RESEARCH



Joint association of the newly proposed dietary index for gut microbiota and sleep disorders with survival among US adult population with diabetes and pre-diabetes



Ke Si¹, Chuanqin Shi², Yajing Huang¹, Chuanfeng Liu¹, Jingwei Chi¹, Lili Xu¹, Ying Chen¹ and Yangang Wang^{1*}

Abstract

Background Diet and sleep disorders are associated with risks of metabolic diseases such as diabetes. The dietary index for gut microbiota (DI-GM) is a newly proposed index designed to assess dietary quality associated with maintaining a healthy gut microbiota. The authors aim to investigate the separate and joint prognostic effect of DI-GM and sleep disorders on the survival of US population with diabetes and pre-diabetes.

Methods Data were from the National Health and Nutrition Examination Survey (NHANES) 2007–2018 at baseline linked to the 2019 National Death Index records.

Dietary recall data were collected to calculate the DI-GM and sleep disorders were assessed by self-reported questionnaires. The Cox proportional hazard model were used to evaluate the associations between separate and joint prognostic effects of DI-GM and sleep disorders with mortality outcomes among diabetic and pre-diabetic patients.

Results A total of 10718 Participants with diabetes and pre-diabetes were ultimately included in this study (weighted population: 67,232,394, weighted mean age [SE]: 57.0 [0.1] years; weighted female proportion: 51.8%). Among these participants, higher DI-GM was more prevalent in those without sleep disorders. During the median follow-up of 13.3 years, 1448 deaths occurred, including 346 participants died from cancer, and 367 died from cardiovascular disease (CVD)...Multivariable models indicated that the joint effects of DI-GM (\geq 6) and no sleep disorders were associated with lower risks for all-cause (HR 0.53, 95% CI: 0.38–0.79) and CVD mortality (HR 0.36, 95% CI: 0.19–0.65).

Conclusions In a nationally representative sample of US population with diabetes and pre-diabetes, high DI-GM combined with no sleep disorders was associated with significantly reduced all-cause and CVD mortality risks.

Keywords Joint association, DI-GM, Sleep disorders, Survival, NHANES

*Correspondence:

Yangang Wang

wangyg1966@126.com

¹ Department of Endocrinology, Affiliated Hospital of Qingdao University, Qingdao 266003, China

² Marine Biomedical Research Institute of Qingdao, School of Medicine and Pharmacy, Key Laboratory of Marine Drugs, Chinese Ministry of Education, Ocean University of China, Qingdao 266003, China

Introduction

Diabetes is a significant public health problem worldwide, which is one of the leading causes of death in the United States. Epidemiological studies have demonstrated the association between diabetes and a variety of physical health conditions including cancer [1], cardiovascular disease (CVD) [2], renal diseases [3] as well as mortality from physical disorders. According to the latest statistics from the Centers for Disease Control and



© The Author(s) 2025. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/.

Prevention (CDC), as of 2021, approximately 97.6 million people aged 18 or older in the United States had pre-diabetes, accounting for 38.0% of the adult population [4]. Pre-diabetes may progress to diabetes, increasing the risk of CVD and death. Growing evidence indicates that disturbed sleep architecture is associated with impaired glucose homeostasis, decreased insulin secretion and an increased risk of diabetes [5, 6]. Recent studies have also found an association between poor subjective sleep and deterioration in blood glucose control [7]. Engeda et al. reported that pre-diabetes was associated with trouble maintaining sleep, waking up too early, and short sleep duration [8]. To our knowledge, the evidence regarding the impact of sleep disorders on the mortality risks of individuals with diabetes and pre-diabetes remains limited.

In recent years, an increasing number of studies have demonstrated that the gut microbiota and their metabolites play a crucial role in the development of diabetes through altering the intestinal mucosal barrier, influencing insulin secretion, and body immune regulation [9–11]. Healthy dietary patterns influence the composition and function of the gut microbiota [12]. Currently, one promising strategy for diabetes treatment is modulating gut microbiota composition by dietary interventions. The Mediterranean diet (MED), significantly alters the composition and function of the gut microbiota, increasing the production of short-chain fatty acids (SCFAs) [13]. The Dietary Approaches to Stop Hypertension (DASH) diet, with its high dietary fiber content [14], stimulates the growth of SCFA-producing bacteria, offering protective effects against diabetes [15]. SCFAs have been proved to enhance insulin sensitivity and reduce postprandial blood glucose levels [16]. A systematic review and dose-response meta-analysis found that high adherence to the DASH diet was significantly associated with a reduced risk of diabetes [17]. The MED intervention has been shown to increase the relative abundances of Lachnospiraceae NK4A136, which could improve patients'blood glucose levels, insulin levels, and insulin resistance [18]. With the deepening of research, recently, Kase and his colleagues developed a novel dietary index for gut microbiota (DI-GM) through a comprehensive literature review of longitudinal studies in order to measure the potential impact of diet on gut microbiota [19]. The DI-GM identified 14 foods or nutrients that have either beneficial or unfavorable effects on gut microbiota, with higher scores reflecting healthier gut microbiota. In addition, DI-GM was associated positively with markers of gut microbiota diversity biomarkers specifically urinary enterodiol and enterolactone, indicating that the dietary patterns captured by the DI-GM may contribute to a more diverse gut microbiota [19]. As a reliable tool, DI-GM can evaluate the impact of specific dietary modifications on the composition, and diversity of the gut microbial community, supporting microbiome-targeted dietary recommendations and personalized nutrition strategies for disease prevention and management. The specific impact of DI-GM on diabetes risk has not yet been thoroughly investigated.

Various evidence has shown that sleep has a complex interaction with microbial communities. The microbiota-gut-brain axis contributes to the regulation of sleep behavior both directly and indirectly and may play a critical role in the etiology and pathogenesis of sleep disorders. As a pathway for bidirectional brain-gut communication, the vagus nerve can regulate and receive information from the digestive system. Lactobacillus rhamnosus JB1 has been proved to ameliorate depression and anxiety through vagus nerve-dependent pathways and modulate GABA receptor expression in the brain [20]. Benzodiazepines targeting GABA receptors such as zolpidem are used to treat insomnia [21]. By modulating bacterial metabolites, neuronal signaling, endocrine signaling, and immune responses, the gut microbiota exerts an influence on sleep-wake behavior, in turn, lack of sleep leads to gut microbiota dysbiosis with changes in gut microbiome composition [1, 22]. Microbiota-targeted interventions may be promising novel strategies for themanagement of sleep disorders. To date, few studies have considered the influence of the joint effect of DI-GM and sleep disorders on the mortality risks of patients with diabetes and pre-diabetes. This study aimed to investigate the independent and joint associations of the newly proposed DI-GM and sleep disorders with all-cause, cancer, and CVD mortality among the US adult population with diabetes and pre-diabetes, utilizing National representative data from the NHANES Study. Our findings may bring benefits to the development of relevant evidencebased guidelines in patients with diabetes and pre-diabetes to improve their quality of life and minimize the risk of severe health outcomes.

