Distinct developmental trajectories of body mass index and diabetes risk: A 5-year longitudinal study of Chinese adults

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Keywords

Body mass index, Diabetes, Trajectory

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ABSTRACT

Aims/Introduction: This longitudinal study aimed to explore whether distinct developmental trajectories of body mass index (BMI) would be predictive of diabetes risk in general Chinese adults.

Materials and Methods: A total of 4,519 participants aged >18 years who were free of diabetes in 2011 (baseline of the current analysis) were enrolled in this study. All participants completed a medical examination every year during 2011–2016, and BMI levels were measured two to six (average 5.6) times. Group-based trajectory modeling was applied to identify BMI trajectories over time. New-onset diabetes was confirmed in 2016. **Results:** During 2011–2016, four distinct BMI trajectories were identified according to BMI range and changing pattern over time: "low" (19.6%), "moderate" (33.4%) and "high" (13.6%). A total of 168 (3.7%) new-onset diabetes cases were confirmed in 2016. Compared with the "low" BMI trajectory, participants in the "high" BMI trajectory were at significantly higher risk for new-onset diabetes (adjusted relative risk 3.24, 95% confidence interval 1.27–8.24). Notably, BMI trajectories based on the first four or three annual BMI tests yielded similar results. By contrast, no significant correlation was found between categories of baseline BMI and new-onset diabetes in 2016 after multivariate adjustment.

Conclusions: The present results show that distinct BMI trajectories, even identified using just four or three annual BMI tests, are significantly associated with new-onset diabetes. Monitoring BMI trajectories over time might provide an important approach to identify subpopulations at higher risk for developing diabetes.

INTRODUCTION

Diabetes is a major and growing health concern worldwide¹. In China, the prevalence of diabetes has increased sharply, from 0.67% in 1980, to 10.9% in 2013, as a result of considerable changes in lifestyles and aging². Meanwhile, the overall mortality, disability-adjusted life-years and economic costs attributable to diabetes are showing rapid growth in China^{3,4}. Previous studies suggested that early detection and treatment of diabetes would substantially reduce the related morbidity and mortal-ity^{5,6}. However, current diagnostic procedures, including fasting blood glucose test, oral glucose tolerance test and hemoglobin A1c test, are not suitable for large-scale population screening in

a developing country, such as China. Therefore, using a simple method to identify subpopulations at higher risk for diabetes would critically inform prevention efforts.

Obesity, commonly classified using a simple index – body mass index (BMI) – is well recognized as the major risk factor for diabetes. However, previous studies were often based on BMI at a certain point, regardless of the effect of long-term BMI changes on diabetes risk^{7,8}. Hence, understanding the heterogeneity of diabetes risk by exploring the distinct patterns of BMI changes might carry new insights into diabetes prevention. In addition, in China, the incidence of diabetes is high, despite a relatively low prevalence of obesity, implying that the effect of BMI on diabetes might differ across various racial/ethnic groups⁹.

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Group-based trajectory modeling (GBTM) is a type of latent class growth analysis that can identify distinct clusters of individuals who are following similar BMI trajectories¹⁰. This method assumes that individual differences in trajectories can be summarized by a finite set of different polynomial functions of time¹¹. In the past years, this method has been successfully used to examine the association between BMI trajectories and the risk of hypertension¹², cancer^{13–15} and all-cause mortality^{16,17}. However, to date, just a few studies have explored the association between BMI trajectories identified by GBTM and diabetes risk^{18,19}. Whether distinct BMI trajectories would be predictive of diabetes risk in general Chinese adults is still unknown. To address the gap, by using repeated measurements of BMI in a longitudinal cohort of Chinese adults, we aimed to examine the association of distinct BMI trajectories with newonset diabetes.

