

Baseline Vitamin D Status, Sleep Patterns, and the Risk of Incident Type 2 Diabetes in Data From the UK Biobank Study

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OBJECTIVE

Circulating vitamin D concentrations have been associated with the risk of type 2 diabetes (T2D), but the results are inconsistent. Emerging evidence suggests that vitamin D metabolism is linked to sleep behaviors. We investigated the prospective association between serum 25-hydroxyvitamin D (250HD) and the risk of incident T2D and whether such association was modified by sleep behaviors.

RESEARCH DESIGN AND METHODS

The study included 350,211 individuals free of diabetes in the UK Biobank. Serum 250HD (nmol/L) concentrations were measured. Five sleep behaviors including sleep duration, insomnia, snoring, chronotype, and daytime sleepiness were included to generate overall sleep patterns, defined by healthy sleep scores. We also calculated genetic risk scores of sleep patterns.

RESULTS

During a median follow-up of 8.1 years, we documented 6,940 case subjects with incident T2D. We found that serum 25OHD was significantly associated with a lower risk of incident T2D, and the multivariate adjusted hazard ratio (HR) (95% CI) per 10 nmol/L increase was 0.88 (0.87–0.90). We found a significant interaction between 25OHD and overall sleep patterns on the risk of incident T2D (*P* for interaction = 0.002). The inverse association between high 25OHD and T2D was more prominent among participants with healthier sleep patterns. Among the individual sleep behaviors, daytime sleepiness showed the strongest interaction with 25OHD (*P* for interaction = 0.0006). The reduced HR of T2D associated with high 25OHD appeared to be more evident among participants with no frequent daytime sleepiness compared with those with excessive daytime sleepiness. The genetic variations of the sleep patterns did not modify the relation between 25OHD and T2D.

CONCLUSIONS

Our study indicates that higher serum 25OHD concentrations are associated with a lower risk of incident T2D, and such relations are modified by overall sleep patterns, with daytime sleepiness being the major contributor.

Vitamin D is a fat-soluble steroid hormone essential for calcium homeostasis and bone health (1,2). Several randomized controlled trials have explored the role of supplemental vitamin D in preventing cancer and cardiometabolic diseases in recent years (3–5). However, the effects of vitamin D supplementation on the risk of type 2 diabetes

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(T2D) were mixed across intervention studies (4-6). For example, the Vitamin D and Type 2 Diabetes (D2d) randomized controlled trial showed that vitamin D₃ supplementation did not lower the risk of T2D among persons with prediabetes (4). In addition, a body of previous prospective studies have associated serum levels of 25hydroxyvitamin D (25OHD), the best measure of vitamin D status, with T2D (7–10). Several meta-analyses of prospective cohort studies showed a reduced risk of T2D associated with high circulating vitamin D levels, yet statistically significant heterogeneity in these observations was detected (11-13). The inconsistent associations might be partly due to the modification effects of varying lifestyle factors related to vitamin D status across the studies.

Emerging evidence has linked various sleep behaviors with vitamin D metabolism (14-16). For example, short sleep duration and obstructive sleep apnea were associated with 250HD deficiency (16,17). Subjects with excessive daytime sleepiness were found to have an increased likelihood of low serum 250HD concentrations (15), possibly due to inadequate outdoor sunlight exposure that reduces the biosynthesis of vitamin D (18). Notably, unhealthy sleep behaviors have also been related to an increased risk of T2D (19,20). Therefore, we hypothesized that sleep behaviors might modify the association between 25OHD and the risk of T2D.

In this prospective study of 350,211 participants from the UK Biobank, we tested the association between serum 250HD and incident T2D and particularly examined the potential modification effect of our recently developed overall sleep patterns, which incorporated sleep duration, chronotype, insomnia, snoring, and excessive daytime sleepiness (21). We also assessed the modification effect of the genetically determined sleep behaviors.

RESEARCH DESIGN AND METHODS

Study Design and Population

The detailed study design and methods of the UK Biobank study have been described elsewhere (22). In brief, UK Biobank recruited >500,000 community-based volunteers aged 40–69 years across the U.K. between 2006 and 2010. Participants provided health information through touchscreen questionnaires and anthropometric measurements. Blood samples were collected for genotyping and biomarker analyses. All participants gave written consent, and ethical approval was obtained from the North West Multi-Centre Research Ethics Committee (London, U.K.).

