

# **Elucidating the Role of Vitamin D in Depression**

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### **Abstract**

Depression is a debilitating and prevalent mental disorder that creates huge societal and economic burden. Vitamin D has been implicated in several mental disorders, but existing studies on the role of circulating vitamin D (25-hydroxyvitamin D [25(OH)D]) in depression have yielded inconsistent findings. Given that vitamin D levels are readily modifiable through dietary supplementation and/or lifestyle changes, understanding the relationship between vitamin D and depression could lead to cost-effective public health strategies for preventing depression. In this dissertation, we aimed to elucidate this relationship by leveraging data from large genome-wide association studies (GWAS), longitudinal registry, and epidemiologic cohorts.

In Chapter 1, we explored the association between 25(OH)D levels and the risk of incident major depressive disorder (MDD) in a general healthcare population (n=233,970). Low 25(OH)D levels were associated with an increased risk for 5-year MDD. Our study also revealed that these associations were stronger among older adults and white or Asian adults. In Chapter 2, we performed a two-sample bidirectional Mendelian randomization analysis, using summary-level data from the largest GWAS studies of 25(OH)D and depression to date, to evaluate the causal relationship between 25(OH)D and depression. There was no causal association between average lifelong 25(OH)D levels and depression, although the possibility of weaker causal effects or threshold effects remains. In Chapter 3, we examined the association between maternal vitamin D status during pregnancy and offspring depression during childhood and adolescence, and further assessed whether polygenic risk for depression (PRS) modified the association. There was no evidence for a strong association between maternal vitamin D status and

offspring depression during either childhood or adolescence. Additionally, there were no interactions between PRS and maternal 25(OH)D on offspring depression.

These findings suggest that vitamin D deficiency is associated with an increased risk for depression, although small changes in 25(OH)D are not causally related to depression. There was little evidence for an association between prenatal 25(OH)D exposures and early life depression. Further work is needed to understand threshold effects of 25(OH)D on health outcomes and whether there are critical windows of vulnerability to vitamin D exposure.

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## **Chapter 1. Serum 25-hydroxyvitamin D and incident major depression in a general healthcare population**

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## Abstract

**Objective:** Emerging evidence suggests vitamin D may be involved in the pathophysiology of depression, however, longitudinal epidemiological studies are sparse and inconclusive. We aimed to investigate the association between serum 25-hydroxyvitamin D [25(OH)D] levels and incident major depression in a large general healthcare population.

**Methods:** Using longitudinal electronic health records from the Partners Healthcare Research Patient Data Registry, we identified adults (ages  $\geq 18$ ) with at least one measurement of 25(OH)D and no ICD codes for major depressive disorder (MDD) prior to the first 25(OH)D measurement (n=233,970). Vitamin D status was classified based on conventional clinical cut-offs: deficient ( $< 20$  ng/mL), insufficient (20-29.9 ng/mL), and normal ( $\geq 30$  ng/mL); MDD case status was defined using ICD codes and treatment records. The association between 25(OH)D and 5-year incident MDD was assessed using adjusted Cox regression models.

**Results:** Over a 5-year study period, 10,152 (4.3%) patients developed MDD. Every 10 ng/mL increase in 25(OH)D was associated, on average, with a decreased risk for MDD (adjusted HR (aHR)=0.89; 95% CI: 0.87, 0.90). Compared to adults with normal 25(OH)D levels, people with insufficient (aHR=1.17; 95% CI: 1.12, 1.23) or deficient (aHR=1.37; 95% CI: 1.30, 1.45) levels of 25(OH)D were at increased risk for MDD. The association was stronger among older age groups, and white and Asian adults.

**Conclusions:** In a large healthcare system population, low 25(OH)D was associated with higher risk for major depression. This association differed by age and race/ethnicity. Further studies to confirm the causal relationship between vitamin D and depression are needed.

**Keywords:** Vitamin D, risk factor, major depression, EHR

## Introduction

Major depressive disorder (MDD) is a highly prevalent and often debilitating mental disorder, estimated to affect 16.6% of adults in the US (1), and currently ranks as the number one leading cause of disability worldwide (2). Major depression has been linked to higher suicide rates, cardiovascular and metabolic diseases, and impaired functioning (3, 4), which cumulatively lead to enormous economic and healthcare costs (3). Given this public health burden, more research is needed to understand the etiology of MDD, which in turn, can inform prevention strategies.

Accumulating evidence suggests that vitamin D, which acts as a neurosteroid hormone, may play a role in etiology of depression (5-7). Although specific biological pathways linking low vitamin D levels to depression are understudied, several have been postulated. First, vitamin D receptors (VDR) and hydroxylases are expressed in the same human brain regions (e.g., hippocampus and amygdala) implicated in depression pathophysiology (8-10). Second, vitamin D plays a role in regulating autoimmune and inflammatory pathways associated with psychological stress and depression (11-13). Third, vitamin D may exert neuroprotective effects through antioxidant mechanisms, stimulation of neurotrophin production, or formation of essential neurotransmitters (12, 14, 15). Such mechanisms have been supported by animal studies. One study revealed that vitamin D-deprived rodents show persistent changes in brain functioning, and VDR knockout mice display depression-like behaviors, such as lowered activity levels and heightened anxiety (12, 16). Furthermore, active vitamin D metabolites in mice have been shown to affect cholinergic and dopaminergic pathways, which have been associated with depression and other psychiatric conditions (8).

Despite ample support from animal models, longitudinal epidemiological studies to date have been limited. Some studies have estimated that lower 25-hydroxyvitamin D (25(OH)D), the main circulating and stable form of vitamin D (17), is associated with up to a 2-fold higher risk for adult depression (5, 18). However, other studies have reported non-significant findings (19-21). Different

results across studies may be attributable to the study population examined (e.g., studying only men (22), elderly populations (20), patients with cardiovascular disease (23)), sample size (ranging from n=656-7358), methods used to assess depression, or the failure to consider sub-group differences based on age, sex, and race/ethnicity. As such, there is a need for larger studies in general clinical populations with more objective measures for depression.

The current study used longitudinal registry data, namely the Partners Healthcare Research Patient Data Registry (RPDR), to examine the association between serum 25(OH)D levels and incident MDD in a general adult healthcare population. This study aimed to provide further insights on this association by addressing some of the limitations of previous studies by focusing on a much larger, younger, and more diverse population with MDD case status defined by clinician diagnosis and/or treatment. Elucidating the link between vitamin D and depression has important public health implications. Circulating 25(OH)D can be easily and inexpensively elevated through supplementation and/or lifestyle interventions that can be implemented at a population level to prevent depression and other health outcomes.

## **Methods**

### **Data Source and Study Design**

This study analyzed electronic health records (EHRs) extracted through the RPDR (24). The RPDR is a central data warehouse for all inpatient and outpatient records of >6.5 million patients, from two major teaching hospitals (Brigham and Women's Hospital and Massachusetts General Hospital), and affiliated community and special hospitals in the Greater Boston area. Data were examined between January 2000 and December 2016.

Adults were eligible for study inclusion if they had one or more measurements of serum 25(OH)D, had no ninth or tenth versions of the *International Classification of Diseases* (ICD-9/ICD-10)

codes for MDD (296.2x – 296.3x/F32.x-F33.x) prior to their first 25(OH)D measurement, and were at least 18 years of age at the time of their first 25(OH)D measurement (**Supplemental Figure 1.1**).

Baseline date was defined as the date of the patient’s first recorded 25(OH)D measurement. Patients with 25(OH)D levels that were three standard deviations (SD) above/below the study population mean and/or had a diagnoses of bipolar disorder (BPD) (ICD-9 codes 296.4x-296.8x/ICD-10 codes F31.0x-F31.9x) or manic disorder (ICD-9 codes 296.0x-296.1x/ICD-10 codes F30.1x-F30.4x, F30.8, F30.9) at any time were also excluded. Follow-up time for each patient began at the first measurement of 25(OH)D until the first occurrence of MDD (defined below) or the last recorded date in the RPDR within five years from baseline. Controls with any ICD codes of 296.9/F39 (i.e., Other and unspecified episodic mood disorder) or 311/F32.9 (i.e., Depressive disorder, not elsewhere classified) were excluded to reduce the likelihood of outcome misclassification. A 5-year study window allowed for sufficient accrual of MDD cases, while maintaining biologically plausible relevance of observed 25(OH)D levels to MDD incidence. Previous studies have reported that the effect of lower vitamin D levels may be diluted with follow-up periods longer than 5 years (25). Institutional Review Board approval was granted by the Partners Human Research Committee.

### **Serum 25(OH)D Measurements**

Within the Partners system, serum 25(OH)D levels were assessed using radioimmunoassays (RIA) before 2008, and high-performance liquid chromatography-tandem mass spectrometry (HPLC-MS) thereafter; prior studies have reported high correlation between the two assays ( $r=0.95$ ) (26). Assays were performed at Clinical Laboratory Improvement Amendment (CLIA)-certified laboratories, and the dates/times of the 25(OH)D assays were available in the EHRs. For patients with multiple measurements taken during the follow-up period, we analyzed only their first 25(OH)D measurement, consistent with EHR studies on 25(OH)D on other health outcomes (27). Of note, among patients with two or more 25(OH)D measures ( $n=136,863$ ), we found low to moderate variability over 5 years (coefficient of

variation=20.7%; IQR: 0-33.5%), suggesting that a single measurement is likely an acceptable estimate of mean 25(OH)D levels over the study period.

Serum 25(OH)D was included in models both continuously and categorically to assess whether the influence of 25(OH)D on MDD were due to incremental changes and/or threshold effects. Vitamin D status was defined by clinical cut-offs (28, 29): normal ( $\geq 30$  ng/mL), insufficient (20-29.9 ng/mL) or deficient ( $< 20$  ng/mL).

### **MDD Case Definition and Validation**

Incident MDD cases were identified using ICD-9/ICD-10 diagnostic codes. MDD cases were patients who, after a 25(OH)D measurement, met at least one of the following criteria: (1) received  $\geq 2$  ICD codes for MDD (296.2x – 296.3x/F32.x-F33.x) at least one month apart; or (2)  $\geq 1$  ICD code(s) for MDD *and* treatment with an antidepressant recorded at  $\geq 2$  visits; or (3)  $\geq 1$  ICD code(s) for MDD *and* treatment with electroconvulsive therapy (ECT). In cases with multiple ICD codes, the date of MDD diagnosis was defined as the earliest recorded ICD code date. Controls were patients who did not fulfill the MDD case definition and did not have any ICD codes for MDD. To ensure the validity of this MDD case definition, two experienced, trained clinicians (JWS, KWC) performed chart reviews on a random set of 50 MDD cases (**Supplementary Methods**). Positive predictive value (PPV), which is the probability that those identified as MDD cases were true cases, was computed to quantify the classification performance of the EHR rule-based MDD definition. Potential cases adjudicated as having insufficient information were removed from the denominator of the PPV calculation. The criteria used to define MDD had a PPV of 92% (95% CI: 84%, 99%).

### **Statistical Analysis**

Kaplan-Meier survival curves and log-rank tests were used to assess the univariate association between 25(OH)D and incident MDD over the five year study period. Multivariable Cox regression models were fit to determine the association between 25(OH)D and incident MDD. Three models were



used to control for a range of potential confounding factors that were shown previously to associate with depression risk and/or vitamin D levels (30, 31) (**Supplementary Methods**). The minimally adjusted model (Model 1) adjusted for age, sex, race/ethnicity, assay, and season of 25(OH)D measurement. Model 2 included all Model 1 covariates and additionally adjusted for vitamin D supplementation, the Charlson-Deyo comorbidity index, and intensity of healthcare use, which could represent a source of ascertainment bias. As vitamin D testing is not systematically performed in healthcare settings, it is possible that those with indications for vitamin D testing are more likely to have both lower vitamin D levels and generally poorer health as well as higher risk for MDD. Therefore, Model 3 built upon Model 2 by including all medical conditions that are indications for vitamin D testing and treatments that may affect circulating 25(OH)D levels. The proportional hazards assumption, which states that the ratio of hazards for any two individuals is constant across time, was tested by including a cross-product interaction term between 25(OH)D and follow-up time in the model. There was no statistical evidence for the violation of the assumption. Interactions between 25(OH)D and age, sex, and race/ethnicity were tested. Results were presented as adjusted hazard ratios (aHR) and 95% confidence intervals (CIs). The statistical significance threshold was set at a two-tailed  $p$ -value  $< 0.05$ . All analyses were performed in STATA 14.0 (StataCorp, College Station, TX) and SAS 9.4 (SAS Institute, Cary, North Carolina).

### **Sensitivity Analysis**

Given that other methods may better account for observed seasonal variations in 25(OH)D levels (**Supplemental Figure 1.2**), we reran our primary analyses using seasonality-adjusted 25(OH)D, representing mean annual 25(OH)D levels, derived using a cosinor model (32). Details about the derivation of mean annual 25(OH)D are described in the **Supplementary Methods**. Briefly, the cosinor model involved fitting raw 25(OH)D measures against sine and cosine transformations of time (i.e., month of blood sampling) in a linear regression model. Each individual's mean annual 25(OH)D was derived from the intercept and residuals yielded from the model. Published studies have reported the

validity of this method for estimating an individual's mean annual 25(OH)D levels, which could potentially reduce biases in 25(OH)D measurements among adults in North America (33).

### **Bias Analysis**

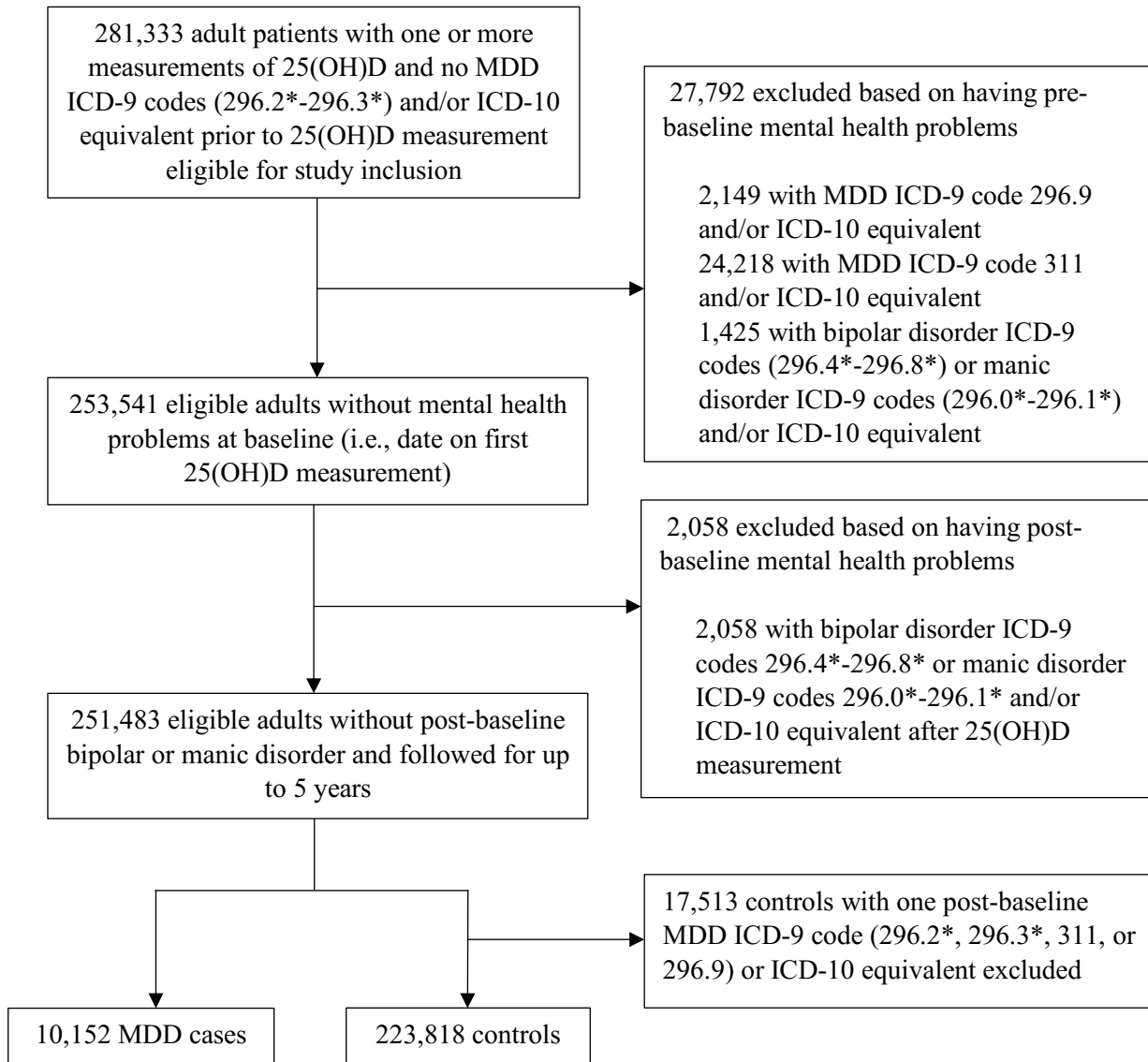
It is possible that there was residual confounding from variables that were not systematically collected in the EHR, including physical activity, body mass index (BMI), socioeconomic status (SES), smoking, and alcohol consumption (34). We therefore performed a bias analysis (35, 36) to adjust for possible unmeasured confounding by these variables. Briefly, bias factors, representing the potential magnitude of confounding by unmeasured confounders, were derived from a sub-sample of the study cohort who were also part of the Partners Healthcare Biobank cohort (37) and had data on the confounders listed above within one year of their 25(OH)D measurement. These bias factors were used to adjust the effect estimates obtained in the full study cohort, yielding HRs corrected for unmeasured confounding. Bias factors that change the original estimates by >30% indicate large confounding bias, while changes <7% are considered negligible confounding bias (38). Full details of the bias analysis are provided in the **Supplementary Methods**.

## **Results**

### **Study Cohort Characteristics**

Of the 233,970 adult patients included in the study cohort, 10,152 patients (4.3%) developed MDD over the study period (**Figure 1.1**). In the full cohort, the mean 25(OH)D was 28.8 ng/mL (SD=12.2), and 55.3% were classified as vitamin D insufficient or deficient. Baseline characteristics are displayed in **Table 1.1**. The study population had a mean age of 50.6 (SD=15.7) and comprised mainly females (66.3%) and whites (74.9%). Compared to the entire RPDR population (n~6.5 million), women (RPDR 52.9% vs. study cohort 66.3%) and patients 60 and older (RPDR 53.2% vs. study cohort 68.5%) were overrepresented, but racial/ethnic distributions were similar. Compared to adults with normal

25(OH)D levels, adults who were either vitamin D insufficient or deficient were younger, more likely to be male, have higher Charlson-Deyo comorbidity scores, have higher healthcare use intensity, and were less likely to be on vitamin D supplementation (**Table 1.1**).



**Figure 1.1 Flow chart for study cohort selection**

**Table 1.1 Study cohort baseline characteristics (n=233,970)**

	Full cohort (N=233,970)		Serum 25(OH)D <sup>1</sup>						p <sup>2</sup>
			Deficient (N=58,021)		Insufficient (N=71,466)		Normal (N=104,483)		
	N	%	N	%	N	%	N	%	
<b>Gender</b>									
Male	78,918	33.7	22,206	38.3	26,215	36.7	30,497	29.2	<0.001
Female	155,052	66.3	35,815	61.7	45,251	63.3	73,986	70.8	
<b>Ethnicity</b>									
Asian	11,177	4.8	4,167	7.2	3,826	5.4	3,184	3.0	<0.001
Black	17,362	7.4	8,603	14.8	5,005	7.0	3,754	3.6	
Hispanic	14,693	6.3	5,548	9.6	5,383	7.5	3,762	3.6	
Other	5,109	2.2	2,061	3.6	1,692	2.4	1,356	1.3	
Unknown	10,375	4.4	2,824	4.9	3,269	4.6	4,282	4.1	
White	175,254	74.9	34,818	60.0	52,291	73.2	88,145	84.4	
	<b>Mean</b>	<b>SD</b>	<b>Mean</b>	<b>SD</b>	<b>Mean</b>	<b>SD</b>	<b>Mean</b>	<b>SD</b>	<b>p<sup>2</sup></b>
<b>Age</b>	50.62	15.74	48.85	16.15	49.54	15.56	52.34	15.45	<0.001
	<b>N</b>	<b>%</b>	<b>N</b>	<b>%</b>	<b>N</b>	<b>%</b>	<b>N</b>	<b>%</b>	<b>p<sup>2</sup></b>
<b>Season of blood draw</b>									
Spring	60,018	25.7	13,408	23.1	18,594	26.0	28,016	26.8	<0.001
Summer	62,938	26.9	17,502	30.2	19,245	26.9	26,191	25.1	
Fall	54,451	23.3	8,797	15.2	15,935	22.3	29,719	28.4	
Winter	56,563	24.2	18,314	31.6	17,692	24.8	20,557	19.7	
<b>Assay</b>									
RIA	40,937	17.5	14,513	25.0	12,006	16.8	14,418	13.8	<0.001
HPLC-MS	193,033	82.5	43,508	75.0	59,460	83.2	90,065	86.2	
<b>Charlson-Deyo index</b>									
0-1	150,485	64.3	34,869	60.1	47,156	66.0	68,460	65.5	<0.001
2-3	49,866	21.3	12,129	20.9	14,849	20.8	22,888	21.9	
4-6	21,298	9.1	6,364	11.0	6,079	8.5	8,855	8.5	
≥7	12,321	5.3	4,659	8.0	3,382	4.7	4,280	4.1	
<b>Past-year vitamin D supplementation use</b>	4,138	1.8	735	1.3	1001	1.4	2402	2.3	<0.001

**Table 1.1 (continued)**

	Full cohort (N=233,970)		Serum 25(OH)D						p <sup>2</sup>
			Deficient (N=58,021)		Insufficient (N=71,466)		Normal (N=104,483)		
	Median	IQR	Median	IQR	Median	IQR	Median	IQR	
<b>Healthcare use density</b>	0.010	<0.01, 0.02	0.011	<0.01, 0.03	0.010	<0.01, 0.02	0.010	<0.01, 0.02	<0.001
	N	%	N	%	N	%	N	%	p <sup>2</sup>
<b>Indicators for vitamin D testing</b>									
History of rickets	20	0.0	6	0.0	5	0.0	9	0.0	0.81
History of osteomalacia	68	0.0	17	0.0	15	0.0	36	0.0	0.27
History of osteopatia	14,660	6.3	2,671	4.6	3,749	5.2	8,240	7.9	<0.001
History of chronic kidney disease	7,084	3.0	2,573	4.4	1,904	2.7	2,607	2.5	<0.001
History of liver disease	6,261	2.7	2,297	4.0	1,857	2.6	2,107	2.0	<0.001
History of pancreatic insufficiency	931	0.4	340	0.6	261	0.4	330	0.3	<0.001
History of malabsorption syndromes	7,571	3.2	1,905	3.3	2,286	3.2	3,380	3.2	0.69
History of hyper-/hypo-thyroidism	6,664	2.8	1,646	2.8	1,954	2.7	3,064	2.9	0.05
History of pathological fractures	2,213	0.9	668	1.2	637	0.9	908	0.9	<0.001
History of chronic bronchitis	1,767	0.8	643	1.1	480	0.7	644	0.6	<0.001
History of type 2 Diabetes	23,561	10.1	8,036	13.9	7,001	9.8	8,524	8.2	<0.001
History of hypercalcemia	2,691	1.2	955	1.6	754	1.1	982	0.9	<0.001
History of antiseizure medication	22,745	9.7	5,983	10.3	6,655	9.3	10,107	9.7	<0.001
History of glucocorticoid medication	14,096	6.0	3,547	6.1	4,267	6.0	6,282	6.0	0.55

<sup>1</sup>Vitamin D status defined by the following 25(OH)D cut-offs: Deficient, <20 ng/mL; Insufficient, 20-29.9 ng/mL; Normal, ≥30 ng/mL

<sup>2</sup>Chi-squared test for categorical variables; ANOVA for normally distributed continuous variables; Kruskal-Wallis for skewed continuous variables

### Serum 25(OH)D and 5-year Incident MDD

Serum 25(OH)D levels were inversely associated with five-year MDD incidence (log-rank test  $p < 0.0001$ ) (Figure 1.2), with the highest and lowest MDD incidence in the deficient and normal categories, respectively. Table 1.2 displays the HR estimates for incident MDD according to continuous and categorical 25(OH)D. After adjusting for covariates, every 10 ng/mL increase in 25(OH)D was associated, on average, with an 11% reduction in MDD risk (HR=0.89, 95% CI: 0.87, 0.90). This 10 ng/mL difference is equivalent to moving from the deficient to insufficient or insufficient to normal category. In addition, compared to patients with normal 25(OH)D levels, patients with insufficient (HR=1.17; 95% CI: 1.12, 1.23;  $p < 0.0001$ ) or deficient levels (HR=1.37; 95% CI: 1.30, 1.45;  $p < 0.0001$ ) had a 17-37% increased risk for MDD. Deficient 25(OH)D levels also conferred an increased risk for MDD compared to insufficient 25(OH)D levels (HR=1.17; 95% CI: 1.11, 1.23;  $p < 0.0001$ ).

