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\*CORRESPONDENCE Yan Li ∑li\_xyan@126.com

<sup>†</sup>These authors share first authorship

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# Risk factor screening and prediction modeling of gastrointestinal adverse reactions caused by GLP-1RAs

Rui Gao<sup>1†</sup>, Yue Li<sup>1,2†</sup>, Anan Li<sup>3</sup>, Penglin Zhou<sup>3</sup>, Huiying Zong<sup>3</sup> and Yan Li<sup>1\*</sup>

<sup>1</sup>Department of Clinical Pharmacy, The First Affiliated Hospital of Shandong First Medical University, Jinan, China, <sup>2</sup>Department of Medicine, Qilu Institute of Technology, Jinan, China, <sup>3</sup>Department of Pharmacy, Shandong University of Traditional Chinese Medicine, Jinan, China

**Objective:** This study aimed to explore the risk factors for gastrointestinal side effects (GISEs) in patients with type 2 diabetes mellitus (T2DM) during treatment with glucagon-like peptide-1 receptor agonists (GLP-1RAs) based on real-world data and to develop a prediction model for GLP-1RA-related GISEs.

**Methods:** A total of 855 patients who attended the First Affiliated Hospital of Shandong First Medical University from January 2020 to May 2023 were selected as the study participants, who were divided into the training set (598 cases) and the validation set (297 cases) using a simple random sampling method at a ratio of 7:3. The general information and biochemical indicators of the participants were collected to assess the risk factors for GLP-1RA-related GISEs, and multifactorial logistic regression analysis was used to obtain the best predictors. A nomogram prediction model was constructed. The Hosmer–Lemeshow test was used to assess the differentiation and calibration of the nomogram model, and decision curve analysis (DCA) was used to evaluate the clinical utility of the model.

**Results:** Age, gender, history of gastrointestinal disorders, and number of combined oral medications were found as risk factors for the occurrence of GISEs in patients with T2DM using GLP-1RAs (p < 0.05). The nomogram prediction model based on these four factors had good discriminability (AUC values of the training and validation sets of 0.855 and 0.836, respectively) and accuracy (Hosmer–Lemeshow test: p > 0.05 for the validation set). DCA showed that the prediction model curve had clinical utility in the threshold probability interval of >5%.

**Conclusions:** The established nomogram model has an excellent predictive effect on GISEs induced by GLP-1RAs in patients with T2DM.

#### KEYWORDS

type 2 diabetes mellitus, glucagon-like peptide-1 receptor agonists, gastrointestinal side effects, risk factors, nomogram

# 1 Introduction

The prevalence of type 2 diabetes mellitus (T2DM) has been on the rise globally in recent years, with data from the International Diabetes Federation (IDF) showing that there will be 783.2 million people (12.2%) living with diabetes by 2045 (1), of which T2DM will account for approximately 90% (2). Glucagon-like peptide-1 receptor agonists (GLP-1RAs) were approved for T2DM treatment in 2005. Now, the American Diabetes Association (ADA) has recommended in the Standards of Care in Diabetes the addition of GLP-1RAs and/or SGLT2 inhibitors (SGLT2i) to the treatment plan for patients with T2DM who have been diagnosed with or are at high risk of atherosclerotic cardiovascular disease (ASCVD) (3). Cardiovascular outcome studies have demonstrated that GLP-1RAs are effective in preventing cardiovascular events (such as acute myocardial infarction or stroke), reducing associated mortality (4-7), and in helping to prevent the progression of renal complications in patients with T2DM (5-8). GLP-1RAs are recommended as the preferred injectable hypoglycemic therapy for T2DM, even superior to insulin therapy (9), due to their definite cardiac and renal benefits.