METHODS

Study population

The Health Management Center of Third Xiangya Hospital, Changsha, Hunan, China is one of the largest medical examination centers in China, mainly servicing employees from hundreds of institutions in Changsha. Of these institutions, 127 continuously chose the center for their annual employee medical examinations during 2011–2016. In the present study, we retrospectively recruited 5,004 participants from the 127 institutions. All participants were aged >18 years and had completed a medical examination every year during 2011–2016 (Figure 1). The present study was in accordance with the guidelines of the Declaration of Helsinki and was approved by the Medical Ethics Committee of Third Xiangya Hospital. All participants



Figure 1 | Timeline of annual visits and study design. At baseline, all participants were free of diabetes. New-onset diabetes was confirmed in 2016. (a) The primary analysis identified distinct body mass index (BMI) trajectories using six BMI tests (during 2011–2016). Regression models were then constructed to examine the association between distinct BMI trajectories and new-onset diabetes. (b,c) Secondary analyses reconstructed the regression models after identifying distinct BMI trajectories using four and three BMI tests.

signed an informed consent form and agreed to share their health information for medical research.

Because diabetes is our primary outcome variable, 94 participants who had missing data on diabetes in 2011, and 288 participants who were diagnosed with diabetes in 2011 were excluded from this study. Meanwhile, 56 participants who had missing data on the diagnosis of diabetes in 2016 were also excluded. In addition, as the calculation of BMI trajectories required at least two BMI tests^{20,21}, we further excluded 47 participants who could not meet this criterion. Finally, 4,519 participants were deemed eligible for the present study.

Diagnosis of diabetes

Venous blood samples were collected in the morning after overnight fasting for 8–12 h and then transfused into ethylenediaminetetraacetic acid-containing vacuum tubes. Blood samples were stored at –20°C until analyzed. The concentration of fasting blood glucose was determined by enzymatic colorimetric assay using an automated analyzer (Hitachi 7600-110; Hitachi, Tokyo, Japan) at the central laboratory of Third Xiangya Hospital. The intra-assay coefficients of variation for blood glucose were <2.5%. Diabetes was defined by the presence of any of the following: (i) self-reported doctor-diagnosed diabetes; (ii) current use of insulin or oral hypoglycemic agents; and (iii) fasting blood glucose \geq 7.0 mmol/L²².

Assessment of BMI

Bodyweight, height and waist circumference were measured to the nearest 0.1 kg or 0.1 cm, with participants wearing light clothes and no shoes. BMI was calculated as bodyweight in kilograms (kg) divided by the square of height in meters (m^2) .

Assessment of potential covariates

Information on demographic variables (age, sex, race, occupation and marital status), lifestyle factors (cigarette smoking and alcohol consumption) and medical history were obtained by well-trained interviewers using standardized questionnaires. A total of 11 racial groups, including Han, Miao, Tujia, Hui, Zhuang, Bai, Dong, Man, Yao, Mongolian and Xibo, were identified in our study sample. For the analyses, just two categories were considered: Han and all other minorities. Occupation was classified into seven groups according to the types of institutions.

Blood pressure (BP) was measured to the nearest 0.1 mmHg in a sitting position after a 10-min rest. Using a corrected mercury sphygmomanometer, two readings were obtained for both systolic and diastolic BP, with a 30-s interval. The mean of the two readings was considered as the participant's BP. If the two readings differed by >5 mmHg, BP was re-measured, and the participant's BP was finally calculated as the average of the three readings. Hypertension was defined as systolic BP \geq 140 mmHg and/or diastolic BP \geq 90 mmHg, or self-reported doctor-diagnosed hypertension, or current use of antihypertensive agents²³.

Other biochemical measurements, including total cholesterol, triglycerides, hemoglobin, creatinine and uric acid, were also taken at the central laboratory of Third Xiangya Hospital. Details about the measurement methods were published previously^{23,24}.

Statistical analysis

All analyses were carried out using Stata version 14.0 (Stata Corp., College Station, TX, USA). The BMI trajectories were determined by GBTM, fitted using the user-written "traj" procedure in Stata^{10,25}. We used censored normal distribution to model BMI. We chose the number and shape (i.e., intercept, linear, quadratic and cubic) of BMI trajectories following the criteria: lowest Bayesian information criterion, high posterior probabilities (>0.7) and \geq 5% of total sample in a trajectory group. In a step-by-step manner, we finally decided on the best-fitting model with four linear trajectories (Figure 2).