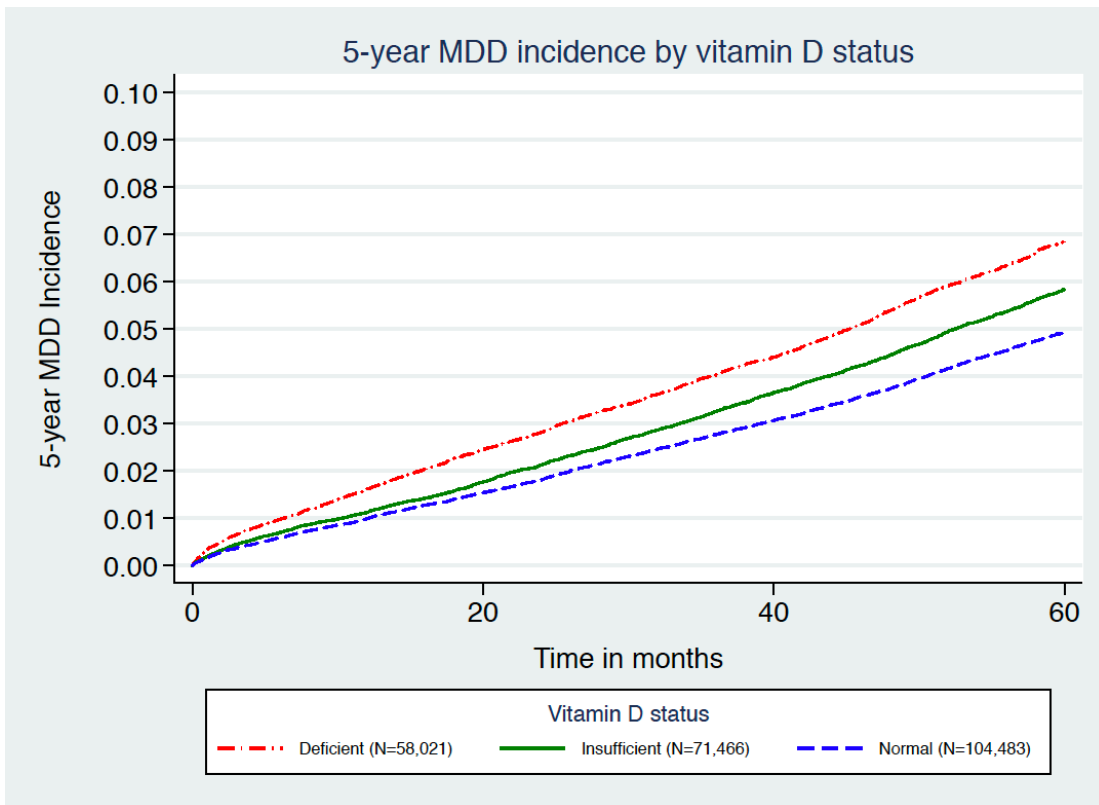


Figure 1.2 Kaplan-Meier 5-year major depressive disorder (MDD) incidence estimates by vitamin D status

**Table 1.2 Association between serum 25(OH)D levels and incident MDD using Cox regression analysis**

Serum 25(OH)D	MDD cases		Model 1			Model 2			Model 3		
	N	%	HR	95% CI	p	HR	95 % CI	p	HR	95 % CI	p
<i>Continuous</i> (per 10 ng/mL increase)	10,152	4.3	0.87	0.85, 0.88	<0.0001	0.88	0.86, 0.90	<0.0001	0.89	0.87 0.90	<0.0001
<i>Clinical Cut-offs</i>											
Normal ( $\geq$ 30 ng/mL)	4,104	3.9	ref	ref	ref	ref	ref	ref	ref		ref
Insufficient (20-29.9 ng/mL)	3,154	4.4	1.19	1.13, 1.25	<0.0001	1.40	1.33, 1.47	<0.0001	1.17	1.12, 1.23	<0.0001
Deficient (<20 ng/mL)	2,894	5.0	1.46	1.39, 1.54	<0.0001	1.18	1.13, 1.24	<0.0001	1.37	1.30, 1.45	<0.0001
<i>Insufficient vs. Deficient</i>											
Insufficient (20-29.9 ng/mL)	3,154	4.4	ref	ref	ref	ref	ref	ref	ref		ref
Deficient (<20 ng/mL)	2,894	5.0	1.23	1.17, 1.30	<0.0001	1.18	1.12, 1.24	<0.0001	1.17	1.11, 1.23	<0.0001

Abbreviations: 25(OH)D: 25-hydroxyvitamin D; CI: confidence interval; HR: hazard ratio; MDD: major depressive disorder

Model 1: Adjusted for age, sex, race/ethnicity, assay, season

Model 2: Model 1 covariates + healthcare use intensity + vitamin D supplementation + Charlson-Deyo comorbidity index

Model 3: Model 2 covariates + medical conditions and therapies indicating vitamin D testing

### **Serum 25(OH)D and 5-year Incident MDD within Subgroups**

We found evidence suggesting that the association between 25(OH)D levels and MDD was modified by age ( $p=0.001$ ) and race/ethnicity ( $p=0.006$ ), but not sex ( $p=0.12$ ). Specifically, the association between vitamin D deficiency and MDD risk was stronger among older adults ( $\text{age} \geq 40$ ;  $\text{HR}=1.37\text{-}1.48$ ) compared to younger adults ( $\text{HR}=1.28$ ) (**Table 1.3**). Furthermore, the association between vitamin D deficiency and MDD risk was only statistically significant in Asian ( $\text{HR}=1.38$ ;  $p=0.04$ ) and white ( $\text{HR}=1.46$ ;  $p<0.0001$ ) individuals, but not black ( $\text{HR}=1.09$ ;  $p=0.37$ ) or Hispanic ( $\text{HR}=1.09$ ;  $p=0.34$ ) individuals.

### **Sensitivity Analysis**

Seasonality-adjusted mean annual 25(OH)D levels were strongly and positively correlated with raw 25(OH)D levels (Pearson's  $r=0.98$ ); the absolute difference between the two measures was small (mean=-0.05, SD=2.18). 9.7%, 17.3%, and 8.9% of those with deficient, insufficient, and normal 25(OH)D levels, respectively, were reclassified either one category up or down. Those in the insufficient group were slightly more likely to be reclassified to the deficient (9.8%) than normal (7.5%) group. However, models fit with the mean annual 25(OH)D levels yielded almost identical estimates to those in the primary analyses (**Supplemental Table 1.1**), suggesting the robustness of the results to different methods of seasonality adjustments.

### **Bias Analysis**

Additional analyses were performed to assess the robustness of the HR estimates to corrections for potential unmeasured confounding. 1,119 individuals in the study cohort were eligible for inclusion in the Partners Biobank sub-cohort used to calculate bias factors. Compared to the full cohort, the sub-cohort was slightly over-represented in including patients with normal 25(OH)D levels (55.0% in sub-cohort vs. 44.7% in the full cohort), whites (sub-cohort, 87.7% vs. full cohort, 74.9%), and vitamin D supplementation users (sub-cohort, 6.6% vs. full cohort, 1.8%). In the sub-cohort, more than half



exercised for less than 2.5 hours per week, were not obese, never smoked, had at least one alcohol drink per week, were employed in the past 5 years, and completed college or graduate school (**Supplemental Table 1.2**).

Bias factors ranged between 0.87 and 1.17, with physical activity showing the largest confounding effects on the association between 25(OH)D and MDD. After correcting for confounding bias induced by physical activity, the aHR estimates attenuated (Vitamin D insufficiency: HR=1.07, 95% CI: 1.03, 1.13; deficiency: HR=1.17, 95% CI: 1.11, 1.24), but remained statistically significant (**Supplemental Table 1.3**). Estimates were also robust to corrections for all other confounders.

**Table 1.3 The association between serum 25(OH)D levels and incident MDD, stratified by age group, sex, and race/ethnicity**

	MDD cases		Serum 25(OH)D <sup>1</sup>												
			Continuous			Normal			Insufficient			Deficient			
			N	%	HR	95% CI	p	HR	95% CI	p	HR	95% CI	p	HR	95% CI
<b>Age group</b>															
<40	3,296	5.4	0.90	0.86, 0.92	<0.0001	ref	ref	ref	1.14	1.04, 1.24	0.004	1.28	1.17, 1.41	<0.0001	
40-59	4,194	4.2	0.86	0.84, 0.89	<0.0001	ref	ref	ref	1.21	1.12, 1.30	<0.0001	1.48	1.36, 1.60	<0.0001	
≥60	2,662	3.6	0.90	0.87, 0.92	<0.0001	ref	ref	ref	1.19	1.09, 1.31	<0.0001	1.37	1.23, 1.51	<0.0001	
<b>Sex</b>															
Females	7,391	4.8	0.90	0.88, 0.91	<0.0001	ref	ref	ref	1.17	1.11, 1.23	<0.0001	1.32	1.24, 1.40	<0.0001	
Males	2,761	3.5	0.85	0.82, 0.89	<0.0001	ref	ref	ref	1.19	1.09, 1.31	<0.001	1.51	1.37, 1.66	<0.0001	
<b>Race/ethnicity</b>															
Asian	308	2.8	0.86	0.77, 0.96	0.01	ref	ref	ref	1.21	0.90, 1.64	0.21	1.38	1.02, 1.86	0.04	
Black	808	4.7	0.91	0.85, 0.98	0.01	ref	ref	ref	0.93	0.75, 1.14	0.47	1.09	0.90, 1.32	0.37	
Hispanic	992	6.8	0.94	0.89, 1.01	0.08	ref	ref	ref	1.08	0.92, 1.27	0.36	1.09	0.92, 1.29	0.34	
White	7,480	4.3	0.87	0.86, 0.90	<0.0001	ref	ref	ref	1.18	1.12, 1.25	<0.0001	1.46	1.37, 1.55	<0.0001	

Abbreviations: 25(OH)D: 25-hydroxyvitamin D; CI: confidence interval; HR: hazard ratio; MDD: major depressive disorder

All models adjusted for assay, season, healthcare use intensity, vitamin D supplementation, Charlson-deyo comorbidity index, and medical conditions and therapies indicating vitamin D testing, and mutually adjusted for the other two tested variables

<sup>1</sup>Vitamin D status defined by the following 25(OH)D cut-offs: Deficient, <20 ng/mL; Insufficient, 20-29.9 ng/mL; Normal, ≥30 ng/mL. Continuous 25(OH)D was analyzed as per 10 ng/mL increment.

## Discussion

In this large, general healthcare cohort, we found that low 25(OH)D levels were associated with incident MDD over a five-year period, independent of potential confounders. Compared to adults with normal 25(OH)D levels, the risk of incident MDD was almost 20-40% higher among adults with insufficient and deficient levels, respectively. The association was robust to corrections for potential confounding bias and was stronger among older age groups (age>40 years), and white and Asian adults.

To our knowledge, this is the largest study to date exploring the association between 25(OH)D and incident MDD (N =233,970), over 30-fold larger than the largest prior study (23). The current study extends existing work on the prospective association between 25(OH)D levels and incident depression (19, 20, 23, 39-41) by focusing on a larger and ethnically/racially diverse cohort with a broader age range. Most existing studies have examined elderly populations (aged  $\geq 55$ ) because of the high prevalence and severe health consequences of vitamin D deficiency in this vulnerable group (42, 43). However, in recent years, vitamin D deficiency has also become a health concern among younger adults, given the increasing proportion of the population with night-shift work and working, living, and exercising indoors, resulting in reduced sunlight exposure (42, 44). Indeed, studies suggest vitamin D deficiency affects 30-50% of children and young adults even in geographical regions that receive ample sunlight (45, 46). There is, however, a lack of studies exploring the association between vitamin D levels on incident depression in younger populations. In this study, we examined the association in a younger cohort comprising mainly individuals aged  $\leq 60$ . Our findings provide preliminary evidence to support that vitamin D deficiency may also increase MDD risk among younger adults.

Findings from our subgroup analysis in older adults (ages >60) add to the literature on vitamin D deficiency and late-life depression, which has so far yielded mixed conclusions. Our findings, which reveal a 1.37-fold higher risk of incident MDD among older adults with vitamin D deficiency, are consistent with that from the most recent and largest epidemiological study to date. In a community-

dwelling Irish population, adults with deficient vitamin D levels had a 1.49-fold higher risk of incident depression, compared to adults with sufficient levels (18). Relative to previous studies in elderly cohorts, our analysis had higher statistical power due to a substantially larger sample (n=73,815). We also had sufficient power to make comparisons across all three vitamin D categories and observed significantly different risks for incident MDD even between the insufficient and deficient groups. Prior studies that collapsed vitamin D categories may have diluted the observed effects, which could partially explain their non-significant findings.

Intriguingly, vitamin D deficiency was associated with higher MDD risk in white and Asian adults, but not in black or Hispanic adults. Similar racial differences have been observed in studies examining the association between serum 25(OH)D levels and other health outcomes including bone mineral density, stroke, and mortality (47-51). For instance, in the Multi-Ethnic Study of Atherosclerosis (MESA), lower 25(OH)D was associated with an increased risk for incident coronary heart disease in white and Chinese participants, but not black or Hispanic participants (50). The reason for racial differences in the association of low 25(OH)D levels with incident depression remains unknown. One explanation is that black individuals have lower levels of vitamin D binding proteins and higher levels of bioavailable 25(OH)D, therefore total 25(OH)D measured using standard clinical assays may not be accurate for defining vitamin D status in these populations (50, 52). Other potential mechanisms such as variation in circulating 1,25-dihydroxyvitamin D (1,25(OH)<sub>2</sub>D) levels and genetic differences in VDR affinity may also explain lower susceptibility to the adverse effects of low serum 25(OH)D in black populations (53-55). Future studies evaluating the prospective association between 25(OH)D and MDD in ethnic minority populations are needed.

Vitamin D may directly impact depression risk through plausible biological mechanisms including the regulation of serotonin synthesis, neuritogenesis, neuromodulation, neuroprotection, or downregulation of inflammatory responses (8). On the other hand, the observed associations could be explained by non-causal or indirect vitamin D effects. First, low vitamin D may be a marker of lifestyle risk factors and chronic illnesses that contribute to the development of depression (20, 22, 56). However,

in the current study, the association between vitamin D and MDD incidence persisted after adjustment for baseline medical comorbidity and when restricted to younger populations with less severe chronic conditions and presumably more active lifestyles. Second, the effect of vitamin D on psychiatric traits may in fact be mediated through calcium suppression and increases in parathyroid hormone (PTH) levels (57). Indeed, prior studies have shown associations between hyperparathyroidism and impaired cognition and depression (58, 59). Additional work is needed to unravel the relationship between vitamin D, PTH, and depression.

Key strengths of this study include a large sample size and examination of 25(OH)D levels collected before MDD diagnosis, minimizing the likelihood of reverse causation. Further, by leveraging data from the subset of participants who were further characterized in the Partners Biobank research cohort, we were able to demonstrate the robustness of the estimates to potential biases induced by unmeasured confounders such as physical activity. However, the study is not without limitations. First, diagnoses were limited to those captured by clinical documentation in the EHR. This may have led to outcome misclassification, although such misclassification would likely have biased our estimates towards the null. Second, while we attempted to account for potential selection bias resulting from differential vitamin D testing due to factors related to both 25(OH)D and MDD, the adjustments may have been incomplete. Third, patients in the study population were all adults, predominantly white, and mainly from the Greater Boston Area, which is located at a higher latitude with sunlight attenuation during winter months. Therefore, these results may not be generalizable to children or adolescents and populations in different geographic locations or with different demographic distributions. Fourth, some covariates were poorly captured in the EHRs, which could have led to residual confounding. For example, only 2% of the study population had recorded past-year vitamin D supplementation use, compared to the 28% past-month vitamin D supplementation use reported in the National Health and Nutrition Examination Survey (NHANES) (60), indicating possible underestimation of dietary supplementation use. It was also not possible to account for changes in vitamin D supplementation use after vitamin D testing, such that patients who were tested to be vitamin D deficient started to take vitamin D

supplements. Nonetheless, in these cases, the results would likely have been biased towards the null, and the true effect sizes would have been larger. Lastly, the Partners Biobank sub-cohort used to derive bias factors to correct for unmeasured confounding had slightly different demographic distributions to the full study population, thus corrections for confounding bias may be incomplete. Nonetheless, comparable associations between these confounders and depression have been reported in other populations (61-63), suggesting the validity of the bias corrections.

Our findings support an association between 25(OH)D levels and MDD, which persisted after adjustments for a range of confounders and correction for confounding bias. However, as with all observational studies, causality cannot be inferred. Trials such as the Vitamin D and Omega-3 Trial-Depression Endpoint Prevention (VITAL-DEP) are underway to evaluate the effect of vitamin D supplementation on depression (64), which could provide firmer conclusions on the causal relationship. If truly causal, the normalization of 25(OH)D in the population may be a cost-effective and safe public health strategy for preventing depression.

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## Appendix

### Supplementary Methods

#### *MDD Chart Reviews*

Chart reviews on 50 random identified MDD cases were performed by two trained and experienced clinicians (JWS, KWC), who reviewed all medical notes in the patient's records to determine diagnosis of MDD or "not enough information" to make a decision. Guidelines for assigning diagnosis were based on DSM-V criteria for MDD, and a confidence level of low or intermediate/high was assigned to patients that were adjudicated to be an MDD case. Cases for which there was uncertainty were discussed with the senior psychiatrist (JWS) until consensus was reached. Two cases did not have sufficient information for classification, and of the remaining 48 patients, 4 were classified as having no evidence of MDD, 6 were classified as possible MDD cases (low confidence), and 38 were classified as probable/definite MDD cases (intermediate/high confidence).

#### *Assessment of Covariates*

Demographic factors including age, sex, and race/ethnicity were retrieved from the closest record prior to the date of study entry. Season of vitamin D testing was derived from the recorded date of blood draw. To account for the general burden of chronic medical conditions, ICD-9 coding algorithms were used to derive the Deyo modification to the Charlson co-morbidity index (1, 2), which has previously been validated in clinical populations (3, 4). This algorithm was implemented in R (5), using the *medicalrisk* package (6). Prescription of vitamin D supplementation, including cholecalciferol (D<sub>3</sub>), calcitriol (1,25-(OH)<sub>2</sub>D<sub>3</sub>), and ergocalciferol (D<sub>2</sub>), was assessed using prescription records within the one-year period prior to the 25(OH)D measurement. To determine if detection of MDD was higher among patients with lower vitamin D levels due to higher healthcare utilization, which could reflect a possible

ascertainment bias, we derived a variable to quantify and control for the intensity of healthcare utilization. Healthcare facility use density was calculated by summing the number of inpatient and outpatient visits and dividing by the amount of time the patient had been in the Partners Healthcare system prior to the 25(OH)D measurement. Similar equations have been used previously to quantify intensity of healthcare utilization in patients in the Partners Healthcare system (7). History of medical conditions that are indications for vitamin D testing (8, 9), including rickets, osteomalacia, osteoporosis, chronic kidney disease, liver disease, pancreatic insufficiency, malabsorption syndromes (e.g., IBD, cystic fibrosis), hyper- or hypo-thyroidism, non-traumatic fractures, chronic bronchitis, type II diabetes, hypercalcemia, were defined by the presence of corresponding ICD codes prior to baseline. Treatments known to affect vitamin D metabolism (9-11), such as anti-seizure and/or glucocorticoid medication, were identified using medication records. Age, Deyo-Charlson comorbidity index, and healthcare facility use were analyzed as continuous variables, and all other covariates were modelled categorically.

### ***Derivation of Mean Annual 25(OH)D***

The cosinor model takes the sine and cosine transformations of time and fits them as predictors of 25(OH)D in a linear regression model (12). The coefficients from the model are then transformed to obtain the amplitude and phase shift of the sine curve. In the cosinor model, 25(OH)D is modeled as a sine wave, characterized by phase shift (location of peak and trough on time axis), height (vertical shift of sine wave), and amplitude (distance between mean to peak or trough location of the curve). The mean annual 25(OH)D concentration in the study sample was estimated by the coefficient for the model intercept which, in combination with the residuals of the models, was used to derive the mean annual 25(OH)D levels for each individual.