Methods

Study design and population

This study utilized data collected from the National Health and Nutrition Examination Survey (NHANES), a series of cross-sectional, complex, multi-stage surveys, conducted by the Centers for Disease Control and Prevention (CDC) [23]. The NHANES study has been conducted on 2-year cycles since 1999 to monitor the health and nutritional status and diet of the US population. The protocol for the NHANES study was approved by the Institutional Review Board of the National Center for Health Statistics (NCHS), and written informed consent was provided by all participants. Participants in this study were invited to engage in a face-to-face interview, undergo a comprehensive series of physical examinations, and participate in laboratory tests in a mobile examination facility [24]. Six survey cycles of data in the NHANES database from 2007 to 2018 were analyzed in this study. Follow-up for mortality status was conducted for all individuals until the end of December 2019. Participants with missing data on glucose and HbA1c, cancer investigation, sleep questionnaire, dietary recall, and incomplete mortality follow-up data were excluded from this study. In total, 10718 participants with diabetes and pre-diabetes aged 20 years or older were eligible in the final analysis (Fig. 1).

Diagnosis of diabetes and pre-diabetes

According to the American Diabetes Association (ADA) criteria [25], diabetes was diagnosed based on the presence of any of the following criteria: (a) fasting plasma glucose (FPG) level \geq 126 mg/dL or HbA1c level \geq 6.5%; (b) a response 'yes' to the question,"Has a doctor ever told you that you have diabetes?"or"Are you currently taking insulin or oral hypoglycemic medication?". Pre-diabetes was diagnosed based on the presence of any of the following criteria: (a) FPG level between 100 mg/dL and 125 mg/dL or HbA1c level between 5.7% and 6.4%; (b) a response 'yes' to

the question,"Has a doctor ever told you that you have pre-diabetes?".

Dietary index for gut microbiota

Kase et al. developed a novel dietary index for gut microbiota (DI-GM), which has been designed to reflect the dietary composition associated with the gut microbiota profiles [19]. They considered 14 foods or nutrients as integral components of the DI-GM including beneficial components like fermented dairy, chickpeas, soybean, fiber, cranberries, whole grains, avocados, broccoli, coffee and green tea, as well as unfavorable components like red meat, processed meat, refined grains, and high-fat diet (mainly soybean oil, $\geq 40\%$ of energy from fat). Studies suggest high-fat diets are associated with reduced microbial abundance and increased gut permeability [26]. High-fat diets rich in meat protein may promote different and less diverse populations of sulphur-metabolising bacteria [27]. Thus high-fat diets act as unfavourable components of DI-GM. The components and scoring criteria for the DI-GM are detailed in the Supplementary Table S1. For beneficial to gut microbiota items, a value of 1 was assigned if the intake exceeded the sex-specific median, otherwise a value of 0 was given. For unfavorable to gut microbiota items, a value of 0 was assigned if the intake exceeded the sex-specific median or 40% (for High-fat diet), otherwise a value of 1 was given. The total DI-GM score ranges from 0 to 13, with beneficial items scoring from 0 to 9 and unfavorable items from 0 to 4. A higher



Fig. 1 Flowchart of the study inclusion and exclusion of participants

score indicating a more favorable gut microbiota profile. In NHANES, The components of the DI-GM were derived from data obtained through two 24-h dietary recalls of all food and drink consumed on the day before the interview (from midnight to midnight), with a time interval varying from 3 to 10 days. The first dietary recall interview was collected face-to-face in the Mobile Examination Center (MEC) and the second interview was collected by telephone 3 to 10 days later. The mean intake of foods, food groups, and nutrients from the two 24-h recalls were used to construct the DI-GM. Previous studies have demonstrated that DI-GM score of ≥ 6 shows a significant negative correlation with adverse clinical outcomes, while the group scoring 0-3 is considered to be at high risk [28, 29]. In the present study, the DI-GM scores were categorized into three groups: 0-3, 4-6, and ≥ 6 .

Sleep disorders

The response to "Have you ever told a doctor or other health professional that you have trouble sleeping?" or "Have you ever been told by a doctor or other health professional that you have a sleep disorder?" in the Sleep Disorder Questionnaire, were used to assess sleep disorder and then participants were classified according to the answers (yes or no). Sleep time was categorized based on the question"How many hours of sleep do you usually get on weeknights or work nights?"The recorded sleep time was divided into short sleep (less than 7 h per night), normal sleep (7–9 h per night), and long sleep (more than 9 h per night).

Ascertainment of mortality

The mortality data in this study were linked to the National Death Index (NDI) provided by NCHS, through December 31, 2019. The cause of death was coded by the 10th International Classification of Diseases (ICD-10). The study outcomes were all-cause mortality, cancer, and cardiovascular disease (CVD) mortality. The duration of follow-up was determined from the baseline interview to the date of death or December 31, 2019.

Covariates

The selection of covariates was based on the knowledge regarding the factors influencing the prognosis of cancer survivors. Sociodemographic characteristics consisted of basic information, including age, gender(male or female), race/ethnicity(Mexican American, other Hispanic, non-Hispanic White, non-Hispanic Black, and other races), and educational level(Less Than 9th Grade, 9-11th Grade, High School Grad, Some College degree, College Graduate or above), family poverty income ratio (total family income divided by the poverty threshold; < 1.3, 1.3 to \leq 3.5, > 3.5). Body mass index(BMI) was calculated by

dividing weight (kg) by height in square meters and was classified as < 30, ≥ 30 kg/m². Total calorie intake on the sum of two days (DR1TOT and DR2TOT) were utilized for analysis. Participants were considered non-smokers if they had either never smoked or had consumed fewer than 100 cigarettes in their lives. Smoking status was categorized, with "Yes" indicating participants who had smoked at least 100 cigarettes in their lives, and "No" for those who had either never smoked or had consumed fewer than 100 cigarettes. For alcohol use, the threshold for classifying"No"and"Yes"was based on having at least 12 alcoholic drinks per year. A history of hypertension and hypercholesterolemia was self-reported by participants who had been diagnosed by a physician or other health professional in a standardized medical condition questionnaire.