Baseline characteristics of the study sample were summarized as the median (interquartile range) for continuous variables, and as the number (percentage) for categorical variables. Comparisons of baseline characteristics by distinct BMI trajectories were assessed using Kruskal–Wallis test for continuous variables, and the χ^2 -test for categorical variables. Trends in baseline characteristics across distinct BMI trajectories were assessed using Spearman's correlation for continuous variables, and the Cochran–Armitage trend test for binary variables. Modified Poisson regression models, using the robust error variance²⁶, were used to estimate the relative risks (RRs) for new-onset diabetes in 2016. All potential covariates in the regression models were collected at baseline. Missing values of potential covariates were assumed to be random, and were handled using multiple imputation with chained equations²⁷. As approximately 30% of



Figure 2 | Group-based trajectory modeling used to determine distinct body mass index (BMI) trajectories. Distinct BMI trajectories were identified using six BMI tests (during 2011–2016).

the study sample had missing values on at least one study variable, we generated 30 imputed datasets in the imputation models²⁷. To optimize the imputation, we added all study variables into the imputation models²⁷.

Statistical interactions between BMI trajectories and sex (male vs female), age (<45 vs \geq 45 years) and impaired fasting glucose (fasting blood glucose <5.6 vs \geq 5.6 mmol/L) in relation to risks of new-onset diabetes were examined using likelihood ratio tests in the regression models. To compare with traditional methods based on baseline BMI, we excluded 196 participants who had missing values on baseline BMI in a secondary analysis. To explore whether BMI trajectories based on fewer BMI tests could predict new-onset diabetes in 2016, we further examined the association of BMI trajectories identified using just four or three annual BMI tests with new-onset diabetes in 2016 (Figure 1).

RESULTS

Baseline characteristics of study participants by distinct BMI trajectories

Of the 4,519 participants, 2,438 (53.9%) were men, and the median age at baseline was 42.0 years (interquartile range 32.0-57.0 years). Based on the BMI range and changing pattern during 2011-2016, four linear BMI trajectories were identified in these participants (Figure 2). Table 1 shows the baseline characteristics of the study participants by distinct BMI trajectories. Participants belonging to the "high" BMI trajectory were more likely to be older, male, current smoker and current alcohol drinker at baseline, and they tended to have an increased prevalence of hypertension, as well as higher levels of BMI, waist circumference, fasting blood glucose, total cholesterol, triglycerides, hemoglobin, creatinine and uric acid at baseline. In addition, the percentages of occupation, married status and use of antihypertensive at baseline were also significantly different across distinct BMI trajectories. However, there was no significant difference in race and family history of diabetes by distinct BMI trajectories.

Association between distinct BMI trajectories and diabetes risk A total of 168 (3.7%) new-onset diabetes cases were identified in 2016. As shown in Table 2, the incidence of new-onset diabetes progressively increased from the "low" to "high" BMI trajectory (0.6, 2.0, 5.1 and 9.1%, respectively). In model 1 adjusting for age and sex at baseline, the RR for new-onset diabetes in the comparison between the "high" and "low" BMI trajectory was 9.17 (95% confidence interval [CI] 3.67-22.89). After further adjusting for all other potential covariates, the risk was attenuated and the RR for new-onset diabetes in the comparison between the "high" and "low" BMI trajectory was 3.24 (95% CI 1.27-8.24). No significant interactions were observed between distinct BMI trajectories and sex, age and impaired fasting glucose in relation to risks of new-onset diabetes (P-values for interactions were 0.520, 0.389 and 0.849, respectively).