However, the occurrence of gastrointestinal side effects (GISEs) has challenged the application of GLP-1RAs, which were clinically characterized by symptoms such as nausea (42.23%), vomiting (21.90%), diarrhea (21.93%), and constipation (8.41%) (10), and the incidence of severe adverse reactions was 15.43% (10). It has been shown that GISEs are associated with central and peripheral glucagonlike peptide-1 receptor (GLP-1R) activation (11). These adverse effects caused by treatment with GLP-1RAs are generally short-lived and gradually subside after a few weeks of treatment and therefore can be tolerated by most patients. Nevertheless, there are a few patients who are unable to tolerate severe GISEs caused by GLP-1RAs, resulting in the discontinuation of GLP-1RA treatment, leading to delays in the disease control and to financial losses. The study results obtained by Zhang et al. (12) showed that GLP-1RAs had the highest risk of intolerable gastrointestinal adverse events among hypoglycemic drugs, with a cumulative percentage area under the ranking curve of 91.8. The severity of GISEs varies among GLP-1RAs (10). Some studies have confirmed that the risk factors for GISEs caused by liraglutide may include the co-application of  $\alpha$ -glucosidase inhibitors or metformin, history of gastrointestinal disorders, gender, age, and elevated levels of thyroid-stimulating hormone (TSH) (13-15). However, until now, there has been a lack of well-designed studies to validate the risk factors for GISEs caused by GLP-1RAs; thus, there is an urgent need to establish a risk prediction model for the identification of the high-risk population. Therefore, the main purposes of this study were to investigate the risk factors for GLP-1RA-related GISEs and to develop an effective prediction model so as to provide a basis for the individualization of GLP-1RAs.

# 2 Methods

### 2.1 Research design

#### 2.1.1 Study participants and data sources

The data used in the study were derived from the healthcare big data platform and the electronic medical record system of the First Affiliated Hospital of Shandong First Medical University (Shandong Provincial Qianfoshan Hospital). Patients with T2DM who received GLP-1RAs for the first time between January 1, 2020, and May 31, 2023, were included. T2DM was diagnosed according to the diagnostic criteria in the Chinese Guidelines for the Prevention and Control of Type 2 Diabetes Mellitus (2020), as shown in Supplementary Table S1. Patient information including the demographic characteristics, co-administered medications, laboratory data, history of gastrointestinal disorders, and clinical diagnosis were collected for further analysis.

#### 2.1.2 Inclusion and exclusion criteria

The inclusion criteria for study participation were: 1) patients with T2DM; 2) aged 18 years and older; 3) patients with poor glycemic control and first-time treatment with GLP-1RAs; and 4) patients with well-established demographics, laboratory data, and medical and medication history. The exclusion criteria were: 1) patients with malignant tumors; 2) patients with a dialysis or renal transplantation history; 3) patients with end-stage diseases; 4) patients with a history of drug abuse or with severe pancreatic diseases; 5) patients with severe gastrointestinal dysfunction; 6) patients with severe hepatic or renal insufficiency; and 7) women who are pregnant or breastfeeding.

# 2.1.3 Determination of the correlation of adverse drug reactions and GLP-1RAs

The symptoms of GISEs include nausea, vomiting, abdominal pain, diarrhea, reflux, flatulence, and other similar conditions. The correlation between the use of GLP-1RAs and the occurrence of GISEs was evaluated according to Naranjo's scale (16) (a Naranjo score  $\geq$ 5 was considered as a GLP-1RA-related GISE). The Naranjo's scoring criteria are shown in Supplementary Table S2.

## 2.2 Data processing

A simple random sampling method was used to divide the study population into a training set (598 cases) and a validation set (297 cases) at a ratio of 7:3 for analysis. All statistical analyses were performed in R software (version 3.6.3). The graphics were drawn using GraphPad Prism 9.0.0 software. Categorical variables were reported as frequencies and percentages, normally distributed continuous variables were described as the mean ± standard deviation, and non-normally distributed continuous variables were described as median (interquartile range). For the data statistics, a t-test was used for normally distributed variables, the Wilcoxon test for non-normally distributed variables, and the chisquare test for categorical variables (i.e., sex, history of gastrointestinal disorders, and smoking history). Binary logistic regression analysis was used to screen the predictors and to construct a nomogram prediction model. Moreover, receiver operating characteristic (ROC) curves were plotted, and the Hosmer-Lemeshow test was used to assess the discriminability and calibration of the prediction model. Finally, decision curve analysis (DCA) was used in order to assess the clinical utility of the model.

