of risk score for forecasting concurrent urinary sepsis in patients was 0.801, with specificity of 73.07%, sensitivity of 79.36% and Youden index of 52.44%.



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#### CONCLUSION

Sex, age, history of lumbar and abdominal pain, operation time, ULEU and UGLU are independent risk factors for urogenic sepsis in diabetic patients with UUTC.

Key Words: Diabetes mellitus; Upper urinary tract calculi; Urinary sepsis; Risk factors; Risk prediction model; Logistic regression; Concurrent urinary sepsis

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**Core Tip:** This study was to determine risk factors of concurrent urinary sepsis in patients with diabetes mellitus comorbid with upper urinary tract calculi and construct a risk prediction model. Gender, age, history of lumbago and abdominal pain, operation time, urinary leukocytes and urinary glucose were independent risk factors for concurrent urinary sepsis. It is helpful to identify high-risk patients at an early stage and implement active and effective intervention measures to reduce complications and improve the prognosis of patients.

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## INTRODUCTION

With the improvement of living standards, the incidence of urinary calculi worldwide is increasing gradually. Its incidence in adults in China is 1%-5%, and upper urinary tract calculi (UUTCs) are a more frequently seen problem[1]. Calculi are caused by several factors, such as socioeconomic status, environmental factors and eating habits<sup>[2]</sup>. The incidence of calculi in northern and southern areas of China differs, with a high incidence in southern areas (5%-10%)[3]. Other factors such as gender and genetic susceptibility can also affect the occurrence of calculi, and the incidence rate among males is three times that among females[4]. In China, the annual incidence of urinary calculi is 150-200 cases per 100000 people, and approximately 25% of patients need hospitalization and surgical treatment. After treatment, the recurrence rate is high at approximately 50% within 10 years[5].

The urethra connects the urinary system with the outside world, providing a way for bacteria and other pathogens to invade the urinary system<sup>[6]</sup>. Usually, the flushing action during urination and urethral mucosa form a natural protective barrier to prevent bacteria from remaining, growing and reproducing[7]. However, various factors hinder the defensive function of the urinary system. For example, obstruction can result in stagnant water above the obstruction, making it easier for pathogenic bacteria to invade and colonize the urinary system, giving rise to infection[8]. Without timely intervention, the infection may develop into urinary sepsis or even septic shock, endangering the life of patients.

Sepsis is a serious disease and an acute physical reaction caused by infection, with associated physiological, pathological and biochemical abnormalities[9]. Sepsis is defined as organ dysfunction due to the host's uncontrolled immune response to infection. Unfortunately, sepsis is a global health threat with high mortality[10]. Without timely treatment, sepsis may develop into septic shock and multiple organ dysfunctions and even cause death[11].

The incidence of diabetes mellitus (DM) is increasing. According to the latest statistics of the International Diabetes Federation, the global prevalence of DM has reached 9.3%. DM can give rise to systemic damage, leading to immune dysfunction and proneness to serious infection [12]. However, currently, the risk factors for concurrent urinary sepsis in patients with UUTCs and DM are still under investigation.

Accordingly, this study aimed to determine the risk factors for concurrent urinary sepsis in patients with DM comorbid with UUTCs to provide a reference for clinical therapy and prevention.

#### MATERIALS AND METHODS

#### Patients

We retrospectively analyzed 384 patients with DM complicated with UUTCs treated in People's Hospital of Jincheng between February 2018 and May 2022. Inclusion criteria were: (1) Imaging results, such as urinary ultrasound, intravenous urography or abdominal computed tomography, suggested the presence of UUTCs; (2) a clear history of DM that met the guidelines for the diagnosis and treatment of senile DM in China[13]; and (3) a complete medical history and laboratory and imaging data. Exclusion criteria were: (1) Age < 18 years; (2) pregnant women; (3) bilateral UUTCs; (4) hematological disease, immune system disease, or malignant tumors; (5) treated with immunomodulatory drugs; and (6) other primary infection, such as lung or abdominal infection. The 384 patients were screened according to the inclusion and exclusion criteria, and 204 patients were enrolled. According to Guidelines for Emergency Treatment of Sepsis/Septic

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Shock in China (2018)[14] and the diagnostic criteria for urinary sepsis[14], patients were assigned to an occurrence group (n = 78) and nonoccurrence group (n = 126).

#### Collection of clinical data

The clinical data were collected through the medical record system of our hospital, including: (1) General information: Gender, age and body mass index; (2) medical history: Lumbago and abdominal pain, hematuria, symptoms of urinary tract irritation, hypertension, and DM; (3) urine examination data: Urinary leukocytes (U-LEU), urinary nitrite (U-NIT), urinary glucose (U-GLU), and urinary occult blood; and (4) imaging examination data: Lateral classification, location, maximum diameter and hydronephrosis of calculi.

#### Statistical analysis

We used R language 4.1.1 software (R Foundation for Statistical Computing, Vienna, Austria) for data cleaning and analysis, and constructed a model. Logistic regression was adopted for screening the risk factors, and receiver operating characteristic (ROC) curve was adopted for value verification. This study used Graph Pad Prism 8.0 for data visualization. P < 0.05 indicated a significant difference.

## RESULTS

#### Analysis of clinical data

The occurrence and nonoccurrence groups did not differ significantly for hematuria and hypertension (P > 0.05) (Table 1), but they did differ significantly for gender, age, history of lumbago and abdominal pain, symptoms of urinary tract irritation and operation time (P < 0.05) (Table 1).

#### Comparison of urinary examination indexes

The occurrence and nonoccurrence groups did not differ significantly for urinary occult blood (P > 0.05) (Table 2), but they did differ significantly for U-LEU, U-NIT, and U-GLU (P < 0.01) (Table 2).

#### Imaging index detection

The occurrence and nonoccurrence groups did not differ significantly for lateral classification of calculi, obstruction position, maximum calculi diameter and severity of hydronephrosis (P > 0.05) (Table 3).

#### Logistic regression analysis

According to the above results, meaningful indicators were assigned (Table 4). The backward logistic regression method was used. Gender, age, history of lumbago and abdominal pain, operation time, U-LEU and U-GLU were independent risk factors for concurrent urinary sepsis (P < 0.05) (Table 5).

#### Construction of risk model

Based on the  $\beta$  coefficient of logistic regression, a risk score for predicting concurrent urinary sepsis was constructed. Risk score = 0.794 × gender + 0.941 × age + 0.901 × history of lumbago and abdominal pain - 1.071 × operation time + 1.972 × U-LEU + 1.541 × U-GLU. According to the comparison results, the occurrence group had notably higher risk scores than the nonoccurrence group (Figure 1A) (P < 0.0001). According to ROC curve analysis, the area under the curve (AUC) of risk score for forecasting concurrent urinary sepsis was 0.801, with specificity of 73.07%, sensitivity of 79.36% and Youden index of 52.44% (Figure 1B).

#### DISCUSSION

Urinary sepsis is a dangerous disease. Without timely diagnosis and treatment, its prognosis is unfavorable[15]. There are approximately 2.8 to 9.8 million new cases of urinary sepsis every year, with 1.6 million deaths[16]. Thus, it is important to quickly identify urinary sepsis and provide effective timely treatment. In the guidelines of the European Association of Urology (2017 edition), the definition of urinary sepsis has been updated, which emphasizes that the disease is more serious than uncomplicated infection and may cause organ dysfunction and become life-threatening[17]. Patients with UUTCs and DM are more likely to have urinary sepsis[18]. This is because calculi may trigger infection, and DM makes patients susceptible to various diseases[19]. Therefore, early identification of high-risk factors in these patients and effective intervention have become the focus of many hospitals, which has also been recognized by the World Health Organization.

Our study retrospectively analyzed the risk factors for concurrent urinary sepsis in patients with DM complicated with UUTCs. Gender, age, history of lumbago and abdominal pain, operation time, U-LEU and U-GLU were independent risk factors for concurrent urinary sepsis. The risk of urinary sepsis was about 2.212 times higher in women than in men. Prior research has revealed that women with ureteral calculi or who undergo endoscopic lithotripsy face an independently increased risk of urinary sepsis[20]. However, one other study has revealed no independent correlation between gender and incidence of urinary sepsis[21]. Kumar *et al*[22] have revealed that the reasons why older women are prone to urinary

Table 1 Analysis of clinical data					
Factors		Occurrence group ( <i>n</i> = 78)	Nonoccurrence group ( <i>n</i> = 126)	χ² value	P value
Gender				9.219	0.002
	Male	30	76		
	Female	48	50		
Age				5.732	0.016
	≥60 yr	45	51		
	< 60 yr	33	75		
BMI				0.292	0.588
	$\geq 25 \text{ kg/m}^2$	28	50		
	$< 25 \text{ kg/m}^2$	50	76		
History of lumbago and abdominal pain				8.459	0.003
	Yes	38	36		
	No	40	90		
Hematuria				0.569	0.450
	Yes	36	65		
	No	42	61		
Symptoms of urinary tract irritation				4.036	0.044
	Yes	60	80		
	No	18	46		
Hypertension				0.533	0.465
	Yes	30	55		
	No	48	71		
Operation time		64.32 ± 15.35	59.94 ± 7.04	4.662	0.001

BMI: Body mass index.



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Figure 1 Value of risk score in predicting urinary sepsis in patients with diabetes mellitus and upper urinary tract calculi. A: Risk score in predicting urinary sepsis; B: Area under the curve of risk score for prediction of urinary sepsis. <sup>d</sup>P < 0.0001.

tract infection and progression to urinary sepsis may include poor perineal hygiene, postmenopausal estrogen deficiency, atrophic vaginitis, uterine and bladder prolapse and the use of vaginal supports.

With the increase of age, the functions of various organs or systems tend to decline, including liver and kidney dysfunction, cardiovascular system defects, and immune system defects[23]. Weakened compensatory ability of organs and systems in patients gives rise to a decline in overall physical function, and patients with DM are more susceptible to

Table 2 Detection of uri	nary i	ndexes			
Factors		Occurrence group ( <i>n</i> = 78)	Nonoccurrence group ( <i>n</i> = 126)	X <sup>2</sup>	P value
Urine occult blood				1.210	0.750
	-	18	35		
	1+	22	28		
	2+	24	38		
	3+	14	25		
U-LEU				15.330	0.002
	-	4	30		
	1+	26	38		
	2+	20	37		
	3+	28	25		
U-NIT				12.499	0.001
	-	33	85		
	+	45	41		
U-GLU				11.596	0.008
	-	7	33		
	1+	17	31		
	2+	40	50		
	3+	14	12		

U-LEU: Urinary leukocytes; U-NIT: Urinary nitrite; U-GLU: Urinary glucose.

Table 3 Comparison of imaging indexes							
Factors		Occurrence group ( <i>n</i> = 78)	Nonoccurrence group ( <i>n</i> = 126)	<b>X</b> <sup>2</sup>	P value		
Lateral classification of calculi				1.494	0.221		
	Left	48	88				
	Right	30	38				
Obstruction position				2.938	0.086		
	Ureter	64	90				
	Kidney	14	36				
Maximum diameter of calculi				0.048	0.825		
	≥ 20 mm	39	65				
	< 20 mm	39	61				
Degree of hydronephrosis				0.400	0.526		
	Yes	11	22				
	No	67	104				

infection in such cases [24]. Urinary tract obstruction can easily give rise to secondary infection, systemic inflammatory reaction and even sepsis[25]. In this study, the risk of urinary sepsis in patients aged > 60 years was 2.563 times that in patients < 60 years old, which is in agreement with previous studies.

Similar to prior research, U-LEU was an independent risk factor for UUTCs complicated with urinary sepsis in our study[26]. Some researchers believe that positive urine bacterial culture can more accurately predict the occurrence of urinary sepsis[27]. However, our study did not include the results of urine bacterial culture as a predictor because of the lag time of urine bacterial culture. Usually, it takes 2-3 d or even longer to achieve the results of urine bacterial culture, which leads to a lag in forecasting ability. Therefore, we mainly adopted U-LEU as an indicator, which suggests purulent

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#### Table 4 meaningful indicators were assigned

Factors	Assignment
Gender	Male = 0, female = 1
Age	$\ge 60 \text{ yr} = 1, < 60 \text{ yr} = 0$
History of lumbago and abdominal pain	Yes = 1, no = 0
Symptoms of urinary tract irritation	Yes = 1, no = 0
Operation time	$\ge 60 \text{ yr} = 1, < 60 \text{ yr} = 0$
U-LEU	- = 0, 1 + -3 + = 1
U-NIT	- = 0, + = 1
U-GLU	- = 0, 1 + -3 + = 1
Occurrence	Occurrence group = 1, nonoccurrence group = 0

U-LEU: Urinary leukocytes; U-NIT: Urinary nitrite; U-GLU: Urinary glucose.

#### Table 5 Logistic multivariate regression

Fasters	p	Standard array		Dualua	OR	95%CI	
ractors	р	Standard error	X	P value		Lower limit	Upper limit
Gender	0.794	0.335	5.603	0.018	2.212	1.146	4.268
Age	0.941	0.346	7.408	0.006	2.563	1.301	5.047
History of lumbago and abdominal pain	0.901	0.348	6.700	0.010	2.462	1.245	4.871
Symptoms of urinary tract irritation	0.628	0.379	2.753	0.097	1.875	0.892	3.939
Operation time	-1.071	0.345	9.619	0.002	0.342	0.174	0.674
U-LEU	1.972	0.602	10.725	0.001	7.182	2.207	23.373
U-NIT	0.491	0.339	2.101	0.147	1.634	0.841	3.172
U-GLU	1.541	0.509	9.171	0.002	4.668	1.722	12.652

U-LEU: Urinary leukocytes; U-NIT: Urinary nitrite; U-GLU: Urinary glucose.

inflammation in the urinary tract, so it can be used to predict the risk of urinary sepsis.

In patients with DM, the reasons for positive U-GLU may include an increase in blood glucose and decrease in renal glucose threshold. Hyperglycemia causes failure of glucose absorption in the renal tubules, so that glucose is excreted in the urine[18]. Additionally, DM can decrease the ability of renal tubules to absorb glucose, and the glucose in urine cannot be completely reabsorbed[28]. Both of these conditions may lead to positive U-GLU. Positive U-GLU may indicate poor control of DM or diabetic nephropathy. Diabetic nephropathy can easily damage the genitourinary system, causing difficulty in controlling urinary tract infection or recurrence. High concentration of U-GLU provides heat for the growth and metabolism of pathogenic bacteria, which in turn leads to disorder of the body's defense mechanisms.

In this study, a history of lumbago and abdominal pain had a strong correlation with concurrent urinary sepsis in patients with DM and UUTCs. Lumbago and abdominal pain are frequent symptoms of UUTCs, and one of the manifestations of many patients with urinary sepsis[29]. UUTCs can give rise to urinary retention, bacterial reproduction and infection, increasing the risk of urinary sepsis. DM complicated with UUTCs is a risk factor for urinary sepsis[30]. Patients with DM are often accompanied by various pathophysiological changes such as decreased immune function and metabolic disorder, which can lead to urinary tract infection. Therefore, patients with DM and UUTCs are at higher risk of urinary sepsis.

Long operation time increases the risk of infection. Long-term exposure of wound tissue increases the probability of infection by surrounding flora, and long operation time also increases bleeding and absorption of perfusion fluid, which increase the risk of infection. Urethral obstruction during surgery may also lead to urinary retention, creating favorable conditions for bacterial reproduction[24,31].