Statistical analysis

Given the complex multistage stratified probability survey design employed by NHANES, all analyses in this study incorporated sample weights, clustering, and stratification to accurately estimate appropriate variance and ensure national representation of results. The baseline characteristics were presented according to DI-GM classification $(0-3,4-6, \geq 6)$, with the survey-weighted means and standard errors (SE) to describe continuous variables and weighted percentages with their 95% CI to describe categorical variables. In addition, to mitigate the impact of missing variables for covariates, we utilized the multiple imputation approach implemented in the"mice"package in R. Multivariable Cox proportional hazard regression model was utilized to determine the hazard ratios (HR) and their 95% CI for the associations of the DI-GM and sleep disorders with all-cause, cancer, and CVD mortality respectively. Fully adjusted models were adjusted for age, sex, race and ethnicity, educational attainment, family poverty income ratio, body mass index, total calorie intake, sleep time, smoking status, alcohol use, hypertension, and hypercholesterolemia. To explore the joint associations, participants were categorized based on the presence of the DI-GM and sleep disorders to assess mortality risks using multivariable Cox proportional hazards regression models with adjustments for the same set of covariates. All analyses were conducted in subpopulations according to age (< 65 years, \geq 65 years), sex (male, female), and race and ethnicity (Mexican American, other Hispanic, non-Hispanic White, non-Hispanic Black and other race), using Cox proportional hazards regression models to examine the stability of the results. Finally, sensitivity analysis was performed by excluding participants who died within the initial 2-year follow-up, and had a history of cancer or CVD disease to lessen the probability of reverse causation.

All analyses were conducted using R version 4.4.1 and SPSS (IBM) version 27, and a two-sided P < 0.05 was set for a significant difference.

Results

Baseline characteristics

A total of 10718 individuals with diabetes and pre-diabetes were ultimately included in this study (weighted population: 67,232,394, weighted mean age [SE]: 57.0 [0.1] years; weighted female proportion: 51.8%). The characteristics of the population categorized according to their DI-GM scores are presented in Table 1. Notably, participants with higher DI-GM (\geq 6) appeared to be more likely to be older, female, non-Hispanic White, have higher calories intake and normal sleep (7–9 h per night), and less likely to have hypertension and sleep disorders. The joint analysis of the prevalence of DI-GM and sleep disorders showed a significant downward trend in the prevalence of sleep disorders among participants with diabetes and pre-diabetes as DI-GM increased (Trend test *P* = 0.016) (Fig. 2).

Relationship between DI-GM, sleep disorders, and mortality

During the follow-up period of up to 13.3 years (median, 6.4 years), 1448 deaths occurred, including 346 participants died from cancer, and 367 died from CVD. Participants with either high DI-GM (≥ 6) or no sleep disorders had reduced risks of all-cause and CVD mortality (Table 2). After adjusting for covariates, the HRs for all-cause, cancer, and CVD mortality among individuals with DI-GM (\geq 6) were 0.71 (95% CI: 0.57–0.93), 0.92 (95% CI: 0.70-1.54) and 0.63 (95% CI: 0.49-0.85), respectively, compared to those with DI-GM (0-3). Each 1-point increase in DI-GM was associated with 6% and 9% reduction in the risk of death from all-cause and CVD mortality, respectively. As shown in Fig. 3, we further considered diabetic and pre-diabetic individuals separately, finding that each 1-point increase in DI-GM was associated with 8% and 10% reduction in the risk of all-cause mortality, respectively. Additionally, the risks of all-cause and CVD mortality among individuals with diabetes and pre-diabetes decreased as the beneficial to gut microbiota increased (HR 0.84, 95% CI: 0.80-0.93; HR 0.81, 95% CI: 0.77-0.92), respectively, but the unfavorable to gut microbiota was not associated with an increased risk of mortality (Table 2). Meanwhile, the HRs for all-cause, cancer, and CVD mortality among individuals with sleep disorders were 1.28 (95% CI: 1.10-1.37), 1.10(95% CI: 0.82-1.47), and 1.44 (95% CI: 1.21-1.84), respectively, compared to those without sleep disorders.

Joint association of DI-GM, sleep disorders with mortality

In the joint analyses, diabetic and pre-diabetic participants with with DI-GM (≥ 6) and no sleep disorders had the lowest risks of all-cause and CVD mortality (Fig. 4). Specifically, compared with participants with DI-GM (0–3) and sleep disorders, the HRs for all-cause and CVD in the participants with DI-GM (≥ 6) and no sleep disorders were 0.53 (95% CI: 0.38–0.79) and 0.36 (95% CI: 0.19–0.65), respectively (Table 3). To ensure the reliability of our findings, a sensitivity analysis was performed. The main results remained similar after excluding the participants with a history of cancer (n = 1269), and participants with a history of CVD (n = 964) (Supplementary Table S2-S4).

Subgroup analyses

In subgroup analyses, the results of the joint analysis of DI-GM and sleep disorders are detailed in Supplementary Table S5 and Supplementary Table S6. The joint effect of DI-GM and sleep disorders on all-cause mortality was significant in participants aged <65 years (HR, 0.45; 95% CI, 0.23–0.81) and ≥65 years (HR, 0.59; 95% CI, 0.38–0.91), both male (HR, 0.54; 95% CI, 0.33–0.87) and female (HR, 0.51; 0.32-0.82) participants, participants with BMI < 30 (HR, 0.51; 95% CI, 0.34-0.79), Mexican-American participants (HR, 0.31; 95% CI, 0.12-0.85), other Hispanic participants (HR, 0.25; 95% CI, 0.11–0.57), non-Hispanic White participants (HR, 0.53; 95% CI, 0.34-0.81), and non-Hispanic Black participants (HR, 0.58; 95% CI, 0.35-0.94) (Supplementary Table S5). The association between DI-GM, sleep disorders, and CVD mortality was more pronounced in participants both aged <65 years (HR, 0.28; 95% CI, 0.11-0.62) and ≥ 65 years (HR, 0.48; 95% CI, 0.28-0.82), both male (HR, 0.50; 95% CI, 0.28-0.89) and female (HR, 0.31; 0.17-0.57) participants, participants with both BMI <30 (HR, 0.44; 95% CI, 0.27–0.73), and BMI ≥ 30 (HR, 0.29; 95% CI, 0.13-0.65), other Hispanic participants (HR, 0.23; 95% CI, 0.08-0.66), non-Hispanic Black participants (HR, 0.26; 95% CI, 0.11-0.64) and other race (HR, 0.13; 95% CI, 0.06-0.75)(Supplementary Table S6). Additionally, the combined impact of DI-GM and sleep disorders on all-cause mortality was similar in both diabetic and pre-diabetic participants. Notably, this joint effect was more strongly associated with CVD mortality in pre-diabetic participants (HR 0.32, 95% CI: 0.16-0.66), compared to those with diabetes (HR 0.49, 95% CI: 0.29-0.81) (Supplementary Table S6).