In the current study, we excluded those with diabetes at baseline (n = 27,258) and those with missing values on serum 25OHD concentrations (n = 51,634) or sleep variables (n = 74,303) at baseline, leaving a total of 350,211 participants in the main analysis. In the genetic analysis, we used the UK Biobank genetic data of 274,356 unrelated individuals of European descent.

Assessment of Serum 250HD Concentrations

A range of biochemistry markers was measured in the blood sample collected at baseline (2006–2010). Serum vitamin D (nanomoles per liter) was measured by chemiluminescence technology analysis on a DiaSorin Ltd. LIAISON XL. Calibration and quality control were conducted by the UK Biobank. In brief, coefficients of variation for low, medium, and high internal quality control level samples for each biomarker were provided. The coefficients of variation for serum 25OHD ranged from 5.0 to 6.1%. In addition, the external quality assurance result was 100% for vitamin D. The detailed information of the measurements is provided on the UK Biobank website (https://biobank .ndph.ox.ac.uk/showcase/showcase/docs/ serum_biochemistry.pdf).

Assessment of Sleep Behaviors

The information on sleep behaviors was based on self-reported records collected from 2006 to 2010 at baseline visit. Details about the sleep behavior questions in the UK Biobank study were described elsewhere (21). Healthy sleep patterns were defined by five individual sleep behaviors (sleep duration, chronotype, insomnia, snoring, and excessive daytime sleepiness) (21). Low-risk sleep behaviors were defined as follows: sleep 7-8 h/day; early chronotype ("morning" or "morning than evening"); reported never or rarely insomnia symptoms; no self-reported snoring; and no excessive daytime sleepiness ("never/rarely" or "sometimes"). For each sleep behavior, low risk or high risk was coded as 1 or 0, respectively. The five scores were summed to obtain a healthy sleep score ranging from 0 to 5, with higher scores indicating a healthier sleep pattern. The participants were classified into three groups of "healthy sleep pattern" (healthy sleep scores of 4 or 5), "intermediate sleep pattern" (healthy sleep scores of 2 or 3), and "poor sleep pattern" (healthy sleep scores of 0 or 1).

Definition of Genetic Risk Score

The genotyping process and arrays used in the UK Biobank study have been reported in detail previously (23). We created the genetic risk score (GRS) for the five sleep behaviors separately using the single nucleotide polymorphisms (SNPs) reaching genome-wide association significance in the recent published genome-wide association studies (GWAS) (24-28). The information on the selected SNPs is provided in Supplementary Table 1. Individual SNP was coded as 0, 1, and 2 according to the number of risk alleles. The effect size (β coefficient) for each SNP was obtained from the reported GWAS data. The GRS was calculated using a weighted method (29,30): weighted $\mathsf{GRS} = (\beta_1 \times \mathsf{SNP}_1 + \beta_2 \times \mathsf{SNP}_2 + \ldots +$ $\beta_n \times SNP_n$ × (*N*/sum of the β coefficients). We determined whether participants were at high or low genetic risk for each sleep behavior according to the median values of the corresponding GRS. Genetic risk for each sleep behavior was coded as 1 for high risk or 0 for low risk and added to calculate GRS for the overall sleep patterns. The sleep patterns GRS ranged from 0 to 5, with higher scores indicating a higher genetic predisposition to unhealthy sleep patterns.

Assessment of Outcomes

We used the reported UK Biobank algorithms to determine T2D status (31). Hospital inpatient records were obtained through linkage to Hospital Episode Statistics for England, Scottish Morbidity Record data for Scotland, and the Patient Episode Database for Wales. Admissions and diagnoses data of the records were used to ascertain incident T2D with the ICD-10 code of E11.