We constructed a risk prediction model based on the regression coefficient. A prediction model is a mathematical model that estimates the probability of a specific event or disease according to the combination of multiple risk factors. Through the analysis and assessment of risk factors, a data-based model can be established to help doctors and researchers better understand the risk factors of a disease and develop better prevention and treatment plans. In this

study, the occurrence group had notably higher risk scores than the nonoccurrence group, and the AUC of risk score in forecasting urinary sepsis was > 0.8, indicating a high value of the risk model in predicting urinary sepsis.

We analyzed the risk factors for concurrent urinary sepsis in patients with DM complicated with UUTCs by logistic regression model, and successfully constructed a prediction model. However, our study had some limitations. First, there was no external verification of our results. This was because in such a single-center study, it was impossible to establish an effective verification set because of the small number of samples collected. Second, as a single-center research model, its universality needs further verification. Therefore, we hope to carry out prospective research and collect more samples in the future to improve the conclusions.

## CONCLUSION

Gender, age, history of lumbago and abdominal pain, operation time, U-LEU and U-GLU were independent risk factors for concurrent urinary sepsis in patients with DM and UUTCs. It is helpful to identify high-risk patients at an early stage and implement effective intervention measures to reduce complications and improve prognosis.

## ARTICLE HIGHLIGHTS

#### Research background

In patients with diabetes mellitus (DM), long-term hyperglycemia can trigger increases in sugar, protein and other substances in urine, promoting formation of calculi. Urinary calculi can lead to urinary tract infection, renal insufficiency and other complications, and even become life-threatening in severe cases, causing a serious impact on the health of patients with DM. However, at present, the risk factors for urinary sepsis are not uniform. The purpose of this study was to analyze the risk factors for urinary sepsis in patients with DM complicated with upper urinary tract calculi (UUTCs) to provide potential indicators for clinical observation.

#### Research motivation

It is helpful to identify high-risk patients at an early stage and implement and effective intervention measures by constructing a prediction model, thus reducing complications and improving prognosis.

#### Research objectives

We successfully predicted high-risk patients by establishing a risk model, which was beneficial to clinical and targeted treatment and prevention.

#### Research methods

We constructed a risk model of urinary sepsis by logistic regression model, which provided an observation model for the prediction of high-risk patients.

#### Research results

Although we successfully established a risk model, due to the small number of patients, it was impossible to carry out external verification, so more data are needed to verify whether the model is universal.

#### Research conclusions

Gender, age, history of lumbago and abdominal pain, operation time, and urinary leukocytes and urinary glucose were independent risk factors for concurrent urinary sepsis in patients with DM and UUTCs, and we predicted high-risk patients using a risk model.

#### Research perspectives

The universality of the model could be verified based on multicenter data, and then extended to clinical practice.

## FOOTNOTES

Author contributions: Gou JJ designed and performed the research and wrote the paper; Zhang C designed the research and supervised the report; Han HS designed the research and contributed to the analysis; Wu HW provided clinical advice; Zhang C, Han HS, and Wu HW supervised the report.

Institutional review board statement: The study was reviewed and approved by the Institutional review board of People's Hospital of Jincheng (Approval No. JCPH.No20230401001).

Informed consent statement: All study participants or their legal guardian provided informed written consent about personal and medical data collection prior to study enrolment.



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**Retrospective Study** 

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ORIGINAL ARTICLE

## Effect of sitagliptin combined with Yiqi yangyin huoxue decoction on clinical efficacy and hemorheology in early diabetic nephropathy

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## Abstract

#### BACKGROUND

Early diabetic nephropathy (DN) is a complication of diabetes mellitus. It mainly affects kidney microvessels and glomerular function, and its timely and effective treatment is critical for early DN. However, the effects of treatments comprising simple Western medicine are not optimal. With the promotion and implementation of integrated Chinese and western medicine treatments, remarkable results have been achieved for many diseases. To this end, we explored the clinical efficacy of integrated traditional Chinese and western medicines for the treatment of early DN.

#### AIM

To investigate the effect of sitagliptin tablets combined with Yiqi yangyin huoxue decoction on clinical efficacy and hemorheology in patients with early DN.

#### **METHODS**

Through a retrospective analysis, 123 patients with early DN were admitted to the endocrinology clinic of the Changzhou NO. 7 People's Hospital from January 2021 to October 2022 and were selected as study subjects. After rigorous screening, 100 patients with early DN were enrolled. The control group (CG, n =50) and the observation group (OG, n = 50) were divided according to the treatment method. The CG were treated with sitagliptin, and the OG were treated with sitagliptin plus the Yiqi yangyin huoxue decoction. Both groups were treated for 3 mo. For both groups, the baseline data and clinical efficacy were compared, and changes in blood glucose levels, lipid levels, renal function, and hematological indicators before (T0) and after (T1) treatment were assessed.

## RESULTS

The total effective rate for the OG was 94.00% and that of the CG was 80.00% (P <



0.05). After treatment (T1), the levels of fasting blood glucose, 2 h postprandial glucose, total cholesterol, triacylglycerol, and low-density lipoprotein cholesterol in OG patients were obviously lower than those in the CG (P < 0.05), and cystatin C, homocysteine, urinary microalbumin, and blood creatinine values in OG patients were also obviously lower than those in the CG (P < 0.05); erythrocyte deposition, plasma viscosity, whole blood high shear viscosity, and whole blood low shear viscosity were markedly lower in OG patients than in the CG (P < 0.05).

#### CONCLUSION

Sitagliptin combined with Yiqi yangyin huoxue decoction has a remarkable effect when used to treat patients with early DN. Further, it is helpful in improving hemorheological indices and controlling disease progression.

Key Words: Sitagliptin; Yiqi yangyin huoxue decoction; Early diabetic nephropathy; Renal function; Clinical effects; Hemorheology

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**Core Tip:** Early diabetic nephropathy (DN) mainly involves kidney microvessels and glomerular functions, and its timely and effective treatment is critical. In clinical practice, the overall treatment effect of medication alone is not ideal. The development of traditional Chinese medicine has demonstrated its superiority for the treatment of many diseases. In the present study, sitagliptin tablets combined with a blood decoction were used to treat patients with early DN. The results showed that sitagliptin combined with Yiqi yangyin huoxue decoction had a significant effect on the treatment of early DN, which was helpful in improving hemorheological indices.

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## INTRODUCTION

Diabetes mellitus (DM) is a metabolic disease characterized by defective insulin secretion and activity. The prevalence of diabetes in China has been increasing, and that of DM in people over 18 years of age is as high as 11.2%[1]. Approximately 20%–40% of patients with diabetes in China have diabetic nephropathy (DN), and 21.8% of type 2 diabetes patients have this condition[2]. Early DN renal injury involves the glomeruli, renal tubules, renal interstitium, and renal blood vessels, with an increase in persistent urinary albumin excretion and a progressive decrease in the glomerular filtration rate, Early DN with glucose and lipid metabolism disorders, insulin resistance, hemodynamic changes, oxidative stress, inflammation, autophagy disorders, and other mechanisms[2-3]. Some patients with Type 2 DM (T2DM) have elevated urinary albumin levels upon diagnosis. If the microalbuminuria period is not treated promptly, it will rapidly progress to a lengthy albuminuria period until end-stage renal disease develops[4]. End-stage treatment of DN is more difficult than that of other renal diseases, with high disability rates. Accordingly, delaying the progression of DN is the focus of current research.

Presently, the key to clinical and early stage DN management is timely and effective treatment, such that disease development can be effectively controlled, which is beneficial for improving prognosis. Western medicine mainly involves lifestyle adjustments, a low-protein diet, lowering glucose and blood pressure, adjusting fat, reducing proteinuria, and improving renal function, as fundamental treatment principles<sup>[5]</sup>. Sitagliptin is a dipeptidyl peptidase-4 inhibitor that effectively improves fasting and postprandial blood glucose<sup>[6]</sup> in patients with T2DM<sup>[6]</sup>. Clinically, sitagliptin attenuates the oxidative stress response in kidney tissues, delays the progression of DN, and prevents the deterioration of renal function<sup>[7]</sup>. However, owing to the poor prognosis of DN, the overall treatment impact of individual medications is not ideal; therefore, other therapeutic drugs must be used in combination to improve the curative effect on early DN. With the development of traditional Chinese medicine (TCM), targeted protection of the potentially involved viscera, the predictable control of disease progress, and intervention-based treatment, significant results have been achieved for many diseases<sup>[8]</sup>. Therefore, this study was based on the basic pathogenesis of early DN with Qi and Yin insufficiency, dampness and heat, the application of sitagliptin tablets with Yiqi yangyin huoxue decoction for the treatment of early DN patients, an observation of its clinical impact on early DN patients, and its influence on blood rheology for early clinical intervention and to delay the progression of DN.

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## MATERIALS AND METHODS

#### **Patient characteristics**

Through a retrospective analysis, 123 patients with early DN admitted to the endocrinology clinic of the Changzhou NO. 7 People's Hospital from January 2021 to October 2022 were selected as the study participants. After rigorous screening, 100 patients with early DN were enrolled. The control group (CG, n = 50) and the observation group (OG, n = 50) were divided according to the treatment method.

The inclusion criteria were as follows: (1) All patients were between 18 and 75 years of age, independent of sex; (2) All patients met the diagnostic criteria for T2DM; and (3) Baseline data, clinical characteristics, laboratory indicators, and other data of all patients were complete. The exclusion criteria were as follows: (1) Patients with type 1 DM or other special types of DM; (2) Patients with infectious diseases, liver disease, cardiovascular disease, urinary system diseases, and kidney diseases; (3) Patients with malignant tumors; (4) Those unable to cooperate in clinical treatment or with mental illness; (5) Pregnant or lactating women; (6) Recent cases of diabetic ketoacidosis; and (7) Patients whose renal lesions were not induced by DM.

Diagnostic criteria for T2DM were as follows, based on the 1999 World Health Organization expert consultation report [9,10]: (1) Typical diabetes symptoms (polydipsia, polyuria, polyphagia, unexplained weight loss) plus random blood glucose  $\geq 11.1 \text{ mmol/L}$ ; (2) Fasting blood glucose (FBG)  $\geq 7.0 \text{ mmol/L}$ ; and (3) Oral glucose tolerance test 2 h blood glucose test  $\geq 11.1 \text{ mmol/L}$ , for patients without diabetes symptoms, which needed to be repeated on another day. Diagnostic criteria for early DN[11] were as follows: Referring to the 2012 updated Kidney Disease Outcomes Quality Initiative Clinical Practice Guidelines for Diabetes and Chronic Kidney Disease, excluding other causes (24 h infection, fever, congestive heart failure) of chronic kidney disease, the urine microalbumin (mg/L)/urine creatinine (g/L) (UACR) was significantly elevated in two of three consecutive examinations (30 mg/g  $\leq$  UACR  $\leq$  300 mg/g). DN staging using the international universal Mogensen staging[12] was performed as follows: Mogensen I, II, and III stages represent early DN; the urinary albumin excretion rate in early DN was examined three consecutive times within 3 mo and was approximately 20–200 µg/min or 30–300 mg/d. Diagnostic criteria for TCM symptoms[13] were as follows: Consistent with Qi and Yin insufficiency and blood stasis syndrome, the main symptoms were turbid urine, dull complexion, hand-footheart heat, fatigue, limb edema, and low back pain; secondary symptoms were frequent urination, palpitations, dry throat, dry mouth, skin nails, and limb numbness; and the tongue pulse was as follows: Tongue thin, red or dark purple, sublingual vein, ecchymosis, and a weak or string pulse.

#### Therapeutic method

All enrolled participants received basic treatment as follows: (1) Diabetes health knowledge education, focusing on explaining the relevant knowledge of early DN to the participants and persuading patients to correctly understand and treat such diseases so that they can maintain a good mood throughout the treatment process; (2) Diet control, developing a high-quality, low-protein diet for each patient, a personalized diet plan, and a low-salt diet for patients with hypertension; (3) Exercise guidance, according to the specific circumstances of the patient, personalized exercise should be arranged for the patient to ensure that aerobic exercise and anaerobic exercises are performed alternately, with the exercise time for each patient being  $\geq$  150 min per week, thus gradually improving the exercise level of the patient; (4) Blood glucose-lowering therapy, based on the actual blood glucose status of the patient, oral medicine or insulin was selected and the blood glucose index was monitored in real time to ensure that it was stable and met the inclusion criteria; FBG was controlled between 5 and 8 mmol/L, and 2 h postprandial glucose (2 h PG) was controlled between 7 and 10 mmol/L; and (5) Symptomatic treatment, other treatments that did not interfere with the purpose of the study and non-angiotensin receptor blockers drugs were used to regulate blood pressure to ensure that this met the inclusion criteria.

The CG patients were treated with sitagliptin (Merck Sharp & Dohme Ltd., Chinese medicine approval word: J20140095, specification: 100 mg, seven tablets and four boards) at 100 mg/time, po, qd, and treatment was performed for 3 mo. OG patients were orally administered sitagliptin combined with Yiqi yangyin huoxue decoction daily in the morning and evening once each, after meals, and treatment was performed for 3 mo. The Yiqi yangyin huoxue decoction proposed by the investigator and the prescription composition are listed in Table 1.

#### **Observational indicators**

(1) General observation measures included sex, age, disease duration, systolic blood pressure (SBP), diastolic blood pressure (DBP), and body mass index (BMI); (2) Based on the Guiding Principles for Clinical Research on the Treatment of Diabetes with New TCM and the Guiding Principles for Clinical Research of New TCM in the Treatment of Chronic Nephritis, complete cure: 24 h urine albumin excretion rate (UAER) normal, renal function is normal; excellent: 24 h UAER reduced by 40%, normal or basically normal renal function; effective, 24 h UAER reduced by < 40%, renal function is normal or improved; invalid: Clinical manifestations and laboratory examination are not improved or aggravated; (3) Improvements in pre-treatment (T0) and posttreatment (T1) blood glucose levels, namely FBG and 2 h PG; (4) The blood lipid contents before treatment (T0) and after treatment (T1), namely total cholesterol, triacylglycerol, and low-density lipoprotein cholesterol (LDL-C), were compared between the two groups; (5) Pre-and post-treatment (T0) renal functions, including improvements in cystatin C (Cys-C), homocysteine (HCY), urinary microalbumin (U-mAlb), and blood creatinine (Cr) levels, were also compared; and (6) Blood rheological indices before (T0) and after treatment (T1), including changes in red blood cell deposition, plasma viscosity, high resection viscosity of whole blood, and low resection viscosity of whole blood, were assessed.

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Table 1 Composition of decoction used for nourishing Qi, nourishing Yin, and promoting blood circulation						
Туре	Dosage, g					
Astragalus	20					
Chinese yam	15					
Poria cocos	10					
Dwarf lilyturf root	10					
Danpi	10					
Common peony root	10					
Prepared Rehmannia root	10					
Dogberry	15					
Sealwort	10					
Root of Zhejiang figwort	10					
Ligusticum wallichii	10					

#### Statistical analysis

Data were analyzed using SPSS statistical software version 25.0. The measurement data were expressed as the mean ± SD and analyzed based on a t-test. Count data are expressed as the frequency percentage (n%) and were compared by performing a  $\chi^2$  test. Statistical significance was set at *P* < 0.05.

#### RESULTS

#### Basic data analysis

The number of patients included and the flow chart of the analysis method are shown in Figure 1. There was no statistically significant difference in gender, age, course of disease, SBP, DBP, and BMI clinical data between the two groups of patients (P > 0.05) are presented in Table 2.

#### Evaluation of therapeutic efficiency

After 3 mo of treatment, the total effective rate was 94.00% for OG patients; the total effective rate was 80.00% for CG patients, the difference between the two groups is statistically significant (P < 0.05; Figure 2).