Discussion

This study is the first to investigate the joint effect of DI-GM and sleep disorders on the mortality risks in a nationally representative US population with diabetes

Table 1 Baseline characteristics of US population stratified by groups of DI-GM, NHANES 2007 to 2018

Characteristics	Total	No. of participants by	s by DI-GM group (weighted %)ª		
		DI-GM 0–3	DI-GM 4–6	DI-GM≥6	
Overall	10718	1970 (17.4)	4958 (43.9)	3790 (38.7)	
Age, mean (SE), year	57.0 (0.1)	56.3 (0.2)	56.2 (0.1)	58.9 (0.2)	
Sex, n (%)					
Men	5279 (48.2)	1058 (52.7)	2462 (48.1)	1759 (46.1)	
Women	5439 (51.8)	912 (47.3)	2496 (51.9)	2031 (53.9)	
Race and ethnicity, n (%)					
Mexican American	1699 (9.0)	264 (8.1)	893 (0.8)	542 (7.3)	
Other Hispanic	1161 (5.7)	210 (5.3)	558 (6.2)	393 (5.2)	
Non-Hispanic White	3948 (51.8)	702 (51.3)	1692 (48.1)	1554 (56.1)	
, Non-Hispanic Black	2751 (25.0)	629 (29.6) 1330 (26.8)		792 (20.8)	
Other Race	1159 (8.6)	165 (5.7)	485 (8.1)	509 (10.5)	
Educational attainment, n (%)					
Less Than 9th Grade	1419 (13.2)	255 (13.0)	631 (13.2)	533 (13.4)	
9-11th Grade	1678 (15.7)	312 (15.4)	753 (15.9)	613 (15.4)	
High School Grad	2419 (22.7)	412 (21.0)	1158 (24.0)	849 (22.2)	
Some College degree	2917 (27.0)	566 (28.1)	1343 (26.5)	1008 (26.7)	
College Graduate or above	2285 (21.4)	425 (22.4)	1073 (20.4)	787 (22.3)	
Eamily poverty income ratio, n (%)		,		()	
< 1.3	3709 (34.6)	688 (34.1)	1719 (34.5)	1302 (34.9)	
> 13 to 35	3698 (33.7)	705 (35.7)	1674 (33.8)	1319 (32.7)	
> 35	3311 (317)	577 (30.1)	1565 (31.7)	1169 (32.5)	
$BMI ka/m^2 n (%)$	5511 (51.7)	577 (50.17	1565 (51.7)	1105 (52.5)	
< 30	7872 (73 7)	1/36 (73 7)	3688 (75.0)	27/18 (72.2)	
> 30	2846 (263)	534 (26 3)	1270 (25.0)	1042 (27.8)	
Total calories intake mean (SE) kcal	3961 28 (15.0)	3903 85 (33 8)	3907 00 (22 9)	4048 65 (24 3)	
Smoking status n (%)	5561.26 (15.6)	5765.65 (55.6)	5507.00 (22.5)	10 10:00 (2 1.0)	
No	5183 (/0.2)	990 (48 5)	2400 (40 3)	1784 (494)	
Vec	5535 (50.8)	980 (51 5)	2540 (57)	2006 (50.6)	
Alcoholic use p (%)	(0.0)	500 (51.5)	2545 (5.7)	2000 (50.0)	
No	4614 (71.2)	850 (73.6)	2130 (71.6)	1634 (69.7)	
Voc	1909 (29.9)	311 (26 4)	2130 (71.0)	663 (30.0)	
Hypertension n (%)	1000 (20.0)	511 (20.4)	054 (20.4)	003 (30.0)	
No	7939 (75 3)	1/28 (73.8)	3617 (74.9)	2703 (76 4)	
Vac	2880 (24.8)	542 (26 2)	13/1 (25.2)	2735 (70.4)	
Humarlinidamia n (04)	2000 (24.0)	542 (20.2)	1541 (25.2)	997 (23.0)	
No	5652(50.6)	1011 (50.8)	2506 (52.0)	2045 (55.0)	
Vac	5067 (43.0)	060 (40 2)	2360 (32.5)	1745 (45.0)	
Sleep disorders p. (%)	5007 (45.5)	500 (45.2)	2502 (47.1)	1745 (45.0)	
	7402 (65 7)	1 207 (62 0)	2425 (64 4)	2600 (60 4)	
No	7402 (05.7)	1207 (03.0)	1522 (25.6)	2000 (08.4)	
Clean time p (04)	5510 (54.5)	065 (57.0)	1525 (55.0)	1110 (51.0)	
sieep time, n (%)	2020 (22.0)	700 (26 1)	1025 (25 1)	1204 (21.6)	
< /	5656 (55.9)	709 (56.1)	1855 (55.1)	1294 (31.6)	
7-9	700 (C 0)	1089 (50.2)	2784 (58.8)	2298 (03.3)	
> y	/U9 (b.U) 4 05 (0.02)	1/2 (7.0)	222 (0.1)	190 (5.1) 6 77(0.00)	
Dependicial to quit microbiotal and (SE)	4.90(0.02)	2.55(U.UZ)	4.51(0.01)	0.77(0.02)	
Demencial to gut microbiota, mean (SE)	2.29 (0.03)	0.97 (0.04)	1.85 (0.03)	5.50 (U.U3)	
Disketes p (%)	1.35 (0.01)	2.42 (U.U2)	1.34 (U.UT)	0.79 (0.01)	
Diabetes, n (%)	4883 (41.3)	902 (46.7)	2310 (42.1)	1011 (38.1)	
Pre-diabetes, n (%)	5835 (58.7)	1008 (53.3)	2648 (57.9)	2179 (61.9)	

All means and SEs for continuous variables and percentages for categorical variables were weighted. The DI-GM score comprises BGMS and UGMS, categorized into three groups: 0–3, 4–5, and \geq 6

Abbreviations: BMI body mass index (calculated as weight in kilograms divided by height in meters squared), DI-GM dietary index for gut microbiota, h/d hours per day, NHANES National Health and Nutrition Examination Survey, SE Standard Error

^a The weighted percentages may not reach exactly 100% due to missing data



Fig. 2 Joint prevalence of DI-GM and sleep disorders in a nationally representative sample of US population with diabetes and pre-diabetes, NHANES 2007 to 2018. Abbreviations: DI-GM, dietary index for gut microbiota; NHANES, National Health and Nutrition Examination Survey. Data were weighted to be nationally representative

and pre-diabetes. Our results revealed that about 38.7% had DI-GM score of more than 6 points, and 30.9% had sleep disorders. Over the follow-up period of 13.3 years (median, 6.4 years), high DI-GM, increase in beneficial to gut microbiota, and no sleep disorders were associated with decreased risks of all-cause and CVD mortality. In the joint analysis, individuals with diabetes and pre-diabetes with DI-GM (0-3) and sleep disorders had a higher risk of all-cause and CVD mortality than those with a single risk factor. The significance of the association was consistently observed across sensitivity and subgroup analyses. The association between the joint effect of DI-GM and sleep disorders and CVD mortality was more pronounced in participants with pre-diabetes than diabetes. These findings underscore the potential of personalized dietary interventions focused on DI-GM coupled with sleep modulation for mortality risks prevention, particularly in diabetes and pre-diabetes population.