Statistical Analysis

Baseline characteristics of the participants were described as means or percentages according to the quintiles of serum 25OHD levels. Follow-up time was calculated from the baseline date to diagnosis of T2D, death, or the censoring date (31 March 2017), whichever came first. Cox proportional hazards model was used to estimate the hazard ratio (HR) and 95% CI. The proportional hazards assumptions for the Cox model were tested using Schoenfeld residuals method. Models were adjusted for age (continuous), sex (male/female), race (White European, mixed, South Asian, Black, and others), UK Biobank assessment center, average total annual household income (<£18,000, £18,000-30,999, £31,000-51,999, £52,000-100,000, >£100,000, and "do not know" or missing), Townsend deprivation index (continuous), alcohol consumption (current, former, or never), smoking status (current, former, or never), BMI (kilograms per meter squared, continuous), physical activity (MET-minutes per week, continuous), healthy diet score (0, 1, 2, 3, 4, or 5), sun exposure time in the summer (hours per day, continuous), season of blood collection (winter: December-February; spring: March–May; summer: June–August; and autumn: September-November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), and family history of diabetes (yes/no). The healthy diet score was calculated by using the following factors: vegetable intake at least four tablespoons each day (median); fruit intake at least three pieces each day (median); fish intake at least twice each week (median); unprocessed red meat intake no more than twice each week (median); and processed meat intake no more than twice each week (median). Each one point was given for each favorable diet factor, with the total diet score ranging from 0 to 5 (32). If the covariate information was missing, we imputed mean values for continuous variables or used a missingindicator approach for categorical variables such as smoking. The linear trend test was performed by treating serum 250HD as a continuous variable. Stratified analyses were performed a priori by treating serum 25OHD in quintiles according to healthy sleep patterns (healthy sleep pattern, intermediate sleep pattern, and poor sleep pattern) as well as individual sleep factors. We also conducted stratified analyses by sleep patterns GRS (low, intermediate, and high). The interaction test between serum 250HD and each category was performed by using the likelihood ratio test comparing models with and without a cross-product term.

We also conducted several sensitivity analyses to confirm the robustness of our results: additionally adjusted for milk consumption, cereal intake, and vitamin D supplement use; additionally adjusted for skin color, sunscreen use, and air pollution (particulate matter with diameters \leq 2.5 μ m concentrations); and further adjusted for latitude, sleep apnea, depressive symptoms, waist circumferences, and serum glucose levels. We also conducted a subsample analysis among participants free of cardiovascular disease (CVD) and cancer at baseline or restricted case subjects with incident T2D to >2 years from the baseline survey. In addition, we conducted a sensitivity analysis among participants with 250HD deficiency (250HD <20 ng/mL) (33).

All analyses were performed using SAS software (version 9.4; SAS Institute Inc., Cary, NC). All *P* values for the tests were two-sided, and *P* values <0.05 were considered as statistically significant.

Data and Resource Availability

This research was conducted using the public UK Biobank Resource.

RESULTS

Table 1 shows the baseline characteristics of the study participants according to the quintiles of serum 250HD concentrations. Participants with higher 25OHD were older, mainly of White descent, and associated with lower Townsend deprivation index. In addition, they were less likely to be current smokers, but more likely to be current drinkers and with lower BMI. They tended to have a healthy diet and more sun exposure time in the summer and were more likely to have higher physical activity levels and healthy sleep scores. Furthermore, a high prevalence of vitamin D supplementation was observed among participants with higher serum 250HD concentrations.

During a median follow-up of 8.1 years (2,804,109 person-years), we documented 6,940 case patients with incident T2D. We found that higher 25OHD concentrations were significantly associated with a lower risk of incident T2D in the age, sex-, and multivariate-adjusted models (Table 2). In the age- and sex-adjusted model, a 10 nmol/L increase of 25OHD levels was associated with a 19% reduction in the risk of incident T2D (95% CI 18–20%). After further adjustment for race, UK Biobank assessment center, house-hold income, Townsend deprivation index, alcohol consumption, smoking status,

BMI, physical activity, healthy diet, sun exposure time in the summer, season of blood collection, antihypertensive medications use, cholesterol medications use, and family history of diabetes, 250HD levels were inversely associated with the risk of incident T2D in a doseresponse fashion. The HR (95% CI) of T2D was 0.88 (0.87–0.90) for a 10 nmol/L increase of 250HD; and a 49% lower risk was observed in the highest quintile compared with the lowest quintile of 250HD (*P* trend <0.001).

We conducted a stratified analysis according to the healthy sleep scores to evaluate whether overall sleep patterns modified the association between 25OHD and the risk of T2D. We found a significant interaction between 250HD and overall sleep patterns on the risk of incident T2D (*P* for interaction = 0.002), in which the protective HR of high 25OHD was more evident among participants with higher healthy sleep scores. The HR (95% CI) of incident T2D associated with a 10 nmol/L increase of 250HD levels was 0.87 (0.84-0.89) among participants with a healthy sleep pattern, 0.89 (0.87-0.90) among participants with an intermediate sleep pattern, and 0.92 (0.88-0.96) among participants with a poor sleep pattern, respectively. Similar interaction patterns were observed in the analysis on the quintiles of 250HD (Fig. 1).