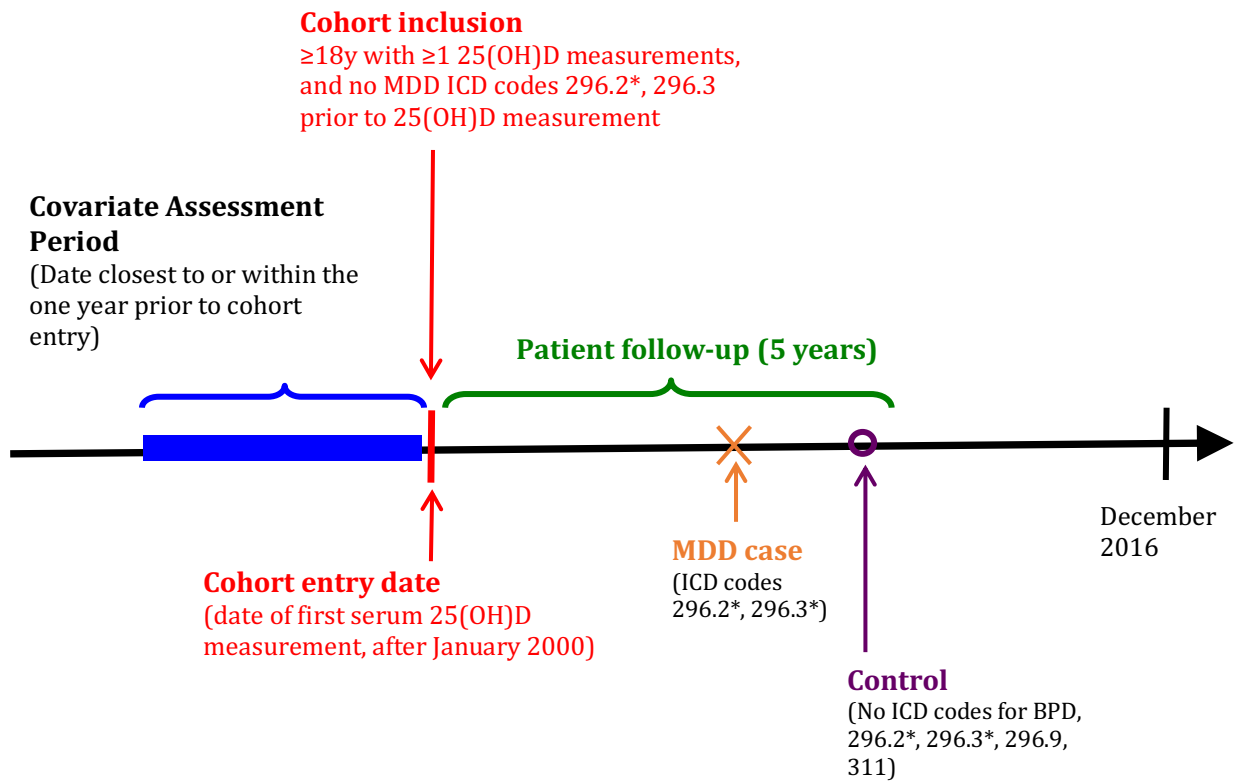
### ***Bias Analysis***

Confounders assessed in the bias analysis were defined and coded as follows: physical activity was defined as the total hours of moderate to intense activity per week and categorized into tertiles (<0.4, 0.4-2.49, >2.5 hours/week); body mass index (BMI; kg/m<sup>2</sup>) was grouped into two categories based on WHO BMI categories (non-obese: BMI<30; obese BMI≥30) (13); socioeconomic factors included employment in the past 5 years (yes or no) and highest educational attainment (high school or less than high school, some college, or college/graduate); smoking status was classified as current, past, and never smokers; and alcohol consumption was grouped into 3 categories (<1, 1-6, or 7-12 drinks/week). In the Partners Biobank sub-cohort, we obtained estimates for the association between these confounders and MDD among those with normal vitamin D levels ( $RR_{dz0}$ ) and the prevalence of different levels of the confounders ( $P_{z0}$ ,  $P_{z1}$ ). Using this information, bias factors for each confounder were calculated (Equation (1)) and used to correct the HR estimates and confidence intervals from the main analysis. Analyses were performed in SAS 9.4.

$$Bias\ factor = \frac{\sum_z RR_{dz0} P_{z1}}{\sum_z RR_{dz0} P_{z0}} \quad (1)$$

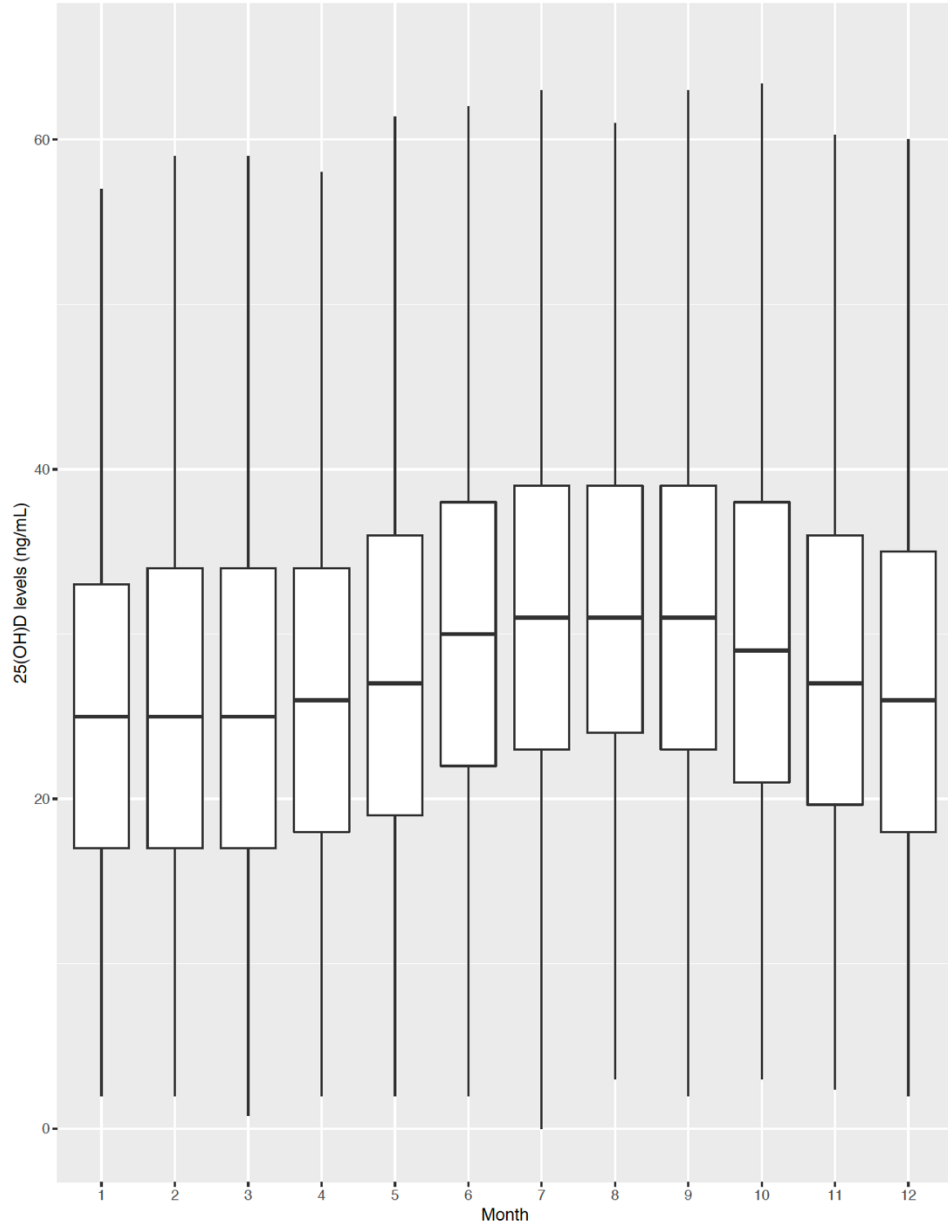
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**Supplemental Figure 1.1 Study design scheme**





**Supplemental Figure 1.2 Patterns of 25(OH)D levels by month in the study population**

**Supplemental Table 1.1 Association between mean annual serum 25(OH)D levels and incident major depressive disorder using Cox regression models**

Serum 25(OH)D	n	MDD cases (%)	Model 1		Model 2		Model 3	
			HR (95 % CI)	p	HR (95 % CI)	p	HR (95 % CI)	p
<i>Continuous</i> (per 10 ng/mL increase)	233,970	10,152 (4.3)	0.87 (0.85, 0.88)	<0.0001	0.88 (0.86, 0.90)	<0.0001	0.89 (0.87, 0.90)	<0.0001
<i>Conventional Clinical Cut-offs</i>								
Normal ( $\geq 30$ ng/mL)	104,483	4,104 (3.9)	ref	ref	ref	ref	ref	ref
Insufficient (20-29.9 ng/mL)	71,466	3,154 (4.4)	1.19 (1.14, 1.25)	<0.0001	1.19 (1.13, 1.25)	<0.0001	1.18 (1.13, 1.24)	<0.0001
Deficient (<20 ng/mL)	58,021	2,894 (5.0)	1.50 (1.42, 1.58)	<0.0001	1.43 (1.36, 1.51)	<0.0001	1.41 (1.34, 1.48)	<0.0001
<i>Insufficient vs. Deficient</i>								
Insufficient (20-29.9 ng/mL)	71,466	3,154 (4.4)	ref	ref	ref	ref	ref	ref
Deficient (<20 ng/mL)	58,021	2,894 (5.0)	1.26 (1.19, 1.32)	<0.0001	1.20 (1.15, 1.27)	<0.0001	1.19 (1.13, 1.26)	<0.0001

Abbreviations: 25(OH)D: 25-hydroxyvitamin D; CI: confidence interval; HR: hazard ratio; MDD: major depressive disorder

Model 1: Adjusted for age, sex, race/ethnicity, assay, season

Model 2: Model 1 covariates + healthcare use intensity + vitamin D supplementation + Charlson-Deyo comorbidity index

Model 3: Model 2 covariates + medical conditions and therapies indicating vitamin D testing

**Supplemental Table 1.2 Distribution of confounders in the Partners Healthcare Biobank sub-cohort**

	<b>N (%)</b>
<b>Total moderate-intense physical activity (hours/week)</b>	
Quantile 1 (<0.4)	377 (34.6)
Quantile 2 (0.4-2.49)	382 (35.0)
Quantile 3 ( $\geq$ 2.5)	332 (30.4)
<b>BMI (kg/m<sup>2</sup>)</b>	
Non-obese (<30)	833 (74.2)
Obese ( $\geq$ 30)	286 (25.8)
<b>Smoking</b>	
Current	67 (6.0)
Past	405 (36.4)
Never	642 (57.6)
<b>Alcohol (drinks/week)</b>	
<1	482 (43.7)
1-6	483 (43.8)
7-12	139 (12.5)
<b>Employment in past 5 years</b>	
Yes	893 (79.8)
No	226 (20.2)
<b>Highest educational attainment</b>	
High school or less	113 (10.2)
Some college	101 (9.0)
College or graduate	898 (80.8)

**Supplemental Table 1.3 Bias factor-corrected hazard ratios (HR) due to potential confounding by physical activity, smoking, alcohol consumption, and SES characteristics**

Vitamin D status <sup>1</sup>	Original aHR (95% CI)	Bias-corrected aHR (95% CI) <sup>2</sup>					
		Adjusted for physical activity (BF1=1.09; BF2=1.17)	Adjusted for obesity (BF1=0.99; BF2=0.98)	Adjusted for smoking (BF1=0.98; BF2= 0.87)	Adjusted for alcohol (BF1=0.99; BF2= 0.99)	Adjusted for education (BF1=1.01; BF2=1.02)	Adjusted for occupation (BF1= 0.98; BF2= 0.96)
Normal	ref	ref	ref	ref	ref	ref	ref
Insufficient	1.17 (1.12, 1.23)	1.07 (1.03, 1.13)	1.18 (1.13, 1.24)	1.19 (1.14, 1.26)	1.18 (1.13, 1.24)	1.16 (1.11, 1.22)	1.19 (1.14, 1.26)
Deficient	1.37 (1.30, 1.45)	1.17 (1.11, 1.24)	1.40 (1.33, 1.48)	1.57 (1.49, 1.67)	1.38 (1.31, 1.46)	1.34 (1.27, 1.42)	1.43 (1.35, 1.51)

Abbreviations: BF = bias factor; BF1 = bias factor for insufficient vitamin D category; BF2 = bias factor for deficient vitamin D category; CI = confidence interval; aHR = adjusted hazard ratio.

<sup>1</sup> Vitamin D status defined by the following 25(OH)D cut-offs: Deficient, <20 ng/mL; Insufficient, 20-29.9 ng/mL; Normal, ≥30 ng/mL

<sup>2</sup> Bias-corrected aHR= (Original aHR)/(BF)

## **Chapter 2. Investigating the causal relationship between serum 25-hydroxyvitamin D and depression: A two-sample bidirectional Mendelian randomization study**

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## Abstract

**Objectives:** Evidence from observational studies regarding the association between serum 25-hydroxyvitamin D [25(OH)D] levels and depression has been inconsistent. Further, the causality and directionality of the relationship remains unclear. To address this gap, we conducted a two-sample bidirectional Mendelian randomization (MR) using summary-level data from the largest genome-wide association studies to date for serum 25(OH)D and depression.

**Methods:** Six genome-wide significant single-nucleotide polymorphisms (SNPs) associated with 25(OH)D levels in 79,366 individuals were used as genetic instruments to investigate the causal effect of serum 25(OH)D on depression, while 81 SNPs that were genome-wide significantly associated with depression in 807,553 individuals were used to assess the causal effect of depression on serum 25(OH)D levels. MR effect estimates were combined using inverse-variance weighted (IVW) meta-analysis, and other methods, including weighted median, likelihood, and MR Egger.

**Results:** We did not find evidence for an effect of 25(OH)D on depression, with an IVW odds ratio of 0.98 (95% CI: 0.95, 1.01;  $p=0.12$ ) for depression per standard deviation increase in serum 25(OH)D. There was also no evidence for the effect of depression on 25(OH)D levels (Change in 25(OH)D levels comparing depression cases to controls:  $\beta=-0.008$ , 95% CI: -0.03, 0.01,  $p=0.50$ ). Results using the likelihood, MR Egger, and weighted median methods yielded consistent estimates.

**Conclusion:** The current MR study provides no evidence for a strong causal relationship between lifelong circulating 25(OH)D levels and risk of depression. However, we cannot rule out the possibility of weaker causal effects and non-linear associations.

**Keywords:** Vitamin D, depression, Mendelian randomization

## Introduction

Depression is a multifactorial disorder that likely results from a combination of genetic, physiologic, and/or environmental factors, such as life stressors, childhood adversities, inflammation, and lack of physical activity (1-5). Growing evidence suggests that vitamin D deficiency, defined by circulating 25-hydroxyvitamin D [25(OH)D], may be implicated in the etiology of depression (6, 7), given reported associations with other neuropsychiatric disorders (8, 9). Vitamin D is a neurosteroid hormone that has important effects on the brain, spanning neurological development and memory-related cognitive functions (10-13). Vitamin D receptors are also present in subcortical limbic brain regions that are implicated in depression risk (8, 14-16). Furthermore, neuromodulating and neuroprotective effects of vitamin D have been reported in animal studies, which reveal plausible biological pathways through which vitamin D may influence vulnerability to depression (8, 14, 16).

In humans, observational studies have generally found that vitamin D deficiency is associated with an increased risk for depression, with hazard ratios ranging between 1.42 and 1.80 (6, 17). However, a number of observational studies have also failed to detect associations (18-20), which suggests that other factors may explain some of the previously observed associations. First, the possibility of residual confounding cannot be excluded. That is, important factors related to both 25(OH)D and depression, including socioeconomic status, diet, physical activity, and other lifestyle characteristics, may not have been fully accounted for in prior studies. Second, some of the true associations may in fact be in the reverse direction. For instance, individuals with depression may change their outdoor activity and/or dietary habits, which may lower their 25(OH)D levels. Curiously, results from published randomized controlled trials do not support the use of vitamin D supplementation as a strategy to reduce depressive symptoms among adults (6, 7, 21, 22). However, these studies have been criticized for having relatively small sample sizes, low doses, and short treatment durations, making it difficult to draw more definitive conclusions about the relationship between vitamin D status and depression risk (6).

Mendelian randomization (MR) has emerged in recent years as a method that can strengthen causal inference and may serve as a bridge between observational and interventional studies. MR is an instrumental variable approach that uses genetic variants, typically single-nucleotide polymorphisms (SNPs), as proxies for the exposure of interest to infer causal relationships between two phenotypes (23). MR analysis is based on the principle that genetic variants are randomly assorted at conception and as a result, two common issues in observational studies can be circumvented, including (1) residual confounding, since genetic variants are unlikely associated with potential exposure-outcome confounding factors, and (2) reverse causation, as bi-directional MR allows assessment of the directionality of effects. A handful of published MR studies have been performed to understand the causal role of 25(OH)D on a range of physical and mental health outcomes (24). These studies have shown that circulating 25(OH)D levels may be causally associated with multiple sclerosis (25, 26), cancers (27, 28), and Alzheimer's disease (29), but do not appear to causally impact risk for schizophrenia (30) or Parkinson's disease (31).

In the current study, we performed a two-sample bidirectional MR analysis to determine whether there is a causal relationship between 25(OH)D levels and risk for depression. Summary-level data for these analyses came from two of the largest genome-wide association study (GWAS) meta-analyses to date: (1) the Study of Underlying Genetic Determinants of Vitamin D and Highly Related Traits (SUNLIGHT) Consortium (32) provided GWAS summary statistics for 25(OH)D; (2) the Howard *et al.* (33) study provided GWAS summary statistics for depression, based on a meta-analysis of results from the Psychiatric Genomics Consortium (PGC) (34), UK Biobank (35), and 23andMe (36). This study expands upon a recent MR analysis of vitamin D and major depressive disorder (MDD) (37) using summary data for depression from a substantially larger cohort (n=807,553 vs. n=173,005) and also examines the effect of depression on 25(OH)D levels.



## Methods

The current analysis was based solely on GWAS summary statistics from previously reported studies that have respectively obtained appropriate ethical approval (32, 34), and no individual-level data was used. Therefore, additional Institutional Review Board approval was not required for this project.

### Data Sources and SNP selection

**Serum 25(OH)D.** Summary-level data was obtained from the SUNLIGHT GWAS meta-analysis (32). This GWAS for 25(OH)D levels comprised 79,366 individuals of European ancestry and identified six independent, genome-wide significant (GWS,  $p < 5 \times 10^{-8}$ ) SNPs, which together explained ~2.84% of serum 25(OH)D variation (30). Four of these variants are located in genes that were previously implicated in the synthesis, metabolism, or transport of vitamin D (38).

**Depression.** Summary-level statistics were obtained from a meta-analysis of the three largest GWAS of depression (33), performed by the Psychiatric Genetics Consortium (PGC-MDD) (34), UK Biobank (35), and 23andMe (36). The study, which included 246,363 depression cases and 561,190 controls, identified 102 SNPs that were GWS-associated with depression. The GWS SNPs, combined in a polygenic risk score (PRS), explained up to 3% of the depression variation on the liability scale. Several of the identified SNPs have previously been linked to neuropsychiatric traits or disorders.

For exposure-related SNPs that were unavailable in the GWAS summary statistics for the outcome, proxy SNPs in high linkage disequilibrium ( $R^2 > 0.8$ ) were identified using LDlink, a web-based platform that utilizes data from the 1000 Genomes Project Phase 3 to explore genomic structures (39). For the 25(OH)D to depression analysis, all 6 SNPs were available in the depression GWAS. For the depression to 25(OH)D analysis, 46 of 102 SNPs were identified, 42 SNPs were matched to proxy SNPs ( $R^2$ : 0.80-1.00), and 14 SNPs were excluded due to a lack of suitable proxies in the 25(OH)D GWAS. An

additional 7 strand ambiguous SNPs (i.e., A/T or G/C SNPs), for which the forward strand could not be inferred based on the minor allele frequency (MAF), were excluded. 81 variants remained for the depression to 25(OH)D MR analysis.

### **SNP validation**

SNPs used as instrumental variables (IVs) for our MR analysis were validated by testing the following three MR assumptions: (1) strong association between the IV and exposure; (2) IV is independent of confounders of the exposure-outcome association; and (3) absence of horizontal pleiotropy, meaning that the IVs exert effects on the outcome only through the exposure and no other independent biological pathways (40, 41). By selecting genetic variants that were GWS-associated with the exposure, we ensured that the first assumption was fulfilled.

To evaluate the second assumption, we examined linkage disequilibrium (LD) patterns between all SNPs selected as genetic instruments for the analysis in 1000 Genomes Project European (CEU) samples (42), recognizing that SNPs highly correlated with other risk loci could lead to confounding (43). For any SNP-pairs with evidence of LD ( $r^2 > 0.1$ ), the variant with the strongest association (i.e., smallest p-value) with the exposure was retained for analysis.

To verify the third MR assumption, we assessed for potential horizontal pleiotropic effects of the selected SNPs that may arise from their effects on confounders of the exposure-outcome association. Pleiotropy is less likely for the selected 25(OH)D IVs, most of which map strongly to 25(OH)D pathways. In addition, prior work in the 1958 British Birth Cohort found no associations between four of the 25(OH)D-related SNPs and potential pleiotropic pathways such as physical activity, sun exposure, and smoking habits (all  $p > 0.05$ ) (44). We also performed a search of associations between the selected IVs and other health outcomes/traits and potential confounders using PhenoScanner, a curated database of a publicly available GWAS data (45) (**Supplemental Tables 2.1 and 2.2**). Additional analyses were performed, as summarized below, to evaluate the effects on the results of removing SNPs with any

evidence of having pleiotropic pathways and evaluate the robustness of our findings to violations of the pleiotropy assumption.

### **Statistical Analysis**

All analyses were performed using the *TwoSampleMR* (version 0.4.16) package in R version 3.3.1 (46). The primary MR analysis was performed using the inverse-variance weighted (IVW) meta-analysis, which takes a weighted mean of ratio estimates from multiple IVs (47). For the 25(OH)D to depression MR analysis, the ratio estimate ( $\exp(\hat{\beta})$ ) is interpreted as the odds ratio (OR) of depression per standard deviation (SD) increase in  $\ln(25(OH)D)$  concentrations. The SD (1 SD=0.33 nmol/L on the log-transformed scale) was derived from a population-based Swedish Mammography cohort (48), and has been used in published 25(OH)D MR analyses (49). For the depression to serum 25(OH)D MR analysis,  $\hat{\beta}$  is interpreted as the difference in  $\ln(25(OH)D)$  levels comparing cases of depression to controls.

The Cochran's Q statistic was calculated to test for statistical heterogeneity, which could indicate the presence of horizontal pleiotropy (50, 51). To evaluate the robustness of the causal estimates, the weighted median, MR-Egger regression, and likelihood methods were also employed (50, 52, 53). Briefly, the weighted median ratio estimate is the median of the weighted empirical distribution function of all SNP ratio estimates (53); this method provides an unbiased estimate of the causal effect even when up to half of the SNPs are invalid instruments. MR Egger involves fitting a weighted linear regression of the SNP-outcome against SNP-exposure effect estimates (50) and provides an unbiased causal effect estimate by adjusting for average pleiotropic bias. The likelihood-based approach calculates a maximum likelihood estimate (MLE), with the SNP-exposure and SNP-outcome associations jointly modeled under a bivariate normal distribution. This method produces valid estimates even when the SNP-exposure and SNP-outcome associations are correlated (47). MR-PRESSO was also implemented to detect and remove horizontal pleiotropic outliers (54).

We also performed sensitivity analyses by excluding SNPs that may potentially have pleiotropic effects. For the 25(OH)D to depression analysis, analyses were performed removing the following SNPs: (1) rs12785878 (*DHCR7*), found to be strongly associated with ancestry (44); (2) rs3755967 (*GC*), which could affect depression through opposing effects on the bioavailability of vitamin D, rather than influencing 25(OH)D levels (55). For the depression to 25(OH)D analysis, we performed additional analyses excluding SNPs that were GWS-associated with potential confounders of the association between 25(OH)D levels and depression, such as educational attainment and body mass index (**Supplemental Table 2.2**).

### Power Calculations

Statistical power for the bidirectional MR analysis was estimated using the online mRnd tool (<http://cnsgenomics.com/shiny/mRnd/>), developed by Brion and colleagues (56). Power estimations were made for a range of effect estimates (OR=0.75-0.98 per SD increase in ln(25(OH)D); beta=0.01-2.0 comparing depression case to control status), with type I error fixed at  $\alpha=0.05$  and assuming no heterogeneity. The study had approximately 80% power to detect an effect size of OR=0.93 for the risk of depression per 1 SD increase in ln(25(OH)D) (nmol/L), and 80% power to detect an effect size of beta=2.0 for the difference in ln(25(OH)D) comparing depression cases to controls (**Supplemental Figure 2.1**).

## Results

### Effect of 25(OH)D on Depression

Of the six 25(OH)D variants included in the MR analysis, the SNP in the *GC* gene (rs3755967) had the strongest association with 25(OH)D levels ( $\beta=0.09$  per-allele difference in ln(25(OH)D),  $p=4.74 \times 10^{-343}$ ) (**Supplemental Table 2.3**). None of the 25(OH)D SNPs were GWS-associated with depression (all  $p \geq 0.01$ ) (**Supplemental Table 2.3**). There was no evidence for an association between

25(OH)D levels on depression using the IVW model (OR of depression per 1 SD increase in 25(OH)D levels=0.98; 95% CI: 0.95, 1.01; p=0.12) (Table 2.1, Figure 2.1).