Variables	Total $(n = 4,519)$	Trajectory 1: Low (n = 884)	Trajectory 2: Moderate (<i>n</i> = 1,511)	Trajectory 3: Moderate-high (<i>n</i> = 1,510)	Trajectory 4: High (<i>n</i> = 614)	<i>P</i> -value*	P for trend ^{\dagger}
Age (years) Male (%) BMI (kg/m ²) VVC (cm)	42.0 (32.0–57.0) 2,438 (53.9) 23.4 (20.9–25.6) 80.0 (71.0–87.0)	32.0 (27.0-43.0) 194 (22.0) 19.4 (18.5-20.1) 67.0 (64.0-71.0)	41.0 (31.0–55.0) 616 (40.8) 22.0 (21.1–22.9) 75.0 (71.0–80.0)	47.0 (37.0–61.0) 1,115 (73.8) 25.1 (24.2–25.9) 85.0 (80.0–88.0)	47.0 (37.0–63.0) 513 (83.6) 28.2 (27.3–29.2) 93.0 (88.0–96.0)	 <0.001 <0.001 <0.001 <0.001 <0.001 	<0.001 <0.001 <0.001 <0.001
Race (%) Han Other minorities	4,350 (96.6) 153 (3.4)	851 (96.3) 33 (3.7)	1,466 (97.5) 38 (2.5)	1,443 (95.9) 62 (4.1)	590 (96.7) 20 (3.3)	0.102	0.577
Occupation (%) Civil servants	1,887 (41.8)	230 (26.0)	574 (38.0)	755 (50.0)	328 (53.4)	<0.001	I
Educators Media workers Medical workers	166 (3.7) 747 (16.5) 1 000 (24.1)	26 (2.9) 168 (19.0) 353 (30.0)	56 (3.7) 244 (16.2) 470 (78.4)	68 (4.5) 238 (15.8) 230 (15.8)	16 (2.6) 97 (15.8) 69 (11.2)		
Bank workers Police officers Others	228 (5.1) 132 (2.9) 260 (6.0)	48 (5.4) 8 (0.9) 51 (5.8)	91 (60) 37 (25) 80 (53)	63 (4.2) 57 (3.8) 60 (6.0)	26 (4.2) 30 (4.9) 48 (7.8)		
Current smoker (%) Current alcohol drinker (%)	200 (6.0) 989 (26.5) 2.034 (60.1)	71 (9.5) 759 (38.3)	239 (19.4) 608 (54.6)	450 (36.5) 807 (71.6)	229 (44.6) 360 (77.4)	<0.001	<0.001
Married status (%)	3,415 (86.3)	535 (70.9)	1,108 (84.7)	1,265 (93.9)	507 (92.9)	<0.001	<0.001
Family history of diapetes (%) Hypertension (%)	347 (7.7) 946 (21.6)	(7.4) 49 (5.8)	201 (7.9) 201 (13.8)	426 (28.9)	46 (7.5) 270 (45.5)	≤0.001	<0.001
Use of antihypertensive (%) Fasting blood glucose (mmol/L)	407 (9.0) 5.0 (4.7–5.3)	24 (2.7) 4.8 (4.5–5.1)	91 (6.0) 4.9 (4.7–5.2)	179 (11.9) 5.1 (4.8–5.5)	113 (18:4) 5.2 (4:8–5.6)	<0.001 <0.001 <0.001	<0.001 <0.001
Total cholesterol (mmol/L) Trialvcerides (mmol/L)	4.7 (4.2–5.3) 1.13 (0.79–1.67)	4.4 (3.9–5.0) 0.79 (0.62–1.06)	4.6 (4.1–5.3) 1.00 (0.73–1.44)	4.8 (4.3–5.5) 1.37 (0.99–1.96)	4.9 (4.3–5.5) 1.62 (1.20–2.43)	<0.001 <0.001	<0.001 <0.001
Hemoglobin (g/L) Creatinine (µmo/L) Uric acid (µmo/L)	133.0 (124.0–144.0) 66.0 (54.0–78.0) 282.0 (221.0–349.0)	125.0 (119.0–132.0) 55.0 (50.0–64.0) 216.0 (183.0–268.0)	129.0 (122.0–140.0) 61.0 (52.0–73.0) 255.0 (209.0–318.0)	140.5 (130.0–148.0) 73.0 (60.0–81.0) 319.0 (261.8–379.0)	143.0 (133.8–150.0) 75.0 (65.0–84.0) 350.0 (295.0–410.0)	<0.001 <0.001 <0.001 	<0.001 <0.001 <0.001
Data were presented as median (ii using Kruskal–Wallis test for contin Cochran–Armitage trend test for b	interquartile range) or n (5 nuous variables, and the χ pinary variables. WC, waist	%). Distinct body mass in 2 ^{,2} -test for categorical vari circumference.	dex (BMI) trajectories we ables. [†] <i>P</i> -values for trenc	ere identified using six BN I are calculated using Spe	Al tests (during 2011–201 earman's correlation for c	6). *P-values al ontinuous vari	e calculated ables, and the