We also tested the interaction between 25OHD and sleep patterns GRS on the risk of incident T2D. The sleep patterns GRS was significantly correlated with healthy sleep scores (β [SE] = -0.04[0.002]; P < 0.001). There was no significant interaction between sleep patterns GRS and 25OHD observed (Fig. 2).

We further assessed the interaction between 250HD and each sleep behavior separately on the risk of T2D. We found that daytime sleepiness showed a significant interaction with 25OHD (P for interaction = 0.0006) (Table 3). A 10 nmol/L higher 250HD level showed more prominent associations with the risk of incident T2D among participants with no frequent daytime sleepiness (HR 0.88 [95% CI 0.87-0.89]) than those with excessive daytime sleepiness (HR 0.92 [95% CI 0.87-0.97]). In particular, the negative HRs of incident T2D for quintiles 2–5 groups compared with quintile 1 group of 250HD were also attenuated among participants with excessive daytime sleepiness (Fig. 3). The interactions between 250HD and other

	Quintile categories of serum 250HD					
Characteristics	Q1	Q2	Q3	Q4	Q5	P*
25OHD (nmol/L)	22.3	35.9	47.5	59.7	80.5	
Ν	69,737	69,873	70,327	70,131	70,143	
Age (years)	54.7	55.8	56.5	57.0	57.1	<0.001
Sex, male (%)	45.3	44.9	45.3	44.7	44.8	0.03
Race, White (%)	88.2	94.7	96.9	98.1	98.8	<0.001
BMI (kg/m ²)	28.1	27.7	27.3	26.8	26.0	<0.001
Townsend deprivation index	-0.7	-1.3	-1.6	-1.8	-1.9	<0.001
Household income \geq £31,000 (%)	46.4	48.9	48.5	47.7	47.6	<0.001
Current drinkers (%)	89.1	92.5	93.6	94.4	94.8	<0.001
Current smokers (%)	15.1	10.7	9.0	8.4	8.3	<0.001
MET-minutes/week	2,310.4	2,512.3	2,678.3	2,837.4	3,076.0	<0.001
Family history of diabetes (%)	22.9	21.3	20.1	19.1	18.0	<0.001
Sun exposure time in summer (hours/day)	3.4	3.6	3.8	3.9	4.2	<0.001
Antihypertensive medications use (%)	19.1	18.6	18.5	18.5	18.5	0.01
Cholesterol-lowering medications use (%)	14.2	14.1	13.9	14.4	16.2	<0.001
Healthy diet score (%)						<0.001
0–1	15.4	12.7	11.1	10.2	9.2	
2–3	51.6	49.6	48.5	47.4	46.4	
4–5	32.9	37.7	40.4	42.5	44.4	
Vitamin D supplementation (%)	2.1	3.5	5.8	7.7	10.3	<0.001
Healthy sleep score (%)						< 0.001
0–1	5.6	4.4	4.1	3.6	3.3	
2–3	59.9	58.5	57.6	57.0	56.3	
4–5	34.5	37.1	38.4	39.4	40.4	

Table 1—Baseline characteristics of participants according to quintiles of serum 25OHD levels in the UK Biobank study

*P values were calculated by ANOVA (continuous variables) or χ^2 test (categorical variables).

sleep behaviors were not statistically significant.

We performed sensitivity analyses, which showed that the results were largely unchanged after further adjustment for milk consumption, cereal intake, and vitamin D supplement use or skin color, sunscreen use, particulate matter with diameters $\leq 2.5 \,\mu$ m concentrations, sleep apnea, depressive symptoms, waist circumferences, and serum glucose levels (Supplementary Tables 2–4). Moreover, we observed a significant but weak association between healthy

sleep patterns and north coordinate with adjustment for age, sex, and 25OHD (β coefficient = -0.10 per 1,000 km-grid; P < 0.001). The association remained unchanged after restricting participants living in the same address for >5 years (β coefficient = -0.09 per 1,000 km-grid; P < 0.001). The results did not change appreciably with further adjustment for the north coordinate (Supplementary Table 4). In addition, after excluding participants with CVD or cancer at baseline or limiting participants with a follow-up time of >2 years, the results did not alter appreciably (Supplementary Tables 5 and 6). Besides, after further adjustment for serum HbA_{1c} levels or excluding participants with estimated glomerular filtration rate <60 mL/min/1.73 m², the results were stable. Furthermore, the results did not alter appreciably among participants with 25OHD deficiency (Supplementary Table 7). Interestingly, we observed a significant interaction between 25OHD and chronotype among participants with vitamin D deficiency, as the negative HR of incident T2D associated with higher serum 25OHD concentrations