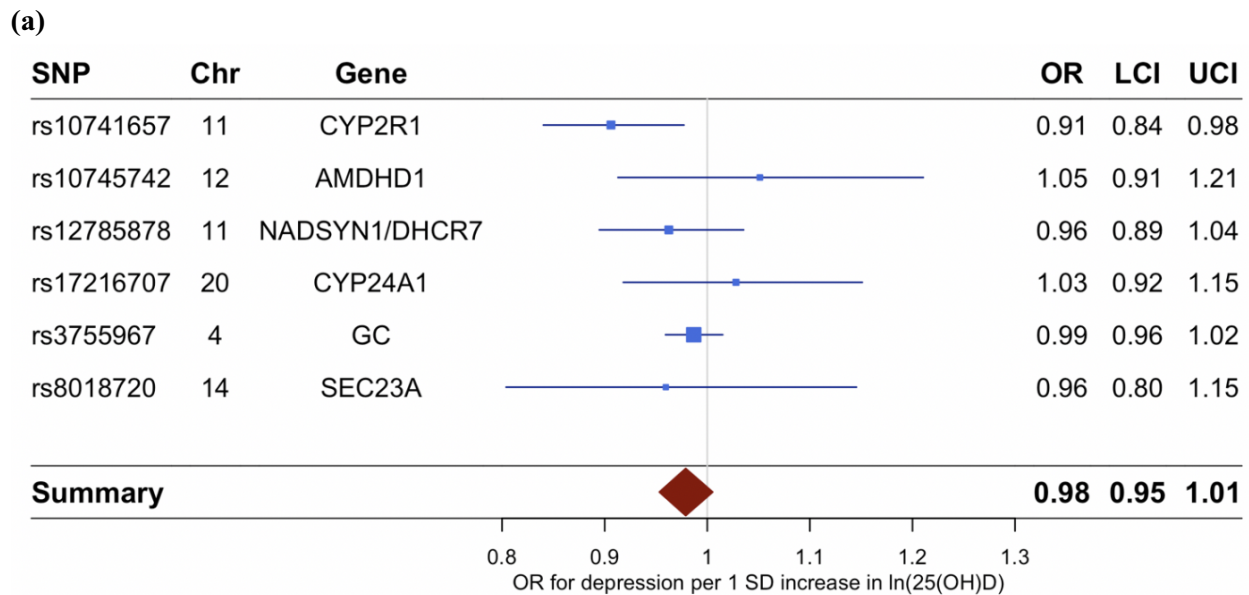
**Table 2.1 MR estimates for the effect of 25(OH)D levels on risk for depression**

Method <sup>a</sup>	No. SNPs	OR <sup>b</sup>	95% CI	p-value
IVW	6	0.98	0.95, 1.01	0.12
Maximum likelihood	6	0.98	0.96, 1.00	0.08
Weighted median	6	0.98	0.96, 1.01	0.19
MR Egger	6	0.98	0.93, 1.04	0.59

IVW: Inverse-variance weighted; MR: Mendelian randomization; OR: odds ratio; SNPs: single-nucleotide polymorphism.

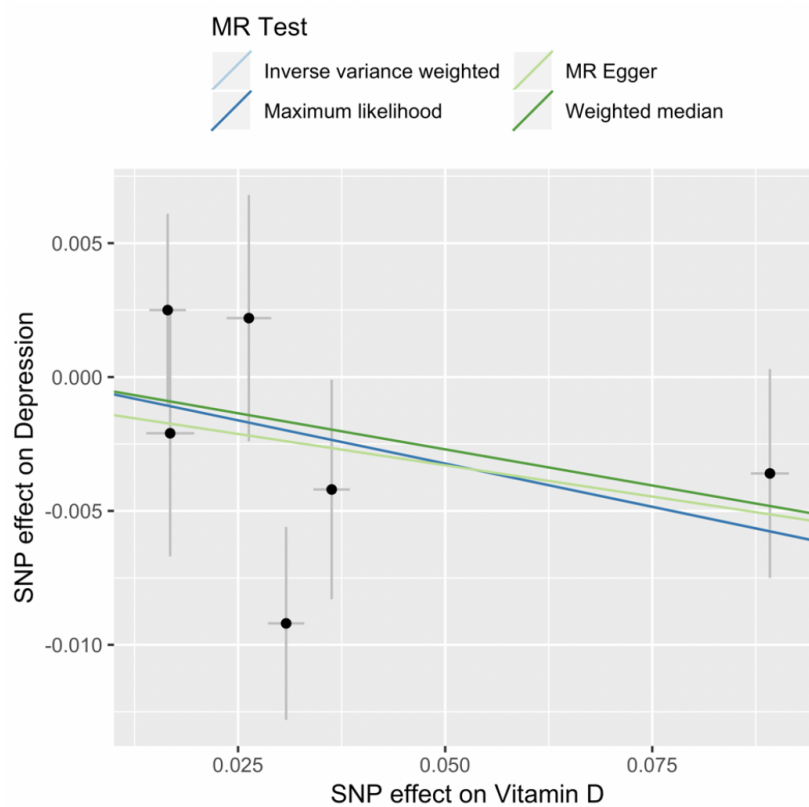
<sup>a</sup>No MR-PRESSO outliers detected

<sup>b</sup>OR=odds ratio for depression per 1 SD increase in ln(25(OH)D) (nmol/L)



**Figure 2.1 Mendelian randomization (MR) plots for the association between 25(OH)D levels and the risk for depression.** (a) Forest plot for the individual and combined SNP effects using the inverse-variance weighted (IVW) model. (b) Scatterplot for the effect of SNPs on ln(25(OH)D) and depression with the slope representing the MR effect estimate for each method.

(b)



**Figure 2.1 (continued)**

There was no evidence of horizontal pleiotropy based on the tests of heterogeneity ( $Q=6.27$ ,  $p=0.28$ ) and Egger regression (Intercept=1.00; 95% CI: 0.99, 1.01;  $p=0.80$ ). MR-PRESSO also did not detect any outliers ( $p=0.31$ ). The causal effect estimates were similar using the likelihood (OR= 0.98; 95% CI: 0.96, 1.00;  $p=0.08$ ), weighted median (OR=0.98; 95% CI: 0.96, 1.01;  $p=0.19$ ), and MR Egger (OR=0.98; 95% CI: 0.93, 1.04;  $p=0.59$ ) methods (**Table 2.1**). Sensitivity analyses excluding the rs12785878 (*DHCR7*) and rs3755967 (*GC*) SNPs did not substantially change the MR effect estimates (**Supplemental Table 2.4**).

### **Effect of Depression on 25(OH)D**

Among the 81 SNPs included in this MR analysis, the SNP that had the strongest association with depression was rs2568958 (OR=1.04;  $p=8.47 \times 10^{-25}$ ), which is found in the uncharacterized RNA gene

*LOC105378797* (**Supplemental Table 2.5**). Twelve SNPs (rs301799, rs7685686, rs11135349, rs3823624, rs3793577, rs58621819, rs57344483, rs4772087, rs9592461, rs7193263, rs143186028, and rs5995992) were nominally associated with serum 25(OH)D ( $p < 0.05$ ), but none reached GWS (**Supplemental Table 2.5**). There was no evidence for causal effects of depression on 25(OH)D levels using the IVW model ( $\beta = -0.008$ ; 95% CI: -0.03, 0.01;  $p = 0.50$ ) (**Table 2.2, Figure 2.2**).

**Table 2.2 MR estimates for the effect of depression case status on 25(OH)D levels.**

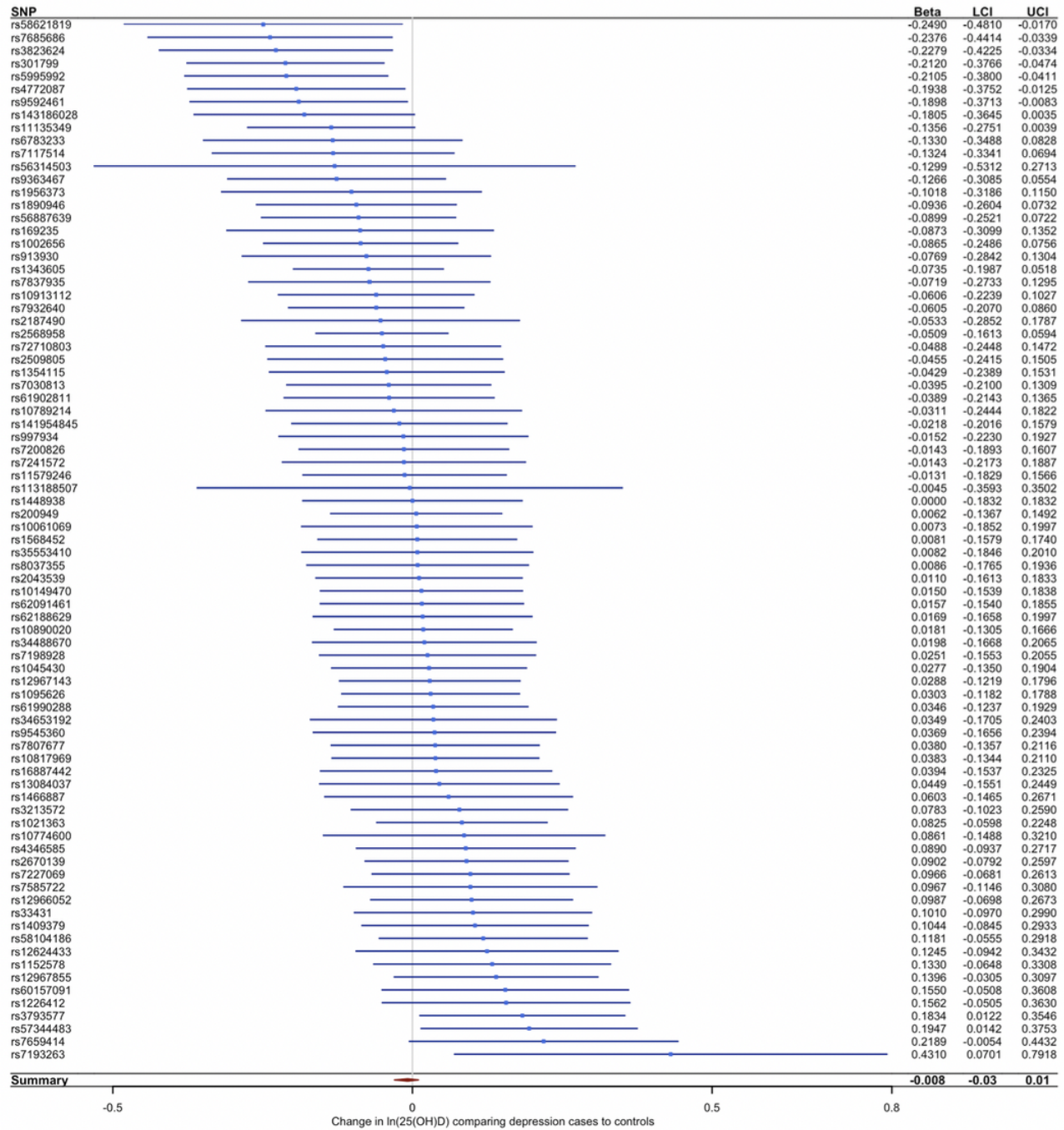
<b>Method<sup>a</sup></b>	<b>No. SNPs</b>	<b>Beta<sup>b</sup></b>	<b>95% CI</b>	<b>p-value</b>
IVW	81	-0.008	-0.03, 0.01	0.50
Maximum likelihood	81	-0.008	-0.03, 0.01	0.45
Weighted median	81	0.008	-0.02, 0.04	0.59
MR Egger	81	-0.027	-0.16, 0.11	0.70

IVW: Inverse-variance weighted; MR: Mendelian randomization; SNPs: single-nucleotide polymorphism.

<sup>a</sup>No MR-PRESSO outliers detected

<sup>b</sup>Beta=Difference in  $\ln(25(OH)D)$  comparing depression cases to controls

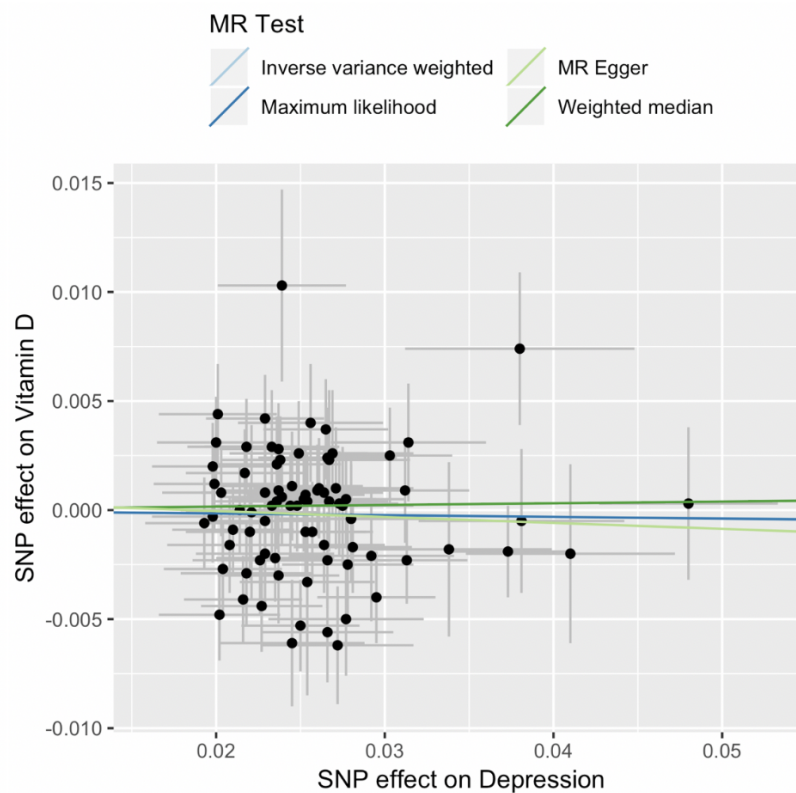
(a)



**Figure 2.2 MR plots for the association between depression case-control status and  $\ln(25(\text{OH})\text{D})$  levels.** (a) Forest plot for the individual and combined SNP effects using the IVW model. (b) Scatterplot for the effect of SNPs on depression and  $\ln(25(\text{OH})\text{D})$  with the slope representing the MR effect estimate for each method.



(b)



**Figure 2.2 (continued)**

There was little evidence for horizontal pleiotropy, with moderate heterogeneity across SNPs ( $Q=100.77$ ;  $p=0.06$ ) and an Egger regression intercept close to the null (Intercept=0.001; 95% CI: -0.006, 0.008;  $p=0.78$ ). MR-PRESSO also did not detect any outliers ( $p=0.09$ ). The causal effect estimates yielded from the likelihood ( $\beta=-0.008$ ; 95% CI: -0.03; 0.01;  $p=0.45$ ), weighted median ( $\beta=0.008$ ; 95% CI: -0.02, 0.04;  $p=0.59$ ), and MR Egger ( $\beta=-0.027$ ; 95% CI: -0.16, 0.11;  $p=0.70$ ) approaches yielded similar results (**Table 2.2**). Sensitivity analysis excluding the 12 SNPs that were associated with potential confounding factors resulted did not change the pattern of effects ( $\beta=-0.002$ ; 95% CI: -0.03, 0.02;  $p=0.85$ ) (**Supplemental Table 2.6**).

## Discussion

Both depression and low vitamin D levels are rising public health issues worldwide, yet their association remains controversial. Understanding whether vitamin D is a causal risk factor for depression would have important implications for depression prevention efforts, since vitamin D levels are easily modifiable through vitamin D supplementation, and lifestyle and/or dietary changes. As such, we performed a MR analysis to examine the causal relationship between 25(OH)D levels and risk for depression. Using genetic instruments from the largest published GWAS meta-analysis for 25(OH)D and depression to date, we employed various MR methods including IVW, likelihood-based MR, weighted median, and MR Egger to explore this association. We did not find evidence for a causal effect of lifelong average 25(OH)D levels on the risk for depression or causal effects in the reverse direction.

Our results are consistent with those reported in a previous MR analysis, which showed no association between lifelong 25(OH)D levels and MDD in European adults (37). The present study also corroborates the findings of a recent longitudinal study that assessed the association between continuous circulating 25(OH)D and incident MDD (19) in older adults, where the authors found a non-significant inverse effect of serum 25(OH)D levels on incident MDD that was of comparable magnitude (HR=0.95, 95% CI: 0.86, 1.05) to that found in the current study (19). Using a bidirectional MR, we were also able to conduct a formal test of whether depression influences 25(OH)D levels, an important analysis in light of concerns about reverse causation in observational studies. While this study may have lacked sufficient power to detect weaker effects, our results suggest that depression did not have substantial effects on 25(OH)D levels, and that prior cross-sectional vitamin D-depression associations may not be explained by the direct effects of depression on vitamin D levels. These results together suggest that modest and incremental changes in 25(OH)D levels are unlikely to reduce the risk for depression.

However, these results should be interpreted in the context of several factors. MR is a powerful statistical method that can be used to establish or refute observational results, but there are important differences between the hypotheses tested under the MR framework compared to those tested in

traditional epidemiological cohort studies. First, 25(OH)D-related SNPs influence lifelong exposures to circulating vitamin D, therefore the MR results are consistent with the lack of a causal relationship between lower lifelong 25(OH)D levels and depression risk. On the other hand, observational studies have conventionally investigated the association between depression and more proximal 25(OH)D levels (i.e., incidence of depression within 1-5 years of vitamin D measurement (57, 58)). Hence, it remains possible that more recent exposures to vitamin D, rather than average lifetime exposures, have causal effects on depression risk. Second, we cannot rule out the existence of specific sensitive periods during which vitamin D levels may have stronger effects on depression risk. Prior studies have found that prenatal 25(OH)D exposures are associated with mental health outcomes, such as schizophrenia (59-61) and neuropsychological scores (62), and that childhood 25(OH)D levels are associated with adolescent depression (63). Therefore, it is possible that early life exposures to 25(OH)D during important periods of brain development have stronger effects on the risk for depression, compared to average lifelong exposures. Third, observational studies have primarily reported the impact of vitamin D deficiency on depression risk; it is plausible that levels below a certain threshold, but not incremental linear changes in circulating vitamin D levels, that are causally related to depression risk (17). Indeed, in a meta-analysis of cohort studies, authors found an increased risk of depression comparing those in the highest vs. lowest vitamin D categories, but no association for 20 nmol/L differences in vitamin D levels (17). While some genetic instruments may be associated with vitamin D deficiency (64), such causal threshold effects could not be tested with currently available MR methods and GWAS data. Further studies are needed to elucidate these potentially complex relationships.

A major strength of this MR study is that we used the largest available GWAS summary statistics for both depression and 25(OH)D levels at the time of writing. Furthermore, participants in both study populations were of European ancestry, reducing potential bias due to population stratification. However, our analysis is not without limitations. First, while the sample sizes were large, the SNPs that we selected as genetic instruments accounted for a relatively small proportion of variation in both 25(OH)D (~2.8%) (32) and depression (~3%) (34). The study therefore lacked statistical power to detect weaker causal

relationships between serum 25(OH)D and depression risk. Second, given unknown functional roles of these tested SNPs, we could not completely eliminate the possibility of horizontal pleiotropy, which could bias the analyses. However, our findings were robust to multiple sensitivity analyses and there was no strong evidence, quantitatively or qualitatively, for the presence of pleiotropic effects. Third, due to a lack of available methods, we could not investigate either non-linear associations between circulating vitamin D and risk of depression nor threshold effects based on conventional clinical cut-offs for vitamin D deficiency (16). MR methods that enable testing for non-linear associations between the exposure and outcome are currently under development (65). Fourth, we could not assess the role of the biologically active form of vitamin D (1,25-dihydroxyvitamin D (1,25(OH)D), for which GWAS have not yet been performed. Therefore, conclusions cannot be drawn about lifelong average levels of 1,25(OH)D on depression risk. Fifth, our null results may be explained by canalization, which are compensatory feedback mechanisms that could buffer harmful effects of lowered vitamin D levels and could bias the association toward the null (43, 66). Lastly, the study was mainly restricted to European adults; thus, results may be less readily generalizable to younger or non-European populations with different prevalence of depression and distributions of circulating vitamin D.

In summary, this MR study does not provide evidence for a causal relationship between lifelong 25(OH)D levels and the risk for depression among individuals of European descent. However, we cannot preclude the possibility of smaller causal associations, sensitive periods, threshold effects, or effects in specific sub-populations. The largest RCT to date (VITAL-DEP), evaluating the effect of vitamin D supplements on preventing depression among approximately 25,000 older adults, is underway and will provide further evidence for the causal effects of vitamin D on depression (67). Future well-designed studies are warranted to determine evidence-based cut-offs for vitamin D deficiency, potentially allowing for stronger conclusions about the vitamin D-depression association. Similar analyses in other ethnic/racial samples could also elucidate the benefit of vitamin D supplementation in other sub-populations.