Table 2 Relative risk and 95% confi	dence interva	al of new-onset	diabetes in 2016, by distir	nct body mass index traje	ectories and categories of	baseline body mass inde	×
Variables	Z	Cases (%)	Model 1 Adjusted RR (95% CI) [†]	Model 2 Adjusted RR (95% CI) [‡]	Model 3 Adjusted RR (95% CI) ^{\$}	Model 4 Adjusted RR (95% CI) [¶]	Model 5 Adjusted RR (95% CI) ^{††}
BMI trajectories	4,519	168 (3.7)					
Trajectory 1: Low	884	5 (0.6)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)
Trajectory 2: Moderate	1,511	30 (2.0)	2.66 (1.04–6.81)	2.48 (0.96–6.41)	2.23 (0.86–5.82)	1.66 (0.64-4.31)	1.48 (0.56–3.88)
Trajectory 3: Moderate-high	1,510	77 (5.1)	5.51 (2.24–13.54)	5.07 (2.04–12.57)	3.65 (1.45–9.14)	2.47 (0.98–6.23)	2.11 (0.84–5.32)
Trajectory 4: High	614	56 (9.1)	9.17 (3.67–22.89)	8.26 (3.28–20.77)	5.35 (2.12–13.52)	3.24 (1.27–8.24)	2.79 (1.09–7.11)
P-value for trend			<0.001	<0.001	<0.001	<0.001	0.001
Baseline BMI (groups) ^{‡‡}	4,323	155 (3.6)					
First group (<20.4)	848	7 (0.8)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	I
Second group (20.4< to <23.6)	1,444	20 (1.4)	1.26 (0.53–2.98)	1.14 (0.47–2.76)	0.94 (0.39–2.28)	0.71 (0.30–1.68)	I
Third group (23.6≤ to <27.0)	1,443	79 (5.5)	3.71 (1.68–8.20)	3.30 (1.47–7.39)	2.16 (0.95-4.94)	1.33 (0.57–3.08)	I
Fourth group (≥27.0)	588	49 (8.3)	5.34 (2.36–12.09)	4.71 (2.07–10.72)	2.72 (1.18–6.25)	1.45 (0.62–3.41)	I
P-value for trend			<0.001	<0.001	<0.001	0.020	
Baseline BMI (quartiles)	4,323	155 (3.6)					
First quartile: (<20.9)	1,083	9 (0.8)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	I
Second quartile: (20.9≤ to <23.4)	1,079	15 (1.4)	1.26 (0.55–2.88)	1.16 (0.50–2.69)	1.04 (0.45–2.43)	0.83 (0.37–1.87)	I
Third quartile: (23.4≤ to <25.6)	1,081	51 (4.7)	3.38 (1.64–6.96)	3.05 (1.46–6.38)	2.28 (1.08-4.80)	1.41 (0.67–2.95)	I
Fourth quartile (≥25.6)	1,080	80 (7.4)	4.92 (2.40–10.05)	4.40 (2.14–9.03)	2.76 (1.33–5.74)	1.73 (0.84–3.55)	Ι
<i>P</i> -value for trend			<0.001	<0.001	<0.001	0.008	
Distinct body mass index (BMI) trajec occupation, current smoker, current a triglycerides, hemoglobin, creatinine a	tories were id Ilcohol drinke and uric acid	dentified using s er and married s at baseline. [¶] Mc	ix BMI tests (during 2011– tatus at baseline. [§] Model 3 odel 4: further adjusted for	-2016). *Model 1: adjustec 3: further adjusted for dia r fasting blood glucose ar	d for age and sex at basel gnosis of hypertension, us nd family history of diaber	ine. [‡] Model 2: further ad se of antihypertensive, to ces at baseline. ^{††} Model <u>5</u>	justed for race, tal cholesterol, s: sensitivity analysis
was carried out by excluding 196 par	ticipants (4,3	23 participants le	eft) who had missing info	rmation on baseline BMI.	#From the first to fourth	group, the proportions (of participants were
19.0, 33.4, 33.4 dria 13.0%, respectivel;	y. Irne aivisio	IN WERE CURSISIE	ant with the proportions c	JI DIVIL ITAJECIOLIES. KK, TELO	auve risk.		