Table 2-Adjusted HRs and 95% CI for serum 250HD levels with T2D in the UK Biobank study

	Serum 250HD (quintiles)					HR (95% CI)	
		for a 10 nmol/L					
	Q1	Q2	Q3	Q4	Q5	increase	P trend
Case subjects/N	2,122/69,737	1,610/69,873	1,370/70,327	1,028/70,131	810/70,143		
Model 1*	1.00	0.70 (0.66–0.75)	0.57 (0.53–0.61)	0.41 (0.38–0.44)	0.32 (0.30–0.35)	0.81 (0.80–0.82)	< 0.001
Model 2 ⁺	1.00	0.85 (0.80–0.91)	0.76 (0.71–0.82)	0.60 (0.56–0.65)	0.51 (0.47–0.56)	0.88 (0.87–0.90)	< 0.001

*Adjusted for age and sex. †Adjusted for age, sex, race (White European, mixed, South Asian, Black, and others), UK Biobank assessment center, average total annual household income (<£18,000,£18,000–30,999,£31,000–51,999,£52,000–100,000, >£100,000, and "do not know" or missing), Townsend deprivation index, alcohol consumption (current, former, never, or missing), smoking status (current, former, never, or missing), BMI, physical activity (MET-minutes), healthy diet score (0, 1, 2, 3, 4, and 5), sun exposure time in the summer (hours per day), season of blood collection (winter: December–February; spring: March–May; summer: June–August; and autumn: September–November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), and family history of diabetes (yes, no, and "do not know" or missing).



Figure 1—The association between serum 25OHD levels and T2D stratified by overall sleep patterns. Results were adjusted for age, sex, UK Biobank assessment center, average total annual household income (<£18,000, £18,000–30,999, £31,000–51,999, £52,000–100,000, >£100,000, and "do not know" or missing), Townsend deprivation index, alcohol consumption (current, former, never, or missing), smoking status (current, former, never, or missing), BMI, physical activity (MET-minutes), healthy diet score (0, 1, 2, 3, 4, and 5), sun exposure time in the summer (hours per day), season of blood collection (winter: December–February; spring: March–May; summer: June–August; and autumn: September–November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), and family history of diabetes (yes, no, and "do not know" or missing). Q, quintile.

was attenuated among participants with late chronotype.

CONCLUSIONS

In this large-scale prospective cohort study, we observed that serum 25OHD concentrations were inversely associated with risk of incident T2D in a doseresponse fashion, independent of traditional risk factors. We also found that the association was significantly modified by overall sleep patterns, while the decreased risk of T2D associated with high serum 25OHD levels appeared to be strengthened among participants with healthier sleep patterns. Among the individual sleep behaviors, daytime sleepiness showed the strongest modification effect.

Our findings of the inverse association between serum 25OHD and T2D are consistent with results from several prospective cohort studies (7–13,34). Even though the mechanisms underlying the observed inverse association between serum 25OHD and the risk of T2D remain unclear, several potential mechanisms may explain the relations. For example, vitamin D might improve β -cell function and subsequently decrease the risk of diabetes (35.36). It was found that 1,25(OH)2D, the bioactive form of 25OHD, might upregulate the expression of insulin receptors, improve insulin responsiveness for glucose transport, and regulate calcium fluxes in the β-cell to stimulate insulin secretion (35). A recent randomized controlled trial also showed that supplementation with high-dose vitamin D₃ in patients with prediabetes and hypovitaminosis D improved insulin sensitivity and reduced diabetes risk (37,38). In addition, a previous study suggests that high 25OHD concentrations may reduce T2D risk through modulation of immune function and inflammatory processes (39). Vitamin D might regulate the proliferation, differentiation, and function of immune and inflammatory cells and, as a consequence, upregulate antiinflammatory pathways and downregulate the activation of these cells (40).