#### Improvement in blood glucose levels

After treatment (T1), the FBG and 2 h PG levels in the OG patients were markedly lower than those in the CG patients (P < 0.05; Figure 3).

#### Improvement in blood lipid levels

At T0, a comparison of TC, TG, and LDL-C contents was not significantly differ between the two groups (P > 0.05), and at T1, the contents of TC, TG, and LDL-C in the OG were markedly lower than those in the CG (P < 0.05; Table 3).

#### Improvement in renal function

At T0, serum Cys-C, HCY, mAlb, and Cr contents was not significantly differ between the two groups (P > 0.05); however, At T1, serum Cys-C, HCY, U-mAlb, and Cr values of patients in the OG were markedly lower than those of patients in the CG (P < 0.05; Table 4).

#### Improvement in hemorheology

At T1, erythrocyte deposition, plasma viscosity, whole blood high shear viscosity, and whole blood low shear viscosity in the OG were markedly lower than those in the CG (P < 0.05; Table 5).

#### DISCUSSION

The basic pathogenesis of diabetes is dryness-heat due to insufficiency. Qi Yin deficiency and meridian obstruction are the pathological causes of diabetes. The early stage of diabetes is dominated by Yin deficiency with heat excess, but when it progresses to the stage of DN, the disease is mostly at the stage of Qi and Yin insufficiency, and blood stasis becomes an important aspect of its pathogenesis<sup>[14]</sup>. Clinically, early DN is commonly characterized by the insufficiency of both Qi



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Table 2 Analysis of basic data between the two groups (mean ± SD)						
Index	OG ( <i>n</i> = 50)	CG ( <i>n</i> = 50)	χ²/t value	<i>P</i> value		
Sex (male/female)	28/22	26/24	0.161	0.688		
Age (yr)	51.24 ± 4.43	$51.28 \pm 4.28$	-0.046	0.963		
Disease course (yr)	$1.48\pm0.47$	$1.49 \pm 0.57$	-0.188	0.851		
SBP (mmHg)	$123.47 \pm 10.28$	124.16 ± 11.33	-0.319	0.750		
DBP (mmHg)	78.26 ± 4.15	79.51 ± 4.36	-1.480	0.142		
BMI (kg/m <sup>2</sup> )	25.21 ± 1.27	$25.19 \pm 1.44$	0.038	0.970		

The course of the disease was diabetic nephropathy. OG: Observation group; CG: Control group; SBP: Systolic blood pressure; DBP: Diastolic blood pressure; BMI: Body mass index.

Table 3 Improvement in blood lipid levels in both groups (mean ± SD)							
Group	Time	OG ( <i>n</i> = 50)	CG ( <i>n</i> = 50)	t-value	<i>P</i> value		
TC (mmol/L)	Т0	$7.05 \pm 1.21$	$7.08 \pm 1.19$	-0.166	0.868		
	T1	$4.97 \pm 0.73$	$6.13\pm0.84$	-7.366	< 0.001		
TC (mmol/L)	Т0	$1.65\pm0.17$	$1.63\pm0.14$	0.637	0.525		
	T1	$1.32 \pm 0.28$	$1.49\pm0.21$	-3.475	< 0.001		
LDL-C (mmol/L)	Т0	$4.13\pm0.68$	$4.09\pm0.71$	0.219	0.827		
	T1	$1.87\pm0.42$	$2.45\pm0.67$	-5.196	< 0.001		

OG: Observation group; CG: Control group; TC: Total cholesterol; TG: Triacylglycerol; LDL-C: Low-density lipoprotein cholesterol; T0: Pre-treatment; T1: Post-treatment

Table 4 Improvement in renal function in the two groups (mean ± SD)							
Group	Time	OG ( <i>n</i> = 50)	CG ( <i>n</i> = 50)	<i>t</i> -value	P value		
Cys-C (mg/L)	Т0	$1.59 \pm 0.43$	$1.57 \pm 0.45$	0.333	0.74		
	T1	$1.57\pm0.45$	$1.48\pm0.39$	-2.566	0.012		
HCY (µmol/L)	Т0	$30.41 \pm 5.73$	$30.44 \pm 5.69$	-0.018	0.986		
	T1	$19.28 \pm 3.15$	22.36 ± 4.28	-4.097	< 0.001		
U-mAlb (mg/L)	Т0	195.67 ± 15.28	195.73 ± 15.14	-0.017	0.987		
	T1	$162.38 \pm 11.54$	$174.63 \pm 13.45$	-4.887	< 0.001		
Cr (µmol/L)	Т0	$102.54 \pm 9.47$	102.71 ± 9.24	-0.096	0.924		
	T1	89.24 ± 7.36	96.53 ± 8.47	-4.591	< 0.001		

OG: Observation group; CG: Control group; Cys-C: Cystatin C; HCY: Homocysteine; U-mAlb: Urinary microalbumin; Cr: Blood creatinine.

and Yin, as well as blood stasis of the venation. Its pathogenesis includes meridian infarction, blocked veins, and deficiency of both Qi and Yin. For DN (see TCM), the following are noted: Urinary turbidity, fatigue, hot hands and feet, dull complexion, swollen limbs, low back pain or frequent urination, palpitations and restlessness, dry throat and mouth, skin and nail lesions, numbness of limbs, tongue thinning and exhibiting a dark red or purple texture, tortuous sublingual veins, ecchymosis, and weak or astringent veins[15]. Therefore, the early detection of DN is beneficial for controlling disease progression. Owing to the poor efficacy of drug treatment alone, we used western medicine and TCM for targeted treatment.

This study shows that the clinical effectiveness of Yiqi yangyin huoxue decoction for the treatment of early DN, as well as that compared with conventional western medicine treatment. We found that the total response rate of patients in the OG was markedly better than that in the CG, indicating that sitagliptin and Yiqi yangyin huoxue decoction, used to treat early diabetic kidney disease, have a good synergistic effect, which can effectively relieve the clinical symptoms of early



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Table 5 Improvement in hemorheology in the two groups (mean ± SD)									
Group	Time	OG ( <i>n</i> = 50)	CG ( <i>n</i> = 50)	<i>t</i> -value	P value				
Red blood cell deposition (%)	Т0	46.88 ± 5.23	46.91 ± 5.19	-0.029	0.977				
	T1	$37.15 \pm 4.13$	$43.96 \pm 4.87$	-7.542	< 0.001				
Plasma viscosity (mPa s)	Т0	$3.26 \pm 0.38$	$3.29 \pm 0.43$	-0.490	0.625				
	T1	$1.52\pm0.25$	$2.27\pm0.31$	-13.465	< 0.001				
Whole blood high shear viscosity (mPa s)	Т0	$5.94 \pm 1.13$	$5.59 \pm 1.07$	1.598	0.113				
	T1	$4.06\pm0.75$	$5.13 \pm 0.81$	-6.854	< 0.001				
Whole blood low shear viscosity (mPa s)	Т0	$10.29 \pm 2.34$	10.33 ± 2.29	-0.108	0.914				
	T1	$7.70\pm1.25$	$8.72 \pm 1.64$	-3.527	0.001				

OG: Observation group; CG: Control group.





#### Figure 1 Number of patients included, and flow chart of analysis method. OG: Observation group; CG: Control group.

diabetic kidney disease, improve renal function, and reverse or delay early kidney injury. Our study also found that after treatment, compared with those before treatment, the levels of serum FBG, 2 h PG, TC, TG, and LDL-C were markedly lower in the OG compared to those in the CG. This shows that sitagliptin combined with Yiqi yangyin huoxue decoction can effectively reduce blood glucose and lipid levels, improve the state of hyperglycemia and hyperlipidemia, inhibit thrombosis formation, and slowed disease progression in patients with early DN. This could be because sitagliptin is a dipeptidyl peptidase-4 inhibitor. It is also a type of hypoglycemic drug that has an inhibitory effect on pancreatic islet α-cell hyperplasia and ultimately increases insulin secretion, such that blood sugar can be effectively regulated[16,17]. There are many drugs in the Yiqi yangyin huoxue decoction that have related effects. For example, when used in combination with astragalus and *Rehmannia*, blood sugar levels can be effectively reduced. By combining the tonifying effect of astragalus with the nourishing and kidney-strengthening effect of *Rehmannia*, the Qi and blood of the patient can also be replenished. *Astragalus membranaceus* can enhance immunity, help to resist oxidation, and have a certain regulatory effect on blood lipid and blood glucose levels in the body[18]. Further, *Rehmannia glutinosa* can improve hormone levels, especially the level of adrenaline, and has a certain improving effect on blood sugar and blood lipids in the body.

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Ling J et al. Therapeutic effect on diabetic nephropathy



Figure 2 Efficacy evaluation based on the two groups. OG: Observation group; CG: Control group. <sup>a</sup>P < 0.05.



**Figure 3 Comparison of blood glucose levels between the two groups.** OG: Observation group; CG: Control group; FBG: Fasting blood glucose; 2 h PG: 2 h postprandial blood glucose; T0: Pre-treatment; T1: Post-treatment. °*P* < 0.001.

Oxidative stress reactions comprise one component of the pathogenesis of DN, and the reactive oxygen species produced by this reaction have an important impact on renal function. CysC is a small-molecule protein that is relatively stable in blood and is mostly used clinically to evaluate renal function. Moreover, it is an independent risk factor for diabetic complications. When renal function is impaired, serum levels increase significantly [19]. HCY is a non-essential amino acid with a sulfur-containing group, demethylated by methionine, and is closely related to renal function and metabolism. Research has shown that HCY can promote the excessive generation of oxygen free radicals and hydrogen peroxide, leading to the aggravation of renal tissue damage and acceleration of disease progression in DN[20]. U-mALB is a negatively charged protein secreted by the liver, and the vast majority of it fails to cross the glomerular filtration membrane charge barrier. An increase in urinary U-mALB content indicates altered glomerular permeability, and glomerular filtration membrane charge barrier is impaired[21]. Serum Cr is also a biochemical index used to evaluate renal function; its increase is related to a decrease in the glomerular filtration rate, and with this, renal function damage or renal failure is considered[22]. The results of our study showed that the levels of CysC, HCY, mALB, and Cr in the OG were markedly lower than those in the CG, indicating that sitagliptin combined with Yiqi yangyin huoxue decoction could reduce albuminuria, improve tubular function, and exert protective effects on renal function. Astragalus in Yiqi yangyin huoxue decoction can improve the balance of water and sodium in the body, change the permeability of blood vessels to reduce proteinuria, and effectively regulate renal function in patients with early DN[23]. Ligusticum wallichii can effectively prevent the production of advanced glycosylation end products in the renal cortex and inhibit the apoptosis of renal cells, thus improving renal function.

Microcirculation disorders are an important pathological basis of DN. Microangiopathy is the pathological basis of typical clinical manifestations, such as proteinuria, in patients with early DN[24]. Hemodynamic abnormalities are also important causes of proteinuria and glomerulosclerosis in patients with DN[25]. The results of this study showed that after treatment, the erythrocyte deposition, plasma viscosity, whole blood high shear viscosity, and whole blood low shear viscosity in the OG were markedly lower than those in the CG. This indicated that sitagliptin combined with Yiqi yangyin huoxue decoction could significantly improve blood rheological indices in patients with early DN. The Yiqi yangyin huoxue decoction uses *A. membranaceus* and prepared *Rehmannia* root as monarch drugs. *Astragalus membranaceus* has a slightly warm and sweet taste, which is beneficial for reducing water-swelling, supplementing Qi, consolidating the surface, strengthening the spleen, and tonifying the middle. The *Rehmannia* root is a processed product of *R. glutinosa*. It has sweet and bitter tastes and is beneficial for the heart, liver, and lung meridians. It nourishes the blood, promotes fluid production, clears heat, and cools the blood. When used in combination, it has the effect of strengthening Qi and Yin. The use of Chinese yam, Dogberry, *Poria cocos*, Sealwort, and Dwarf lilyturf root as medicinal herbs has a role in nourishing



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Qi in patients, and it is the main treatment for patients with spleen deficiency caused by fatigue and lumbar debility. The roots of Zhejiang figwort, Danpi, *L. wallichii*, and common peony root act as adjuvants, prevent thrombosis, improve hypercoagulability, promote blood circulation, and resolve blood stasis; after entering the body, they can improve blood viscosity, speed up blood flow, and reduce glomerular filtration, nourish the kidney, repair the kidney, and work together to complement each other, and thus, they have a marked influence on the treatment of diseases. Limitations of this study are as follows. Owing to time and sample size limitations, the selected subjects were all admitted to our hospital. Next, we will expand the sample size in clinical practice or conduct multicenter research and further explore the mechanism underlying the effects of Yiqi yangyin huoxue decoction for the treatment of early DN.

## CONCLUSION

In conclusion, treatment with sitagliptin combined with Yiqi yangyin huoxue decoction in patients with Qi insufficiency and early DN has a significant effect, which is helpful in improving blood glucose, blood lipids, renal function, and hematological indicators and helping control disease progression, which is worthy of further clinical research.

## **ARTICLE HIGHLIGHTS**

#### Research background

Early diabetic nephropathy (DN) is a major complication of diabetes, a disease induced by glomerular sclerosis caused by a long-term glucose metabolism disorder. Progressive renal damage, edema, polyuria, and proteinuria are the main clinical symptoms of this disease. The main treatment plan comprising Western medicine for early DN is lipid-lowering and blood pressure regulation; however, the treatment effect is not good. Therefore, we used traditional Chinese medicine (TCM) to control disease progression from the root cause.

#### **Research motivation**

With the development of TCM, we already know that the main pathogenic mechanism of early DN is "Kidney deficiency and blood stasis, Qi and Yin insufficiency." Therefore, we used western medicine combined with a TCM program for targeted treatment.

#### **Research objectives**

This study aimed to use sitagliptin tablets combined with Yiqi yangyin huoxue decoction to treat patients with early diabetic nephropathy, to observe the clinical efficacy and blood rheology in these patients, and to provide a basis for early clinical intervention to delay the progression of this disease.

#### **Research methods**

Using a retrospective approach, patients with early DN were randomized into control group (CG) (n = 50) and observation group (OG) (n = 50). CG patients were treated with sitagliptin, and OG patients were treated with sitagliptin plus the Yiqi yangyin huoxue decoction. Both groups were treated for 3 mo.

#### **Research results**

The total response rate of the patients in the OG was 94.00%, and the total response rate of those in the CG was 80.00%. After treatment (T1), fasting blood glucose, 2 h postprandial glucose, total cholesterol, triacylglycerol, low-density lipoprotein cholesterol, cystatin C, homocysteine, urinary microalbumin, and blood creatinine levels in OG patients were markedly lower than those in the CG; further, red blood cell deposition, plasma viscosity, and hyperviscosity of whole blood in OG patients were markedly lower than those in the CG (all P < 0.05).

#### **Research conclusions**

Sitagliptin combined with Yiqi yangyin huoxue decoction has achieved remarkable results in patients with early DN, helping to improve blood glucose, blood lipid, renal function, and hematological indicators.

#### Research perspectives

In this retrospective study, patients with early DN were randomized into the CG and OG; the CG patients were treated with sitagliptin, and The OG patients were treated with sitagliptin plus Yiqi yangyin huoxue decoction; both groups were treated for 3 mo. Comparing the baseline data and clinical efficacy between the two groups, changes in blood glucose, blood lipid, renal function, and the hematology index were observed before (T0) and after treatment (T1). Thus, we explored the efficacy of sitagliptin in combination with Yiqi yangyin huoxue decoction for early DN treatment and its effect on blood rheology.