# **3** Results

## 3.1 Demographic characteristics of the study participants

A total of 855 patients with T2DM were included in the present study, of whom 636 (74.39%) received injections of liraglutide, 137 (16.02%) of polyethylene glycol loxenatide, 44 (5.15%) of semaglutide, 23 (2.69%) of exenatide, 3 (0.35%) of dulaglutide, and 12 (1.40%) of lixisenatide. The research group included 624 (72.98%) men and 231 (27.02%) women, aged 54 years (44–62 years), with duration of diabetes mellitus of 4.00 years (0.15–10.00 years). The glycosylated hemoglobin (HbA<sub>1c</sub>) level was 8.40% (7.30%–9.90%), while the fasting blood glucose was 7.46 ±.75 mmol/L. A total of 526 (61.52%) of all enrolled patients developed GISEs, of which 14 (1.64%) discontinued GLP-1RA treatment due to intolerance to GISEs. The participant inclusion process is shown in Figure 1.

### 3.2 Univariate analysis of the clinical variables

Only the history of gastrointestinal disorders and diastolic blood pressure (DBP) were statistically different between the two groups in the training and validation sets (p < 0.05) (Table 1). The results of unifactorial screening based on the training set data are shown in Table 2. Significant differences were found between patients developing GISEs and those without GISEs in terms of age, gender, duration of diabetes, history of gastrointestinal disorders, smoking history, number of combined oral drugs (the types of combined oral medications are shown in Supplementary Table S3), body mass index (BMI), DBP, free triiodothyronine (FT<sub>3</sub>), free thyroxine (FT<sub>4</sub>), alanine aminotransferase (ALT), total bilirubin (TBIL), uric acid (UA), and estimated glomerular filtration rate (eGFR) (p < 0.05). However, family history of gastrointestinal disorders, history of alcoholism, heart rate (HR), systolic blood pressure (SBP), fasting blood glucose (FBG), glycated hemoglobin (HbA<sub>1c</sub>), total cholesterol (TC), triglycerides (TG), high-density lipoprotein cholesterol (HDL-C), low-density lipoprotein cholesterol (LDL-C), thyroid-stimulating hormone (TSH), fasting insulin (FINS), aspartate aminotransferase (AST), lipid profile, creatinine (enzyme assay) (Cr),  $\beta_2$ -microglobulin ( $\beta_2$ -M), red blood cell count (RBC), white blood cell count (WBC), and platelet count (PLT) were not significantly correlated with GISEs (p > 0.05).

# 3.3 Multifactor logistic regression analysis of the significant variables

The stepwise method of binary logistic regression was used to assess whether the significant variables in the univariate analysis were associated with GISEs (Table 3). Preliminary analysis of logistic regression showed that age [adjusted OR = 1.06 (1.02–1.09)], gender [adjusted OR = 2.86 (1.47–5.60)], number of combined oral drugs [adjusted OR = 1.94 (1.63–2.31)], history of gastrointestinal disorders [adjusted OR = 4.92 (2.39–10.12)], eGFR [adjusted OR = 1.02 (1.00–1.04)], and BMI [adjusted OR = 1.06 (0.99–1.14)]. The *p*-values for eGFR and BMI were greater than 0.05 and not statistically significant. In the final regression model, age [adjusted OR = 1.05 (1.04–1.07)], gender [adjusted OR = 2.32 (1.42–3.80)], number of combined oral drugs [adjusted OR = 1.84 (1.62–2.10)], and history of gastrointestinal disorders [adjusted OR = 6.22 (3.49–11.11)] were associated with GISEs caused by GLP-1RAs.