For the first time, we found that overall sleep patterns significantly modified the relations between 250HD and T2D risk. We recently developed an index approach to jointly evaluate overall sleep patterns (21), and we found that healthy sleep patterns, including early chronotype, sleep 7–8 h/day, never or rarely insomnia, no snoring, and no frequent excessive daytime sleepiness, were associated with a reduced risk of CVD. Our findings indicate that overall sleep patterns may better capture the relation between sleep behaviors and human health. In addition, among individual sleep behaviors, we found that daytime sleepiness showed a significant interaction with 250HD with regard to T2D risk, but not for other sleep behaviors. Notably, significant heterogeneity in the associations between 250HD and T2D across studies was observed in previous meta-analyses. Randomized clinical trials also showed inconsistent results on whether vitamin D supplementation might lower the risk of diabetes (4-6). The D2d trial yielded a nonsignificant effect of vitamin D supplementation on the reduction in the incidence of T2D (4), but a recent meta-analysis of randomized clinical trials found a lower risk of T2D with vitamin D supplementation (6). Our findings suggest that varying sleep behaviors may at least partly account for such heterogeneity.



Figure 2—The association between serum 25OHD levels and T2D stratified by genetically determined sleep patterns. Results were adjusted for age, sex, UK Biobank assessment center, average total annual household income (<£18,000, £18,000–30,999, £31,000–51,999, £52,000–100,000, >£100,000, and "do not know" or missing), Townsend deprivation index, alcohol consumption (current, former, never, or missing), smoking status (current, former, never, or missing), BMI, physical activity (MET-minutes), healthy diet score (0, 1, 2, 3, 4, and 5), sun exposure time in the summer (hours per day), season of blood collection (winter: December–February; spring: March–May; summer: June–August; and autumn: September–November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), family history of diabetes (yes, no, and "do not know" or missing), genotyping batch, and the first 10 genetic principal components. Q, quintile.

Our findings of the interactions between the sleep patterns and 25OHD are biologically plausible. Vitamin D is endogenously synthesized via skin exposure to ultraviolet B radiation (41). Behavioral factors such as sleep behaviors may influence the amount of skin surface exposure to sunlight and reduce skin synthesis

Table 3—Adjusted HRs* and 95% CI for serum 250HD levels with T2D, stratified by sleep behaviors

Sleep behaviors		Case subjects/N	HR (95% CI)	P for interaction
Sleep duration	Low risk	3,990/240,824	0.87 (0.86–0.89)	0.07
	High risk	2,950/109,387	0.90 (0.88–0.92)	
Chronotype	Low risk	4,391/220,373	0.88 (0.86–0.89)	0.25
	High risk	2,549/129,838	0.89 (0.87–0.92)	
Insomnia	Low risk	1,356/86,779	0.87 (0.84–0.90)	0.35
	High risk	5,584/263,432	0.89 (0.87–0.90)	
Snoring	Low risk	3,476/221,562	0.88 (0.86–0.89)	0.27
	High risk	3,464/128,649	0.89 (0.87–0.91)	
Daytime sleepiness	Low risk	6,523/341,273	0.88 (0.87–0.89)	0.0006
	High risk	417/8,938	0.92 (0.87–0.97)	

*Adjusted for age, sex, race (White European, mixed, South Asian, Black, and others), UK Biobank assessment center, average total annual household income (<£18,000, £18,000–30,999, £31,000–51,999, £52,000–100,000, >£100,000, and "do not know" or missing), Townsend deprivation index, alcohol consumption (current, former, never, or missing), smoking status (current, former, never, or missing), BMI, physical activity (MET-minutes), healthy diet score (0, 1, 2, 3, 4, and 5), sun exposure time in the summer (hours/day), season of blood collection (winter: December–February; spring: March–May; summer: June–August; and autumn: September–November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), and family history of diabetes (yes, no, and "do not know" or missing).

of vitamin D (2). For example, time spent outdoors, as a proxy for sunlight exposure, has been associated with vitamin D status in previous studies (18,42). Notably, unhealthy sleep behaviors such as excessive daytime sleepiness may be related to less outside activities and sun exposure, thus reducing 25OHD in human bodies. Furthermore, the functions and synthesis of melatonin and vitamin D are contrary to each other (43). A lack of light exposure related to unhealthy sleep behaviors may enhance melatonin synthesis, which may compete with vitamin D₃ to bind to the vitamin D receptor (43,44). The hypothesis was supported by the previous observations that serum vitamin D was inversely associated with daytime sleepiness (15,45). In particular, of the individual sleep behaviors, the strongest interaction was observed between serum 250HD and daytime sleepiness in the current analysis. Therefore, our data suggest that daytime sleepiness may be the major contributor to the interactions between overall sleep patterns and vitamin D. Additionally, although no significant interaction was detected for