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## Appendix

**Supplemental Table 2.1 Association between 25(OH)D SNPs and other phenotypes as identified by genome-wide association studies**

<b>SNP</b>	<b>Chr</b>	<b>Gene</b>	<b>Associated traits in European populations (<math>p &lt; 5E-08</math>)</b>
rs3755967	4	GC	Blood cell counts (white blood cell, myeloid white cell, basophil, eosinophil, neutrophil, granulocyte)
rs10741657	11	CYP2R1	Height
rs12785878	11	NADSYN1/ DHCR7	-
rs17216707	20	CYP24A1	Glomerular filtration rate, creatinine
rs10745742	12	AMDHD1	-
rs8018720	14	SEC23A	-

**Supplemental Table 2.2 Association between depression SNPs and other phenotypes as identified by genome-wide association studies**

SNP	Chr	Gene	Associated traits in European populations (p<5E-08)
rs2568958	1	LOC105378797	BMI, obesity, overweight, weight, gene expression, age at menarche, arm fat, basal metabolic weight, impedance of leg, leg fat, education, self-reported non-cancer illness, trunk fat, sodium in urine, seen doctor for nerves/anxiety/depression
rs10890020	1	LOC105378800	-
rs10913112	1	RFWD2	-
rs301799	1	RERE	Eosinophil/basophil/lymphocyte count, neutrophil granulocytes, allergic disease, arm fat, asthma, body fat, DBP/SBP, allergy, bone mineral density, leg fat, hypertension, trunk fat, vascular/heart problems
rs1002656	1	-	Neuroticism
rs1890946	1	NRDC	-
rs72710803	1	-	-
rs11579246	1	ELAVL4	-
rs113188507	1	-	Body fat, leg fat, trunk fat, waist circumference
rs169235	1	CACNA1E	Neuroticism
rs1466887	1	-	-
rs10789214	1	SGIP1	-
rs1568452	2	LOC105377628	Impedance of leg
rs62188629	2	-	No medication for pain relief, constipation, heartburn
rs1226412	2	LINC01876	Neuroticism
rs7585722	2	RNF103-CHMP3	-
rs1095626	3	RSRC1	BMI, forced vital capacity (FVC), height
rs4346585	3	-	-
rs141954845	3	FHIT	-
rs13084037	3	KLHDC8B	Age at menarche, hemoglobin, frequency of tiredness/lethargy, impedance of arm/whole body, miserableness, overall health rating, pulse rate, education

**Supplemental Table 2.2 (continued)**

SNP	Chr	Gene	Associated traits in European populations (p<5E-08)
rs6783233	3	LOC101926953	Age at menarche, seen doctor for nerves/anxiety/tension/depression
rs35553410	4	-	-
rs7659414	4	-	Impedance of leg
rs7685686	4	HTT	Alcohol intake frequency, tiredness/lethargy frequency
rs11135349	5	-	Hip circumference
rs10061069	5	FAM172A	-
rs60157091	5	-	Mean platelet volume
rs200949	6	HIST1H1B	Blood cell counts (eosinophil, granulocyte, hematocrit, reticulocyte, lymphocyte, monocyte, myeloid white cell, neutrophil, basophil, neutrophil, white blood cell), hemoglobin concentration, IgA deficiency, DBP, FEV-1, FVC, guilty feelings, hearing problems, intestinal malabsorption, medication for pain relief/constipation/heart burn, dental problems, potassium in urine, seen doctor/psychiatrist for nerves/anxiety/tension/depression, thyroid disorders, sarcoidosis, treatment with insulin
rs9363467	6	-	-
rs2043539	7	TMEM106B	Miserableness, mood swings, neuroticism, seen doctor for nerves/anxiety/tension/depression
rs58104186	7	-	Tiredness/lethargy frequency
rs7807677	7	CTTNBP2	Smoking
rs3823624	7	MAD1L1	Age at first live birth, arm/leg fat, body fat, BMI, impedance of leg, irritability, miserableness, self-reported non-cancer illnesses, overall health rating, education, prescription medications, trunk fat, physical activity
rs16887442	7	-	-
rs7837935	8	CYP7B1	-
rs7030813	9	PAX5	-
rs10817969	9	ASTN2	-



**Supplemental Table 2.2 (continued)**

<b>SNP</b>	<b>Chr</b>	<b>Gene</b>	<b>Associated traits in European populations (p&lt;5E-08)</b>
rs3793577	9	ELAVL2	Depressed mood, guilty feelings, neuroticism, waist circumference
rs2670139	9	DENND1A	-
rs34653192	9	LOC105376009	-
rs1354115	9	-	-
rs913930	9	-	-
rs1021363	10	SORCS3	Seen doctor for nerves/anxiety/tension/depression
rs997934	10	LOC105376345	-
rs7932640	11	GRM5	Seen doctor for nerves/anxiety/tension/depression
rs61902811	11	LOC105369501	Guilty feelings, miserableness, mood swings, neuroticism, seen doctor for nerves/anxiety/tension/depression, worry, neuroticism
rs1448938	11	DCDC1	Red blood cell count, impedance of arm/whole body
rs7117514	11	SHANK2	-
rs2509805	11	-	Red wine intake, walking, neuroticism, physical activity, worry
rs58621819	11	LTBP3	Platelet count, red cell distribution, arm/leg/trunk fat/mass, basal metabolic rate, body fat percentage, BMI, bone mineral density, impedance of arm, pulse rate, waist circumference, weight
rs57344483	11	-	-
rs2187490	11	-	-
rs56314503	12	LOC107984536	-
rs3213572	12	SPPL3	Allergic disease, height
rs10774600	12	ATP2A2	Red blood cell count, eosinophil/basophil counts
rs1343605	13	-	Sleeplessness/insomnia
rs4772087	13	STK24	-
rs9592461	13	PCDH9	-
rs1409379	13	-	-

**Supplemental Table 2.2 (continued)**

SNP	Chr	Gene	Associated traits in European populations (p<5E-08)
rs9545360	13	-	-
rs10149470	14	-	Monocyte count, height, cognitive ability, intelligence, trunk mass
rs61990288	14	LRFN5	-
rs1045430	14	AREL1	Nervous feelings, neuroticism, worrier/anxious feelings
rs1152578	14	ESR2	-
rs1956373	14	RTN1	-
rs8037355	15	-	-
rs34488670	15	SEMA6D	Tobacco smoking, education
rs56887639	16	-	Depressive symptoms
rs7200826	16	SHISA9	-
rs7198928	16	RBFOX1	Neuroticism, worrying
rs7193263	16	RBFOX1	Depressive symptoms
rs12967143	18	TCF4	Height, irritability, neuroticism, seen doctor for nerves/anxiety/tension/depression
rs12967855	18	CELF4	Fed-up feelings, guilty feelings, mood swings, neuroticism, overall health, education, seen doctor for nerves/anxiety/tension/depression, sensitivity/hurt feelings, TV watching, worrying
rs12966052	18	LOC105372125	Mood swings
rs7227069	18	DCC	Fed-up feelings, depressed mood, tiredness/lethargy, low enthusiasm/disinterest, miserableness, mood swings, neuroticism, sitting height, tense/highly strung
rs7241572	18	KCNG2	Wheeze or whistling in chest
rs62091461	18	RAB27B	Arm/leg mass, basal metabolic rate, BMI, body size, hip circumference, weight, body fat
rs33431	19	ZNF536	-
rs143186028	20	EMILIN3	-

**Supplemental Table 2.2 (continued)**

<b>SNP</b>	<b>Chr</b>	<b>Gene</b>	<b>Associated traits in European populations (p&lt;5E-08)</b>
rs12624433	20	SLC12A5	Rheumatoid arthritis
rs5995992	22	EP300	Irritability, mood swings, neuroticism, neutrophil percentage of granulocytes

**Supplemental Table 2.3 Characteristics of genome-wide significant 25(OH)D SNPs used as genetic instruments in the 25(OH)D to depression MR analysis**

SNP	Chr	Gene	Vitamin D-related pathway	EA <sup>c</sup>	Serum 25(OH)D results <sup>a</sup>		Depression results <sup>b</sup>	
					Effect on ln(25OHD) (nmol/L)	p-value	OR <sup>d</sup>	p-value
rs3755967	4	GC	Metabolism	C	0.089	4.74E-343	1.00	0.36
rs10741657	11	CYP2R1	Synthesis	A	0.031	2.05E-46	0.99	0.01
rs12785878	11	NADSYN1/ DHCR7	Synthesis	T	0.036	3.80E-62	1.00	0.31
rs17216707	20	CYP24A1	Metabolism	T	0.026	8.14E-23	1.00	0.64
rs10745742	12	AMDHD1	-	T	0.017	1.88E-14	1.00	0.49
rs8018720	14	SEC23A	-	G	0.017	4.72E-09	1.00	0.65

25(OH)D: 25-hydroxyvitamin D; Chr: chromosome; EA: effect allele; OR: odds ratio of depression per effect allele increase; SNP: single-nucleotide polymorphism.

<sup>a</sup> Summary statistics obtained from the SUNLIGHT GWAS meta-analysis (n=79,366)

<sup>b</sup> Summary statistics obtained from the PGC-MDD2, UKB, 23andMe GWAS meta-analysis (Howard *et al.*; n=807,553)

<sup>c</sup> 25(OH)D-increasing allele

<sup>d</sup> Effect size estimates represents the OR of depression per effect allele, based on the results from the PGC-MDD2, UKB, 23andMe GWAS meta-analysis.

**Supplemental Table 2.4 Sensitivity analyses MR estimates for the effect of 25(OH)D levels on the risk of depression**

IVW Model	No. SNPs	OR	95% CI	p-value
Excluding rs12785878 ( <i>DHCR7</i> )	5	0.98	0.95, 1.01	0.22
Excluding rs3755967 ( <i>GC</i> )	5	0.96	0.91, 1.01	0.11

IVW: Inverse-variance weighted; OR: odds ratio for depression per 1 SD increase in ln(25(OH)D); SNPs: single-nucleotide polymorphism

**Supplemental Table 2.5 Characteristics of genome-wide significant depression SNPs used as genetic instruments in the depression to 25(OH)D MR analysis**

SNP	Chr	Gene	Depression results <sup>a</sup>				Serum 25(OH)D results <sup>b</sup>					
			EA <sup>c</sup>	EAF	OR	p	Proxy SNP <sup>d</sup>	R2	EA <sup>e</sup>	Effect on ln(25(OH)D) <sup>f</sup>	SE	p
rs2568958	1	LOC105378797	A	0.62	1.04	8.47E-25	-	-	-	-0.002	0.00	0.37
rs10890020	1	LOC105378800	A	0.52	0.97	4.03E-15	rs10789363	1.00	A	-5.00E-04	0.00	0.82
rs10913112	1	RFWD2	T	0.38	0.97	3.40E-13	-	-	-	0.002	0.00	0.47
rs301799	1	RERE	T	0.57	0.98	1.36E-12	-	-	-	0.005	0.00	0.01
rs1002656	1	-	T	0.70	0.97	3.74E-12	-	-	-	0.002	0.00	0.29
rs1890946	1	NRDC	T	0.47	0.98	2.68E-11	-	-	-	0.002	0.00	0.28
rs72710803	1	-	A	0.91	0.96	5.29E-11	rs11579578	0.96	T	-0.002	0.00	0.63
rs11579246	1	ELAVL4	A	0.91	1.04	5.71E-10	-	-	-	-5.00E-04	0.00	0.87
rs113188507	1	-	A	0.28	1.02	1.87E-08	rs35020936	0.95	A	1.00E-04	0.00	0.98
rs169235	1	CACNA1E	A	0.75	0.98	2.98E-08	rs199930	0.92	T	-0.002	0.00	0.46
rs1466887	1	-	T	0.55	0.98	4.12E-08	rs1542886	0.80	A	-0.001	0.00	0.57
rs10789214	1	SGIP1	T	0.57	1.02	4.44E-08	-	-	-	-6.00E-04	0.00	0.79
rs1568452	2	LOC105377628	T	0.39	1.03	8.12E-12	-	-	-	2.00E-04	0.00	0.91
rs62188629	2	-	A	0.31	1.02	7.13E-10	rs7586277	0.99	A	4.00E-04	0.00	0.84
rs1226412	2	LINC01876	T	0.79	1.03	3.46E-09	rs2697360	0.98	A	0.004	0.00	0.13
rs7585722	2	RNF103-CHMP3	T	0.85	0.97	2.68E-08	-	-	-	-0.003	0.00	0.37
rs1095626	3	RSRC1	T	0.58	0.97	7.13E-14	rs827113	1.00	T	8.00E-04	0.00	0.69
rs4346585	3	-	T	0.70	0.98	7.13E-10	rs13058822	1.00	A	0.002	0.00	0.35

**Supplemental Table 2.5 (continued)**

SNP	Chr	Gene	Depression results <sup>a</sup>				Serum 25(OH)D results <sup>b</sup>					
			EA <sup>c</sup>	EAF	OR	p	Proxy SNP <sup>d</sup>	R2	EA <sup>e</sup>	Effect on ln(25(OH)D) <sup>f</sup>	SE	p
rs141954845	3	FHIT	A	0.39	1.02	8.15E-10	rs1916801	0.99	A	5.00E-04	0.00	0.81
rs13084037	3	KLHDC8B	A	0.77	0.98	7.08E-09	-	-	-	-0.001	0.00	0.65
rs6783233	3	LOC101926953	T	0.28	1.02	2.90E-08	-	-	-	-0.003	0.00	0.22
rs35553410	4	-	T	0.75	0.98	1.42E-09	rs1022079	0.97	A	2.00E-04	0.00	0.94
rs7659414	4	-	A	0.58	0.98	1.20E-08	-	-	-	-0.004	0.00	0.06
rs7685686	4	HTT	A	0.58	1.02	2.57E-08	-	-	-	-0.005	0.00	0.02
rs11135349	5	-	A	0.47	0.97	6.04E-17	rs10035449	0.82	T	0.004	0.00	0.05
rs10061069	5	FAM172A	C	0.22	0.97	8.15E-11	-	-	-	-2.00E-04	0.00	0.93
rs60157091	5	-	T	0.52	1.02	1.42E-08	rs10939933	1.00	C	0.003	0.00	0.14
rs200949	6	HIST1H1B	A	0.87	1.05	2.53E-19	-	-	-	3.00E-04	0.00	0.94
rs9363467	6	-	T	0.60	1.02	6.44E-11	rs9351533	0.93	A	0.003	0.00	0.18
rs2043539	7	TMEM106B	A	0.42	1.03	9.89E-15	-	-	-	3.00E-04	0.00	0.91
rs58104186	7	-	A	0.47	1.02	1.82E-11	rs11561993	0.99	T	0.003	0.00	0.18
rs7807677	7	CTTNBP2	T	0.55	1.02	1.82E-11	-	-	-	9.00E-04	0.00	0.66
rs3823624	7	MAD1L1	T	0.81	1.03	1.99E-09	-	-	-	-0.006	0.00	0.02
rs16887442	7	-	T	0.43	1.02	8.62E-09	rs4556040	1.00	T	-8.00E-04	0.00	0.70
rs7837935	8	CYP7B1	T	0.15	0.97	3.34E-09	-	-	-	0.002	0.00	0.48

Supplemental Table 2.5 (continued)

SNP	Chr	Gene	Depression results <sup>a</sup>				Serum 25(OH)D results <sup>b</sup>					
			EA <sup>c</sup>	EAF	OR	p	Proxy SNP <sup>d</sup>	R2	EA <sup>e</sup>	Effect on ln(25(OH)D) <sup>f</sup>	SE	p
rs7030813	9	PAX5	T	0.37	1.03	3.07E-12	rs1329572	0.94	A	-0.001	0.00	0.66
rs10817969	9	ASTN2	T	0.72	1.03	3.11E-11	rs2418449	1.00	T	0.001	0.00	0.67
rs3793577	9	ELAVL2	A	0.47	0.98	8.41E-11	-	-	-	-0.004	0.00	0.03
rs2670139	9	DENND1A	T	0.76	0.97	1.21E-10	-	-	-	-0.002	0.00	0.30
rs34653192	9	LOC105376009	C	0.32	0.98	2.23E-09	rs1414019	0.91	A	8.00E-04	0.00	0.75
rs1354115	9	-	A	0.62	1.02	7.08E-09	-	-	-	-9.00E-04	0.00	0.68
rs913930	9	-	A	0.64	0.98	2.42E-08	-	-	-	0.002	0.00	0.45
rs1021363	10	SORCS3	A	0.35	1.03	4.41E-16	-	-	-	0.003	0.00	0.24
rs997934	10	LOC105376345	T	0.38	1.02	4.81E-08	-	-	-	-3.00E-04	0.00	0.90
rs7932640	11	GRM5	T	0.44	1.03	1.62E-15	-	-	-	-0.002	0.00	0.41
rs61902811	11	LOC105369501	A	0.37	0.97	1.40E-12	rs17602038	0.98	T	-0.001	0.00	0.67
rs1448938	11	DCDC1	A	0.42	1.02	1.30E-09	-	-	-	0.000	0.00	0.99
rs7117514	11	SHANK2	A	0.54	0.98	7.29E-09	rs11237238	0.97	A	0.003	0.00	0.20
rs2509805	11	-	T	0.32	1.02	9.17E-09	rs682503	0.81	C	-0.001	0.00	0.64
rs58621819	11	LTBP3	A	0.79	0.98	1.57E-08	rs947791	0.99	A	-0.006	0.00	0.04
rs57344483	11	-	A	0.93	0.96	1.82E-08	rs7115915	1.00	T	-0.007	0.00	0.04
rs2187490	11	-	T	0.91	0.97	3.82E-08	-	-	-	0.002	0.00	0.66

Supplemental Table 2.5 (continued)

SNP	Chr	Gene	Depression results <sup>a</sup>				Serum 25(OH)D results <sup>b</sup>					
			EA <sup>c</sup>	EAF	OR	p	Proxy SNP <sup>d</sup>	R2	EA <sup>e</sup>	Effect on ln(25(OH)D) <sup>f</sup>	SE	p
rs56314503	12	LOC107984536	T	0.75	0.97	2.95E-10	rs41466449	0.96	A	0.003	0.01	0.53
rs3213572	12	SPPL3	A	0.47	1.02	7.61E-10	-	-	-	0.002	0.00	0.39
rs10774600	12	ATP2A2	T	0.17	0.97	3.39E-08	rs1265775	0.96	A	-0.002	0.00	0.49
rs1343605	13	-	A	0.38	1.03	6.23E-18	rs2806949	0.99	A	-0.002	0.00	0.26
rs4772087	13	STK24	T	0.37	1.02	3.91E-10	-	-	-	-0.004	0.00	0.03
rs9592461	13	PCDH9	A	0.49	1.02	9.10E-10	-	-	-	-0.004	0.00	0.04
rs1409379	13	-	T	0.76	1.03	1.67E-09	-	-	-	0.003	0.00	0.27
rs9545360	13	-	A	0.18	0.97	5.02E-09	rs9531035	0.99	A	-0.001	0.00	0.73
rs10149470	14	-	A	0.49	0.97	3.72E-14	-	-	-	-4.00E-04	0.00	0.85
rs61990288	14	LRFN5	A	0.51	0.97	1.68E-13	rs17781288	0.99	C	9.00E-04	0.00	0.66
rs1045430	14	AREL1	T	0.48	0.98	7.31E-13	-	-	-	-7.00E-04	0.00	0.72
rs1152578	14	ESR2	T	0.44	0.98	6.36E-10	-	-	-	-0.003	0.00	0.18
rs1956373	14	RTN1	T	0.74	0.98	2.06E-08	-	-	-	0.002	0.00	0.35
rs8037355	15	-	T	0.56	0.98	3.94E-11	rs16964908	0.88	A	-2.00E-04	0.00	0.92
rs34488670	15	SEMA6D	T	0.79	0.98	6.03E-09	rs12901436	1.00	C	-5.00E-04	0.00	0.83
rs56887639	16	-	A	0.73	0.97	1.51E-12	rs4456522	0.93	A	-0.003	0.00	0.28
rs7200826	16	SHISA9	T	0.26	1.03	3.74E-12	rs12921037	0.84	C	4.00E-04	0.00	0.87
rs7198928	16	RBFOX1	T	0.62	1.02	4.45E-11	rs3785238	0.91	A	-6.00E-04	0.00	0.78



**Supplemental Table 2.5 (continued)**

SNP	Chr	Gene	Depression results <sup>a</sup>				Serum 25(OH)D results <sup>b</sup>					
			EA <sup>c</sup>	EA <sup>f</sup>	OR	p	Proxy SNP <sup>d</sup>	R2	EA <sup>e</sup>	Effect on ln(25(OH)D) <sup>f</sup>	SE	p
rs7193263	16	RBFOX1	A	0.67	0.98	4.33E-10	rs7188257	0.88	T	-0.010	0.00	0.02
rs12967143	18	TCF4	C	0.70	0.97	3.70E-16	-	-	-	-9.00E-04	0.00	0.70
rs12967855	18	CELF4	A	0.33	1.03	1.18E-12	-	-	-	0.004	0.00	0.10
rs12966052	18	LOC105372125	C	0.18	0.97	1.25E-11	-	-	-	-0.003	0.00	0.25
rs7227069	18	DCC	A	0.43	1.02	1.50E-11	rs7232543	0.96	A	-0.002	0.00	0.25
rs7241572	18	KCNG2	A	0.20	1.03	2.70E-10	-	-	-	-4.00E-04	0.00	0.88
rs62091461	18	RAB27B	T	0.23	0.97	1.95E-09	rs12456731	0.81	T	-4.00E-04	0.00	0.85
rs33431	19	ZNF536	T	0.61	1.02	4.81E-08	-	-	-	0.002	0.00	0.34
rs143186028	20	EMILIN3	T	0.18	1.03	2.29E-09	rs6065338	0.95	A	-0.005	0.00	0.05
rs12624433	20	SLC12A5	A	0.26	1.02	7.44E-09	-	-	-	0.003	0.00	0.26
rs5995992	22	EP300	T	0.72	0.97	1.30E-11	rs11090039	0.85	A	-0.006	0.00	0.02

25(OH)D: 25-hydroxyvitamin D; Chr: chromosome; EA: effect allele; OR: odds ratio of depression per effect allele increase; SNP: single-nucleotide polymorphism.

<sup>a</sup> Summary statistics obtained from the PGC-MDD2, UKB, 23andMe GWAS meta-analysis (Howard *et al.*; n=807,553)

<sup>b</sup> Summary statistics acquired from the SUNLIGHT GWAS meta-analysis (n=79,366)

<sup>c</sup> Allele associated with the increased risk of depression

<sup>d</sup> Proxy SNP in SUNLIGHT GWAS

<sup>e</sup> Effect allele in the SUNLIGHT GWAS (prior to data harmonization)

<sup>f</sup> Effect size estimates represent the change in log-transformed 25(OH)D (nmol/L) per effect allele, based on the beta coefficients in the SUNLIGHT GWAS

**Supplemental Table 2.6 Sensitivity analyses MR estimates for the effect of depression case status on 25(OH)D levels**

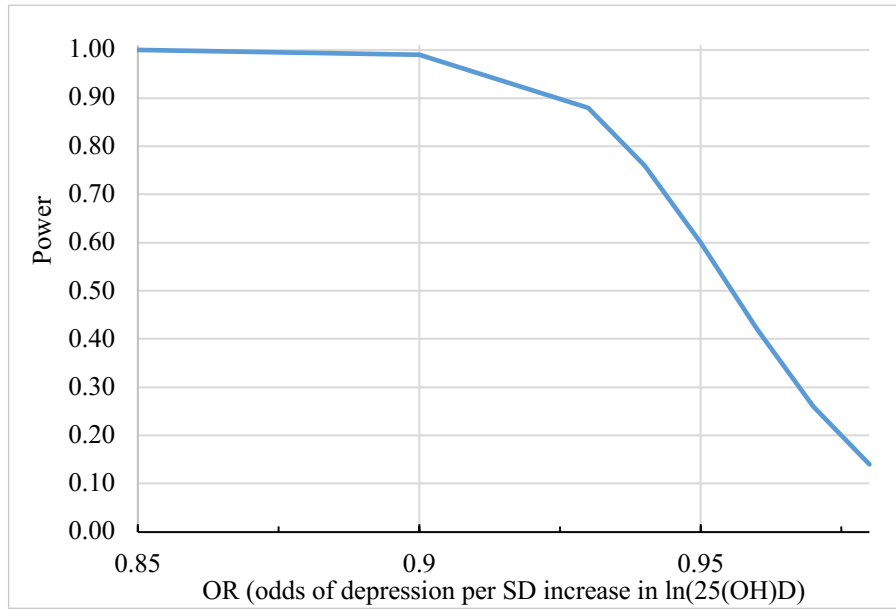
<b>IVW Model</b>	<b>No. SNPs</b>	<b>Beta<sup>a</sup></b>	<b>95% CI</b>	<b>p-value</b>
Excluding 12 SNPs with potential associations with confounding factors <sup>b</sup>	69	-0.002	-0.03, 0.02	0.85

CI: confidence interval; IVW: inverse-variance weighted; SNPs: single-nucleotide polymorphism.

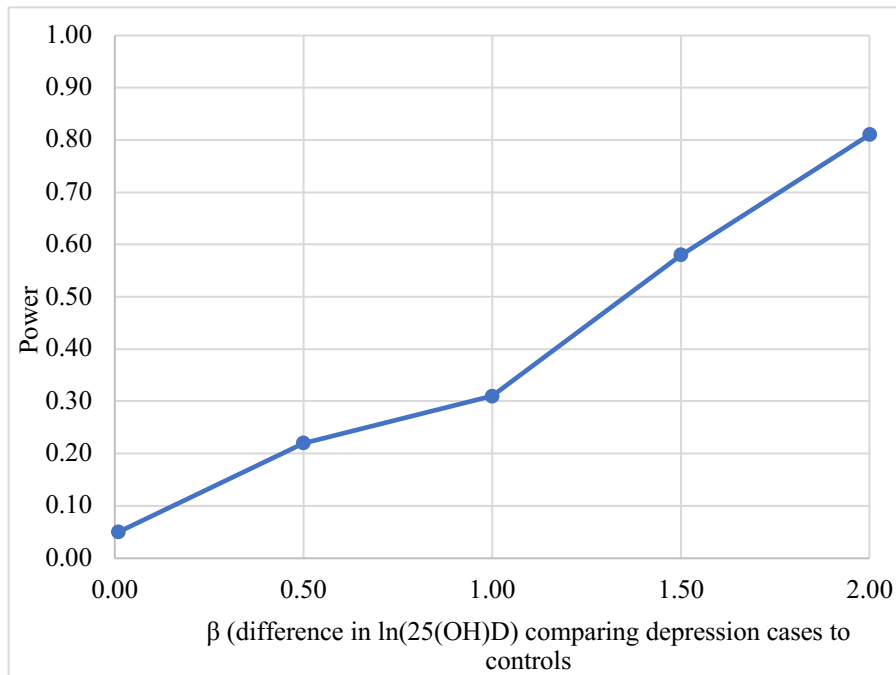
<sup>a</sup> Beta: Change in ln(25(OH)D) comparing depression case to control status

<sup>b</sup> Excluded SNPs: rs2568958, rs301799, rs113188507, rs1095626, rs13084037, rs7807677, rs3823624, rs2509805, rs58621819, rs34488670, rs12967855, and rs62091461

(a)



(b)



**Supplemental Figure 2.1 Power calculation plots for varying effect estimates for the MR analyses.**  
(a) ln(25(OH)D) (nmol/L) to depression and (b) depression to ln(25(OH)D) (nmol/L).