#### TABLE 1 Results of the baseline feature comparison between the training and validation sets.

|   | Training set ( <i>n</i> = 598) | Validation set ( $n = 257$ ) | <i>p</i> -value |
|---|--------------------------------|------------------------------|-----------------|
| Age (years)   | 53 (45-63)                     | 53 (44-60)                   | 0.104           |
| Gender (% women)                                    | 168 (28.09)                    | 63 (24.51)                   | 0.280           |
| Duration of diabetes (years)                        | 4.00 (0.15-10.00)              | 5.00 (0.42-10.00)            | 0.811           |
| Family history of gastrointestinal disorders, n (%) | 30 (5.02)                      | 12 (4.67)                    | 0.829           |
| History of gastrointestinal disorders, n (%)        | 136 (22.74)                    | 41 (15.95)                   | 0.025*          |
| Smoking history, n (%)                              | 230 (38.46)                    | 113 (43.97)                  | 0.190           |
| History of alcoholism, <i>n</i> (%)                 | 191 (31.94)                    | 92 (35.80)                   | 0.531           |
| No. of combined oral drugs                          | 2 (2-4)                        | 2 (2-4)                      | 0.328           |
| SBP (mmHg)  | 132.41 ± 14.55                 | 134.10 ± 13.91               | 0.118           |
| DBP (mmHg)  | 79.96 (73.41-86.17)            | 81.16 (75.97-87.19)          | 0.034*          |
| HR (bpm)  | 82.87 ± 16.96                  | 83.50 ± 13.44                | 0.961           |
| BMI (kg/m <sup>2</sup> )                            | 28.04 (25.93-30.48)            | 28.15 (26.05-30.69)          | 0.693           |
| FBG (mmol/L)  | 7.47 ± 1.86                    | 7.44 ± 1.49                  | 0.894           |
| FINS (µIU/ml)                                       | 8.70 (4.62–15.21)              | 9.77 (5.47–16.16)            | 0.350           |
| HbA <sub>1c</sub> (%)                               | 8.40 (7.30-9.80)               | 8.45 (7.38-10.10)            | 0.398           |
| TC (mmol/L)   | 13.86 (2.10-20.17)             | 14.68 (2.27–19.86)           | 0.560           |
| TG (mmol/L)   | 2.86 (0.24-4.56)               | 0.43 (0.23-4.65)             | 0.945           |
| HDL-C (mmol/L)                                      | 1.03 (0.91–1.17)               | 1.04 (0.90–1.17)             | 0.969           |
| LDL-C (mmol/L)                                      | 2.58 ± 0.86                    | 2.58 ± 0.87                  | 0.966           |
| TSH (µIU/ml)  | 1.83 (1.29–2.63)               | 1.63 (1.24–2.37)             | 0.060           |
| FT <sub>3</sub> (pmol/L)                            | 4.65 (4.11-5.08)               | 4.62 (4.24-5.10)             | 0.361           |
| FT <sub>4</sub> (pmol/L)                            | 18.52 (16.88-20.25)            | 18.38 (16.66–20.20)          | 0.668           |
| ALT (U/L)   | 25.10 (18.50-38.70)            | 26.10 (18.30-40.15)          | 0.927           |
| AST (U/L)   | 18.70 (14.70-24.82)            | 19.30 (14.70–25.20)          | 0.660           |
| TBIL (µmol/L)                                       | 10.70 (8.10-14.62)             | 11.30 (8.30–14.20)           | 0.619           |
| Cr (µmol/L)   | 67.00 (58.00-77.00)            | 66.00 (57.00-74.50)          | 0.214           |
| Urea/Cr (ratio)                                     | 0.08 (0.07-0.10)               | 0.08 (0.07-0.10)             | 0.866           |
| $\beta_2$ -M (mg/L)                                 | 1.63 (1.47–1.91)               | 1.63 (1.44–1.91)             | 0.259           |
| eGFR (ml/min/1.73 m <sup>2</sup> )                  | 107.59 (98.07–117.80)          | 109.14 (102.03–118.16)       | 0.163           |
| UA (μmol/L)   | 340.00 (278.00-392.00)         | 331.50 (283.00-397.50)       | 0.914           |
| RBC (×10 <sup>12</sup> /L)                          | 4.14 (2.00-5.13)               | 4.19 (2.00-5.00)             | 0.633           |
| WBC (×10 <sup>9</sup> /L)                           | 6.34 (5.40-7.56)               | 6.37 (5.32–7.68)             | 0.993           |
| PLT (×10 <sup>9</sup> /L)                           | 221.00 (184.00-262.00)         | 214.00 (183.00-252.00)       | 0.375           |

Variables are expressed as the mean  $\pm$  standard deviation or percentage. *SBP*, systolic blood pressure; *DBP*, diastolic blood pressure; *HR*, heart rate; *BMI*, body mass index; *FBG*, fasting blood glucose; *FINS*, fasting insulin; *HbA*<sub>1,c</sub>, glycated hemoglobin; *TC*, total cholesterol; *TG*, triglycerides; *HDL*-*C*, high-density lipoprotein cholesterol; *LDL*-*C*, low-density-lipoprotein cholesterol; *TSH*, thyroid-stimulating hormone; *FT*<sub>3</sub>, free triiodothyronine; *FT*<sub>4</sub>, free thyroxine; *ALT*, alanine aminotransferase; *AST*, aspartate aminotransferase; *eGFR*, estimated glomerular filtration rate; *Cr*, creatinine; *Urea/Cr*, urea-to-creatinine ratio;  $\beta_2$ -*M*,  $\beta_2$ -microglobulin; *TBIL*, total bilirubin; *UA*, uric acid; *RBC*, red blood cell count; *WBC*, white blood cell count; *PLT*, platelet count.