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## FOOTNOTES

Author contributions: This article was written by Ling J, who independently completed the research design and data analysis; and Yang YH provided important guidance for solving difficult and complex problems.

Institutional review board statement: The study was reviewed and approved by the Biomedical Research Ethics Committee, Changzhou NO. 7 People's Hospital.

Informed consent statement: This is a retrospective study and has applied for exemption from informed consent.

Conflict-of-interest statement: All the authors report no relevant conflicts of interest for this article.

Data sharing statement: Data used in this study were obtained from the corresponding author at yyh0728@163.com.

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META-ANALYSIS

## Effectiveness and safety of traditional Chinese medicine for diabetic retinopathy: A systematic review and network meta-analysis of randomized clinical trials

Hong-Dian Li, Ming-Xuan Li, Wen-Hua Zhang, Shu-Wen Zhang, Yan-Bing Gong

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reviewed.	Wen-Hua Zhang, Shu-Wen Zhang, Yan-Bing Gong, Dongzhimen Hospital, Beijing University of
Peer-review model: Single blind	Chinese Medicine, Beijing 100700, China
Peer-review report's scientific quality classification	<b>Corresponding author:</b> Yan-Bing Gong, PhD, Dean, Dongzhimen Hospital, Beijing University of Chinese Medicine, No. 5 Haiyuncang Hutong, Dongcheng District, Beijing 100700, China.
Grade A (Excellent): A	gyb_1226@163.com
Grade B (Very good): B	
Grade C (Good): C	Abstract
Grade D (Fair): 0	ADSILICI
Grade E (Poor): 0	BACKGROUND
	Diabetic retinopathy (DR) is currently recognized as one of the most serious
<b>P-Reviewer:</b> Rahmati M, Iran;	diabetic microangiopathies and a major cause of adult blindness. Commonly used
Salceda R, Mexico; Horowitz M,	clinical approaches include etiological control, microvascular improvement, and
Australia	surgical intervention, but they are ineffective and have many side effects. Oral
Received: April 17, 2023	Chinese medicine (OCM) has been used for thousands of years to treat DR and is still widely used today, but it is unclear which OCM is more effective for DR
Peer-review started: April 17, 2023	suit where used today, but it is unclear which oew is note encenve for DR.
First decision: June 1, 2023	AIM
<b>Revised:</b> June 14, 2023	To estimate relative effectiveness and safety profiles for different classes of OCMs
Accepted: July 29, 2023	for DR, and provide rankings of the available OCMs.
Article in press: July 29, 2023	METHODS
Published online: September 15,	The search time frame was from the creation of the database to January 2023

The search time frame was from the creation of the database to January 2023. RevMan 5.3 and Stata 14.0 software were used to perform the systematic review and Network meta-analyses (NMA).

## **RESULTS**

A total of 107 studies and 9710 patients were included, including 4767 cases in the test group and 4973 cases in the control group. Based on previous studies and clinical reports, and combined with the recommendations of Chinese guidelines for the prevention and treatment of DR, 9 OCMs were finally included in this study, namely Compound Xueshuantong Capsules, Qiming Granules, Compound



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Danshen Dripping Pills, Hexue Mingmu Tablets (HXMM), Qiju Dihuang Pills (QJDH), Shuangdan Mingmu Capsules (SDMM), Danggui Buxue Decoction (DGBX), Xuefu Zhuyu Decoction and Buyang Huanwu Decoction. When these nine OCMs were analyzed in combination with conventional western medicine treatment (CT) compared with CT alone, the NMA results showed that HXMM + CT has better intervention effect on the overall efficacy of DR patients, HXMM + CT has better effect on improving patients' visual acuity, SDMM + CT has better effect on inhibiting vascular endothelial growth factor, DGBX + CT has better effect on reducing fundus hemorrhage area, HXMM + CT has better effect on reducing fasting blood glucose, and QJDH + CT has better effect on reducing glycated hemoglobin. When there are not enough clinical indicators for reference, SDMM + CT or HXMM + CT treatments can be chosen because they are effective for more indicators and demonstrate multidimensional efficacy.

#### CONCLUSION

This study provides evidence that combining OCMs with CT leads to better outcomes in all aspects of DR compared to using CT alone. Based on the findings, we highly recommend the use of SDMM or HXMM for the treatment of DR. These two OCMs have demonstrated outstanding efficacy across multiple indicators.

Key Words: Diabetic retinopathy; Network meta-analysis; Traditional Chinese medicine; Therapeutic effect; Systematic review

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**Core Tip:** To our knowledge, this study represents the first network meta-analysis (NMA) examining the effectiveness of traditional Chinese medicine in treating diabetic retinopathy (DR). Notably, this NMA includes the largest number of original studies, subjects, and variety of Chinese medicines to date. While the efficacy of Chinese medicine for DR has been widely recognized in China, no previous studies have systematically evaluated which Chinese medicine treatment is the most effective. Therefore, this study fills an important gap in the field.

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## INTRODUCTION

Diabetic retinopathy (DR) is currently recognized as one of the serious diabetic microangiopathies and is the leading cause of blindness in adults. According to the International Diabetes Federation, it is estimated that the number of people with diabetes will reach 642 million worldwide in 2040, and 34.6% of these patients will have DR[1]. DR causes irreversible visual impairment, including abnormal vision, blurred vision, and even blindness. In addition, the presence of DR implies an increased risk of life-threatening systemic vascular complications[2]. Currently, the main treatments for DR include retinal laser photocoagulation, pharmacotherapy, hormonal therapy, and surgery. However, these treatments may lead to adverse effects such as increased angiogenesis, increased intraocular pressure, and retinal hemorrhage[3,4]. Studies have shown that age is a key factor affecting DR, and the number of DR patients in the elderly population will reach new highs as the world ages[5]. In view of the current situation, the pathogenesis of DR is being actively explored around the world and effective therapeutic drugs are being explored.

In fact, traditional Chinese medicine (TCM) has long been considered a promising complementary therapy that dates back more than 1000 years. In China, many facts have proven that herbal medicine can effectively improve the fundus condition of patients, relieve the pain of the disease, and obtain a better quality of life for patients through multi-target and multi-path interventions in DR[6]. Many Oral Chinese medicine (OCM), including proprietary Chinese medicine preparations and herbal granules are widely used in the treatment of DR. For example, Qiming Granules (QM), the first OCM approved by the State Food and Drug Administration for the treatment of DR, whose main ingredients are *Hedysarum Multijugum Maxim*. (Huangqi in chinese), *Radix Puerariae* (Gegen in chinese), *Lycii Fructus* (Gouqizi in chinese), and *Cassiae Semen* (Juemingzi in chinese), *etc.*, were shown to alleviate retinal hypoxia and ischemia by increasing retinal blood flow and improving blood circulation in a multicenter, randomized, parallel controlled clinical trial[7]. Then for example, Compound Xueshuantong Capsules (XST), an OCM commonly used for DR, was shown to effectively improve the disorder of retinal structure and edema in streptozotocin-induced type 2 diabetic rats by activating the PPAR signaling pathway, reversing the reduction in retinal thickness and retinal ganglion cell number, and reducing the apoptotic index of retinal cells[8]. Compound Danshen dripping pills, an oral proprietary Chinese medicine containing Danshen, was found to improve vision and clinical symptoms and reduce the incidence of macular edema compared to captopril in a retrospective study[9].

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In addition to these well known OCMs, there are many lesser known OCMs that are widely used in clinical practice. However, the selection of these OCMs remains a challenge for patients with different disease states. Network metaanalyses (NMA) allow for the comparison of multiple treatments (*i.e.* three or more) using both direct comparisons of interventions within randomized controlled trials (RCT) and indirect comparisons across trials based on a common comparator[10]. To date, there are no studies comparing different types of OCMs used for the treatment of DR. Therefore, we searched all RCTs of OCMs used for the treatment of DR and initiated this NMA to compare the efficacy between them, hoping to provide some suggestions for clinical practice.

## MATERIALS AND METHODS

The study protocol was registered on PROSPERO (International Prospective Register of Systematic Reviews). Registration No. CRD42022352250 (https://www.crd.york.ac.uk/PROSPERO/#myprosperoID=CRD42022352250). This program was developed in accordance with the Preferred Reporting Items For Systematic Review And Meta-analysis Protocols (PRISMA-P)[11]. The PRISMA Extension Statement is used to ensure that all aspects of the methods and results are reported[12].

#### Eligibility criteria

The Population-Intervention-Comparators-Outcomes-Studydesign framework was adopted as the eligibility criteria for the review as following.

#### Study type

The RCT is the original study that we agreed to include. We did not place any restrictions on the language, country, publication date, or phase of the RCT. Duplicate publications, summaries of personal experience, purely theoretical studies, reviews, animal or cellular experiments, and original studies with incorrect or incomplete data in the literature should also be excluded.

#### Population

Patients with DR who meet the standard diagnosis rely on fundus fluorescence angiography and fundus signs to detect microangiomas, exudates, hemorrhages, neovascularization, and other fundus changes. No restriction of age, gender, occupation and region. No concomitant ocular diseases caused by non-hyperglycemic factors, such as primary glaucoma and senile cataract; no acute metabolic diseases, such as diabetic ketoacidosis, within a short period of time before enrollment.

#### Intervention

Regarding the RCT of OCM for DR, the blinding and language are not limited. The control group received only oral western medicine, conventional western medicine treatment (CT) mainly included hypoglycemic drugs, lipid-lowering drugs and antihypertensive drugs, among which hypoglycemic drugs could include subcutaneous injection of insulin, in addition, other injectable drugs were not acceptable; antioxidant and microcirculation improvement drugs, such as calcium dobesilate, pancreatic kininogenase; drugs to promote retinal metabolism and nerve nutrition of the eye, such as lecithin complex iodine, etc. The test group was added to the control group with OCM. By combining previous studies and actual clinical observations, and also referring to the latest published Guideline for the prevention and treatment of type 2 diabetes mellitus in China (2020 edition)[13], we decided to study the 9 most commonly used OCMs which are XST (composed of Panax Notoginseng (Burk.) F. H. Chen Ex C. Chow/Figwort Root/Hedysarum Multijugum Maxim./ Radix Salviae), Compound Danshen Dripping Pills (DS, composed of Radix Salviae/Borneolum Syntheticum/Panax Notoginseng (Burk.) F. H. Chen Ex C. Chow), Qiming Granules (QM, composed of Hedysarum Multijugum Maxim./Radix Puerariae/Rehmanniae Radix Praeparata/Lycii Fructus/Leonuri Fructus/Cassiae Semen/Pollen Typhae/Whitm.ania Pigra Whitman), Hexue Mingmu Tablets (HXMM, Pollen Typhae/Rehmanniae Radix Praeparata/Radix Salviae/Ecliptae Herba/ Chrysanthemi Flos/Scutellariae Radix/Cassiae Semen/Plantaginis Semen/Leonuri Fructus/Fructus Ligustri Lucidi/Prunellae Spica/Gentianae Radix Et Rhozima/Curcumae Radix/Equiseti Hiemalis Herba/Radix Paeoniae Rubra/Cortex Moutan/Angelicae Sinensis Radix/Chuanxiong Rhizoma), Qiju Dihuang Pills (QJDH, Lycii Fructus/Rehmanniae Radix Praeparata/Chrysanthemi Flos/Cornus Officinalis Sieb. Et Zucc./Rhizoma Dioscoreae/Poria Cocos(Schw.) Wolf./Alisma Orientale (Sam.) Juz./Cortex Moutan), Shuangdan Mingmu Capsules (SDMM, Ecliptae Herba/Fructus Ligustri Lucidi), Danggui Buxue Decoction (DGBX, Hedysarum Multijugum Maxim./ Angelicae Sinensis Radix), Xuefu Zhuyu Decoction (XFZY, Persicae Semen/Carthami Flos/Radix Paeoniae Rubra/Chuanxiong Rhizoma/Radix Bupleuri/Platycodon Grandiforus/Licorice/Angelicae Sinensis Radix/ Rehmanniae Radix Praeparata/Achyranthis Bidentatae Radix/Aurantii Fructus) and Buyang Huanwu Decoction (BYHW, Hedysarum Multijugum Maxim./Angelicae Sinensis Radix/Radix Paeoniae Rubra/Chuanxiong Rhizoma/Persicae Semen/ Carthami Flos/Pheretima). To obtain the most accurate efficacy results for OCM, all studies performing non-oral treatments such as laser photocoagulation, surgery, injectable fluids, acupuncture, tui-na, and traditional Chinese gongfu were excluded from this study.

#### Outcome measures

The main efficacy indicators in this study encompass total clinical effectiveness and visual acuity. Following the international DR efficacy determination standard, the efficacy after treatment was categorized into three groups: Significantly



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effective, effective, and ineffective. Significantly effective cases were identified based on two criteria: (1) Improvement in visual acuity by  $\geq$  2 lines; and (2) improvement in two or more of the three fundus indices (microvessel count, hemorrhage, and exudation), or significant improvement in one or more of the three indices without deterioration in the remaining indices. Effective cases were determined by improvement in visual acuity and improvement in at least one of the three fundus indices without deterioration in the remaining indices. Ineffective cases were those that did not meet the criteria for effectiveness based on the indices. For assessing visual acuity, the international standard visual acuity table was utilized, where the counting included 2 rows for no light perception to light perception, and 1 row for each interval including light perception, manual, index, 0.02, 0.04, 0.06, 0.08, and 0.1, while refractive error measurements accounted for corrected visual acuity. The efficacy index considered the total clinical efficiency (comprising significantly effective and effective populations), reflecting the actual determination of clinical effects. Secondary indicators include fundus hemorrhagic area (FHA), fasting blood glucose (FBG), glycated hemoglobin (HbA1c) and vascular endothelial growth factor (VEGF).

#### Data sources and search strategy

A total of seven databases were searched by computer: China national knowledge infrastructure, Wanfang Database, Weipu Journal Database, Chinese Biomedical Literature Database, PubMed, Cochrane Library, and Web of Science database, and the search time of each database was built until January 2023. In addition, references to the literature were incorporated retrospectively to supplement access to relevant literature. The search takes a combination of subject terms and free words. English search terms include: "Diabetic Retinopathy", "Diabetic Retinopathies", "Traditional Chinese medicine", "Compound Xueshuantong Capsules", "Compound Danshen Dripping Pills", "Qiju Dihuang Pills", "Qiming Granules", "Hexue Mingmu Tablets", "Randomized Controlled Trial", *etc.* The detailed search strategy is described in Supplementary Table 1.

#### Data screening and quality evaluation

Two investigators (LMX and LHD) performed literature screening and data extraction, excluding duplicates and then first read the title and abstract to exclude literature that clearly did not meet the requirements, and then read the remaining literature in full to clarify whether it met the inclusion criteria, and if there was disagreement, a third investigator (WZ) had to be consulted for a decision after discussion. Data extraction included: (1) Basic information: Article title, first author, publication time, country/region, *etc.*; (2) study characteristics: Interventions in the trial and control groups, number of cases, age, duration of intervention, and adverse effects in the study subjects; (3) key information required for risk of bias evaluation in the literature; and (4) outcome indicators included in the test and control groups.

Quality evaluation of RCTs was performed usingRevMan 5.3 (Cochrane Collaboration, Copenhagen, The Nordic Cochrane Centre). Assessed by 2 investigators (HL and ML) using the tool for assessing risk of bias recommended in the Cochrane systematic reviewers' handbook 5.1[14], including the following 7 aspects: Random-sequence generation (selection bias); allocation concealment (selection bias); blinding of participants and personnel (performance bias); blinding of outcome assessment (detection bias); incomplete outcome data (attrition bias); selective reporting (reporting bias); others bias. Each aspect can be further categorized as "low risk of bias", "unclear risk of bias", and "high risk of bias". When there is a disagreement, it can be decided by mutual discussion or by consulting 3 investigators (SZ).