### **Chapter 3. Maternal vitamin D status during pregnancy and childhood/adolescent depression: Results from the Avon Longitudinal Study of Parents and Children (ALSPAC)**

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## Abstract

**Background:** Low maternal vitamin D levels [serum 25-hydroxyvitamin D (25(OH)D)] during pregnancy have been linked to offspring neuropsychiatric outcomes such as schizophrenia and autism, but studies on depression are lacking. This study examined the association between maternal vitamin D status during pregnancy and offspring depression during childhood and adolescence, and whether any associations were modified by offspring genetic risk for depression.

**Methods:** Mother-singleton offspring pairs in the Avon Longitudinal Study of Parents and Children (ALSPAC) that had maternal 25(OH)D measurements, offspring genetic data, and offspring depression measures collected in childhood (mean age=10.6 years; n=2938) and/or adolescence (mean age=13.8 years; n=2485) were included in the analyses. Using multivariable logistic regression models, we assessed relations of maternal vitamin D status and offspring polygenic risk score (PRS) for depression to childhood/adolescent depression risk.

**Results:** There was little evidence for an association between maternal vitamin D status during pregnancy and offspring depression in childhood (p-trend=0.72) or adolescence (p-trend=0.07), after adjusting for potential confounders. Offspring depression PRS were independently associated with childhood depression (p-trend=0.003) but did not interact with maternal vitamin D status. These results were robust to adjustments for potential confounders and different cut-offs for vitamin D insufficiency/deficiency.

**Conclusion:** These findings suggest that maternal vitamin D status during pregnancy does not affect the offspring's risk for early life depression.

**Keywords:** Vitamin D, polygenic risk, gene-environment interactions, depression

## Introduction

Vitamin D deficiency among pregnant women is highly prevalent globally, and particularly in geographical regions at higher latitudes, such as Northern Europe and North America (1, 2), where it affects up to 80% of dark-skinned women and 47% of light-skinned women (2-5). Maintaining sufficient vitamin D levels during pregnancy is important, given that it is the sole source of fetal vitamin D, which may play a crucial role in fetal programming and brain development. Since vitamin D levels can be easily increased by lifestyle modifications and supplementation, understanding the influence of maternal vitamin D levels on offspring health outcomes can inform early prevention efforts.

Possible mechanisms linking gestational vitamin D exposure to neurodevelopment include the role of vitamin D in neuronal growth and the signaling and regulation of endocrine functions (6). Animal studies reported that vitamin D deficiency during pregnancy resulted in alterations in brain morphology and functioning of rat offspring with long-lasting effects on behavior spanning into adulthood (6-9). Epidemiological research in humans has provided some evidence suggesting that maternal vitamin D status, defined by circulating levels of 25-hydroxyvitamin D [25(OH)D], during gestation could influence neurocognitive and mental health outcomes in offspring (10-13) and the subsequent development of neuropsychiatric disorders including schizophrenia, autism, and attention deficit/hyperactivity disorder (ADHD) (9, 14-17). As such, early life vitamin D deficiency in offspring may also be a risk factor for the subsequent onset of depression. In a prospective cohort study of children, higher serum 25(OH)D levels during childhood were associated with lower depressive symptoms in adolescence (18). However, studies examining the relationship between maternal 25(OH)D levels during gestation and offspring depression are lacking. To our knowledge, only one study has assessed the effect of low gestational vitamin D levels on the risk of offspring depression (13). However, the relatively small sample size of 850 mother-child pairs and relatively crude definition for depression rendered findings inconclusive.

It is well-established that genetic factors contribute to the risk for depression in both children and young adults (19, 20). Understanding the interactions between maternal vitamin D status during pregnancy and polygenic risk scores (PRS) for depression could potentially have important implications on public health interventions. For instance, the presence or absence of gene-environment interactions (GxE) could inform whether preventive strategies should take on a targeted or universal approach (21).

In the current study, we used data from the Avon Longitudinal Study of Parents and Children study (ALSPAC) (22), a birth cohort study based in South West England, to examine the prospective association between maternal vitamin D status during pregnancy and offspring depression in childhood and adolescence, and to investigate whether the effects of maternal vitamin D status were modified by offspring genetic risk for depression.

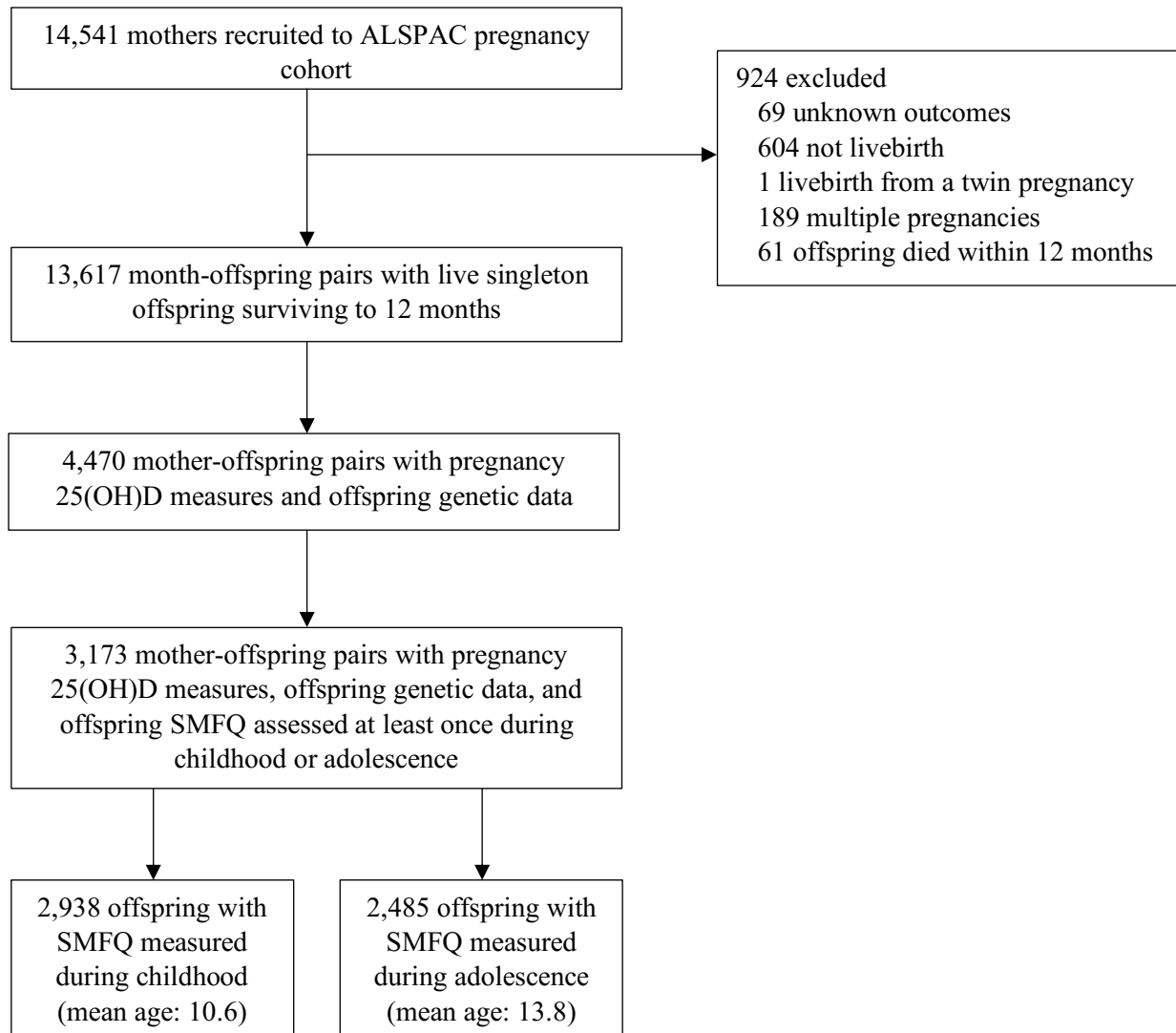
## **Methods**

### **Study Population**

ALSPAC is an ongoing population-based birth cohort from South West England. Detailed information about ALSPAC has been described previously (22, 23) and is available on the study website (<http://www.bristol.ac.uk/alspac>), which also includes a searchable data dictionary. In brief, the study enrolled 14,541 pregnant women with estimated delivery dates between April 1, 1991 and December 31, 1992, which resulted in 14,062 live births (24). Ethical approval was obtained from the ALSPAC Law and Ethics Research Committee and National Health Service (NHS) local research ethics committee, and all participants provided written informed consent at enrolment. Data on health, physical, and psychological development were collected from the parents and children at regular clinic visits and via self-reported questionnaires.

The eligible study sample included mother-offspring pairs with singleton births and complete data on maternal 25(OH)D levels during pregnancy, offspring genetic data, and offspring depressive

symptoms collected during childhood and/or adolescence. Since less than 3% of the ALSPAC cohort were of black or mixed race/ethnicity (25), only white participants were included. Mothers were excluded if their total 25(OH)D levels were three standard deviations (SD) above or below the study population mean (n=39). This yielded analytic sample sizes of 2,938 (child subsample) and 2,485 (adolescent subsample) (**Figure 3.1**).



**Figure 3.1 Study sample flow chart**



## **Offspring Depression**

Depressive symptoms were measured using the Short Moods and Feelings Questionnaire (SMFQ) by a trained interviewer during childhood (mean: 10.62; SD: 0.25) and adolescence (mean: 13.83; SD: 0.21). The SMFQ consists of 13 items that ascertain depressive symptoms in the past 2 weeks. Total SMFQ scores were obtained by summing across all items (possible score range: 0-26), with higher scores corresponding to higher depressive symptoms. The SMFQ has been validated in individuals aged between 6 and 18, and correlates highly with both the Children's Depression Inventory scores and past-year Diagnostic Interview for Children depression scores (26). Since the raw SMFQ scores were positively skewed (skewness: 1.36 for childhood subsample, 1.46 for adolescent sample) and remained skewed after log-transformations, the SMFQ was dichotomized, with depression defined as  $SMFQ \geq 11$ . This threshold has been shown to have high sensitivity and specificity for depression defined by the revised Diagnostic and Statistical Manual of Mental Disorders, Third Edition (DSM-III-R) (27), and has been used in prior ALSPAC studies of depression (28, 29).

## **Serum 25(OH)D Measurements**

Maternal serum 25(OH)D<sub>2</sub> and 25(OH)D<sub>3</sub>, were measured on non-fasting blood samples taken for routine pregnancy tests at any point during pregnancy. Blood samples were assayed using high-performance liquid chromatography tandem mass spectrometry (HPLC/MS) in accordance with Vitamin D External Quality Assessment Scheme requirements. Details about sampling, storage, and processing are described in detail elsewhere (30). For the few mothers who had multiple measurements taken during pregnancy (4.8%) (31), the last result available was used, in line with similar studies that have assessed the influence of maternal 25(OH)D concentrations during pregnancy on child health outcomes (30, 32). Total 25(OH)D was calculated by summing 25(OH)D<sub>2</sub> and 25(OH)D<sub>3</sub> levels.

As total maternal 25(OH)D levels displayed sinusoidal seasonal variation (**Supplemental Figure 3.1**), a seasonality-adjusted 25(OH)D variable was derived to represent mean annual levels of 25(OH)D. Similar trigonometric models have been previously applied to adjust for seasonal variation in 25(OH)D in ALSPAC (30, 32), and has been shown to have good validity (33). Further details of the method are described in the **Supplementary Methods**. Briefly, 25(OH)D was modeled against sine and cosine transformations of month of blood sampling in a linear regression model. The mean annual 25(OH)D concentration for each mother was then estimated using the residuals of the model. The seasonality-adjusted and unadjusted 25(OH)D levels were strongly correlated (Spearman's  $r=0.88$ ).

## **Genetic Data**

9,912 children in the ALSPAC cohort were genotyped on the Illumina HumanHap550-Quad genome-wide single nucleotide polymorphisms (SNPs) genotyping platform. Standard quality control (QC) measures were performed to exclude individuals on the basis of gender mismatch, minimal or excessive heterozygosity, individual genotyping call rates <97%, cryptic relatedness (IBD >10%), and non-European ancestry (assessed using multidimensional scaling analysis and compared to HapMap II). SNPs were excluded based on the following criteria: minor allele frequency (MAF) <1%; missing rate >5%; and significant deviation from Hardy-Weinberg Equilibrium (HWE) ( $p < 5 \times 10^{-7}$ ). 500,527 directly genotyped SNPs and 8,365 children remained after quality control (QC). Imputation was conducted using Impute V2.2.2 (34) against the 1000 genomes reference panel (Phase 1, version 3) (35), with 2186 reference haplotypes (including non-Europeans).

## **Polygenic Risk Scores (PRS)**

PRS for depression were generated using methods described by Purcell *et al.* in PLINK v.1.90 (36). The Psychiatric Genomics Consortium wave 2 (PGC-MDD2) (37) was used as the discovery sample

(meta-analytic subsample excluding 23andMe: n=173,005) and the ALSPAC cohort was used as the training sample. Additional details about the data sources and QC procedures prior to PRS construction are provided in the **Supplementary Methods**. We generated a single PRS for each offspring based on a threshold of  $p < 0.05$ , which was the threshold that maximized the variance explained in major depression (37). PRS was generated by summing the risk alleles (0, 1, or 2) for each SNP, weighted by the natural log-transformed odds ratio (OR) for its association with major depressive disorder in the PGC-MDD2. PRS scores were then standardized using the z-transformation and categorized into three risk groups (low risk: quartile 1; medium risk: quartiles 2 and 3; high risk: quartile 4) (**Supplemental Figure 3.2**).

### **Missing Data**

In the analytic samples, a substantial proportion of participants had missing data on one or more of the covariates. For any single covariate, the amount of missingness ranged between 0.3% and 20.9%, and 71.2% and 72.9% had complete covariate data for the child and adolescent subsamples, respectively. Missing covariate information was imputed using multiple imputation by chained equations (MICE) in the *mice* (version 2.30) R package (38) (**Supplementary Methods**). The distribution of covariates with missing data in the observed data was similar to that in the imputed data (**Supplemental Table 3.1**). All primary analyses were performed in the imputed data sets and effect estimates were pooled across the 20 generated imputed data sets using Rubin's rule (39).

### **Statistical Analysis**

Comparisons of covariate distributions were made across categories of maternal 25(OH)D during pregnancy using the analysis of variance (ANOVA), Kruskal-Wallis rank sum test, or  $\chi^2$  test. Similar comparisons were made between the eligible sample and the excluded sample with maternal 25(OH)D and offspring genetic data, but missing offspring depression measures.

Since continuous maternal 25(OH)D was non-linearly associated with the log odds of depression at both time points (**Supplemental Figure 3.3**), 25(OH)D was modeled as a categorical variable, based on clinical cut-offs defined by the Endocrine Society (i.e., deficient <20 ng/mL; insufficient 20-29.9 ng/mL; normal  $\geq 30$  ng/mL) (4). PRS was modeled categorically by polygenic risk groups, as described above, with the reference group set as low PRS. Logistic regression models were used to assess: (1) the main effects of PRS and maternal 25(OH)D on offspring depression during childhood and/or adolescence, separately (models 1-2) and together (model 3); (2) the interaction effect between PRS and maternal 25(OH)D on offspring depression (model 4). All models were adjusted for gestational age at 25(OH)D measurement, maternal age at delivery, maternal pre-pregnancy BMI, maternal education, maternal occupation, parity, smoking during the first trimester, maternal depression during pregnancy, and child sex (**Supplementary Methods**). Models including PRS were additionally adjusted for the top three principal components (PCs); additional details about the population structure analysis are provided in the **Supplementary Methods**. Models with multiplicative interaction terms were further adjusted for all significant covariate\*maternal 25(OH)D and covariate\*PRS interactions, based on recommendations by Keller *et al.* (40). Effect estimates were presented as odds ratios (ORs) and 95% confidence intervals (CI). Multiplicative interaction terms between maternal 25(OH)D and PRS were tested separately using the Wald test and globally using the likelihood ratio test. Statistical significance was set at two-tailed  $p < 0.05$  for all analyses. All analyses were performed in R (version 3.3.1) (41).

Based on the power calculations (**Supplementary Methods**), the study had 80% power to detect  $OR = 1.4-1.7$  for the association between vitamin D deficiency and child/adolescent depression, and sufficient power to detect GxE effects of magnitudes greater than  $OR_{gxe} = 1.7$ .

## **Additional Analysis**

We conducted several sensitivity analyses to assess the robustness of the findings. First, we fit logistic regression models that further adjusted for covariates that may be affected by or affect 25(OH)D levels during pregnancy, including vitamin D intake, calcium intake, vitamin D supplementation, and oily fish intake at 32 weeks of gestation. As the cut-offs for vitamin D deficiency remain controversial, we additionally conducted analyses using lower cut-offs set by the Institute of Medicine (i.e., deficient <10 ng/mL; insufficient 10-19.9 ng/mL; normal  $\geq$ 20 ng/mL) (44). Also, given that vitamin D may have different effects on fetal brain development at different stages of gestation, we repeated the analyses stratified by the trimester of 25(OH)D measurement. Finally, we performed a complete case analysis on the subsample of participants with complete observed data on all variables (child subsample: n=2,091, adolescent subsample: n=1,812).

We also examined the prospective association between maternal vitamin D status and the risk of offspring depressive symptoms using negative binomial regression models. Negative binomial regression allows for the modeling of the highly skewed and over-dispersed counts of depressive symptoms and could potentially provide greater statistical power to detect associations (45). Estimates were presented as rate ratios (RR) and 95% CI, representing the association between maternal vitamin D status, PRS, and the risk for offspring depressive symptoms.

## **Results**

Among the eligible mother-offspring pairs (n=3,173), the mean maternal 25(OH)D was 27.38 ng/mL (SD=11.97), and 63.5% of mothers were vitamin D insufficient or deficient during pregnancy based on clinical cut-offs defined by the Endocrine Society. There were no significant differences between the number of 25(OH)D measurements taken across the seasons (Winter: 23.9%; Spring: 28.7%; Summer: 24.6%; Fall: 22.8%). The median gestational week at 25(OH)D measurement was 29.4 weeks

(IQR: 12.7, 33.1), with most available measurements collected in the third trimester (58.5%). **Table 3.1** displays the distribution of mother and offspring characteristics by maternal vitamin D status during pregnancy.

**Table 3.1 Maternal and offspring characteristics by maternal vitamin D status during pregnancy, among eligible mother-offspring pairs with complete data on maternal 25(OH)D, offspring genetic data, and at least one SMFQ measure**

	All (n=3173)	Maternal 25(OH)D <sup>1</sup>			p <sup>2</sup>
		<20 ng/mL (n=988)	20-29.9 ng/mL (n=1027)	≥30 ng/mL (n=1158)	
<b>Maternal age at delivery, n (%)</b>					
15-19	46 (1.4)	18 (39.1)	18 (39.1)	10 (21.7)	<b>0.02</b>
20-35	2857 (90.0)	906 (31.7)	915 (32.0)	1036 (36.3)	
>35	270 (8.5)	64 (23.7)	94 (34.8)	112 (41.5)	
<b>Maternal BMI (kg/m<sup>2</sup>), median [IQR]</b>	22.18 [20.53, 24.38]	22.18 [20.53, 24.52]	22.35 [20.53, 24.51]	22.02 [20.47, 23.88]	
<b>Maternal education, n (%)</b>					
Lower than O-levels	577 (19.3)	203 (52.3)	180 (31.2)	194 (33.8)	0.09
O-levels	1080 (36.1)	331 (30.6)	359 (33.2)	390 (36.1)	
Higher than O-levels	1332 (44.6)	390 (29.3)	428 (32.1)	514 (38.6)	
<b>Maternal occupation, n (%)</b>					
Manual	427 (16.0)	154 (36.1)	131 (30.7)	142 (33.2)	<b>0.03</b>
Non-manual	2237 (84.0)	659 (29.5)	747 (33.4)	831 (37.1)	
<b>Parity, n (%)</b>					
0	1446 (46.8)	482 (33.3)	477 (33.0)	487 (33.7)	<b>0.007</b>
≥1	1647 (53.2)	483 (29.3)	524 (31.8)	640 (38.9)	
<b>Gestational week of 25(OH)D measurement, median [IQR]</b>	29.43 [12.71, 33.14]	28.29 [11.29, 32.86]	28.71 [11.43, 32.86]	31.14 [18.75, 33.29]	<b>&lt;0.001</b>
<b>Season of 25(OH)D measurement, n (%)</b>					
Winter	758 (23.9)	343 (45.3)	256 (33.8)	159 (20.9)	<b>&lt;0.001</b>
Spring	912 (28.7)	412 (45.2)	297 (32.6)	203 (22.3)	
Summer	781 (24.6)	100 (12.8)	218 (27.9)	463 (59.3)	
Fall	722 (22.8)	133 (18.4)	256 (35.5)	333 (46.1)	
<b>Tobacco use during 1st trimester, n (%)</b>					
No	2569 (82.2)	752 (29.3)	831 (32.4)	984 (38.3)	<b>&lt;0.001</b>
Yes	557 (17.8)	221 (39.7)	181 (32.5)	155 (27.8)	

Table 3.1 (continued)

	All (n=3173)	Maternal 25(OH)D <sup>1</sup>			p <sup>2</sup>
		<20 ng/mL (n=988)	20-29.9 ng/mL (n=1027)	≥30 ng/mL (n=1158)	
<b>Oily fish intake at 32 weeks, n (%)</b>					
≥1 times/week	813 (26.8)	214 (26.3)	261 (32.1)	338 (41.6)	<b>&lt;0.001</b>
<1 times/week	1045 (34.5)	318 (30.4)	341 (32.6)	386 (37.0)	
Never/rarely	1173 (38.7)	408 (34.8)	379 (32.3)	386 (32.9)	
<b>Vitamin D supplementation at 32 weeks, n (%)</b>					
No	3035 (95.7)	961 (31.7)	985 (32.4)	1089 (35.9)	<b>0.001</b>
Yes	138 (4.3)	27 (19.6)	42 (30.4)	69 (50.0)	
<b>Vitamin D intake at 32 weeks (ug), median [IQR]</b>	3.43 [2.44, 5.40]	3.21 [2.28, 4.84]	3.39 [2.46, 5.46]	3.69 [2.59, 5.72]	<b>&lt;0.001</b>
<b>Calcium intake at 32 weeks (mg), median [IQR]</b>	938.54 [762.25, 1131.15]	935.50 [762.92, 1129.98]	940.01 [763.07, 1129.33]	939.08 [759.17, 1134.02]	0.87
<b>Maternal depression during pregnancy<sup>3</sup>, n (%)</b>					
No	2066 (75.6)	637 (30.8)	642 (31.1)	787 (38.1)	0.10
Yes	667 (24.4)	211 (31.6)	231 (34.6)	225 (33.8)	
<b>Breastfeeding, n (%)</b>					
No	1067 (34.9)	366 (34.3)	356 (33.4)	345 (32.3)	<b>0.001</b>
Yes	1994 (65.1)	585 (29.4)	633 (31.7)	776 (38.9)	
<b>Child sex, n (%)</b>					
Male	1592 (50.2)	483 (30.3)	520 (32.7)	589 (37.0)	0.62
Female	1581 (49.8)	505 (31.9)	507 (32.1)	569 (36.0)	

<sup>1</sup> Unadjusted 25(OH)D levels

<sup>2</sup> P-value calculated using the Chi-squared test for categorical variables and the ANOVA or Kruskal-Wallis rank sum test for normally and non-normally distributed continuous variables, respectively

<sup>3</sup> Maternal depression at 18 and/or 32 weeks of gestation

Maternal age at delivery, oily fish intake, and vitamin D intake during pregnancy increased with maternal 25(OH)D; the proportion of mothers with non-manual occupations, more than one child, were non-smokers during the first trimester, were on vitamin D supplementation, and breastfed increased from lower to higher maternal 25(OH)D levels. 6.6% and 13.0% of the offspring were classified as depression cases during childhood and adolescence, respectively.