|   | GISE(–) ( <i>n</i> = 220) | GISE(+) (n = 378)      | <i>p</i> -value |  |
|---|---------------------------|------------------------|-----------------|--|
| Age (years)   | 48 (39–57)                | 57 (51–65)             | <0.001*         |  |
| Gender (% women)                                    | 40 (18.18)                | 128 (33.86)            | <0.001*         |  |
| Duration of diabetes (years)                        | 1.00 (0.00-8.00)          | 6.00 (0.50-12.00)      | <0.001*         |  |
| Family history of gastrointestinal disorders, n (%) | 7 (3.18)                  | 23 (6.08)              | 0.117           |  |
| History of gastrointestinal disorders, $n$ (%)      | 21 (9.55)                 | 115 (30.42)            | <0.001*         |  |
| Smoking history, n (%)                              | 98 (44.55)                | 132 (34.92)            | 0.013*          |  |
| History of alcoholism, n (%)                        | 82 (37.27)                | 109 (28.84)            | 0.065           |  |
| No. of combined oral drugs                          | 1 (0-3)                   | 3 (2–5)                | <0.001*         |  |
| BMI (kg/m <sup>2</sup> )                            | 28.68 (26.14-31.09)       | 27.72 (25.66-30.17)    | 0.005*          |  |
| HR (bpm)  | 83.44 ± 28.35             | 82.67 ± 13.94          | 0.950           |  |
| SBP (mmHg)  | 132.26 ± 14.69            | 132.50 ± 14.49         | 0.848           |  |
| DBP (mmHg)  | 81.47 (74.31-87.70)       | 79.00 (73.30-85.00)    | 0.013*          |  |
| FBG (mmol/L)  | 7.41 ± 1.85               | 7.52 ± 1.87            | 0.710           |  |
| HbA <sub>1c</sub> (%)                               | 8.40 (7.30-9.90)          | 8.40 (7.32-9.70)       | 0.811           |  |
| TG (mmol/L)   | 3.28 (0.24-4.65)          | 2.67 (0.24-4.46)       | 0.625           |  |
| HDL-C (mmol/L)                                      | 1.04 (0.91–1.16)          | 1.03 (0.92–1.17)       | 0.812           |  |
| LDL-C (mmol/L)                                      | 2.60 ± 0.78               | 2.57 ± 0.91            | 0.745           |  |
| TC (mmol/L)   | 13.34 (2.10–19.80)        | 14.09 (2.09–20.28)     | 0.748           |  |
| TSH (µIU/mL)  | 1.74 (1.25–2.62)          | 1.90 (1.30-2.63)       | 0.451           |  |
| FT <sub>3</sub> (pmol/L)                            | 4.76 (4.23-5.28)          | 4.56 (4.02-4.96)       | <0.001*         |  |
| FT <sub>4</sub> (pmol/L)                            | 18.95 (17.18-20.43)       | 18.24 (16.74-20.10)    | 0.046*          |  |
| FINS (µIU/mL)                                       | 9.20 (4.15–15.46)         | 8.52 (5.00-15.15)      | 0.824           |  |
| ALT (U/L)   | 26.40 (20.22-44.48)       | 24.50 (17.60-4.40)     | 0.028*          |  |
| AST (U/L)   | 19.50 (15.75-27.10)       | 18.00 (14.25-23.70)    | 0.053           |  |
| eGFR (ml/min/1.73 m <sup>2</sup> )                  | 110.98 (100.38-120.94)    | 104.93 (96.97–114.70)  | <0.001*         |  |
| Cr (µmol/L)   | 69.00 (58.00-79.00)       | 67.00 (57.00-75.00)    | 0.224           |  |
| Urea/Cr (ratio)                                     | 0.08 (0.06-0.10)          | 0.08 (0.07-0.09)       | 0.190           |  |
| β <sub>2</sub> -M (mg/L)                            | 1.60 (1.42–1.88)          | 1.68 (1.49–1.96)       | 0.068           |  |
| TBIL (µmol/L)                                       | 12.00 (8.65–15.40)        | 10.30 (8.03-13.80)     | 0.024*          |  |
| UA (µmol/L)   | 360.50 (291.00-408.00)    | 320.00 (271.00-383.50) | 0.013*          |  |
| RBC (×10 <sup>12</sup> /L)                          | 4.35 (2.00-5.11)          | 4.00 (2.00-5.14)       | 0.461           |  |
| WBC (×10 <sup>9</sup> /L)                           | 6.25 (5.41-7.69)          | 6.42 (5.39-7.43)       | 0.829           |  |
| PLT (×10 <sup>9</sup> /L)                           | 221.00 (190.00-255.50)    | 220.50 (183.25-263.00) | 0.733           |  |