Previous studies have explored the relationship among sleep disorders, diabetes, and mortality [30-32]. A study by Schantz et al. [33] found that the presence of both diabetes and frequent sleep disturbances was associated with a higher risk of all-cause mortality compared to either condition alone. A large single-center cohort study reported that sleep duration was a significant indicator of all-cause, extended CVD, and non-extended CVD mortality among patients with diabetes, with the lowest risks observed for patients with 5-7 h of sleep [34]. Another cohort study of the UK Biobank discovered that healthy sleep patterns could reduce the negative impact of metabolic control on CVD death in people with diabetes [32]. People with diabetes and sleep disorders are susceptible to cardiac hypoxia, which worsens the severity of heart failure [35]. Our study further validated the detrimental impact of sleep disorders on the increased risk of all-cause and CVD mortality in individuals with diabetes independently, especially considering prediabetic patients. Similar results were not observed for cancer mortality. Laboratory studies in healthy adults have shown that sleep deprivation is associated with a 40% reduction in glucose clearance rate and increased sympathetic activity compared to a refreshed state, and the latter may deteriorate the status of insulin resistance [36, 37]. Sleep deprivation in people with diabetes is likely to worsen complications and interfere with blood glucose control and management, leading to an excess risk of mortality. Therefore, these findings indicate that sleep disorders could elevate the risk of mortality among patients with diabetes and prediabetes.

The influence of diet on the gut microbiota and its potential role in diabetes and pre-diabetes has been a significant focus of research. Fiber, defined beneficial component in DI-GM, has been reported to increase the proportion of beneficial gut microbes, and improve blood glucose homeostasis, serum metabolomics and systemic inflammation in patients with diabetes [38]. A high-fat diet, having unfavorable effects on gut microbiota within DI-GM, may contribute to diabetes through impaired glucose and lipid metabolism, which impairs the function of key metabolic organs such as the liver, pancreas, and adipose tissue [39]. The Mediterranean Diet (MED) is a plant-based eating pattern that includes a rich variety of vegetables, fruits, whole grains, legumes, and nuts, which are beneficial components in DI-GM. A study has shown that upgraded adherence to a MED pattern reduced the proportion of people with diabetes at high risk of CVD [40]. Recently, adults with pre-diabetes were randomly assigned to follow MED or gut microbiome modulates

Table 2 Association of DI-GM and sleep disorders with all-cause, cancer and CVD mortality among US population with diabetes and pre-diabetes, NHANES 2007 to 2018

Hazard ratio (95% CI), <i>P</i> value	Mortality outcome All-cause mortality								
	0–3	4–6	≥6	Per 1 point increase	Beneficial to gut microbiota	Unfavorable to gut microbiota	No	Yes	
	Death/No	311/1970	669/4958	468/3790	NA	NA	NA	954/7402	494/3316
Age adjusted ^a	Reference	0.86(0.70-1.06)	0.69(0.53-0.88)	0.90(0.85–0.95)	0.83(0.78–0.88)	1.05(0.98-1.13)	Reference	1.28(1.09–1.33)	
Р		0.165	< 0.001	< 0.001	< 0.001	0.153		< 0.001	
MV model 1 ^b	Reference	0.89(0.75-1.11)	0.71(0.57-0.92)	0.92(0.88–0.96)	0.83(0.79-0.91)	1.06(0.98-1.14)	Reference	1.27(1.09–1.35)	
Р		0.302	< 0.001	< 0.001	< 0.001	0.054		< 0.001	
MV model 2 ^c	Reference	0.91(0.76-1.13)	0.71(0.57–0.93)	0.94(0.86-0.97)	0.84(0.80-0.93)	1.08(1.02-1.19)	Reference	1.28(1.10-1.37)	
Р		0.315	< 0.001	< 0.001	< 0.001	0.073		< 0.001	
	Mortality	Mortality outcome							
Hazard ratio	Cancer mo	ortality							
(95% CI), <i>P</i> value	DI-GM group						Sleep disorders		
	0–3	4–6	≥6	Per 1 point increase	Beneficial to gut microbiota	Unfavorable to gut microbiota	No	Yes	
Death/No	70/1970	159/4958	111/3790	NA	NA	NA	234/7402	112/3316	
Age adjusted ^a	Reference	0.93(0.70-1.54)	0.85(0.66-1.22)	0.92(0.86-1.02)	0.97(0.79-1.26)	1.04(0.74–1.39)	Reference	1.15(0.93-1.50)	
Ρ		0.682	0.236	0.097	0.136	0.523		0.279	
MV model 1 ^b	Reference	0.96(0.75-1.69)	0.89(0.70-1.39)	0.96(0.91-1.14)	0.97(0.80-1.29)	1.03(0.73-1.38)	Reference	1.11(0.86-1.48)	
Р		0.793	0.415	0.226	0.212	0.623		0.424	
MV model 2 ^c	Reference	0.98(0.77-1.74)	0.92(0.70-1.54)	0.97(0.91-1.18)	0.98(0.83-1.38)	1.03(0.71-1.39)	Reference	1.10(0.82-1.47)	
Р		0.902	0.589	0.297	0.447	0.764		0.472	
	Mortality outcome								
Hazard ratio	CVD mortality								
(95% CI), P value	DI-GM group						Sleep disorders		
	0–3	4–6	≥6	Per 1 point increase	Beneficial to gut microbiota	Unfavorable to gut microbiota	No	Yes	
Death/No	82/1970	176/4958	109/3790	NA	NA	NA	224/7402	143/3316	
Age adjusted ^a	Reference	0.83(0.66-1.04)	0.61(0.47-0.79)	0.89(0.84-0.95)	0.81(0.76-0.86)	1.08(0.82-1.02)	Reference	1.39(1.11–1.78)	
Р		0.117	< 0.001	< 0.001	< 0.001	0.121		< 0.001	
MV model 1 ^b	Reference	0.85(0.68-1.08)	0.63(0.49-0.83)	0.90(0.85-0.96)	0.81(0.77-0.89)	1.10(0.98–1.20)	Reference	1.41(1.17–1.85)	
Р		0.179	< 0.001	< 0.001	< 0.001	0.053		< 0.001	
MV model 2 ^c	Reference	0.85(0.68-1.11)	0.63(0.49-0.85)	0.91(0.86-0.97)	0.81(0.77-0.92)	1.09(0.95-1.19)	Reference	1.44(1.21-1.84)	
Р		0.184	< 0.001	< 0.001	< 0.001	0.052		< 0.001	

Abbreviations: DI-GM dietary index for gut microbiota, NHANES the National Health and Nutrition Examination Survey, CI confidence interval

^a Adjusted for age

^b Multivariable adjusted model additionally adjusted for sex, race and ethnicity, educational attainment, family poverty income ratio, body mass index, total calories intake, sleep time, smoking status, and alcohol use