Compared to the excluded participants, the eligible sample comprised mothers with higher 25(OH)D levels during pregnancy and offspring with lower PRS (**Supplemental Table 3.2**). Further, mothers in the eligible sample also tended to be older when they gave birth, had higher educational attainment, were in non-manual occupations, had no previous pregnancies, were non-smokers during the first trimester, and showed no evidence of depression during pregnancy. However, the absolute difference between maternal 25(OH)D (i.e., 1.3 ng/mL higher in the eligible sample) and offspring PRS (i.e., 0.07 standardized units lower in the eligible sample) in the eligible and excluded samples was small, hence the results are unlikely to be affected by selection bias.

### ***Maternal Vitamin D Status and PRS on Offspring Depression***

Associations between maternal 25(OH)D during pregnancy, PRS, and offspring depression are presented in **Table 3.2**. There were no associations between maternal 25(OH)D and offspring depression during childhood, although there was suggestive evidence that offspring exposed to deficient vitamin D levels during gestation had higher odds of depression during adolescence, compared to offspring exposed to normal vitamin D levels (OR=1.32; 95% CI: 0.98, 1.79; p=0.07).



**Table 3.2 Association between maternal vitamin D status during pregnancy, offspring polygenic risk scores (PRS), and offspring depression during childhood or adolescence**

	Childhood depression (n=2938)					Adolescent depression (n=2485)				
	n	OR	95% CI	p	P-trend	n	OR	95% CI	p	P-trend
<b>Model 1: Maternal 25(OH)D only</b>										
Normal ( $\geq 30$ ng/mL)	985	<i>ref</i>	<i>ref</i>	<i>ref</i>		841	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	1090	1.02	0.71, 1.48	0.90	0.72	907	1.12	0.84, 1.51	0.44	0.07
Deficient (<20 ng/mL)	863	1.07	0.73, 1.58	0.72		737	1.32	0.98, 1.79	0.07	
<b>Model 2: PRS only</b>										
PRS-Low	735	<i>ref</i>	<i>ref</i>	<i>ref</i>		622	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1471	1.37	0.90, 2.09	0.14	<b>0.003</b>	1242	0.99	0.73, 1.34	0.93	0.06
PRS-High	732	<b>1.94</b>	<b>1.24, 3.03</b>	<b>0.004</b>		621	1.34	0.96, 1.87	0.09	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>										
Normal ( $\geq 30$ ng/mL)	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/mL)	-	1.03	0.71, 1.49	0.87	0.70	-	1.13	0.84, 1.51	0.43	0.07
Deficient (<20 ng/mL)	-	1.08	0.73, 1.59	0.70		-	1.33	0.98, 1.81	0.06	
PRS-Low	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	-	1.36	0.90, 2.09	0.15	<b>0.003</b>	-	0.98	0.73, 1.33	0.92	0.07
PRS-High	-	<b>1.94</b>	<b>1.24, 3.03</b>	<b>0.004</b>		-	1.34	0.96, 1.87	0.09	

Table 3.2 (continued)

	Childhood depression (n=2938)					Adolescent depression (n=2485)				
	n	OR	95% CI	p	P-inter.	n	OR	95% CI	p	P-inter.
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)</b>										
Normal ( $\geq 30$ ng/mL)	-	<i>ref</i>	<i>ref</i>	<i>ref</i>	-	-	<i>ref</i>	<i>ref</i>	<i>ref</i>	-
Insufficient (20-29.9 ng/mL)	-	0.65	0.29, 1.45	0.29	-	-	1.15	0.62, 2.11	0.66	-
Deficient (<20 ng/mL)	-	0.38	0.13, 1.06	0.06	-	-	1.45	0.78, 2.69	0.24	-
PRS-Low	-	<i>ref</i>	<i>ref</i>	<i>ref</i>	-	-	<i>ref</i>	<i>ref</i>	<i>ref</i>	-
PRS-Intermediate	-	0.74	0.63, 1.46	0.38	-	-	1.21	0.71, 2.06	0.49	-
PRS-High	-	1.26	0.37, 2.53	0.52	-	-	1.06	0.58, 1.96	0.84	-
Insufficient*PRS-Intermediate	-	1.97	0.74, 5.26	0.18	-	-	0.80	0.38, 1.66	0.54	-
Deficient*PRS-Intermediate	-	4.09	1.26, 13.3	0.02	0.19	-	0.68	0.32, 1.45	0.32	0.36
Insufficient*PRS-High	-	1.65	0.59, 4.58	0.34	-	-	1.38	0.61, 3.15	0.44	-
Deficient*PRS-High	-	2.92	0.86, 10.0	0.09	-	-	1.39	0.60, 3.22	0.44	-

Abbreviations: 25(OH)D: 25-hydroxyvitamin D; CI: confidence intervals; OR: odds ratio; p-inter.: p for interaction

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal occupation, parity, smoking during 1st trimester, maternal depression during pregnancy, child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + significant PRS\*covariate, maternal 25(OH)D\*covariate interactions

Offspring PRS was positively associated with the odds of depression during childhood ( $p$  for trend=0.003), with high PRS conferring an approximately 2-fold higher odds of childhood depression compared to low PRS (OR=1.94; 95% CI: 1.24, 3.03;  $p$ =0.0004); higher PRS was associated with non-significantly higher odds of depression during adolescence ( $p$  for trend=0.06) (model 2). When modeled together (model 3), the effect estimates for both maternal vitamin D status and PRS were almost identical to those yielded from their separate models. In model 4, there was no evidence for significant interactions between maternal vitamin D status and PRS on childhood ( $p$ =0.19) or adolescent ( $p$ =0.36) depression.

### *Additional Analyses*

Further adjustments for nutritional intake at 32 weeks of gestation (model 5) (**Supplemental Table 3.3**) resulted in estimates similar to those in the main analyses. We also investigated the robustness of the results to alternative, lower cut-offs for vitamin D insufficiency and deficiency. These lower cut-offs resulted in large imbalances in the numbers across categories, resulting in a small number of mothers classified as vitamin D deficient during pregnancy ( $n$ =55-71). Nonetheless, the direction and magnitude of effects were generally similar to those using higher cut-offs for deficiency (**Supplemental Table 3.4**). The results did not change substantially when stratified by trimester of 25(OH)D measurement, although some of the effect directions were reversed in the small subsamples with measurements in the second trimester ( $n$ =381-438) (**Supplemental Table 3.5**). Complete case analysis also showed similar patterns of trends to those from the primary analyses on the imputed data (**Supplemental Table 3.6**), but the associations between PRS and childhood/adolescent depression were attenuated to non-significance, likely due to large reductions in sample sizes.

Results from the negative binomial regression models showed similar associational patterns as those produced from the logistic regression models (**Supplemental Table 3.7**). There were no significant main effects of maternal 25(OH)D nor interactions with PRS on the risk of offspring depressive

symptoms during childhood or adolescence. The only notable difference was the presence of a statistically significant positive association between PRS and the risk of depressive symptoms during adolescence (p-trend=0.03).

## Discussion

In this large, prospective birth cohort, we found little evidence for an association between maternal pregnancy vitamin D status (i.e., blood 25(OH)D level) and offspring depression in childhood or adolescence. In this population, PRS was positively associated with risk for offspring depression at both time points but did not interact with maternal 25(OH)D. These findings were robust to adjustments for a range of potential confounders and different cut-offs for vitamin D insufficiency/deficiency.

Although several studies have reported the influence of maternal and/or cord blood 25(OH)D on the offspring's risk for neuropsychiatric or neurodevelopmental disorders/traits, including schizophrenia (16), ADHD (17), and autism-related traits (46), the same effects do not appear to be present for depression. Results from the current study are consistent with the only existing study, to our knowledge, that assessed the association between maternal 25(OH)D during pregnancy and offspring depression. In the Strom *et al.* study (13), maternal vitamin D deficiency during pregnancy was not associated with offspring risk for depression over 22 years of follow-up. Studies that assessed features similar to depression in childhood, such as emotional problems and/or internalizing symptoms, also did not report significant associations with maternal 25(OH)D (11, 47, 48). Taken together, these findings do not support influences of maternal vitamin D status during pregnancy on offspring depression or related symptoms in early life.

It has been postulated that there may be sensitive developmental periods during which the effects of fetal 25(OH)D on neurodevelopment are particularly salient (9). The precise timing of the

neurodevelopmental effects of vitamin D is unclear, but may have importance at any stage of pregnancy: early gestation, when structures important for behavioral regulation and dopaminergic neurons develop (47, 49); mid-gestation, which is characterized by neural circuit formation and myelination (50, 51); late gestation, when prolific brain development and growth occurs (52-54). When we stratified our analysis by trimester of 25(OH)D measurement, the overall results did not differ from those yielded from the primary analysis in the full cohort. However, the stratified analyses had limited statistical power given the small sample sizes within some of the subgroups. Hence, the possibility for sensitive periods for vitamin D effects remains, and longitudinal studies sampling 25(OH)D in all three trimesters are needed to establish stronger conclusions. Alternatively, the lack of gestational 25(OH)D effects on child depression may be due to critical window(s) of vulnerability to vitamin D exposure occurring postnatally. In the Tolppanen *et al.* study (18), similarly based on children in the ALSPAC cohort, childhood 25(OH)D<sub>3</sub> levels were found to be associated with adolescent depressive symptoms. It is plausible that the protective effects of vitamin D on depression and mental health are exerted through neuroprotective actions, such as moderating inflammatory processes or modulating neurotrophic factors, after birth. Further studies are necessary to characterize the precise mechanisms of vitamin D on the developing brain.

Our study has notable strengths. First, this study was conducted in a large, population-based sample that was more than three times larger than the previous study examining the same associations. Second, the longitudinal study design allowed us to prospectively investigate the effects of maternal 25(OH)D on offspring depression measured during both childhood and adolescence, while adjusting for a wide range of potential confounding factors.

The study also has some limitations. First, only one measurement of 25(OH)D, taken at any time during pregnancy, was available for all mothers in the study sample, which may not have been representative of average 25(OH)D levels for the duration of the pregnancy. However, studies have shown that single measurements may be reasonable proxies for vitamin D status throughout pregnancy, given the strong correlation in 25(OH)D concentrations over time (55, 56). Second, the SMFQ only

captures depressive symptoms in the past 2 weeks, which may not represent depression over longer periods. This may have led to outcome misclassification, particularly for adolescent depression, which has been shown to be highly episodic in nature (57); such misclassification may have biased the results towards the null. Third, it is possible that an association between low maternal vitamin D levels and offspring depression is only detectable at very low levels of maternal 25(OH)D, and that the distribution of 25(OH)D levels in this population was not sufficiently wide to capture this. Compared to prior studies that detected significant associations (16, 46), mothers in the current study were less diverse ethnically/racially and/or socioeconomically, and on average, had higher 25(OH)D levels (27.38 ng/mL compared to 15.3-23.6 ng/mL in other studies) with only a very small proportion having 25(OH)D levels <10 ng/mL (2%). Fourth, as with most longitudinal studies, there was considerable attrition over time, which could introduce bias when both maternal 25(OH)D and offspring depression were associated with loss to follow-up. Nevertheless, since the difference between the average maternal 25(OH)D levels in the excluded and included samples was small in magnitude, we do not expect attrition in this case to substantially influence the results. Fifth, the study was based in a white population, thus the results may not be generalizable to other racial/ethnic groups or populations with different prevalence of vitamin D deficiency and/or depression. Finally, we may have lacked power to detect weaker effects of maternal 25(OH)D on offspring depression and had even lower power to detect GxE. Larger sample sizes are needed to definitively rule out the presence of smaller associations and gene-environment interactions.

In sum, our findings do not support an association between maternal vitamin D status during pregnancy and offspring depression during childhood and adolescence. This suggests that interventions aimed at increasing 25(OH)D levels during pregnancy are unlikely to reduce the risk of offspring depression during childhood or adolescence, although such efforts may still be beneficial for other child health outcomes (58). Larger prospective studies in more racially/ethnically diverse populations with lower average values and/or broader ranges of levels of 25(OH)D during pregnancy are needed to confirm the current findings.

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## Appendix

### Supplementary Methods

#### *Derivation of seasonality-adjusted maternal 25(OH)D measurements*

The cosinor model is a linear regression model that models 25(OH)D against sine and cosine transformations of time (i.e., months) (1). The transformed model coefficients can be used to derive the amplitude (distance between mean to peak or trough location of the curve) and phase shift (location of peak and trough on time axis) of the sine curve. The coefficient for the model intercept is used to obtain the mean annual 25(OH)D concentration in the study population, and the residuals are used to derive the mean annual 25(OH)D levels for each individual. Full details of this method has been described elsewhere (1).

#### *Polygenic risk score construction*

Polygenic risk scores (PRS) for major depressive disorder (MDD) were constructed using summary-level data from the GWAS meta-analysis of MDD from the Psychiatric Genomics Consortium wave 2 (PGC-MDD2; 130,664 cases and 330,470 controls) (2). The PGC-MDD2 consisted of 35 cohorts, including the 29 PGC-MDD anchor cohorts (16,823 cases, 25,632 controls) (2) and 6 additional cohorts: GERA (7,162 cases, 38,307 controls) (3), deCODE (1,980 cases, 9,536 controls) (2), GenScotland (997 cases, 6,358 controls) (4, 5), iPsych (18,629 cases, 17,841 controls) (2), UK Biobank (14,260 cases, 15,480 controls) (6, 7), and 23andMe (75,607 cases, 231,747 controls) (8). Due to general access constraints, full summary statistics were only available for the meta-analytic subsample (n=173,005) excluding the 23andMe cohort. Prior to score construction, SNPs with imputation quality metric score (INFO)<0.8, MAF<1%, call rate<95%, and HWE  $p < 1 \times 10^{-6}$  were filtered out. SNPs were pruned for linkage disequilibrium (LD) using p-value informed clump-based pruning in PLINK version 1.90 (<https://www.cog-genomics.org/plink2>) using the following parameters: --clump-p1 1 --clump-p2 1 --

clump-r2 0.2 --clump-kb 500). Within a specified window, LD clumping preferentially keeps the most significant SNPs and removes all other SNPs in high LD.

### *Assessment of covariates*

Data on maternal pre-pregnancy body mass index (BMI, kg/m<sup>2</sup>), parity, maternal socioeconomic position indicators, tobacco use, nutritional intake, and vitamin D supplementation were provided by mothers through self-administered questionnaires at 18 and/or 32 weeks of pregnancy. Maternal depression during pregnancy was derived using the Edinburgh Postnatal Depression Scale (9), with depressed mood defined as scores  $\geq 13$ . Maternal age at delivery, month/season and gestational age at 25(OH)D measurement (weeks) were obtained from obstetric records. Child sex was obtained from birth records, while information on breastfeeding was collected postnatally from mother-completed questionnaires.

Maternal pre-pregnancy BMI and gestational age at 25(OH)D measurement were included in the models as continuous variables. All other variables were included as categorical variables: maternal age at child birth (age 15-19 years, 20-35 years, or >35 years), education (less than O-levels or equivalent, O-levels, or higher than O-levels), occupation (manual or non-manual), parity (0 or  $\geq 1$  child), first trimester tobacco use (yes or no), oily fish intake at 32 weeks of gestation ( $\geq$ once/week, <once/week, or never/rarely), vitamin D supplementation at 32 weeks of gestation (yes or no), vitamin D intake at 32 weeks of gestation (ug), calcium intake at 32 weeks of gestation (mg), maternal depression during pregnancy (yes or no), breastfeeding (yes or no), child sex (male or female). The reference group for all categorical variables were chosen on the basis of sample size and consistency with the literature.

### *Population structure analysis*

Principal components (PCs) reflect a population's genetic structure and can be used to adjust for population stratification (10). Using genome-wide SNPs in the directly genotyped data, the top 20 PCs

were estimated in PLINK using methods described by Price and colleagues (10). In line with the PGC ricopili pipeline (<https://sites.google.com/a/broadinstitute.org/ricopili/pca>) the following quality control (QC) procedures were performed: MAF>5%; HWE>1.0E-03; missing rate<2%; no strand ambiguous SNPs (AT/GC); no MHC (6:25-35Mb); no Chromosome 8 inversions (8:7-13Mb). 462,838 SNPs remained after QC and were pruned if their  $r^2>0.2$ , within a 200 SNP window. After pruning, 90,414 SNPs were available for principal component analysis. 20 PCs were generated from the analysis and the top 3 PCs were included in the analytic models based on examination of the eigenvalues and a scree plot.

### ***Multivariable multiple imputation***

All exposure, covariate, and outcome variables that were included in the final analysis were included in the imputation model for imputing covariate data, since failure to do so may result in a biased analytic model (11). Variables that were strongly correlated with the covariates, such as home crowding and home ownership, were also included in the imputation model because this has been shown to reduce bias and improve precision (11). For continuous variables, predictive mean matching was used for imputation, while logistic and polynomial regression was used to impute dichotomous and categorical variables, respectively. Twenty imputed data sets were generated using this method.

Several diagnostic tests were performed to assess proper imputation. First, convergence plots were inspected to determine whether convergence was achieved. Second, density plots were examined to compare the distribution of the imputed data to observed data. **Table S1** presents the percentage of missing data for each covariate, as well as a comparison of the distribution of the variables in the observed data versus imputed data.

### ***Power analysis***

Power calculations were performed for both the main effects of maternal 25(OH)D and GxE on offspring depression. Power analysis for the main effects of maternal 25(OH)D on offspring depression, was implemented in an online power calculator for logistic regression



(<http://www.dartmouth.edu/~eugened/power-samplesize.php>), developed using algorithms derived by Demidenko (12, 13). Power analysis for GxE was performed using the *powerGWASinteraction* package in R. All power calculations assumed  $\alpha=0.05$  and a two-sided test; other parameters were derived from the literature or directly estimated in the study cohort.

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**Supplemental Table 3.1 Comparison of the distribution of main covariates in the observed data set and the imputed data set, for mother-offspring pairs with (A) SMFQ in childhood; (B) SMFQ in adolescence**

**(A)**

Covariates with missing data	N missing (%)	Mean (SD) or n (%) in sample	
		Observed data	Imputed data
Maternal pre-pregnancy BMI (kg/m <sup>2</sup> )	264 (5.9)	22.87 (3.63)	22.84 (3.58)
Maternal education			
<O-level		534 (19.0)	570 (19.2)
O-level	166 (3.7)	1016 (36.2)	1077 (36.2)
>O-level		1259 (44.8)	1328 (44.6)
Parity			
0	71 (1.6)	1347 (46.4)	1382 (46.5)
1+		1557 (53.6)	1593 (53.5)
Maternal occupation			
Manual	464 (10.4)	398 (15.9)	504 (16.9)
Non-manual		2113 (84.1)	2471 (83.1)
Tobacco use during the first trimester			
No	40 (0.9)	2434 (82.9)	2466 (82.9)
Yes		501 (17.1)	509 (17.1)
Maternal depression during pregnancy			
No	405 (9.1)	1960 (76.3)	2266 (76.2)
Yes		610 (23.7)	709 (23.8)

**(B)**

Covariates with missing data	N missing (%)	Mean (SD) or n (%) in sample	
		Observed data	Imputed data
Maternal pre-pregnancy BMI (kg/m <sup>2</sup> )	191 (7.6)	22.89 (3.74)	22.89 (3.74)
Maternal education			
<O-level		417 (17.4)	438 (17.4)
O-level	116 (4.6)	848 (35.4)	891 (35.5)
>O-level		1130 (47.2)	1182 (47.1)
Parity			
0	57 (2.3)	1200 (48.9)	1226 (48.8)
1+		1254 (51.1)	1285 (51.2)
Maternal occupation			
Manual	353 (14.1)	322 (14.9)	409 (16.3)
Non-manual		1846 (85.1)	2102 (83.7)
Tobacco use during the first trimester			
No	32 (1.3)	2072 (83.6)	2101 (83.7)
Yes		1254 (16.4)	410 (16.3)
Maternal depression during pregnancy			
No	320 (12.7)	1677 (76.5)	1910 (76.1)
Yes		514 (23.5)	601 (23.9)

**Supplemental Table 3.2 Comparison of covariate distribution among eligible sample and participants excluded due to missing offspring SMFQ**

	<b>Excluded sample (n=1297)</b>	<b>Eligible sample (n=3173)</b>	<b>p<sup>1</sup></b>
<b>Maternal 25(OH)D (ng/mL), median [IQR]<sup>2</sup></b>	24.14 [17.68, 32.82]	25.45 [18.89, 33.42]	<b>0.003</b>
<b>Standardized MDD PRS, mean (SD)</b>	0.03 (0.99)	-0.04 (1.01)	<b>0.04</b>
<b>Maternal age at birth (years), n (%)</b>			
15-19	77 (5.9)	46 (1.4)	<b>&lt;0.001</b>
20-35	1150 (88.7)	2857 (90.0)	
>35	70 (5.4)	270 (8.5)	
<b>Maternal BMI (kg/m<sup>2</sup>), median [IQR]</b>	22.02 [20.47, 24.51]	22.18 [20.53, 24.38]	0.79
<b>Maternal education, n (%)</b>			
Lower than O-levels	301 (29.7)	577 (19.3)	<b>&lt;0.001</b>
O-levels	390 (38.5)	1080 (36.1)	
Higher than O-levels	323 (31.9)	1332 (44.6)	
<b>Maternal occupation, n (%)</b>			
Manual	195 (22.4)	427 (16.0)	<b>&lt;0.001</b>
Non-manual	676 (77.6)	2237 (84.0)	
<b>Parity, n (%)</b>			
0	500 (41.9)	1446 (46.8)	<b>0.005</b>
≥1	693 (58.1)	1647 (53.2)	
<b>Gestational week of 25(OH)D measurement, median [IQR]</b>	29.71 [13.00, 33.29]	29.43 [12.71, 33.14]	0.26
<b>Season of 25(OH)D measurement, n (%)</b>			
Winter	303 (23.4)	758 (23.9)	0.78
Spring	381 (29.4)	912 (28.7)	
Summer	305 (23.5)	781 (24.6)	
Fall	308 (23.7)	722 (22.8)	
<b>Tobacco use during 1st trimester, n (%)</b>			
No	832 (68.5)	2569 (82.2)	<b>&lt;0.001</b>
Yes	383 (31.5)	557 (17.8)	
<b>Maternal depression during pregnancy, n (%)</b>			
No	656 (70.0)	2066 (75.6)	<b>0.001</b>
Yes	281 (30.0)	667 (24.4)	