TABLE 2 Demographic information and metabolic and biochemical indices of patients in the training set with and without gastrointestinal adverse effects during the use of glucagon-like peptide-1 receptor agonists.

Variables are expressed as the mean  $\pm$  standard deviation or percentage.

SBP, systolic blood pressure; DBP, diastolic blood pressure; HR, heart rate; BMI, body mass index; FBG, fasting blood glucose; FINS, fasting insulin; HbA<sub>1c</sub>, glycated hemoglobin; TC, total cholesterol; TG, triglycerides; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; TSH, thyroid-stimulating hormone; FT<sub>3</sub>, free triiodothyronine; FT<sub>4</sub>, free thyroxine; ALT, alanine aminotransferase; AST, aspartate aminotransferase; eGFR, estimated glomerular filtration rate; Cr, creatinine; Urea/Cr, urea-to-creatinine ratio;  $\beta_2$ -M,  $\beta_2$ -microglobulin; TBIL, total bilirubin; UA, uric acid; RBC, red blood cell count; WBC, white blood cell count; PLT, platelet count.

|                                       | Primary model |                   | Final model     |      |                   |                 |
|---------------------------------------|---------------|-------------------|-----------------|------|-------------------|-----------------|
|                                       | β             | OR (95%CI)        | <i>p</i> -value | β    | OR (95%CI)        | <i>p</i> -value |
| Gender (% women)                      | 1.05          | 2.86 (1.47-5.60)  | 0.002           | 0.84 | 2.32 (1.42-3.80)  | < 0.001         |
| History of gastrointestinal disorders | 1.59          | 4.92 (2.39-10.12) | < 0.001         | 1.83 | 6.22 (3.49–11.11) | <0.001          |
| Age (years)                           | 0.05          | 1.06 (1.02–1.09)  | < 0.001         | 0.05 | 1.05 (1.04–1.07)  | < 0.001         |
| No. of combined oral drugs            | 0.66          | 1.94 (1.63–2.31)  | < 0.001         | 0.61 | 1.84 (1.62–2.10)  | < 0.001         |
| eGFR (ml/min/1.73 m <sup>2</sup> )    | 0.02          | 1.02 (1.00-1.04)  | 0.128           |      |                   |                 |
| BMI (kg/m <sup>2</sup> )              | 0.06          | 1.06 (0.99–1.14)  | 0.089           |      |                   |                 |

TABLE 3 Multifactorial logistic regression analysis of the gastrointestinal adverse reactions caused by glucagon-like peptide-1 receptor agonists.

## 3.4 Model development

The GLP-1RA-related GISEs in patients with T2DM were predicted based on four factors (i.e., age, gender, number of combined oral medications, and history of gastrointestinal disorders), which were plotted as a nomogram (Figure 2). According to the nomogram model, the effect of each variable on GISEs was reflected in the respective row lengths and corresponding scores, and the total score of the model was obtained by adding the scores of each factor. The probability that the reference total score corresponds to the risk of GISEs was the risk of GISEs for a specific patient.

## 3.5 Evaluation of the predictive efficiency and clinical applicability of the model

The prediction performance of the nomogram was evaluated using ROC curves (the AUC values for the training and validation sets were 0.855 and 0.836, respectively), as shown in Figures 3A, B, respectively, indicating that the column-line diagram prediction model had good prediction ability. The calibration curves revealed that the predicted probabilities were generally consistent with the observed probabilities, indicating that the model had good goodness-of-fit (Hosmer-Lemeshow test: p = 0.095) (Figure 4A). The DCA curves demonstrated that screening with the GISE risk nomogram model at thresholds greater than 5% resulted in a high net clinical benefit (Figure 4B).