#### Statistical analysis

Stata 14.0 (MRC Biostatistics Unit, Cambridge, United Kingdom) software was used to implement the statistical process. The odds ratio (OR) and its 95% confidence interval (CI) were used for statistical data, and the mean difference and its 95%CI were used for measurement data. Data were preprocessed using the "Network" command, and two comparisons were made between different interventions, according to this case there is no closed loop, so the consistency model is used. The efficacy was ranked according to the surface under the cumulativeranking curve (SUCRA), and the results of the NMA analysis were finally presented in tabular form to obtain the relatively best interventions. Considering the inclusion of multiple observations, we combined the individual observations two by two and used a multivariate approach to determine the dependencies between the results. Use the "Clustering" command to obtain a clustering ranking chart[15]. Sensitivity analysis is performed for factors that may affect the stability of the results. Finally, "correction-comparison" funnel plots were drawn for publication bias assessment.

## RESULTS

#### Literature search results and basic characteristics

A total of 1667 relevant papers were identified and screened for inclusion in 107 randomized controlled trials, involving a total of 9710 patients (4767 in the trial group and 4973 in the control group). Baseline characteristics were carefully matched between the groups. The individual sample sizes of the trials ranged from 30 to 256 individuals, and the observation periods varied from 4 wk to 24 wk. The trials encompassed 9 different OCMs, namely XST (26 studies[16-41]), QM (23 studies[42-64]), DS (21 studies, references[65-85]), HXMM (9 studies[86-94]), QJDH (6 studies[95-100]), SDMM (5 studies[101-105]), DGBX (6 studies[106-111]), XFZY (7 studies[112-118]), and BYHW (4 studies[119-122]). All studies evaluated at least one of the two primary efficacy measures, namely total clinical effectiveness and visual acuity. It should be noted that all included studies were conducted exclusively in China. The literature screening process is shown in Figure 1, the characteristics of the literature are shown in Table 1, and the OCMs involved in the study and their details

## Table 1 Basic characteristics of included randomized controlled trials

Ref.	Sample size	Random	Interventions		Period of	Age	Outcomes	Adverse reactions
	C/E	method	C	E	- treatment	C/E		C/E
Men[16], 2020	40/40	TRD	CT + Calcium Dobesilate	C + XST	8 wk	67.46 ± 2.52/63.12 ± 2.21	1, 2, 3, 4	-
Bai[ <mark>17</mark> ], 2016	38/38	-	CT + Calcium Dobesilate	C + XST	24 wk	51.08 ± 4.73/50.63 ± 5.51	1, 2, 7	3/2
An[ <mark>18</mark> ], 2021	30/30	-	Calcium Dobesilate	C + XST	20 wk	69.51 ± 5.19/70.32 ± 5.39	1, 2	-
Yan and Song[ <mark>19</mark> ], 2020	46/46	TRD	Calcium Dobesilate	C + XST	12 wk	47.4 ± 4.6/48.5 ± 4.9	1,7	22/8
Li <mark>[20]</mark> , 2021	20/20	-	Calcium Dobesilate	C + XST	20 wk	59.67 ± 1.78/58.51 ± 1.31	1,3	-
Jin <mark>[21</mark> ], 2020	40/40	-	Calcium Dobesilate	C + XST	20 wk	66.12 ± 3.45/66.58 ± 3.16	2, 3	-
Wu <mark>[22</mark> ], 2015	50/50	TRD	СТ	C + XST	12 wk	54.1 ± 6.6/54.6 ± 6.2	1	-
Qu[23], 2018	35/35	-	CT + Calcium Dobesilate	C + XST	20 wk	59.7 ± 6.3/60.4 ± 7.2	1	-
Sui[ <mark>24</mark> ], 2017	47/49	-	CT + Calcium Dobesilate	C + XST	4 wk	-	1	-
Jiang <i>et al</i> [ <mark>25</mark> ], 2019	46/46	-	Calcium Dobesilate	C + XST	20 wk	58.72 ± 2.20/58.69 ± 2.15	1, 2, 4	-
Zhu <i>et al</i> [ <mark>26</mark> ], 2016	48/48	-	CT	C + XST	12 wk	56.38 ± 12.19/56.15 ± 12.21	2	-
Sun[ <mark>27</mark> ], 2021	18/18	-	СТ	C + XST	12 wk	61.38 ± 3.69/62.47 ± 3.01	1	-
Liu <i>et al</i> [ <mark>28]</mark> , 2018	45/45	TRD	CT + Calcium Dobesilate	C + XST	12 wk	48.56 ± 7.64/48.34 ± 6.49	1	-
Li <mark>[29]</mark> , 2019	49/49	TRD	CT + Calcium Dobesilate	C + XST	12 wk	$\begin{array}{c} 66.41 \pm \\ 4.11/66.82 \pm 4.03 \end{array}$	1, 2, 4	-
Xiao <mark>[30</mark> ], 2016	110/110	-	CT + Calcium Dobesilate	C + XST	12 wk	50.2 ± 6.4/49.5 ± 5.9	1,7	0/0
Chen <mark>[31</mark> ], 2019	39/39	-	Calcium Dobesilate	C + XST	12 wk	58.17 ± 3.82/59.34 ± 3.27	1, 3, 4	-
Zhao[ <mark>32</mark> ], 2020	43/44	-	Calcium Dobesilate	C + XST	24 wk	53.71 ± 5.52/53.66 ± 5.49	3	-
An[ <mark>33</mark> ], 2020	35/35	-	Calcium Dobesilate	C + XST	12 wk	52.12 ± 15.76/51.17 ± 17.83	1, 4, 7	1/2
Zhang[ <mark>34</mark> ], 2019	19/19	-	Calcium Dobesilate	C + XST	8 wk	51.86 ± 1.92/53.28 ± 2.64	1,7	0/0
Wei <i>et al</i> [ <mark>35</mark> ], 2017	34/34	TRD	СТ	C + XST	35 wk	62.94 ± 3.48/61.31 ± 3.54	1, 3, 4, 7	0/0
Meng <i>et al</i> [36], 2012	38/40	-	CT + Calcium Dobesilate	C + XST	24 wk	49.2 ± 7.8/48.8 ± 6.7	1	-
Zhu and Sui [ <mark>37</mark> ], 2022	76/76	TRD	CT + Calcium Dobesilate	C + XST	12 wk	55.89 ± 4.17/55.94 ± 4.13	1, 2, 3, 7	12/13
Li[ <mark>38</mark> ], 2017	33/33	-	Calcium Dobesilate	C + XST	12 wk	-	2	-
Wang et al [39], 2020	42/44	TRD	CT + Calcium Dobesilate	C + XST	20 wk	68.35 ± 6.82/69.52 ± 7.11	1, 3, 7	4/3
Hu[ <mark>40</mark> ],	30/30	-	Calcium	C + XST	20 wk	55.30 ±	1, 3, 7	0/0



2017			Dobesilate			$2.15/55.67 \pm 2.08$		
Xu and Ru [41], 2020	46/46	-	Calcium Dobesilate	C + XST	12 wk	52.3 ± 3.2/52.1 ± 3.6	1,3	-
Wang[ <mark>42</mark> ], 2016	38/38	-	Iodized Lecithin	C + QM	8 wk	$43 \pm 6/42 \pm 5$	1	-
Wang[ <mark>43</mark> ], 2018	44/44	TRD	Calcium Dobesilate	C + QM	12 wk	58.4 ± 7.5/57.8 ± 6.2	1,7	12/5
Wang et al [44], 2020	32/32	-	Amlodipine besylate	C + QM	12 wk	38.94 ± 4.89/39.87 ± 5.13	1, 2	-
Chen[ <mark>45</mark> ], 2016	45/45	TRD	СТ	C + QM	12 wk	62.05 ± 5.47/63.11 ± 5.64	1	-
Wang <i>et al</i> [ <mark>46]</mark> , 2015	38/41	-	Calcium Dobesilate	C + QM	24 wk	52.1 ± 5.6/52.5 ± 5.3	1, 2, 7	0/0
Sui <i>et al</i> [ <b>4</b> 7], 2014	43/43	-	Calcium Dobesilate	C + QM	12 wk	50.53 ± 11.28/50.22 ± 14.82	1, 2, 5, 6, 7	0/0
Huang[ <mark>48</mark> ], 2017	63/63	-	CT + Calcium Dobesilate	C + QM	12 wk	55.9 ± 4.1/55.6 ± 4.2	1	-
Feng <i>et al</i> [49], 2016	41/42	Lottery -	Calcium Dobesilate	C + QM	12 wk	55.89 ± 6.13/55.26 ± 6.29	1, 2, 5, 6	-
Ge[ <mark>50</mark> ], 2018	53/53	-	CT + Calcium Dobesilate	C + QM	24 wk	50.87 ± 3.71/51.25 ± 3.64	1	-
Yan <mark>[51</mark> ], 2020	38/46	Lottery	CT + Calcium Dobesilate	C + QM	8 wk	56.96 ± 4.59/56.65 ± 4.02	1, 2, 5, 6, 7	8/2
Meng <i>et al</i> [52], 2016	21/21	-	Calcium Dobesilate	C + QM	24 wk	-	1	-
Zhang <i>et al</i> [ <mark>53]</mark> , 2016	46/45	-	СТ	C + QM	24 wk	-	1	-
Zhang[ <mark>54</mark> ], 2013	34/34	-	СТ	C + QM	36 wk	-	1	-
Wu et al [55], 2022	50/50	Lottery	Pancreatic Kinino- genase	C + QM	24 wk	53.82 ± 5.42/54.06 ± 4.93	1, 2, 4, 7	3/5
Zhou and Femng <mark>[56]</mark> , 2018	60/60	-	CT + Calcium Dobesilate	C + QM	12 wk	58.5 ± 6.7/57.9 ± 6.2	1, 2	-
Wang et al [57], 2019	48/52	-	CT + Calcium Dobesilate	C + QM	24 wk	66.8 ± 6.3/66.7 ± 6.2	1,7	0/0
Wang[ <mark>58</mark> ], 2017	47/47	-	CT + Calcium Dobesilate	C + QM	12 wk	54.3 ± 4.9/54.5 ± 4.8	1, 4, 7	0/0
Yin[ <mark>59]</mark> , 2018	46/50	-	Calcium Dobesilate	C + QM	12 wk	55.27 ± 5.42/54.63 ± 5.28	1, 2, 7	0/0
Xin <i>et al</i> [ <mark>60]</mark> , 2019	38/38	-	Epalrestat	C + QM	12 wk	55.1 ± 3.3/55.5 ± 3.2	1, 3, 7	0/0
Zhang <mark>[61</mark> ], 2017	39/39	-	Calcium Dobesilate	C + QM	12 wk	56.8 ± 2.5/56.9 ± 2.1	1, 2, 5, 6, 7	0/0
Yue <mark>[62</mark> ], 2016	38/57	-	СТ	C + QM	24 wk	49.82 ± 6.17/50.67 ± 5.23	1	-
Zheng <i>et al</i> [ <mark>63</mark> ], 2014	15/15	-	СТ	C + QM	24 wk	50.4 ± 3.1/55.2 ± 4.7	1	-
Yang <i>et al</i> [ <b>64</b> ], 2013	36/35	-	СТ	C + QM	24 wk	50.94 ± 8.01/50.23 ± 7.15	1	-
Huang[ <mark>65</mark> ], 2020	20/20	-	Calcium Dobesilate	C + DS	16 wk	52.16 ± 2.45/53.16 ± 2.26	1, 2, 3, 5, 6	-
Zheng and Ji[ <mark>66</mark> ], 2021	43/44	-	СТ	C + DS	8 wk	57.52 ± 6.41/58.21 ± 6.35	1, 4	-
Meng <i>et al</i> [67], 2011	28/30	-	CT + Calcium Dobesilate	C + DS	24 wk	51.20 ± 7.90/50.60 ± 8.70	1,7	0/0



Wang[ <mark>68</mark> ], 2004	16/28	-	CT + Calcium Dobesilate	C + DS	16 wk	50.5 ± 9.36/50.4 ± 8.70	1	-
Zhao[ <mark>69</mark> ], 2019	53/53	TRD	CT	C + DS	8 wk	57.5 ± 14.8/56.8 ± 13.4	1, 3, 4, 7	0/0
Ma et al <mark>[70]</mark> , 2016	34/48	-	СТ	C + DS	24 wk	59.16 ± 9.73/59.01 ± 10.58	5, 6, 7	0/0
Chen <i>et al</i> [71], 2007	25/25	-	CT	C + DS	8 wk	60.56/62.42	2, 3	-
Xu <b>[72]</b> , 2019	43/43	TRD	Calcium Dobesilate	C + DS	16 wk	53.06 ± 4.39/53.11 ± 4.41	1, 3, 7	0/0
Huang et al [73], 2021	45/45	TRD	CT + Calcium Dobesilate	C + DS	24 wk	67.3 ± 5.1/67.5 ± 5.3	2, 4, 7	3/4
Jiao[ <mark>74</mark> ], 2018	75/75	-	CT + Calcium Dobesilate	C + DS	8 wk	56.31 ± 2.19/56.24 ± 3.86	1	-
Li[ <mark>75</mark> ], 2017	89/89	-	CT + Calcium Dobesilate	C + DS	8 wk	55.8 ± 6.8/56.5 ± 7.2	1, 5, 6	-
Yan and Yuan[ <mark>76]</mark> , 2014	20/60	-	СТ	C + DS	24 wk	68.8/65.6	2, 3	-
Zhou[ <mark>77</mark> ], 2008	18/28	-	Calcium Dobesilate	C + DS	24 wk	50.50 ± 9.36/50.40 ± 8.70	1	-
Miao[ <mark>78]</mark> , 2020	24/34	TRD	Calcium Dobesilate	C + DS	16 wk	57.46 ± 4.41/57.33 ± 4.26	2, 3	-
Ruan <i>et al</i> [79], 2017	35/35	-	Calcium Dobesilate	C + DS	16 wk	52.8 ± 1.7/52.5 ± 1.1	1, 3, 5, 6, 7	0/0
Yin <i>et al</i> [80], 2013	50/50	-	CT + Calcium Dobesilate	C + DS	8 wk	59.7/57.9	1, 3	-
Zhu[ <mark>81</mark> ], 2018	57/57	-	СТ	C + DS	12 wk	64.12 ± 1.36/64.17 ± 1.38	1, 2, 3	-
Yang <i>et al</i> [ <mark>82</mark> ], 2013	32/33	-	СТ	C + DS	8 wk	54.2 ± 10.8/55.4 ± 12.1	5, 6	-
Bai <mark>[83</mark> ], 2017	38/38	TRD	CT + Calcium Dobesilate	C + DS	16 wk	-	1, 2, 3, 4, 7	0/0
Guo[ <mark>84</mark> ], 2015	35/100	-	СТ	C + DS	24 wk	59.6 ± 9.7/59.0 ± 10.6	5, 6	-
Liu[ <mark>85</mark> ], 2018	89/89	TRD	СТ	C + DS	4 wk	$54.97 \pm 4.88/55.02 \pm 5.01$	5, 6	-
Liu[ <mark>86</mark> ], 2019	41/42	-	Pancreatic Kinino- genase	C + HXMM	12 wk	-	3,7	2/2
Zhao and Liu[ <mark>87]</mark> , 2021	30/30	-	CT + Calcium Dobesilate	C + HXMM	12 wk	64.84 ± 4.26/65.09 ± 4.37	1,7	4/2
Gao et al [ <mark>88</mark> ], 2020	128/128	TRD	CT + Calcium Dobesilate	C + HXMM	12 wk	57.65 ± 7.82/58.14 ± 7.63	1, 3, 7	4/3
Li[ <mark>89</mark> ], 2021	34/34	Lottery	CT + Calcium Dobesilate	C + HXMM	12 wk	68.49 ± 4.62/67.84 ± 4.57	1, 3	-
Wang et al [90], 2018	100/100	-	Pancreatic Kinino- genase	C + HXMM	12 wk	$50.62 \pm 6.91/50.96 \pm 6.71$	1, 3, 7	4/2
Zhang and Wang <mark>[91]</mark> , 2018	39/39	-	СТ	C + HXMM	12 wk	-	1, 2	-
Ye <i>et al</i> [ <mark>92]</mark> , 2019	88/88	TRD	Calcium Dobesilate	C + HXMM	12 wk	60.9 ± 13.4/60.5 ± 13.4	1, 5, 6	-
Yu[ <mark>93</mark> ], 2019	30/30	-	Epalrestat	C + HXMM	12 wk	-	1,7	2/3
Du <i>et al</i> [ <mark>94</mark> ], 2015	26/25	TRD	СТ	C + HXMM	12 wk	53.39 ± 4.96/52.13 ± 5.01	1,7	0/0