Comparison with traditional methods based on baseline BMI

Previous studies predicting diabetes risk usually divided baseline BMI into groups. In the present study, we first divided baseline BMI into four groups according to the proportions of BMI trajectories. Then, we divided baseline BMI into quartiles, which were in accordance with many previous studies^{28,29}. As shown in Table 2, the incidence of new-onset diabetes progressively increased with higher BMI groups (0.8, 1.4, 5.5 and 8.3%, respectively), as well as with higher BMI quartiles (0.8, 1.4, 4.7 and 7.4%, respectively). However, in models adjusting for all potential covariates, no significant RRs for new-onset diabetes were found in the comparison between the fourth and first group (P = 0.393), as well as in the comparison between the fourth and first quartile (P = 0.137).

By contrast, as described above, there was a significant RR for new-onset diabetes in the comparison between "high" and "low" BMI trajectory (P = 0.014). In a sensitivity analysis, by excluding 196 participants who had missing data on baseline BMI, the RR for new-onset diabetes in the comparison between the "high" and "low" BMI trajectory remained significant (P = 0.032; Table 2).

Association between BMI trajectories identified using fewer BMI tests and diabetes risk

Finally, we explored whether BMI trajectories identified using the first four or three annual BMI tests were associated with new-onset diabetes in 2016 as well. When using the first four annual BMI tests, we also identified four linear BMI trajectories through the GBTM (Figure S1). As shown in Table 3, the incidence of new-onset diabetes progressively increased from the "low" to "high" BMI trajectory (0.7, 2.4, 5.1 and 10.1%, respectively). In models adjusting for all potential covariates, the RR for new-onset diabetes in the comparison between "high" and "low" BMI trajectory was 3.10 (95% CI 1.30–7.38). In a sensitivity analysis, by further excluding 114 participants who were diagnosed with diabetes or had no information about diabetes in 2014, the RR for new-onset diabetes in the comparison between "high" and "low" BMI trajectory remained significant (P = 0.028; Table 3).

Similarly, when using the first three annual BMI tests, we identified four linear BMI trajectories through the GBTM (Figure S1). The incidence of new-onset diabetes progressively increased from the "low" to "high" BMI trajectory (0.6, 2.2, 5.3 and 10.2%, respectively; Table 3). In models adjusting for all potential covariates, the RR for new-onset diabetes in the comparison between the "high" and "low" BMI trajectory was 2.71 (95% CI 1.19–6.19). In a sensitivity analysis, by further excluding 87 participants who were diagnosed with diabetes or had no information about diabetes in 2013, the RR for new-onset diabetes in the comparison between "high" and "low" BMI trajectory remained significant (P = 0.015; Table 3). These results suggest that BMI trajectories based on just four or three annual BMI tests were significantly associated with subsequent new-onset diabetes.