^c Additionally adjusted for hypertension, and hypercholesterolemia

 d The DI-GM scores were categorized into three groups: 0–3, 4–6, and $\geq\!6$



Fig. 3 Dose–response association of DI-GM with all-cause mortality among US populations with diabetes and pre-diabetes (A), diabetes (B), and pre-diabetes (C), NHANES 2007 to 2018. The solid and dashed lines illustrate the estimated hazard ratios along with their respective 95% confidence intervals. Abbreviations: DI-GM, dietary index for gut microbiota; NHANES, the National Health and Nutrition Examination Survey. Adjusted for age, sex, race and ethnicity, educational attainment, family poverty income ratio, body mass index, total calories intake, sleep time, smoking status, alcohol use, hypertension, and hypercholesterolemia

the effects of a personalized postprandial-targeting (PPT) diet in a 6-month dietary intervention. Studies indicated that the PPT diet induced more pronounced changes in the composition of the gut microbiome and improved glycemic control significantly more than a MED diet, which further affected cardiometabolic markers in prediabetes [41, 42]. Currently, available dietary indicators show inconsistent results in reflecting the diversity and abundance of the gut microbiota [43, 44]. DI-GM, as a newly proposed indicator, was designed to reflect the diversity of the gut microbiota and to measure diet quality about gut microbiome health purposely. Our study indicated that high DI-GM significantly reduced the risk of all-cause and CVD mortality in patients with diabetes and pre-diabetes. Based on a comprehensive dietary pattern rather than a single food or nutrient evaluation might provide stronger evidence of the rational dietary recommendation for mortality prevention.

The singular effects of diet or sleep disorders on the prevention of diabetes and pre-diabetes have been wellestablished, however, no studies have yet examined their joint effects on these conditions. Joint analysis allows us to provide more comprehensive survival guidance for patients with diabetes and pre-diabetes by exploring the



Fig. 4 Joint association of DI-GM and sleep disorders with all-cause, cancer, and CVD mortality participants with diabetes and pre-diabetes, NHANES 2007 to 2018. Hazard ratios (solid symbols) with 95% CIs (error bars) of joint association between DI-GM and sleep disorders for all-cause (A), cancer (B), and CVD (C) mortality calculated using multivariable Cox regression models adjusted for age, sex, race and ethnicity, educational attainment, family poverty income ratio, body mass index, total calories intake, sleep time, smoking status, alcohol use, hypertension, and hypercholesterolemia. Abbreviations: DI-GM, dietary index for gut microbiota; NHANES, the National Health and Nutrition Examination Survey; CI, confidence interval; CVD, cardiovascular disease

unique and combined impact of each factor on mortality outcomes. Our results revealed that the combination of high DI-GM (\geq 6) and no sleep disorders could significantly decrease the risks of all-cause and CVD mortality among patients with diabetes and pre-diabetes as compared with either factor alone. According to our subgroup analyses, the stratification of the study population into two subgroups based on diabetes status did not significantly influence the primary outcomes. Notably, this joint effect was more strongly associated with CVD mortality in pre-diabetic participants compared to those with diabetes. The Da Qing Diabetes Study indicated that lifestyle interventions, including diet alone and diet plus exercise, over 6 years reduced the risk of CVD mortality and CVD

Mortality outcome	DI-GM group	Death/No	Hazard ratio (95% CI), P value					
			Age adjusted ^a		MV model 1 ^b		MV model 2 ^c	
All-cause mortality								
Sleep disorders	0–3	113/683	Reference		Reference		Reference	
	4–5	235/1523	0.83(0.58–1.17)	0.280	0.83(0.58–1.16)	0.255	0.84(0.59–1.19)	0.287
	≥6	146/1110	0.66(0.46-0.90)	< 0.001	0.69(0.48–0.93)	0.009	0.69(0.49–0.96)	0.012
No sleep disorders	0–3	198/1287	0.71(0.51-1.01)	0.054	0.70(0.51-0.99)	0.041	0.71(0.52–0.99)	0.048
	4–5	434/3435	0.65(0.48-0.85)	< 0.001	0.68(0.49–0.88)	< 0.001	0.69(0.49–0.92)	0.003
	≥6	322/2680	0.52(0.36-0.75)	< 0.001	0.54(0.39–0.75)	< 0.001	0.53(0.38–0.79)	< 0.001
P for trend				< 0.001		< 0.001		< 0.001
Cancer mortality								
Sleep disorders	0–3	25/683	Reference		Reference		Reference	
	4–5	52/1523	0.87(0.46-1.38)	0.306	0.86(0.50-1.46)	0.541	0.91(0.52–1.59)	0.742
	≥6	30/1110	0.74(0.44-1.24)	0.253	0.82(0.47-1.42)	0.472	0.88(0.50-1.53)	0.616
No sleep disorders	0–3	45/1287	0.91(0.50-1.56)	0.712	0.91(0.50–1.54)	0.870	0.93(0.54-1.61)	0.745
	4–5	107/3435	0.76(0.45-1.20)	0.240	0.83(0.51-1.35)	0.447	0.82(0.50-1.33)	0.413
	≥6	81/2680	0.65(0.41-1.03)	0.065	0.72(0.4-1.18)	0.193	0.76(0.46-1.25)	0.256
P for trend				0.114		0.207		0.258
CVD mortality								
Sleep disorders	0–3	37/683	Reference		Reference		Reference	
	4–5	69/1523	0.55(0.32-0.94)	0.030	0.56(0.34–0.97)	0.036	0.57(0.32-1.03)	0.054
	≥6	37/1110	0.47(0.26-0.84)	0.010	0.48(0.28-0.87)	0.014	0.49(0.28-0.86)	0.012
No sleep disorders	0–3	45/1287	0.48(0.27-0.87)	0.017	0.50(0.29-0.88)	0.017	0.52(0.32-0.93)	0.020
	4–5	107/3435	0.46(0.24-0.80)	< 0.001	0.48(0.28-0.81)	< 0.001	0.49(0.33-0.83)	< 0.001
	≥6	72/2680	0.34(0.19–0.62)	< 0.001	0.32(0.20-0.60)	< 0.001	0.36(0.19–0.65)	< 0.001
P for trend				< 0.001		< 0.001		< 0.001

Table 3 Joint association of DI-GM and sleep disorders with all-cause, cancer, and CVD mortality among US population with diabetes and pre-diabetes, NHANES, 2007–2018

Abbreviations: DI-GM dietary index for gut microbiota, NHANES the National Health and Nutrition Examination Survey, CI confidence interval, CVD cardiovascular disease

^a Adjusted for age

^b Multivariable adjusted model additionally adjusted for sex, race and ethnicity, educational attainment, family poverty income ratio, body mass index, total calories intake, sleep time, smoking status, and alcohol use

^c Additionally adjusted for hypertension and hypercholesterolemia

events by about 30–50% in people with pre-diabetes over the subsequent 30 years [45]. The English National Health Service Diabetes Prevention Programme showed lifestyle intervention led to significant reductions in weight and HbA1c in pre-diabetic people [46]. These findings emphasize the importance of lifestyle interventions in preventing the risk of long-term death in patients with pre-diabetes, in which personalised gut–microbiota focused dietary interventions combined with sleep modulation may be a promising strategy to reduce the risk of death.