Song[ <mark>95</mark> ], 2013	40/40	-	СТ	C + QJDH	12 wk	-	1,7	0/0
Li <mark>[96</mark> ], 2019	40/40	-	Mecobalamin	C + QJDH	24 wk	61.25 ± 6.75/60.85 ± 6.57	1, 3, 4, 7	0/0
Li and Wei [ <mark>97</mark> ], 2019	54/54	-	CT + Calcium Dobesilate	C + QJDH	20 wk	56.1 ± 3.7/55.4 ± 3.1	1, 3, 4, 5, 6, 7	0/0
Wu <mark>[98]</mark> , 2018	29/29	TRD	Pancreatic Kinino- genase	C + QJDH	12 wk	52.4 ± 10.8/52.3 ± 9.2	1, 5, 7	0/0
Guan[ <mark>99</mark> ], 2017	40/40	-	CT + Calcium Dobesilate	C + QJDH	24 wk	40.0 ± 3.1/41.1 ± 2.0	1, 5, 6, 7	6/5
Ainu <i>et al</i> [ <mark>100</mark> ], 2019	50/50	-	CT + Calcium Dobesilate	C + QJDH	4 wk	53.02 ± 5.39/52.61 ± 5.39	2, 3, 4	-
Fu[ <mark>101</mark> ], 2019	40/40	-	Calcium Dobesilate	C + SDMM	24 wk	-	5,7	7/2
Ji and Liu [ <mark>102</mark> ], 2022	52/52	Lottery	Calcium Dobesilate	C + SDMM	12 wk	56.53 ± 4.09/56.63 ± 4.02	1, 3, 4, 7	7/5
Jin and Zhang[ <mark>103</mark> ], 2019	71/71	-	CT + Calcium Dobesilate	C + SDMM	16 wk	62.39 ± 8.34/63.07 ± 8.08	1, 7	9/2
Liu <i>et al</i> [ <mark>104</mark> ], 2019	60/60	TRD	CT + Calcium Dobesilate	C + SDMM	16 wk	57.10 ± 9.26/57.54 ± 8.11	1, 3, 4, 7	0/0
Pang[ <mark>105</mark> ], 2015	40/40	-	СТ	C + SDMM	16 wk	49.6 ± 5.3/49.4 ± 5.7	1	-
Deng <i>et al</i> [ <mark>106</mark> ], 2018	40/40	TRD	Calcium Dobesilate	C + DGBX	12 wk	59.48 ± 8.22/59.62 ± 8.30	1, 3, 7	0/0
Wang and Chen[107], 2020	75/75	-	Calcium Dobesilate	C + DGBX	12 wk	62.38 ± 2.00/62.40 ± 2.02	1,4	-
Wu[ <mark>108</mark> ], 2013	33/34	-	СТ	C + DGBX	12 wk	-	1, 2, 5, 6, 7	0/0
Sun <i>et al</i> [ <mark>109</mark> ], 2019	90/92	-	CT	C + DGBX	12 wk	55.3 ± 3.7/55.9 ± 3.5	1, 3, 7	10/6
Yu <mark>[110</mark> ], 2020	28/28	-	CT + Calcium Dobesilate	C + DGBX	12 wk	$60.01 \pm 8.26/60.48 \pm 8.11$	1, 2	-
Xu[ <mark>111</mark> ], 2018	43/43	TRD	СТ	C + DGBX	12 wk	56.2 ± 7.3/56.6 ± 7.1	1,7	4/0
Huang [ <mark>112</mark> ], 2017	54/54	TRD	Iodized Lecithin	C + XFZY	8 wk	62.15 ± 11.80/61.84 ± 12.11	2	-
Zhu[ <mark>113</mark> ], 2010	36/36	TRD	СТ	C + XFZY	8 wk	55.42 ± 5.35/55.73 ± 5.10	1, 5, 6, 7	0/0
Xiong and Chen[ <mark>114</mark> ], 2019	43/43	TRD	Calcium Dobesilate	C + XFZY	16 wk	53.14 ± 7.25/54.28 ± 7.13	2, 5, 6	-
Liu[ <mark>115</mark> ], 2020	57/57	Lottery	СТ	C + XFZY	16 wk	50.38 ± 3.67/50.47 ± 3.22	2	-
Gong et al [ <mark>116</mark> ], 2014	40/40	-	СТ	C + XFZY	12 wk	57.24 ± 10.60/55.36 ± 9.28	1,5	-
Hao[ <mark>117</mark> ], 2012	84/66	-	СТ	C + XFZY	4 wk	-	1	-
Huang [ <mark>118</mark> ], 2004	50/70	-	СТ	C + XFZY	12 wk	64.5/65	1	-
Zhang[ <mark>119</mark> ], 2019	46/46	-	СТ	C + BYHW	8 wk	64.37 ± 9.13/66.46 ± 9.90	1	-
Yang[ <mark>120]</mark> , 2019	40/40	TRD	CT + Calcium Dobesilate	C + BYHW	8 wk	48.66 ± 8.82/50.35 ± 9.06	1, 2, 5	-
Tian[ <mark>121</mark> ],	27/27	-	Metformin	C + BYHW	12 wk	53.05 ±	1	-

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2019						1.18/53.12±1.12		
Qu and Yao 3 [122], 2009	30/32	-	CT + Iodized Lecithin	C + BYHW	12 wk	-	1	-

A total of 6 studies[95-100] used Qiju Dihuang Pills (QJDH) (consisting of Lycii Fructus/Gouqizi in chinese; Rehmanniae Radix Praeparata/Dihuang in chinese; Chrysanthemi Flos/Juhua in chinese; Cornus Officinalis Sieb. Et Zucc./Shanzhuyu in chinese; Rhizoma Dioscoreae/Shanyao in chinese; Poria Cocos(Schw.) Wolf./Fuling in chinese; Alisma Orientale (Sam.) Juz./Zexie in chinese; Cortex Moutan/Mudanpi in chinese). Four[95-97,100] of these six studies added Paeoniae Radix Alba (Baishao in Chinese), Tribulifructus (Jili in Chinese), Cassiae Semen (Juemingzi in Chinese), and Angelicae Sinensis Radix (Danggui in Chinese). Although there are some changes in components, the whole is still dominated by QJDH, which has the same efficacy as QJDH, so in this study, we classified these 6 studies as using QJDH. The composition of all Oral Chinese medicine (OCM) in this study is shown in Supplementary Table 2, and the source of OCMs, quality control report and chemical analysis report are shown in Supplementary Table 3. TRD: Table of random digit; CT: Conventional treatment, including blood glucose control, blood pressure lowering, lipid regulation and other conventional treatments; E: Experimental group; C: Control group; QM: Qiming Granules; DS: Compound Danshen Dripping Pills; XFZY: Xuefu Zhuyu Decoction; DGBX: Danggui Buxue Decoction; HXMM: Hexue Mingmu Tablets; SDMM: Shuangdan Mingmu Capsules; BYHW: Buyang Huanwu Decoction; QJDH: Qiju Dihuang Pills; XST: Compound Xueshuantong Capsule; -: Not mentioned; 1: Total effective rate; 2: Visual acuity; 3: Fundus hemorrhage area; 4: Vascular endothelial growth factor; 5: Fasting blood glucose; 6: Glycated hemoglobin; 7: Adverse reactions.



Figure 1 The PRISMA flow diagram for search and selection processes of the meta- analysis. PRISMA: Preferred Reporting Items for Systematic Reviews and Meta- Analyses; RCT: Randomized controlled trials.

are shown in Supplementary Tables 2 and 3, respectively.

#### Risk of bias evaluation results

The quality of the included literature was evaluated using the "Risk Assessment Tool" recommended by the Cochrane Collaboration: 33 studies[16,19,22,28,29,35,37,39,43,45,49,51,55,69,72,73,78,83,85,88,89,92,94,98,102,104,106,111-115,120] mentioned the specific randomization method used and therefore assessed as "Low risk". The other 74 studies only mentioned the randomized grouping without mentioning the specific method used for allocation and were therefore evaluated as "Unclear risk". None of the included studies mentioned allocation concealment and blinding, and were evaluated as "Unclear risk". All studies had clear outcome indicators and were evaluated as "Low risk"; no duplicate publications or published biases were found in any of the studies and were evaluated as "Low risk"; other biases were unknown and were evaluated as "Unclear risk". All data were reported completely and were comparable between groups (Figure 2 and Supplementary Figure 1).

#### Results of reticulated meta-analysis

Mesh relationship diagram and consistency testing: The reticulation between the nine included OCM is shown in Figure 3. The total number of arms in the 107 papers totals 214. Lines between nodes indicate direct comparative evidence



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#### Figure 3 Reticulation of diabetic retinopathy treated with socio-comportemental and medical indicator. CT: Conventional western medicine treatment.

between the two interventions, no lines indicate no direct comparison, indirect comparisons can be made through reticulated Meta-analysis. The thickness of the line represents the number of included studies comparing each treatment, and the circular area represents the sample size of the population using the measure. All interventions involved in this study did not form a closed loop and did not require consistency testing.

Total effective rate: A total of 89 studies [16-20,22-25,27-31,33-37,39-69,72,74,75,77,79-81,83,87-99,102-111,113,116-122] were included to compare the total effective rate after OCM + CT treatment. The evidence diagram is shown in Figure 4A. The difference was statistically significant (P < 0.05) for the total effective rate for these nine OCM + CT treatments compared with CT (Figure 5A). The nine OCMs + CT total effective rate in descending order are: DS (SUCRA = 75.0%) > DGBX (SUCRA = 72.8%) > SDMM (SUCRA = 71.4%) > QM (SUCRA = 66.4%) > QJDH (SUCRA = 62.4%) > XST = HXMM (SUCRA = 52.5%) > BYHW (SUCRA = 26.8%) > XFZY (SUCRA = 20.1%) > CT (SUCRA = 0.0%). The probability ranking is shown in Figure 6A.

Visual acuity: A total of 34 studies [16-18,21,23,25,26,29,37,38,44,46,47,49,51,55,56,59,60,65,71,73,76,78,81,83,91,100,108,110, 112,114,115,120] with 8 OCMs were included to compare visual acuity after OCM + CT treatment. The evidence diagram is shown in Figure 4B. In terms of visual acuity, supplemental XST, QM, DS, HXMM, QJDH, DGBX and XFZY treatments were statistically significant compared with CT treatment alone (P < 0.05) (Figure 5B). The eight OCMs + CT in order of highest to lowest improvement in visual acuity are: HXMM (SUCRA = 76.3%) > XST (SUCRA = 67.4%) > DS (SUCRA = 67.2%) > DGBX (SUCRA = 64.8%) > BYHW (SUCRA = 50.3%) > XFZY (SUCRA = 44.4%) > QM (SUCRA = 40.1%) > QJDH (SUCRA = 36.2%) > CT (SUCRA = 3.4%). The probability ranking is shown in Figure 6B.

VEGF: A total of 18 studies [16,25,29,31,33,35,55,58,66,69,73,83,96,97,100,102,104,107] with 6 OCMs were included to compare VEGF after OCM + CT treatment. The evidence diagram is shown in Figure 4C. In terms of VEGF, supplemental XST, DS, QJDH and SDMM were statistically significant compared with CT treatment alone (P < 0.05) (Figure 5C). The seven OCMs + CT in order of VEGF reduction from highest to lowest are: SDMM (SUCRA = 97.7%) > DS (SUCRA = 74.4%) > QJDH (SUCRA = 57.9%) > DGBX (SUCRA = 45.9%) > XST (SUCRA = 44.8%) > QM (SUCRA = 24.7%) > CT

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Figure 4 Network diagrams of comparisons on different outcomes of treatments in different groups (Oral Chinese medicines + conventional western medicine treatment) of patients with diabetic retinopathy. A: Total effective rate; B: Visual acuity; C: Vascular endothelial growth factor; D: Fundus hemorrhage area; E: Fasting blood glucose; F: Glycated hemoglobin. CT: Conventional western medicine treatment.

(SUCRA = 4.7%). The probability ranking is shown in Figure 6C.