DISCUSSION

In the present longitudinal study carried out during 2011–2016, we identified four distinct developmental trajectories of BMI by the GBTM. We found that distinct BMI trajectories were significantly associated with new-onset diabetes in 2016. In addition, we found that BMI trajectories identified using the first four or three annual BMI tests were still significantly associated with new-onset diabetes in 2016. By contrast, no significant correlation was found between categories of baseline BMI and new-onset diabetes in 2016 after multivariate adjustment.

Although few previous studies have explored the association between BMI trajectories determined by GBTM and diabetes risk, the majority of them focused on the effect of childhood BMI trajectories on adult diabetes³⁰⁻³² or the effect of BMI trajectories before and during early pregnancy on gestational diabetes³³. For example, in a longitudinal cohort consisting of 2,449 USA residents, Tao et al.³¹ found that childhood (4-19 years) BMI trajectories have a significant impact on adult (20-51 years) diabetes. In an analysis of 8,009 Australian women aged 18-36 years, Kakoly et al.33 identified three distinct BMI trajectories across six surveys, and found that women in a high-rising trajectory were independently associated with gestational diabetes, as compared with those in a low-stable trajectory (OR 2.50, 95% CI 1.80-3.48). However, these studies carried out with children or pregnant women could not provide direct evidence for the primary prevention of diabetes in general adults.

For those studies directly carried out with general adults, three of them³⁴⁻³⁶ only showed the developmental trajectories of BMI before the diagnosis of diabetes, without examining the relative risk of diabetes across distinct BMI trajectories. In another study consisting of 24,875 Australian adults, Peter et al.37 showed that the incidence of diabetes was different across distinct BMI trajectories; however, this study did not adjust for potential covariates in the regression models. Therefore, whether distinct BMI trajectories predict diabetes risk could not be concluded from these studies. To date, there are just two studies^{18,19} that have examined the relative risk of diabetes by distinct BMI trajectories in general adults. Nevertheless, both studies spanned decades. It remains uncertain whether distinct BMI trajectories have an impact on diabetes risk, even in a shorter period, such as several years. In addition, we could not apply the conclusions to inform diabetes prevention in China, as both studies were carried out in Western countries. Thus, more evidence is really needed.

To our knowledge, the present study is the first to describe the association between distinct BMI trajectories and diabetes risk in general Chinese adults. We found that distinct BMI trajectories, even identified using the first four or three annual BMI tests, were significantly associated with new-onset diabetes. The findings suggest that distinct BMI trajectories might be an effective tool to predict future diabetes. In addition, no significant interactions between distinct BMI trajectories and sex, age and impaired fasting glucose in relation to risks of new-onset diabetes were observed in the present study, implying that the