# 4 Discussion

The main finding of the present study is that age, gender, number of combined oral medications, and history of gastrointestinal disorders are risk factors for GISEs caused by GLP-1RAs. A nomogram model was used to establish a risk prediction model for the clinical application of GLP-1RAs, which is of great significance for the safety and individualized administration of GLP-1RAs in patients with T2DM.

A previous study found that aging is associated with an increased incidence of liraglutide-related GISEs (14), which is consistent with the outcomes of this study. The reasons may be that, with increased age, the gastrointestinal tract undergoes the following physiological degenerative changes: a) decreased resistance of the gastric mucosa to damages, which leads to an increased risk of gastrointestinal ulcer disease; b) gradually decreased secretion of gastric acid and pepsin, resulting in a decline in digestive function and, thus, an increased risk of functional digestive malfunction; and c) reduced water absorption capacity of the large intestine and rectum and a weakened motility, thus making it susceptible to swelling and constipation (17). Therefore, the use of GLP-1RAs should be taken with caution in elderly patients.



Nomogram including age, gender, number of combined oral drugs, and history of gastrointestinal disorders for the risk of gastrointestinal side effects in patients using glucagon-like peptide-1 receptor agonists.



#### FIGURE 3

(A) Receiver operating characteristic (ROC) curves for the training set of the prediction model. (B) ROC curves for the validation set of the prediction model.



The number of combined oral medications is also a risk factor for the occurrence of GISEs, probably due to the fact that the greater the number of combined medications, the more complex the drug interactions in vivo (18). In addition, more oral medications have a superimposed gastrointestinal irritant effect that increases the risk of developing GISEs. Some studies have reported that 6%-30% of adverse drug reactions are caused by drug-drug interactions (19), particularly 5%-9% of adverse drug reactions in hospitalized patients (19), and that inappropriate combinations of medications could also lead to the enhanced side effects of specific medications (18), which could increase the risk of gastrointestinal adverse reactions in patients with T2DM treated with GLP-1RAs. A history of gastrointestinal disorders could also contribute to the development of GISEs, although gastrointestinal disorders are not a contraindication to the use of GLP-1RAs and comprised the risk factors for the development of GISEs in our study (p < 0.001). This might be due to the fact that GLP-1RAs have a delaying effect on gastrointestinal emptying, which could exacerbate gastrointestinal disorders or predispose to the development of prior gastrointestinal disorders. Therefore, patients with a history of gastrointestinal disorders are advised to use GLP-1RAs with caution in order to avoid poor tolerance and compliance.

In this study, the first nomogram prediction model was constructed to predict the risk of GLP-1RA-associated GISEs in patients with T2DM. The AUC values of both the training and test sets indicated that the model had high validity and accuracy in predicting GISEs, which had certain practical application value. The calibration curve showed that the predicted probability of the model was in good conformity with the actual probability. However, this study still has some limitations. As a cross-sectional study, we used data from patients during their hospitalization to analyze the status for a short period after the use of the target medication, and there was no information on the long-term follow-up of outpatients and inpatients, which could have some confounding factors. In addition, documentation of the GISEs was taken from the medical records, which might have some factual bias.

In summary, this study explores and establishes a convenient and practical nomogram model for predicting the risk of GISEs after the use of GLP-1RAs in patients with T2DM, which provides early warning information for patients to understand the risk of developing GISEs as early as possible and to take timely and appropriate interventions.

## Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

## Ethics statement

Ethical approval was not required for the study involving humans in accordance with the local legislation and institutional requirements. Written informed consent to participate in this study was not required from the participants or the participants' legal guardians/next of kin in accordance with the national legislation and the institutional requirements.

## Author contributions

RG: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Software, Visualization, Writing – original draft. YuL: Investigation, Methodology, Visualization, Writing – original draft. AL: Investigation, Methodology, Visualization, Writing – original draft. PZ: Data curation, Writing – original draft. HZ: Data curation, Writing – original draft. YaL: Conceptualization, Funding acquisition, Resources, Supervision, Writing – review & editing.

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# Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

# **Generative AI statement**

The author(s) declare that no Generative AI was used in the creation of this manuscript.

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## Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2024. 1502050/full#supplementary-material

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