**FHA:** A total of 32 studies[16,20,21,31,32,35,37,39-41,60,65,69,71,72,76,78-81,83,86,88-90,96,97,100,102,104,106,109] with 7 OCMs were included to compare FHA after OCM + CT treatment. The evidence diagram is shown in Figure 4D. These seven OCM + CT were statistically significant (P < 0.05) compared with CT for FHA, respectively (Figure 5D). The seven OCMs + CT reduce FHA in the following order from highest to lowest: DGBX (SUCRA = 87.5%) > SDMM (SUCRA =

Α									
Compound									
Xueshuanton									
Capsules									
0.90	Qiming	]							
(0.62,1.29)	Granules								
0.82 (0.52,1.30)	0.92 (0.59,1.44)	Compound Danshen Dripping Pills		_					
1.01 (0.63,1.64)	1.13 (0.71,1.81)	1.23 (0.71,2.12)	Hexue Mingmu Tablets						
0.91	1.02	1.11	0.90	Qiju					
(0.44,1.90)	(0.49,2.11)	(0.51,2.42)	(0.41,1.98)	<b>Dihuang Pills</b>					
0.82 (0.38,1.77)	0.91 (0.43,1.97)	0.99 (0.44,2.24)	0.81 (0.35,1.84)	0.90 (0.33,2.42)	Shuangdan Mingmu Capsules				
0.84	0.94	1.02	0.83	0.92	1.02	Danggui			
(0.52.1.36)	(0.58.1.51)	(0.59.1.77)	(0.47.1.46)	(0.41.2.03)	(0.45.2.34)	Buxue			
(0.52,1.50)	(0.56,1.51)	(0.55,1.77)	(0.47,1.40)	(0.41,2.03)	(0.45,2.54)	Decoction			
1.57 (0.89,2.78)	1.75 (1.00,3.08)	1.91 (1.02,3.57)	1.55 (0.81,2.94)	1.72 (0.74,4.01)	1.92 (0.79,4.63)	1.87 (0.98,3.58)	Xuefu Zhuyu Decoction		
1.43 (0.74,2.78)	1.60 (0.83,3.09)	1.74 (0.85,3.55)	1.41 (0.68,2.92)	1.57 (0.63,3.91)	1.75 (0.68,4.50)	1.71 (0.82,3.55)	0.91 (0.41,2.01)	Buyang Huanwu Decoction	
3.69 (2.83,4.80)	4.11 (3.20,5.30)	4.48 (3.08,6.50)	3.63 (2.44,5.40)	4.03 (2.04,7.98)	4.50 (2.18,9.27)	4.40 (2.93,6.60)	2.35 (1.42,3.89)	2.57 (1.40,4.72)	СТ

В

(	Compound								
Xu	ieshuantong								
	Capsules								
(	1.05 (0.97,1.14)	Qiming Granules							
	1.00 (0.91,1.10)	0.95 (0.87,1.04)	Compound Danshen Dripping Pills						
(	0.96 (0.78,1.17)	0.91 (0.74,1.12)	0.96 (0.77,1.18)	Hexue Mingmu Tablets					
	1.08 (0.90,1.29)	1.02 (0.85,1.23)	1.08 (0.89,1.30)	1.13 (0.87,1.47)	Qiju Dihuang Pills				
	1.00 (0.86,1.16)	0.95 (0.82,1.11)	1.00 (0.85,1.17)	1.05 (0.82,1.33)	0.93 (0.74,1.16)	Danggui Buxue Decoction			
	1.05 (0.93,1.18)	1.00 (0.89,1.12)	1.05 (0.92,1.19)	1.10 (0.88,1.37)	0.97 (0.80,1.19)	1.05 (0.88,1.25)	Xuefu Zhuyu Decoction		
	1.03 (0.85,1.25)	0.98 (0.81,1.19)	1.04 (0.85,1.26)	1.08 (0.83,1.42)	0.96 (0.75,1.24)	1.04 (0.82,1.30)	0.99 (0.80,1.22)	Buyang Huanwu Decoction	
(	1.18 (1.11,1.25)	1.12 (1.06,1.19)	1.18 (1.10,1.27)	1.23 (1.01,1.50)	1.09 (0.92,1.30)	1.18 (1.02,1.36)	1.13 (1.02,1.25)	1.14 (0.95,1.37)	СТ

## С

Compound						
Xueshuantong						
Capsules						
0.51	Qiming					
(0.10,2.72)	Granules					
2.04 (0.74,5.61)	3.97 (0.70,22.41)	Compound Danshen DrippingPills				
1.36 (0.45,4.13)	2.66 (0.44,15.90)	0.67 (0.20,2.23)	Qiju Dihuang Pills			
5.69 (1.57,20.65)	11.09 (1.65,74.54)	2.80 (0.71,11.00)	4.17 (0.99,17.62)	Shuangdan Mingmu Capsule		
0.98 (0.19,5.10)	1.90 (0.22,16.65)	0.48 (0.09,2.67)	0.72 (0.12,4.22)	0.17 (0.03,1.14)	Danggui Buxue Decoction	
0.28 (0.15,0.54)	0.55 (0.12,2.58)	0.14 (0.06,0.31)	0.21 (0.08,0.51)	0.05 (0.02,0.15)	0.29 (0.06,1.33)	СТ



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D							
Compound Xueshuantong Capsules							
0.84 (0.52,1.38)	Qiming Granules						
0.88 (0.71,1.10)	1.05 (0.64,1.71)	Compound Danshen Dripping Pills					
0.64 (0.49,0.85)	0.76 (0.45,1.29)	0.730 (0.55,0.97)	Hexue Mingmu Tablets				
0.95 (0.69,1.30)	1.12 (0.66,1.93)	1.07 (0.78,1.47)	1.47 (1.03,2.11)	Qiju Dihuang Pills			
1.09 (0.75,1.57)	1.29 (0.73,2.29)	1.23 (0.85,1.78)	1.69 (1.12,2.55)	1.15 (0.75,1.77)	Shuangdan Mingmu Capsules		
1.18 (0.82,1.70)	1.40 (0.80,2.48)	1.34 (0.93,1.93)	1.84 (1.23,2.75)	1.25 (0.82,1.91)	1.09 (0.68,1.74)	Danggui Buxue Decoction	
0.44 (0.38,0.51)	0.52 (0.33,0.83)	0.50 (0.42,0.58)	0.68 (0.54,0.87)	0.46 (0.35,0.61)	0.40 (0.29,0.57)	0.37 (0.27,0.52)	СТ

Ε

Qiming Granules								
1.39 (0.77,2.48)	Compound Danshen Dripping Pills							
3.48 (1.17,10.33)	2.51 (0.90,6.98)	Hexue Mingmu Tablets						
1.93 (0.95,3.95)	1.39 (0.75,2.59)	0.56 (0.18,1.68)	Qiju Dihuang Pills					
2.99 (0.97,9.19)	2.16 (0.75,6.23)	0.86 (0.21,3.50)	1.55 (0.50,4.84)	Shuangdan Mingmu Capsules				
1.06 (0.41,2.75)	0.76 (0.32,1.84)	0.30 (0.09,1.09)	0.55 (0.21,1.45)	0.35 (0.10,1.30)	Danggui Buxue Decoction			
1.10 (0.56,2.19)	0.80 (0.45,1.42)	0.32 (0.11,0.94)	0.57 (0.28,1.16)	0.37 (0.12,1.13)	1.04 (0.40,2.70)	Xuefu Zhuyu Decoction		
1.06 (0.37,3.02)	0.76 (0.29,2.04)	0.30 (0.08,1.17)	0.55 (0.19,1.59)	0.35 (0.09,1.40)	1.00 (0.29,3.46)	0.96 (0.34,2.73)	Buyang Huanwu Decoction	
0.86 (0.53,1.40)	0.62 (0.45,0.85)	0.25 (0.09,0.65)	0.44 (0.26,0.75)	0.29 (0.10,0.79)	0.81 (0.36,1.84)	0.78 (0.48,1.26)	0.81 (0.32,2.05)	СТ

F

Qiming Granules						
0.97 (0.48,1.96)	Compound Danshen Dripping Pills		_			
1.43 (0.45,4.56)	1.47 (0.50,4.35)	Hexue Mingmu Tablets				
2.77 (1.07,7.15)	2.85 (1.21,6.69	1.94 (0.55,6.81)	Qiju Dihuang Pills			
0.96 (0.29,3.11)	0.99 (0.33,2.97)	0.67 (0.16,2.83)	0.35 (0.10,1.24)	Danggui Buxue Decoction		
1.08 (0.42,2.78)	1.11 (0.48,2.60)	0.76 (0.22,2.65)	0.39 (0.14,1.12)	1.13 (0.32,4.02)	Xuefu Zhuyu Decoction	
0.64 (0.36,1.14)	0.66 (0.44,0.99)	0.45 (0.16,1.23)	0.23 (0.11,0.49)	0.67 (0.24,1.87)	0.59 (0.28,1.25)	ст

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Figure 5 Pooled estimates of the network meta-analysis. A: Pooled risk d ratios (95% credible intervals) for the total effective rate; B: Pooled risk d ratios (95% credible intervals) for visual acuity; C: Pooled risk d ratios (95% credible intervals) for vascular endothelial growth factor; D: Pooled risk d ratios (95% credible intervals) for fundus hemorrhage area; E: Pooled risk d ratios (95% credible intervals) for fasting blood glucose; F: Pooled risk d ratios (95% credible intervals) for glycated hemoglobin. CT: Conventional western medicine treatment.

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**Figure 6 Surface under the cumulativeranking curve for outcomes.** A: Total effective rate; B: Visual acuity; C: Vascular endothelial growth factor; D: Fundus hemorrhage area; E: Fasting blood glucose; F: Glycated hemoglobin. CT: Conventional western medicine treatment; XST: Xueshuantong Capsules; QM: Qiming Granules; DS: Compound Danshen Dripping Pills; HXMM: Hexue Mingmu Tablets; QJDH: Qiju Dihuang Pills; SDMM: Shuangdan Mingmu Capsules; DGBX: Danggui Buxue Decoction; XFZY: Xuefu Zhuyu Decoction; BYHW: Buyang Huanwu Decoction.

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77.7%) > XST (SUCRA = 67.9%) > QJDH (SUCRA = 58.8%) > DS (SUCRA = 45.9%) > QM (SUCRA = 45.1%) > HXMM (SUCRA = 17.0%) > CT (SUCRA = 0.0%). The probability ranking is shown in Figure 6D.

**FBG:** A total of 21 studies[47,49,51,61,65,70,75,79,82,84,85,92,97-99,101,108,113,114,116,120] with 8 OCMs were included to compare FBG after OCM + CT treatment. The evidence diagram is shown in Figure 4E. In terms of FBG, supplemental DS, HXMM, QJDH and SDMM treatments were statistically significant compared with CT treatment alone (P < 0.05) (Figure 5E). The eight OCMs + CT lower FBG in the following order from highest to lowest: HXMM (SUCRA = 91.1%) > SDMM (SUCRA = 86.4%) > QJDH (SUCRA = 73.8%) > DS (SUCRA = 54.2%) > XFZY (SUCRA = 36.1%) > BYHW (SUCRA = 34.0%) > DGBX (SUCRA = 33.2%) > QM (SUCRA = 27.8%) > CT (SUCRA = 13.5%). The probability ranking is shown in Figure 6E.

**HbA1c:** A total of 17 studies[47,49,51,61,65,70,75,79,82,84,85,92,97,99,108,113,114] with 6 OCMs were included to compare HbA1c after OCM + CT treatment. The evidence diagram is shown in Figure 4F. In terms of HbA1c, supplemental DS and QJDH treatments were statistically significant compared with CT treatment alone (P < 0.05) (Figure 5F). The six OCMs + CT lower HbA1c in the following order from highest to lowest: QJDH (SUCRA = 95.5%) > HXMM (SUCRA = 65.3%) > XFZY (SUCRA = 50.8%) > QM (SUCRA = 45.0%) > DS (SUCRA = 43.6%) > DGBX (SUCRA = 42.2%) > CT (SUCRA = 7.6%). The probability ranking is shown in Figure 6F.

**Cluster analysis and meta-analysis two-by-two comparison results:** The key outcome indicators included in this study were cluster analyzed to derive the intervention of different OCMs + CT for two outcome indicators at the same time. In terms of total effective rate and visual acuity, HXMM, XST, DS and DGBX in the upper right corner of Figure 7A performed better; in terms of total effective rate and VEGF, SDMM and DS performed better (Figure 7B); in terms of total effective rate and FHA, DGBX and SDMM in the upper right corner of Figure 7C performed better; in terms of FBG and HbA1c, QJDH and HXMM in the upper right corner of Figure 7D performed better.

A two-by-two comparison of the nine OCMs + CT and CT was performed at each index. The total clinical efficiency of DR treatment with all 9 OCMs + CT was found to be higher than that of CT alone; CT in combination with XST, QM, DS, HXMM, DGBX and XFZY were superior to CT alone in improving visual acuity, respectively; in anti-VEGF, CT in combination with XST, DS, QJDH and SDMM were better than CT alone, respectively; CT in combination with XST, QM, DS, HXMM, QJDH, SDMM and DGBX were superior to CT alone in reducing FHA, respectively; CT in combination with DS, HXMM, QJDH and SDMM were superior to CT alone in lowering FBG, respectively; CT in combination with DS and QJDH were superior to CT alone in lowering HbA1c, respectively. All results are plotted as forest plots shown in Supplementary Figure 2.

#### Sensitivity analysis

To verify the stability of the above results, we performed NMA with sample size and duration of treatment as sensitivity factors that may affect the results. Of the total 107 studies, 54 studies[16-18,20,21,23,27,31-36,38,40,42,44,46,52,54,60,61,63-68,71,76-79,82,83,87,89,91,93-96,98,99,101,105,106,108,110,113,116,120-122] with a case load of no more than 80 were included in the sensitivity analysis. The results revealed that there was no significant difference in total effective rate for CT + SDMM compared with CT (OR: 0.94, 95%CI: 0.94-23.98, P > 0.05), unlike the original NMA; for HbA1c, DGBX + CT was effective in reducing HbA1c compared with CT (OR: 0.67, 95%CI: 0.48-0.94, P < 0.05), unlike the original NMA. There was no significant change in the remaining indicators, so the sample size was considered as a possible factor influencing the results (Supplementary Figure 3). When using duration of treatment as a sensitivity factor, we divided all studies according to 12 wk of treatment, and a total of 65 studies[16,19,22,24,26-31,33,34,37,38,41-45,47-49,51,56,58-61,66,69,71,72,75,80-82,85-95,98,100,102,106-113,116-122] with no more than 12 wk of treatment were included in the sensitivity analysis. Results found that DGBX + CT was more effective than CT in terms of VEGF (OR: 0.29, 95%CI: 0.39-1.16, P > 0.05), unlike the original NMA; for HbA1c, the effect of DS + CT was not significantly different from CT (OR: 0.67, 95%CI: 0.39-1.16, P > 0.05), unlike the original NMA; for HbA1c, the effect of DS + CT was not significantly different from CT (OR: 0.77, 95%CI: 0.38-1.57, P > 0.05), unlike the original NMA. Consideration of sample size may have influenced this result (Supplementary Figure 4).

#### The small sample effect and publication bias

The total effective rate (significantly effective + effective), visual acuity, VEGF, FHA, FBG and HbA1c were used as evaluation indicators to produce a comparative corrected funnel plot of the study to assess the small sample effect, see Figure 8. The results showed that the total effective rate and FBG comparison corrected funnel plots showed basic symmetry, with studies roughly symmetrically distributed on both sides of the midline, suggesting that a small sample effect is less likely. The poor symmetry of the corrected funnel plot for visual acuity, VEGF, FHA, and HbA1c comparisons suggests the possibility of a small sample effect. The reasons may be related to the mixed quality of included studies, small sample size, inconsistent treatment regimens, and different pathological stages of study subjects. Further analysis of publication bias using Begg's and Egger's tests revealed the presence of publication bias for total clinical effect-iveness ( $P_{Begg} < 0.001$ ,  $P_{Egger} < 0.001$ ). Additionally, there was a potential publication bias for FBG ( $P_{Begg} = 0.01$ ;  $P_{Egger} = 0.151$ ) and FHA ( $P_{Begg} = 0.224$ ,  $P_{Egger} = 0.041$ ). However, no publication bias was observed for visual acuity, VEGF, and HbA1c ( $P_{Begg} > 0.05$ ;  $P_{Egger} > 0.05$ ). The presence of bias in these particular outcomes may be attributed to various factors, including the mixed quality of the included studies, small sample sizes, variations in treatment regimens, and differences in the pathological stages of the study subjects. These factors could contribute to heterogeneity and potential reporting biases within the literature.



Figure 7 Cluster analysis for outcome indicators. A: Total effective rate and vision; B: Total effective rate and VEGF; C: Total effective rate and fundus hemorrhage area; D: Fasting blood glucose and HbA1c. CT: Conventional western medicine treatment; XST: Xueshuantong Capsules; QM: Qiming Granules; DS: Compound Danshen Dripping Pills; HXMM: Hexue Mingmu Tablets; QJDH: Qiju Dihuang Pills; SDMM: Shuangdan Mingmu Capsules; DGBX: Danggui Buxue Decoction; XFZY: Xuefu Zhuyu Decoction; BYHW: Buyang Huanwu Decoction.