Table 3 Relative risk and 95% of	confidence in	iterval of new-on	iset diabetes in 2016, by b	oody mass index trajectorie	s identified using fewer b	ody mass index tests	
Variables	\$.	Cases (%)	Model 1 Adjusted RR (95% Cl) [‡]	Model 2 Adjusted RR (95% CI) [§]	Model 3 Adjusted RR (95% Cl) [¶]	Model 4 Adjusted RR (95% Cl) ^{††}	Model 5 Adjusted RR (95% CI) ^{‡‡}
BMI trajectories (four tests) Traiectory 1-1 ow	4,475 1.031	167 (3.7) 7 (0.7)	1 00 (Ref)	1 (M) (Ref)	1 00 (Ref)	1 00 (Ref)	100 (Ref)
Trajectory 2: Moderate	1,487	35 (2.4)	2.66 (1.11–6.33)	2.47 (1.03–5.94)	2.07 (0.85–5.00)	1.60 (0.67–3.82)	3.65 (0.83–16.09)
Trajectory 3: Moderate-high	1,440	73 (5.1)	5.13 (2.22–11.88)	4.75 (2.03–11.10)	3.28 (1.39–7.79)	2.27 (0.96–5.38)	4.14 (0.95–18.13)
Trajectory 4: High	517	52 (10.1)	9.50 (4.05–22.26)	8.59 (3.64–20.27)	5.22 (2.20–12.43)	3.10 (1.30–7.38)	5.46 (1.20-24.82)
P-value for trend			<0.001	<0.001	<0.001	<0.001	0.018
BMI trajectories (three tests)	4,408	166 (3.8)					
Trajectory 1: Low	984	6 (0.6)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)
Trajectory 2: Moderate	1,493	33 (2.2)	2.49 (1.11–5.59)	2.32 (1.02–5.27)	2.04 (0.89-4.66)	1.52 (0.67–3.45)	2.35 (0.69-8.05)
Trajectory 3: Moderate-high	1,433	76 (5.3)	4.37 (2.00–9.56)	4.06 (1.84–8.95)	2.85 (1.27–6.39)	1.93 (0.86-4.34)	3.44 (1.00–11.84)
Trajectory 4: High	498	51 (10.2)	8.27 (3.73–18.36)	7.51 (3.36–16.82)	4.81 (2.13–10.87)	2.71 (1.19–6.19)	4.67 (1.35–16.15)
P-value for trend			<0.001	<0.001	<0.001	0.001	0.001
¹ For body mass index (BMI) traje larly, for BMI trajectories identified adjusted for race, occupation, cu total cholesterol, triglycerides, her Patients with diabetes might loss trajectories identified using four F identified using three BMI tests, E this diverter.	ctories identi d using three rrent smoker, moglobin, cr e or control t BMI tests, 114 87 participant	fied using four B BMI tests, 111 p current alcohol eatinine and uric their bodyweight 4 participants wh ts who were diag	MI tests, 44 participants w articipants were excluded drinker and married status acid at baseline. ^{1†} Model : To avoid the potential in o were diagnosed with di gnosed with diabetes or h	vere excluded from the tot from the total population s at baseline. "Model 3: fur 4: further adjusted for fast npact of new-onset diabet iabetes or had no informat iab no information about (al population due to the [*] Model 1: adjusted for ac ther adjusted for diagnosi ing blood glucose and far es during follow up on BN tion about diabetes in 201 diabetes in 2013 were furt	analysis requiring at least ge and sex at baseline. [§]) s of hypertension, use of mily history of diabetes at MI, sensitivity analysis was A were further excluded. her excluded. CI, confide	two BMI tests. Simi- Aodel 2: further antihypertensive, : baseline. ^{‡‡} Model 5: carried out. For BMI For BMI trajectories nce interval; RR, rela-
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effect of distinct BMI trajectories on diabetes risk was consistent across these factors.

The strengths of the present study included a longitudinal design, a large sample size, the availability of repeated measurements of BMI over time and the use of GBTM. However, some limitations should also be noted. First, the diagnosis of diabetes was based on fasting blood glucose without using the oral glucose tolerance test or hemoglobin A1c test, which is due to limited data collection for such a large cohort. Second, we did not collect information about the type of diabetes but, as the incident rate of type 1 diabetes is extremely low in Chinese adults³⁸, the majority of new-onset cases in our cohort should be type 2 diabetes. Third, approximately 30% of our study sample had missing values on at least one study variable. We thus used multiple imputation to increase the power of statistical tests. Multiple imputation is well recognized as a flexible and reliable tool for handling missing data, and has been widely used in clinical studies³⁹. Fourth, the current study included only Chinese adults. As the BMI level and its implications for diabetes differ widely across various racial/ethnic groups⁹, the present findings might not be generalizable to other populations.

In conclusion, the present results show that distinct BMI trajectories, even identified using just four or three annual BMI tests, are significantly associated with new-onset diabetes. The findings suggest that monitoring BMI trajectories over time might provide an important approach to identify subpopulations at higher risk for diabetes in general Chinese adults.

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DISCLOSURE

The authors declare no conflict of interest.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

Figure $S1 \mid$ Distinct body mass index trajectories identified using (a) four body mass index tests and (b) three body mass index tests.