There are several possible biological explanations for our findings. The Brain-Gut-Microbiome Axis plays a central role in the pathophysiology of diabetes. Individual sleep status has an impact on metabolism by affecting the microbiota composition, which in turn can alter insulin metabolism and cognitive function. Sleep disorders cause proliferation of pathogenic bacteria,

Gryllostomatidae and Enterobacteriaceae, as well as a decrease in bacteria producing SCFAs [47]. SCFAs stimulate the production and secretion of GLP-1, which directly promotes insulin secretion and inhibits glucagon secretion through interactions with pancreatic β cells [48]. Additionally, microbiota can also modulate the inflammatory response that leads to cognitive decline in a model of sleep deprivation (SD) [49]. Colonisation of the SD-associated microbiome induces gut leakage with higher circulation of LPS and other bacterial toxins. These microbial products and metabolites can activate inflammasomes, triggering systemic inflammation and impaired glucose metabolism [50]. Therefore, the DI-GM and sleep disorders have adverse biological effects on individuals with diabetes and pre-diabetes, inflammation and metabolic disorders may explain the combined effect of these two risk factors.

Our study has several strengths. To our knowledge, this is the first study to evaluate the joint effect of DI-GM and sleep disorders on mortality outcomes in a nationally representative sample of the population with diabetes and pre-diabetes. Sample weights, clustering and stratification were used to assess appropriate variance and ensure national representation of the US population. Additionally, the nationally representative sample included multiple races to increase generalisability to other populations with diabetes and pre-diabetes. Sensitivity and stratified analyses to evaluate the robustness of the main findings. The consistent findings after excluding the participants who died within 24 months or had a history of cancer or CVD add robustness to our main conclusions and reduce concerns about reverse causation.

Admittedly, some limitations should be noted in our study. Firstly, as a cross-sectional study, this research design cannot establish causal relationships between DI-GM and diabetes. Future prospective cohort studies and randomized controlled trials will be required to verify potential causality. Secondly, the data on dietary intake and sleep disorders were self-reported, and were subject to recall biases and misclassification. Thirdly, the selection of components in the DI-GM was constrained by the limited literature available on each type of food, while foods that have not been thoroughly investigated in relation to the gut microbiota were not included in the index. Finally, the type of sleep disorder was not clearly defined.

Conclusion

In summary, this population-based cohort study reveals that the newly proposed DI-GM is negatively correlated with the prevalence of diabetes and pre-diabetes. Furthermore, diabetic and pre-diabetic individuals with high DI-GM and no sleep disorders are associated with significantly reduced all-cause and CVD mortality risks.

These results highlight the importance of considering both gut–microbiotafocused dietary interventions and sleep status in developing targeted intervention strategies to improve survival outcomes among diabetic and prediabetic individuals.

Supplementary Information

The online version contains supplementary material available at https://doi. org/10.1186/s12937-025-01162-0.

Supplementary Material 1. Supplementary Material 2. Supplementary Material 3. Supplementary Material 4.

Supplementary Material 5. Supplementary Material 6.

Acknowledgements

We thank all participants in the NHANES databases.

Trial registration

The data in this study was sourced from the NHANES database, and the research was conducted as a retrospective analysis.

Clinical trial number

Not applicable.

Authors' contributions

KS conducted the database search, performed the statistical analysis and wrote the first draft of the manuscript. CS and YH extracted data and contributed to the discussion. CL, JC, LX, and YC contributed to the review and editing. YW revised the final manuscript. All authors contributed to the article and approved the submitted version.

Funding

There is no funding statement in this article.

Data availability

The data sets generated and analyzed during this study are available at NHANES website: https://www.cdc.gov/nchs/nhanes/index.htm.

Declarations

Ethics approval consent to participate

The NCHS Research Ethics Review Board approved the study protocol, and written informed consent was provided by all participants.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

Received: 26 December 2024 Accepted: 6 June 2025 Published online: 18 June 2025

References

- Yang D, Huang W, Wu CW, et al. Acute sleep deprivation exacerbates systemic inflammation and psychiatry disorders through gut microbiota dysbiosis and disruption of circadian rhythms. Microbiol Res. 2023;268:127292.
- Kim MK, Lee KN, Han K, et al. Diabetes Duration, cholesterol levels, and risk of cardiovascular diseases in individuals with type 2 diabetes. J Clin Endocrinol Metab. 2024;109(12):e2317–23.
- Shan S, Luo Z, Yao L, et al. Cross-country inequalities in disease burden and care quality of chronic kidney disease due to type 2 diabetes mellitus, 1990–2021: findings from the global burden of disease study 2021. Diabetes Obes Metab. 2024;26(12):5950–9.
- 4. National Diabetes Statistics Report. Prevention, Centers For Disease Control, 2023.
- Antza C, Kostopoulos G, Mostafa S, et al. The links between sleep duration, obesity and type 2 diabetes mellitus. J Endocrinol. 2022;252(2):125–41.
- 6. Mao Y. Sleep architecture changes in diabetes. J Clin Med. 2024;13(22):6851.
- Zhang Y, Liu C, Xu Y, et al. The relationship between sugar-sweetened beverages, sleep disorders, and diabesity. Front Endocrinol. 2023;13:1041977.
- Engeda J, Mezuk B, Ratliff S, et al. Association between duration and quality of sleep and the risk of pre-diabetes: evidence fromNHANES. Diabet Med. 2013;30(6):676–80.
- Pöysti S, Silojärvi S, Brodnicki TC, et al. Gut dysbiosis promotes isletautoimmunity by increasing T-cell attraction in islets via CXCL10 chemokine[J]. J Autoimmun. 2023;140:103090.