#### Adverse reactions

Of the total 107 RCTs, 47 mentioned adverse reactions, but only 20 of these studies[17,19,33,37,39,43,51,55,73,86-88,90,93, 99,101-103,109,111] had patients with adverse reactions, and the other 26 studies[30,34,35,40,46,47,57-61,67-70,72,79,83,94-98,104,106,108,113] in which all patients had no adverse reactions. A total of 76 patients in the experimental group had adverse reactions during the treatment, and 127 patients in the control group had adverse reactions. A total of 7 OCMs were involved, and the results are shown in Table 2.

# DISCUSSION

To the best of our knowledge, this study is the first NMA focused on the combination of OCM with conventional Western medicine for the treatment of DR, and is the NMA with the largest number of original studies included, the largest number of subjects, and the largest variety of OCM included to date. Only five NMAs[123-127] have reported the therapeutic effect on DR, and only one[125] of them is about the efficacy of herbal injections for DR. This study comprehensively collected RCTs involving 9 commonly used OCMs in China and included 6 clinical indicators commonly used to evaluate the efficacy of DR, the largest number of included indicators we are aware of to date in a similar report.

Although surgical treatment and intravitreal drug injection for DR are becoming popular, there are still some problems that are difficult to solve in a short period of time. First, invasive therapies have a limited audience. Total retinal photocoagulation is the primary treatment for proliferative DR and is not advocated for the treatment of non-proliferative DR; intraocular injections of VEGF inhibitors are more effective primarily in the treatment of DR with macular edema. In some forms of non-clearing vitreous hemorrhage, vitrectomy has been shown to remain the only method for removing fibrous proliferation and relieving traction detachment, with mixed results[128]. Second, safety issues need to be kept in mind, for example, one study found that although intraocular steroid injections led to rapid regression of dimethyl ether, however, this improvement did not persist and was associated with a significant increase in the incidence of elevated intraocular pressure and cataracts[129]. Furthermore, we must take into account the additional financial burden incurred by this type of treatment. A study from Canada reported that Grid laser therapy adds an additional cost benefit per

Table 2 Occurrence of adverse reactions of Oral Chinese medicine															
OCMs compound	Group	Sample	Hypoglycemia	Stomach upset ( <i>n</i> )	Loss of appetite	Insomnia	Fever	Dizzy	Nausea	Liver damage	Kidney damage	Macular edema	Corneal damage	ltchy skin	Fatigue
Xueshuantong Capsules	С	42	0	14	5	0	0	0	6	8	9	0	0	0	0
	Е	28	0	8	8	0	0	0	8	2	2	0	0	0	0
Qiming Granules	С	23	0	1	0	0	0	0	0	3	2	8	7	1	1
	Е	12	0	2	0	0	0	0	0	1	0	3	3	1	2
Compound Danshen Dripping Pills	С	3	0	3	0	0	0	0	0	0	0	0	0	0	0
	Е	4	0	4	0	0	0	0	0	0	0	0	0	0	0
Hexue Mingmu Tablets	С	16	2	4	3	0	0	0	3	0	0	0	0	3	1
	Е	12	0	6	2	0	0	0	1	0	0	0	0	2	1
Qiju Dihuang Pills	С	6	0	0	0	0	0	0	6	0	0	0	0	0	0
	Е	5	0	0	0	0	0	0	5	0	0	0	0	0	0
Shuangdan Mingmu Capsules	С	23	0	5	0	0	5	2	8	0	0	0	0	0	3
	Е	9	0	2	0	0	2	2	3	0	0	0	0	0	0
Danggui Buxue Decoction	С	14	2	4	0	2	0	4	0	0	0	0	2	0	0
	Е	6	2	2	0	0	0	0	1	0	0	0	0	0	1

OCM: Oral Chinese medicine; E: Experimental group; C: Control group; The numbers in the table represent the number of patient cases.

quality of life adjusted year[130]. In contrast, OCM is not only increasingly proving to have surprising clinical efficacy [131], is affordable[132], and is indicated for patients in almost all stages of DR with a broad universal indication. Therefore, a systematic and comprehensive evaluation of the therapeutic efficacy of OCM for DR is essential.

This study found that DS showed excellent efficacy in improving visual acuity levels and total clinical effectiveness. DS is composed of three herbs *Radix Salviae* (Danshen in Chinese), *Panax notoginseng (Burkill) F. H. Chen ex C. H.* (Sanqi in Chinese) and *borneol* (bingpian in Chinese). According to Chinese medicine, Danshen and Sanqi have the effect of activating blood circulation and dispelling blood stasis, and are commonly used herbs for treating diseases of blood stasis and obstruction; Bingpian is obtained from the stem of *Blumea balsamifera* (*L.*) *DC*. or the leaves of *Cinnamomum camphora* by water steam distillation and recrystallization, and is documented in the famous Chinese medical work *Annotation of Materia Medica* from the Tang Dynasty (about 659 AD) as a treatment for eye diseases. Each of the three herbs has its own characteristics and at the same time exerts the effect of activating blood circulation and removing blood stasis, which is in line with modern pharmacological research. It was found[133] that Tanshinone IIa (the main active component of Danshen) promoted phosphorylation of AMP-activated protein kinase AMPK at T172 in retinal pigment epithelial cells and inhibited monolayer permeability of human retinal epithelial cells under high glucose conditions, similar to that



Figure 8 Risk of bias funnel chart. A: Total effective rate; B: Visual acuity; C: Vascular endothelial growth factor; D: Fundus hemorrhage area; E: Fasting blood glucose; F: Glycated hemoglobin. a: CT: Conventional treatment, including blood glucose control, blood pressure lowering, lipid regulation and other conventional treatments; b: Compound Xueshuantong Capsules; c: Qiming Granules; d: Compound Danshen Dripping Pills; e: Hexue Mingmu Tablets; f: Qiju Dihuang Pills; g: Shuangdan Mingmu Capsules; h: Danggui Buxue Decoction; i: Xuefu Zhuyu Decoction; j: Buyang Huanwu Decoction.

under normal glucose, while apparently preventing co-localization of NF-KB and p300 and inhibiting their binding, thereby reducing ARPE-19 cell monolayer permeability. A study by Fan et al [134] found that Tanshinone IIa significantly downregulated the expression levels of VEGF and intercellular adhesion molecule-1 (ICAM-1) in a dose-dependent manner under HG conditions, probably by mediating proliferation, migration and inhibition of angiogenesis in human retinal endothelial cells. Sanqi and its extracts have anti-inflammatory[135-138], antioxidant, inhibit platelet aggregation, regulate blood glucose[139,140], regulate blood pressure[141-143], improve insulin resistance[144,145], inhibit neuronal apoptosis[146-148] and neuronal protection[149-152]. In particular, ginsenoside Rb1 (the main active compound extracted from the rhizome of Sanqi) has been widely demonstrated to be promising as an antidiabetic and its complications, improving diabetes-related complications by regulating oxidative stress, apoptosis, inflammatory response, enhancing insulin sensitivity, improving leptin resistance, activating the activation of lipocalin signaling pathway, and inhibiting fibronectin expression[153-161]. In addition to showing good clinical treatment rates, DGBX has shown excellent performance in improving the fundus hemorrhage area. According to TCM theory, DGBX is mainly used for treating diseases caused by fatigue and internal injury, blood deficiency and qi weakness. Although there are only two herbs in the formula, Radix Astragali Mongolici (Huangqi in Chinese) and Angelicae Sinensis Radix (Danggui in Chinese), the

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formula is short but powerful and can promote the production of tangible blood from invisible qi, which is a classic OCM with the effect of nourishing Qi and promoting blood circulation. DGBX was found to affect lipid metabolism in the early stages of atherosclerosis in diabetic Goto-Kakizaki rats, and the mechanism may be related to the regulation of intravascular lipid metabolism genes[162]. Astragalus polysaccharides are the main active components of Huangqi, can reduce the levels of tumor necrosis factor- $\alpha$ , ICAM-1, vascular VEGF and p-Akt in the retina of diabetic rats, and affect the Akt-VEGF signaling pathway by anti-inflammatory and reducing the adhesion of leukocytes to the diabetic retina[163], while inhibiting peripapillary cell apoptosis and basement membrane thickening. Dangui and its active ingredients were able to inhibit the VEGF- $\alpha$  pathway to improve the inflammatory response and apoptosis in the retina of diabetic rats, butyrylcholinesterase and $\beta$ -site amyloid precursor protein cleaving enzyme 1 to exert antidiabetic effects[165].

An interesting finding is that SDMM, QJDH and HXMM are all OCMs with the main effect of nourishing the Yin of the liver and kidney, and they all showed good results in lowering glucose. Chinese medicine theory believes that diabetes is a disease with Yin deficiency as the fundamental pathogenesis, and DR develops from diabetes, so the treatment should focus on replenishing Yin. Ligustrum lucidum (Nvzhenzi in Chinese) and Ecliptae Herba (Mohanlian in Chinese) are common components of HXMM and SDMM, which are widely used in China for the treatment of liver-kidney yin deficiency syndrome[166] (A TCM pathological diagnostic pattern caused by the imbalance of yin and yang[167,168], which is closely related to the development of diabetes). They were found to increase insulin sensitivity, enhance the function of islet β-cells INS-1 and β-tc-6[169], reduce retinal oxidative stress levels, repair diabetes-induced abnormal transcriptome[170], improve retinal cell apoptosis[171], inhibit NLRP3 inflammasome and autophagy signaling pathways [172], regulate homocysteine pathway, reduce lipid peroxidation and scavenge free radicals[173], thereby reducing fundus microangiopathy and protecting normal retinal barrier function in diabetic mice. Meanwhile, SDMM was confirmed to be the most effective complementary and alternative drug for inhibiting VEGF in this study, exerting anti-VEGF effects by inhibiting VEGF-induced RF/6A cell tube formation[174], inhibiting NF-κB activity to regulate advanced glycosylation end-product accumulation, oxidative stress and mitochondrial function, and thus improving retinal cell apoptosis by downregulating PKCô, P47phox and ERK1/2[175,176]. QJDH is a very well-known Chinese OCM for the treatment of eye diseases, which is composed of Liuwei Dihuang Pills (an ancient remedy with very good treatment of diabetes and its complications) plus Chrysanthemi Flos and Lycii Fructus, and has been shown to possibly inhibit the development of DR by interfering with multiple biological pathways such as regulation of response to insulin, glucose homeostasis, and angiogenesis[177]. It was found that Lycii Fructus extract and the active ligand taurine dosedependently enhanced cell viability, reduced apoptosis, downregulated caspase-3 protein expression and caspase-3 enzymatic activity, and downregulated mRNA encoding pro-inflammatory mediators of MMP-9 and fibronectin as well as COX-2 and iNOS protein expression in human retinal epithelial cell lines after HG treatment in order to achieve a protective effect on human retinal epithelial cell lines under HG exposure, thereby delaying the progression of DR[178, 179]. Several network pharmacological and experimental validations found that chemical components such as luteolin, kaempferol, beta-sitosterol, and thymol were able to improve apoptosis-related protein expression by regulating the NLRP/NOX4 signaling pathway, downregulate network hub genes of tumor necrosis factor, and other multibiological pathways, inhibit VEGF-induced RF/6A cell tube formation, and slow down the DR process[171,174,180,181], and these chemicals were also found in Chrysanthemi Flos.

It is noteworthy that metabolic diseases, such as hyperglycemia, hyperlipidemia, and hypertension, serve as crucial risk factors contributing to the development of DR[2]. Effective management of these risk factors is imperative in the prevention and treatment of DR. In this regard, incorporating exercise training and neuromuscular electrical stimulation [182,183] can prove to be a valuable strategy. The guidelines established by the American College of Sports Medicine and the American Diabetes Association emphasize the significance of initial guidance from a qualified exercise training for individuals with type 2 diabetes[184]. These guidelines advocate for the implementation of appropriate exercise training to optimize outcomes related to glycemic control, blood pressure, lipid levels, and cardiovascular risk management.

Despite the clear strengths of this study, there are some limitations. First, the quality of the included studies in this study warrants improvement, as most of them were short-term, single-center, and had small sample sizes. Second, clinical effectiveness, being a commonly used measure in clinical practice, may vary slightly in its definition across different randomized controlled trials, leading to a certain degree of heterogeneity in the results. And then, due to the relatively strict inclusion criteria applied in this study, some high-quality individualized TCM treatment studies were excluded, which may limit the comprehensive representation of the characteristics of TCM. Finally, the included literature lacked direct comparisons between different TCMs, and a closed loop was not formed in the evidence network, thereby allowing only indirect comparisons to assess the efficacy advantages and disadvantages of different interventions. Despite these limitations, the present study remains one of the most comprehensive studies available and holds significant clinical reference value.

This study aims to optimize patient outcomes by tailoring treatment strategies to each patient's unique condition. We recommend the utilization of SDMM + CT or HXMM + CT for treatment due to their demonstrated efficacy across multiple indicators. Specifically, HXMM + CT has shown greater effectiveness in improving patients' visual acuity, while SDMM + CT exhibits stronger inhibitory effects on VEGF. Furthermore, DGBX + CT has proven to be more effective in reducing FHA, HXMM + CT excels in reducing FBG, and QJDH + CT demonstrates superior efficacy in reducing HbA1c. Additionally, we suggest combining OCMs with western drugs for the treatment of DR, as this combination has been shown to yield superior outcomes compared to interventions with western drugs alone. Hence, it is crucial to select appropriate treatment methods in clinical practice based on the individual circumstances of patients with DR to attain maximum benefits from combined Chinese and Western medicine interventions.

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# CONCLUSION

This study provides evidence that combining OCMs with western drugs leads to better outcomes in all aspects of DR compared to using western drugs alone. Based on the findings, we highly recommend the use of SDMM or HXMM for the treatment of DR. These two OCMs have demonstrated outstanding efficacy across multiple indicators.

# **ARTICLE HIGHLIGHTS**

## Research background

Diabetic retinopathy (DR) is one of the most important factors in adult blindness, yet rationalized DR treatment protocols are currently not systematically updated.

# Research motivation

Current traditional Chinese medicine treatment options for DR need to be re-evaluated.

# Research objectives

To investigate which complementary alternative treatment with herbs is the most effective for the different clinical characteristics of DR patients.

### Research methods

Alternative treatment options to traditional Chinese medicine were incorporated and assessed by employing a mesh meta-analysis to prioritize the therapeutic effects of these options based on various clinical observations.

### Research results

When these nine Oral Chinese medicines were analyzed in combination with conventional western medicine treatment (CT) compared with CT alone, the results showed that Hexue Mingmu Tablets has better intervention effect on the overall efficacy, visual acuity and reducing fasting blood glucose, Shuangdan Mingmu Capsules has better effect on inhibiting vascular endothelial growth factor, Danggui Buxue Decoction has better effect on reducing fundus hemorrhage area, and Qiju Dihuang Pills has better effect on reducing glycated hemoglobin.

# Research conclusions

Shuangdan Mingmu Capsules or Hexue Mingmu Tablets in combination with western drugs for DR may be the ideal treatment option.

#### Research perspectives

Bringing guidance to the clinical use of DR, as well as providing direction to basic experiments.

# FOOTNOTES

Author contributions: Li HD and Li MX conceived and designed the study, performed the initial search and screening of the literature; Zhang SW was involved in the decision of literature inclusion; Zhang WH assessed the quality of the literature; Li HD and Li MX entered the literature data, including author information, year of publication, outcome indicators, and the profile of the study population; Li MX performed the network meta-analysis and produced the figures; Li HD performed the first draft of the manuscript; Zhang WH and Zhang SW performed the preliminary check and revision of the first draft; Gong YB performed the final revision of the manuscript and critically reviewed the results of the data; In addition, all personnel were familiar with the content of the manuscript; Li HD and Li MX contributed equally to this study.

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