Figure 3—The association between serum 25OHD levels and T2D stratified by daytime sleepiness. Results were adjusted for age, sex, UK Biobank assessment center, average total annual household income (<£18,000, £18,000–30,999, £31,000–51,999, £52,000–100,000, >£100,000, and "do not know" or missing), Townsend deprivation index, alcohol consumption (current, former, never, or missing), smoking status (current, former, never, or missing), BMI, physical activity (MET-minutes), and healthy diet score (0, 1, 2, 3, 4, and 5), sun exposure time in the summer (hours per day), season of blood collection (winter: December–February; spring: March–May; summer: June–August; and autumn: September–November), antihypertensive medications use (yes/no), cholesterol medications use (yes/no), and family history of diabetes (yes, no, and "do not know" or missing). Q, quintile.

other sleep behaviors, the negative association of high serum 250HD with T2D appeared to be more marked among participants with a low risk of sleep duration, chronotype, insomnia, or snoring symptoms. Short sleep duration has been reported to change outdoor activities and was significantly associated with low serum 250HD levels (46,47). Previous studies also showed that obstructive sleep apnea, characterized by snoring symptoms, played a vital role in serum vitamin D deficiency (16,17). The study also showed that subjects with more severe obstructive sleep apnea tended to present lower 250HD related to increased prevalence of diabetes (17).

Previous studies suggest that genetic susceptibility may modify the relations between environmental factors and human health outcomes (48,49). However, we did not observe a statistically significant interaction between serum 25OHD and genetically determined sleep behaviors in the study. The GRSs of the sleep patterns were calculated based on genetic variants that accounted for a small proportion of the variance in the sleep behaviors; this may partly explain the null interaction between the genetic scores and 25OHD on T2D.

Strengths and Limitations

To the best of our knowledge, this is the first prospective study to assess the interaction between 250HD and sleep behaviors on incident T2D risk. The major strengths of this study include the large sample size, the prospective design, and the well-validated measures of serum 250HD. More importantly, a wide range of covariates, including lifestyles, dietary habits, and sun exposure, was collected, which allowed for rigorous confounding adjustment. Several potential limitations, however, should be addressed. First, some case subjects with incident T2D were based on the secondary diagnosis; thus, the incident time of T2D used in the current analysis may be later than the actual onset time. Second, a single measurement of serum 250HD at baseline was used in the study, which did not take into account the changes of the biomarker during the follow-up time. However, the use of a single serum 25OHD measurement in epidemiologic studies is supported by a population-based study showing that baseline 250HD values could predict 50% of follow-up values 14 years later (50). Besides, a recent GWAS showed that \sim 30% of serum 250HD values were heritable (51). Third, misclassification of exposures might exist since we adopted self-reported sleep data in this study. Fourth, other sleep behaviors, such as restless legs syndrome, that could interact with vitamin D on the risk of T2D were not included in the healthy sleep scores. Furthermore,

causality could not be determined because of the observational nature of this study, and randomized clinical trials are needed to confirm our findings. In addition, although we carefully controlled for a number of potential confounders including demographic and diet factors, lifestyles, sun exposure, as well as medical history in the analyses, residual confounding could still be present. Moreover, the current study only included participants aged 40-69 years. Since sleep behaviors and 250HD may differ across ages, the relations among 250HD, sleep behaviors, and T2D in other age-groups need further studies. Finally, the current study was based on national data, and most of the participants were of European descent; thus, the generalization of the results to other populations should be interpreted with caution.

Conclusion

In summary, our study indicates that higher serum 25OHD concentrations are associated with a lower risk of incident T2D, and such relations are modified by overall sleep patterns, with daytime sleepiness being the major contributor. Our findings highlight the importance to consider lifestyle factors such as sleep behaviors in investigation of the relation between vitamin D and human diseases. Our findings, if confirmed by replications, may have implications for the development of T2D prevention strategies targeting improvement of vitamin D supplementation among people with sleep disorders, especially daytime sleepiness.

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Duality of Interest. No potential conflicts of interest relevant to this article were reported. **Author Contributions.** M.W. and L.Q. conceived and designed the study, interpreted the data, and drafted and critically revised the manuscript. M.W. and T.Z. performed the statistical analysis. All authors contributed to the interpretation of the results and critical revision of the manuscript. All authors approved the final manuscript. L.Q. is the guarantor of this work and, as such, had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

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