- Yuan M, Sun T, Zhang Y, et al. Quercetin alleviates insulin resistance and repairs intestinal barrier in db/db mice by modulating gut microbiota. Nutrients. 2024;16(12):1870.
- Rosell-Mases E, Santiago A, Corral-Pujol M, et al. Mutual modulation of gut microbiota and the immune system in type 1 diabetes models. Nat Commun. 2023;14(1):7770.
- Dong H, Zhuang H, Yu C, et al. Interactions between soluble dietary fibers from three edible fungi and gut microbiota. Int J Biol Macromol. 2024;278(Pt 3):134685.
- Ross FC, Patangia D, Grimaud G, et al. The interplay between diet and the gut microbiome: implications for health and disease. Nat Rev Microbiol. 2024;22(11):671–86.
- Toi PL, Anothaisintawee T, Chaikledkaew U, et al. Preventive role of diet interventions and dietary factors in type 2 diabetes mellitus: an umbrella review. Nutrients. 2020;12(9):2722.
- Li YJ, Chen X, Kwan TK, et al. Dietary fiber protects against diabetic nephropathy through short-chain fatty acid-mediated activation of G protein-coupled receptors GPR43 and GPR109A. J Am Soc Nephrol. 2020;31(6):1267–81.
- Vitale M, Giacco R, Laiola M, et al. Acute and chronic improvement in postprandial glucose metabolism by a diet resembling the traditional Mediterranean dietary pattern: Can SCFAs play a role? Clin Nutr. 2021;40(2):428–37.
- Quan X, Shen X, Li C, et al. Adherence to the dietary approaches to stop hypertension diet reduces the risk of diabetes mellitus: a systematic review and dose-response meta-analysis. Endocrine. 2024;86(1):85–100.
- Galié S, García Gavilán J, Camacho Barcía L, et al. Effects of the mediterranean diet or nut consumption on gut microbiota composition and fecal metabolites and their relationship with cardiometabolic risk factors. Mol Nutr Food Res. 2021;65(19):e2000982.
- Kase BE, Liese AD, Zhang J, et al. The development and evaluation of a literature-based dietary index for gut microbiota. Nutrients. 2024;16(7):1045.
- Bravo JA, Forsythe P, Chew MV, et al. Ingestion of Lactobacillus strain regulates emotional behavior and central GABA receptor expression in a mouse via the vagus nerve. Proc Natl Acad Sci U S A. 2011;108(38):16050–5.
- Nutt DJ, Stahl SM. Searching for perfect sleep: the continuing evolution of GABAA receptor modulators as hypnotics. J Psychopharmacol. 2010;24(11):1601–12.
- 22. Wang X, Wang Z, Cao J, et al. Gut microbiota-derived metabolites mediate the neuroprotective effect of melatonin in cognitive impairment induced by sleep deprivation. Microbiome. 2023;11(1):17.
- National Center for Health Statistics. National Health and Nutrition Examination Survey, 2015.
- Zipf G, Chiappa M, Porter KS, et al. National health and nutrition examination survey: plan and operations, 1999-2010. Vital Health Stat 1. 2013;56:1–37.
- ElSayed NA, Aleppo G, Aroda VR, et al. 2. Classification and diagnosis of diabetes:standards of care in diabetes—2023. Diabetes Care. 2023;46(Supplement_1):S19–40.
- Malesza IJ, Malesza M, Walkowiak J, et al. High-fat, Western-style diet, systemic inflammation, and gut microbiota: a narrative review. Cells. 2021;10(11):3164.
- Nguyen LH, Ma W, Wang DD, et al. Association between sulfur-metabolizing bacterial communities in stool and risk of distal colorectal cancer in men. Gastroenterology. 2020;158(5):1313–25.
- Liu J, Huang S. Dietary index for gut microbiota is associated with stroke among US adults[J]. Food Funct. 2025;16(4):1458–68.
- 29. Zhang X, Yang Q, Huang J, et al. Association of the newly proposed dietary index for gut microbiota and depression: the mediation effect of phenotypic age and body mass index. Eur Arch Psychiatry Clin Neurosci, 2024.
- Wang Y, Huang W, O Neil A, et al. Association between sleep duration and mortality risk among adults with type 2 diabetes: a prospective cohort study. Diabetologia. 2020;63(11):2292–304.
- Gu K, Min SH, Cho J. Sleep duration and mortality in patients with diabetes: results from the 2007–2015 Korea national health and nutrition examination survey. Diabetes Metab. 2022;48(3):101312.
- 32. Li J, Yin J, Luo Y, et al. Association of healthy sleep pattern with the risk of cardiovascular disease and all-cause mortality among people

- von Schantz M, Ong JC, Knutson KL. Associations between sleep disturbances, diabetes and mortality in the UK Biobank cohort: a prospective population-based study. J Sleep Res. 2021;30(6):e13392.
- 34. Li C, Lin C, Liu C, et al. Sleep duration predicts subsequent long-term mortality in patients with type 2 diabetes: a large single-center cohort study. Cardiovasc Diabetol. 2022;21(1):60.
- Mansor LS, Mehta K, Aksentijevic D, et al. Increased oxidative metabolism following hypoxia in the type 2 diabetic heart, despite normal hypoxia signalling and metabolic adaptation. J Physiol. 2016;594(2):307–20.
- Spiegel K, Leproult R, Van Cauter E. Impact of sleep debt on metabolic and endocrine function. The Lancet. 1999;354(9188):1435–9.
- Sakamoto K, Butera MA, Zhou C, et al. Overnutrition causes insulin resistance and metabolic disorder through increased sympathetic nervous system activity. Cell Metab. 2025;37(1):121–37.
- Chen L, Liu B, Ren L, et al. High-fiber diet ameliorates gut microbiota, serum metabolism and emotional mood in type 2 diabetes patients. Front Cell Infect Microbiol. 2023;13:1069954.
- Prasad M, Rajagopal P, Devarajan N, et al. A comprehensive review on high -fat diet-induced diabetes mellitus: an epigenetic view. J Nutr Biochem. 2022;107:109037.
- 40. Martínez-González MA, Montero P, Ruiz-Canela M, et al. Yearly attained adherence to Mediterranean diet and incidence of diabetes in a large randomized trial. Cardiovasc Diabetol. 2023;22(1):262.
- Ben-Yacov O, Godneva A, Rein M, et al. Gut microbiome modulates the effects of a personalised postprandial-targeting (PPT) diet on cardiometabolic markers: a diet intervention in pre-diabetes. Gut. 2023;72(8):1486–96.
- Ben-Yacov O, Godneva A, Rein M, et al. Personalized postprandial glucose response-targeting diet versus mediterranean diet for glycemic control in prediabetes. Diabetes Care. 2021;44(9):1980–91.
- Cotillard A, Cartier-Meheust A, Litwin NS, et al. A posteriori dietary patterns better explain variations of the gut microbiome than individual markers in the American Gut Project. Am J Clin Nutr. 2022;115(2):432–43.
- 44. Maskarinec G, Hullar MAJ, Monroe KR, et al. Fecal microbial diversity and structure are associated with diet quality in the multiethnic cohort adiposity phenotype study. J Nutr. 2019;149(9):1575–84.
- 45. Yu L, Wang J, Gong Q, et al. Influence of a diet and/or exercise intervention on long-term mortality and vascular complications in people with impaired glucose tolerance: da qing diabetes prevention outcome study. Diabetes Obes Metab. 2024;26(4):1188–96.
- Valabhji J, Barron E, Bradley D, et al. Early outcomes from the english national health service diabetes prevention programme. Diabetes Care. 2020;43(1):152–60.
- Li Y, Shao L, Mou Y, et al. Sleep, circadian rhythm and gut microbiota: alterations in Alzheimer's disease and their potential links in the pathogenesis. Gut Microbes. 2021;13(1):1957407.
- Zhao L, Zhang F, Ding X, et al. Gut bacteria selectively promoted by dietary fibers alleviate type 2 diabetes. Science. 2018;359(6380):1151–6.
- Wang Z, Chen WH, Li SX, et al. Gut microbiota modulates the inflammatory response and cognitive impairment induced by sleep deprivation. Mol Psychiatry. 2021;26(11):6277–92.
- Guo Z, Pan J, Zhu H, et al. Metabolites of gut microbiota and possible implication in development of diabetes mellitus. J Agric Food Chem. 2022;70(20):5945–60.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.