<sup>1</sup> P-value calculated using Chi-squared test for categorical variables and the student's t-test or Kruskal-Wallis test for continuous variables

<sup>2</sup> Mean annual 25(OH)D levels

**Supplemental Table 3.3 Association between maternal vitamin D status during pregnancy, PRS, and child/adolescent depression, additionally adjusting for other covariates (Model 5)**

	Childhood depression (n=2938)				Adolescent depression (n=2485)			
	OR	95% CI	p	p-interaction	OR	95% CI	p	p-interaction
<b>Model 5: PRS and Maternal 25(OH)D</b>								
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	0.63	0.28, 1.42	0.27	-	1.14	0.62, 2.10	0.66	-
Deficient (<20 ng/mL)	0.35	0.12, 0.99	0.05		1.46	0.79, 2.71	0.23	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	0.74	0.37, 1.48	0.40	-	1.22	0.72, 2.08	0.46	-
PRS-High	1.28	0.64, 2.59	0.48		1.09	0.59, 2.00	0.79	
Insufficient*PRS-Intermediate	1.94	0.73, 5.19	0.19		0.80	0.38, 1.67	0.55	
Insufficient*PRS-High	4.06	1.25, 13.19	0.02	0.20	0.68	0.32, 1.44	0.31	0.37
Deficient*PRS-Intermediate	1.61	0.86, 4.48	0.36		1.38	0.60, 3.15	0.44	
Deficient*PRS-High	2.95	0.58, 10.12	0.08		1.37	0.59, 3.16	0.46	

Abbreviations: CI: confidence intervals; OR: odds ratio; PRS: polygenic risk score

Model 5: Model 4 covariates + oily fish intake + vitamin D supplementation + vitamin D intake + Calcium intake

**Supplemental Table 3.4 Association between maternal vitamin D status during pregnancy, PRS, and child/adolescent depression, using thresholds of <10, 10-19.99, and ≥20 ng/mL**

	Childhood depression (n=2938)					Adolescent depression (n=2485)				
	n	OR	95% CI	p	p-trend	n	OR	95% CI	p	p-trend
<b>Model 1: Maternal 25(OH)D</b>										
Normal (≥20 ng/mL)	2075	<i>ref</i>	<i>ref</i>	<i>ref</i>		1748	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (10-19.9 ng/mL)	792	1.08	0.77, 1.51	0.66	0.83	682	1.26	0.97, 1.64	0.09	0.14
Deficient (<10 ng/mL)	71	0.87	0.31, 2.46	0.80		55	1.05	0.46, 2.39	0.92	
<b>Model 2: PRS</b>										
PRS-Low	735	<i>ref</i>	<i>ref</i>	<i>ref</i>		622	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1471	1.37	0.90, 2.09	0.14	<b>0.003</b>	1242	0.99	0.73, 1.34	0.93	0.06
PRS-High	732	<b>1.94</b>	<b>1.24, 3.03</b>	<b>0.004</b>		621	1.34	0.96, 1.87	0.09	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>										
Normal (≥20 ng/mL)	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (10-19.9 ng/mL)	-	1.08	0.77, 1.51	0.67	0.82	-	1.27	0.97, 1.65	0.08	0.14
Deficient (<10 ng/mL)	-	0.89	0.32, 2.50	0.82		-	1.04	0.45, 2.38	0.93	
PRS-Low	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	-	1.37	0.90, 2.09	0.15	<b>0.003</b>	-	0.99	0.73, 1.34	0.93	0.07
PRS-High	-	<b>1.93</b>	<b>1.23, 3.03</b>	<b>0.004</b>		-	1.34	0.96, 1.87	0.09	

Supplemental Table 3.4 (continued)

	Childhood depression (n=2938)					Adolescent depression (n=2485)				
	n	OR	95% CI	p	p-trend	n	OR	95% CI	p	p-trend
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)</b>										
Normal ( $\geq 20$ ng/mL)	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (10-19.9 ng/mL)	-	0.40	0.14, 1.17	0.10	-	-	1.41	0.82, 2.41	0.21	-
Deficient (<10 ng/mL)	-	1.16	0.15, 9.26	0.89		-	0.70	0.09, 5.60	0.74	
PRS-Low	-	<i>ref</i>	<i>ref</i>	<i>ref</i>		-	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	-	1.04	0.64, 1.69	0.86	-	-	1.07	0.74, 1.55	0.71	-
PRS-High	-	1.60	0.96, 2.67	0.07		-	1.27	0.84, 1.92	0.25	
Insufficient*PRS-Intermediate	-	3.46	1.08, 11.07	0.04		-	0.72	0.37, 1.39	0.32	
Deficient*PRS-Intermediate	-	0.71	0.06, 8.97	0.79	0.25	-	2.03	0.20, 20.32	0.55	0.39
Insufficient*PRS-High	-	2.68	0.79, 9.08	0.11		-	1.18	0.58, 2.42	0.64	
Deficient*PRS-High	-	0.72	0.04, 13.52	0.83		-	0.75	0.04, 14.63	0.85	

Abbreviations: CI: confidence intervals; OR: odds ratio; p-inter.: p for interaction; PRS: polygenic risk score

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal occupation, parity, smoking during 1st trimester, maternal depression during pregnancy, and child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + significant PRS\*covariate, maternal 25(OH)D\*covariate interactions



**Supplemental Table 3.5 Association between maternal vitamin D status during pregnancy, PRS, and child/adolescent depression, stratified by trimester of 25(OH)D measurement**

	Trimester 1									
	Childhood depression (n=781)					Adolescent depression (n=652)				
	OR	LCI	UCI	p	P-trend	OR	LCI	UCI	p	P-trend
<b>Model 1: Maternal 25(OH)D</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	1.56	0.69	3.52	0.28	0.34	1.12	0.61	2.05	0.71	0.09
Deficient (<20 ng/ml)	1.55	0.67	3.56	0.30		1.62	0.90	2.92	0.11	
<b>Model 2: PRS</b>										
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.40	0.64	3.08	0.40	0.72	0.97	0.53	1.77	0.93	0.27
PRS-High	1.19	0.46	3.07	0.72		1.39	0.71	2.70	0.34	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	1.58	0.70	3.57	0.27	0.34	1.14	0.62	2.08	0.68	0.09
Deficient (<20 ng/ml)	1.55	0.68	3.58	0.30		1.64	0.91	2.98	0.10	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.42	0.64	3.14	0.38	0.70	0.97	0.53	1.77	0.93	0.29
PRS-High	1.22	0.47	3.14	0.69		1.40	0.71	2.73	0.33	
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.51	0.10	2.71	0.43	-	0.65	0.15	2.84	0.57	-
Deficient (<20 ng/ml)	0.54	0.10	3.03	0.49		2.38	0.65	8.69	0.19	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	0.43	0.09	2.08	0.29	-	1.01	0.29	3.49	0.99	-
PRS-High	0.54	0.08	3.51	0.52		1.30	0.33	5.16	0.71	
Insufficient*PRS-Intermediate	5.00	0.65	38.35	0.12		1.69	0.31	9.22	0.54	
Deficient*PRS-Intermediate	4.30	0.53	34.61	0.17	0.62	0.65	0.14	3.01	0.58	0.46
Insufficient*PRS-High	2.90	0.25	33.14	0.39		2.56	0.40	16.29	0.32	
Deficient*PRS-High	3.00	0.25	35.22	0.38		0.56	0.10	3.17	0.51	

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal social class, parity, smoking during 1st trimester, maternal psychopathology during pregnancy, child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + any significant PRS\*covariate and maternal 25(OH)D\*covariate interactions

**Supplemental Table 3.5 (continued)**

	Trimester 2									
	Childhood depression (n=438)					Adolescent depression (n=381)				
	OR	LCI	UCI	p	P-trend	OR	LCI	UCI	p	P-trend
<b>Model 1: Maternal 25(OH)D</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.78	0.30	2.03	0.61	0.05	1.30	0.63	2.66	0.48	0.91
Deficient (<20 ng/ml)	0.21	0.04	1.03	0.05		1.03	0.48	2.19	0.95	
<b>Model 2: PRS</b>										
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.41	0.36	5.51	0.62	0.14	0.64	0.30	1.36	0.24	0.71
PRS-High	2.60	0.65	10.43	0.18		0.79	0.34	1.84	0.59	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.74	0.28	1.98	0.55	0.04	1.33	0.64	2.77	0.44	0.80
Deficient (<20 ng/ml)	0.20	0.04	0.95	0.04		1.10	0.51	2.37	0.81	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.38	0.34	5.55	0.65	0.12	0.64	0.30	1.36	0.25	0.64
PRS-High	1.76	0.67	11.33	0.16		0.80	0.35	1.87	0.61	
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)*</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	-	-	-	-	-	-	-	-	-	-
Deficient (<20 ng/ml)	-	-	-	-	-	-	-	-	-	-
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	-	-	-	-	-	-	-	-	-	-
PRS-High	-	-	-	-	-	-	-	-	-	-
Insufficient*PRS-Intermediate	-	-	-	-	-	-	-	-	-	-
Deficient*PRS-Intermediate	-	-	-	-	-	-	-	-	-	-
Insufficient*PRS-High	-	-	-	-	-	-	-	-	-	-
Deficient*PRS-High	-	-	-	-	-	-	-	-	-	-

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal social class, parity, smoking during 1st trimester, maternal psychopathology during pregnancy, child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Insufficient sample size to run GxE analysis

Supplemental Table 3.5 (continued)

	Trimester 3									
	Childhood depression (n=1719)					Adolescent depression (n=1452)				
	OR	LCI	UCI	p	P-trend	OR	LCI	UCI	p	P-trend
<b>Model 1: Maternal 25(OH)D</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.91	0.56	1.47	0.71	0.53	1.13	0.76	1.66	0.55	0.19
Deficient (<20 ng/ml)	1.17	0.72	1.92	0.52		1.31	0.87	1.98	0.19	
<b>Model 2: PRS</b>										
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.27	0.73	2.21	0.39	<b>0.004</b>	1.12	0.75	1.69	0.58	0.06
PRS-High	<b>2.15</b>	<b>1.22</b>	<b>3.78</b>	<b>0.008</b>		1.54	0.98	2.40	0.06	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.92	0.57	1.49	0.74	0.50	1.10	0.75	1.63	0.62	0.23
Deficient (<20 ng/ml)	1.20	0.73	1.96	0.47		1.30	0.86	1.96	0.22	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.27	0.73	2.21	0.40	<b>0.005</b>	1.12	0.74	1.69	0.58	0.06
PRS-High	<b>2.15</b>	<b>1.22</b>	<b>3.79</b>	<b>0.008</b>		1.54	0.98	2.41	0.06	
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)*</b>										
Normal ( $\geq 30$ ng/ml)	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-30 ng/ml)	0.71	0.26	1.95	0.51	-	1.12	0.51	2.45	0.78	-
Deficient (<20 ng/ml)	0.26	0.06	1.22	0.09		1.10	0.48	2.54	0.82	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	0.72	0.30	1.72	0.46	-	1.31	0.67	2.56	0.43	-
PRS-High	1.35	0.57	3.17	0.50		1.00	0.45	2.21	0.99	
Insufficient*PRS-Intermediate	1.28	0.36	4.61	0.70		0.78	0.30	2.03	0.61	
Deficient*PRS-Intermediate	7.25	1.31	40.10	0.02	0.10	0.80	0.28	2.24	0.67	0.24
Insufficient*PRS-High	1.61	0.46	5.71	0.46		1.48	0.80	4.39	0.48	
Deficient*PRS-High	4.64	0.81	26.50	0.08		2.47	0.50	7.69	0.12	

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal social class, parity, smoking during 1st trimester, maternal psychopathology during pregnancy, child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + any significant PRS\*covariate and maternal 25(OH)D\*covariate interactions

**Supplemental Table 3.6 Complete case analysis for the association between maternal vitamin D status during pregnancy, PRS, and child/adolescent depression**

	Childhood depression (n=2091)				Adolescent depression (n=1812)			
	OR	95% CI	p	p-trend	OR	95% CI	p	p-trend
<b>Model 1: Maternal 25(OH)D</b>								
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	1.19	0.77, 1.84	0.43	0.57	1.01	0.71, 1.43	0.97	0.15
Deficient (<20 ng/mL)	1.14	0.72, 1.80	0.57		1.30	0.91, 1.85	0.14	
<b>Model 2: PRS</b>								
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.39	0.89, 2.24	0.16	0.12	0.85	0.60, 1.20	0.34	0.10
PRS-High	1.51	0.90, 2.56	0.12		1.36	0.93, 1.99	0.11	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>								
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	1.19	0.77, 1.84	0.44	0.55	1.03	0.73, 1.47	0.86	0.15
Deficient (<20 ng/mL)	1.14	0.72, 1.80	0.58		1.32	0.92, 1.88	0.13	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.38	0.88, 2.23	0.17	0.12	0.85	0.60, 1.20	0.34	0.10
PRS-High	1.51	0.90, 2.56	0.12		1.36	0.93, 2.00	0.11	
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)</b>								
	OR	95% CI	p	p-inter.	OR	95% CI	p	p-inter.
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	0.49	0.17, 1.32	0.16	-	0.71	0.35, 1.43	0.35	-
Deficient (<20 ng/mL)	0.29	0.09, 0.84	0.03		1.28	0.65, 2.54	0.47	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	0.74	0.36, 1.57	0.43	-	4.50	1.19, 17.7	0.03	-
PRS-High	0.61	0.24, 1.47	0.28		3.73	0.77, 19.3	0.11	
Insufficient*PRS-Intermediate	2.70	0.90, 0.87	0.08		0.85	0.35, 2.05	0.72	
Insufficient*PRS-High	2.89	0.92, 0.99	0.08	0.12	0.72	0.31, 1.66	0.44	0.18
Deficient*PRS-Intermediate	4.70	1.35, 1.78	0.02		2.36	0.90, 6.30	0.08	
Deficient*PRS-High	3.39	0.88, 1.41	0.08		1.47	0.57, 3.86	0.43	

Abbreviations: CI: confidence intervals; OR: odds ratio; p-inter.: p for interaction; PRS: polygenic risk score

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal occupation, parity, smoking during 1st trimester, maternal depression during pregnancy, and child sex

Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + significant PRS\*covariate, maternal 25(OH)D\*covariate interactions

**Supplemental Table 3.7 Negative binomial regression results for the association between maternal vitamin D status during pregnancy, PRS, and child/adolescent depressive symptoms**

	Childhood depression (n=2938)				Adolescent depression (n=2485)			
	RR	95% CI	p	P-trend	RR	95% CI	p	P-trend
<b>Model 1: Maternal 25(OH)D</b>								
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	1.02	0.94, 1.10	0.68	0.48	1.02	0.94, 1.11	0.63	0.15
Deficient (<20 ng/mL)	1.03	0.95, 1.12	0.48		1.07	0.98, 1.17	0.15	
<b>Model 2: PRS</b>								
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.10	1.01, 1.18	0.03	<b>0.002</b>	1.01	0.93, 1.10	0.86	<b>0.03</b>
PRS-High	1.15	1.06, 1.26	0.002		1.11	1.01, 1.23	0.04	
<b>Model 3: PRS and Maternal 25(OH)D (Main effects)</b>								
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	1.02	0.94, 1.10	0.68	0.49	1.02	0.94, 1.11	0.63	0.15
Deficient (<20 ng/mL)	1.03	0.95, 1.11	0.50		1.07	0.98, 1.17	0.15	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.09	1.01, 1.18	0.03	<b>0.002</b>	1.01	0.92, 1.10	0.87	<b>0.04</b>
PRS-High	1.15	1.06, 1.26	0.002		1.11	1.01, 1.23	0.04	
<b>Model 4: PRS and Maternal 25(OH)D (Interaction)*</b>								
	RR	95% CI	p	P-inter.	RR	95% CI	p	P-inter.
Normal ( $\geq 30$ ng/mL)	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
Insufficient (20-29.9 ng/mL)	0.95	0.82, 1.11	0.52	-	1.06	0.89, 1.26	0.49	-
Deficient (<20 ng/mL)	0.94	0.80, 1.11	0.49		0.99	0.83, 1.19	0.94	
PRS-Low	<i>ref</i>	<i>ref</i>	<i>ref</i>		<i>ref</i>	<i>ref</i>	<i>ref</i>	
PRS-Intermediate	1.02	0.90, 1.17	0.72	-	1.03	0.88, 1.19	0.72	-
PRS-High	1.08	0.93, 1.26	0.32		1.04	0.88, 1.24	0.63	
Insufficient*PRS-Intermediate	1.09	0.91, 1.32	0.34		0.91	0.74, 1.12	0.39	
Deficient*PRS-Intermediate	1.11	0.91, 1.36	0.28	0.79	1.05	0.84, 1.31	0.66	0.41
Insufficient*PRS-High	1.09	0.88, 1.34	0.43		1.02	0.81, 1.30	0.84	
Deficient*PRS-High	1.13	0.90, 1.42	0.28		1.21	0.94, 1.56	0.14	

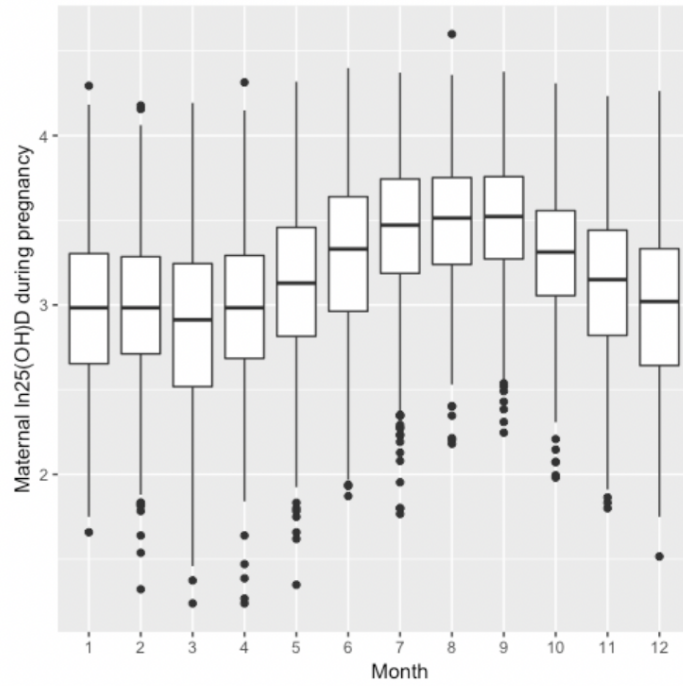
Abbreviations: CI: confidence intervals; RR: risk ratio; p-inter.: p for interaction; PRS: polygenic risk score

Model 1: Adjusted for gestational age of 25(OH)D measurement, maternal age, maternal pre-pregnancy BMI, maternal education, maternal occupation, parity, smoking during 1st trimester, maternal depression during pregnancy, and child sex

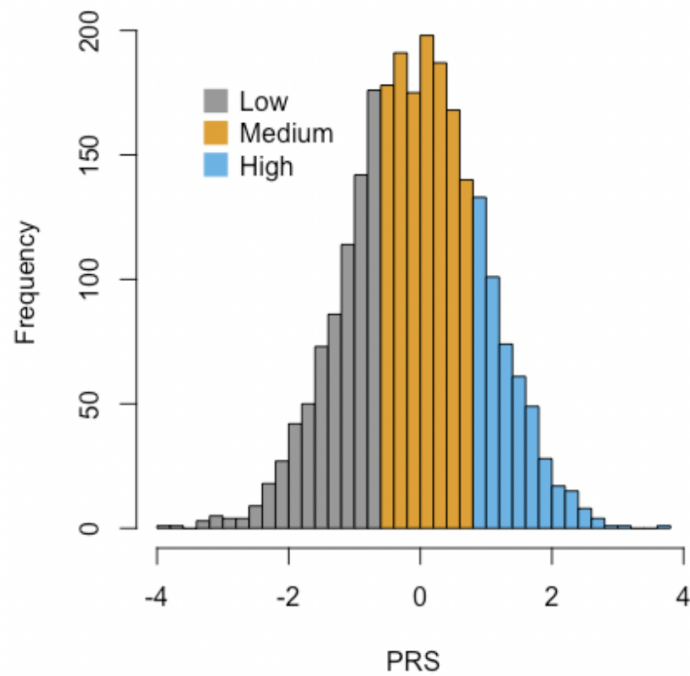
Model 2: Model 1 confounders + 3 PCs

Model 3: Model 2 confounders

Model 4: Model 3 confounders + significant PRS\*covariate, maternal 25(OH)D\*covariate interactions

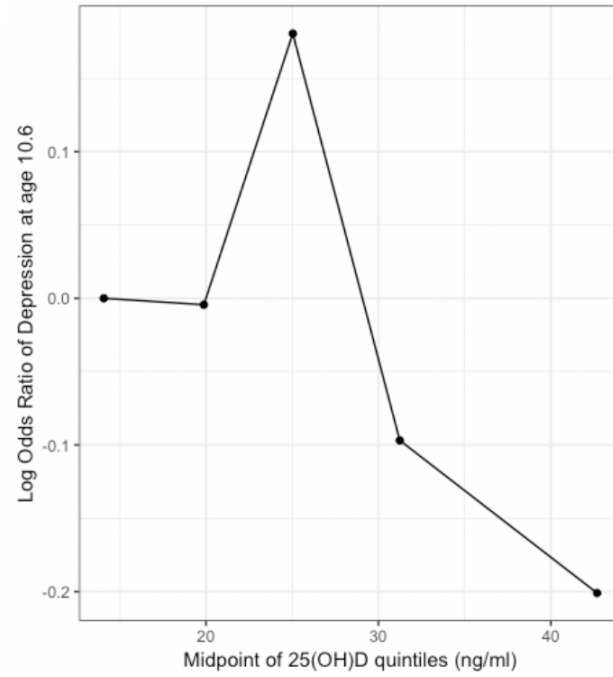


**Supplemental Figure 3.1 Patterns of maternal 25(OH)D levels during pregnancy by month in the study population**

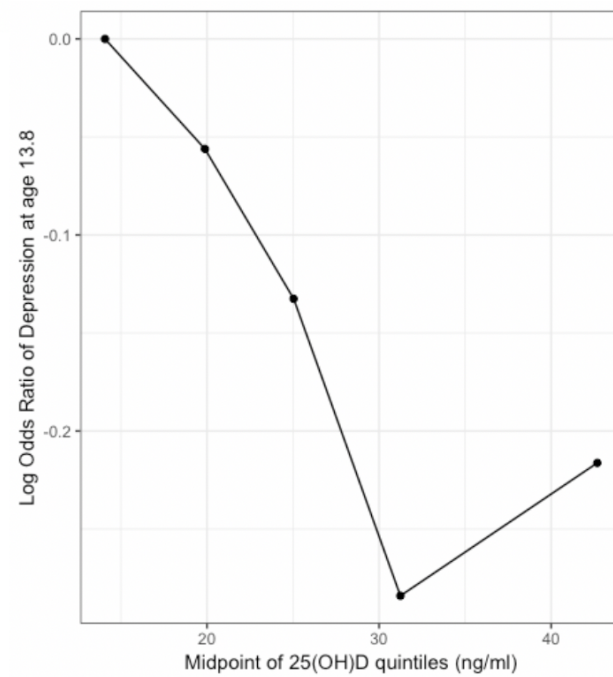


**Supplemental Figure 3.2 Distribution of offspring MDD PRS and categorizations (low, medium, and high polygenic risk) in the study population**

(A)



(B)



**Supplemental Figure 3.3 The association between quintiles of maternal 25(OH)D, compared to the lowest quintile, and the log odds of offspring depression in (A) childhood and (B) adolescence**