

Causal roles of daytime sleepiness in cardiometabolic diseases and osteoporosis

M. GUO¹, T. FENG², M. LIU³, Z. HUA², Y. MA², J.-P. CAI², X.-J. LI⁴

¹Department of Clinical Nutrition, The First Affiliated Hospital of Chongqing Medical University, Chongqing, China

²Institute of Traumatology & Orthopedics, Nanjing University of Chinese Medicine, Nanjing, Jiangsu, China

³Department of Orthopedic Surgery, Lianyungang Second People's Hospital, Lianyungang, Jiangsu, China

⁴Department of Gerontology, The First Branch of The First Affiliated Hospital of Chongqing Medical University, Chongqing, China

Abstract. – OBJECTIVE: Daytime sleepiness has some association with cardiometabolic diseases and osteoporosis, but it is unknown whether their relationship is causal. This two-sample Mendelian randomization (MR) study aims to explore their causal relationship.

MATERIALS AND METHODS: We included the largest genome-wide association studies (GWASs) associated with daytime sleepiness, cardiometabolic diseases and osteoporosis. 34 single nucleotide polymorphisms (SNPs) were used as the instrumental variables of daytime sleepiness.

RESULTS: Genetic predisposition to excessive daytime sleepiness was strongly associated with increased risk of coronary artery disease (beta-estimate: 0.610, 95% confidence interval [CI]: 0.128 to 1.093, standard error [SE]: 0.246, p -value=0.013) and may increase the incidence of type 2 diabetes (beta-estimate: 0.614, 95% CI: 0.009 to 1.219, SE: 0.309, p -value=0.047). We found no causal influence of daytime sleepiness on heart failure, atrial fibrillation, cerebral ischemia, intracerebral hemorrhage, forearm bone mineral density (FA-BMD), femoral neck BMD (FN-BMD), and lumbar spine BMD (LS-BMD).

CONCLUSIONS: This study suggested that excessive daytime sleepiness was causally associated with increased risk of coronary artery disease, which may benefit to prevent this disease.

Key Words:

Daytime sleepiness, Cardiometabolic diseases, Osteoporosis, Mendelian randomization study.

colepsy and circadian rhythm disorders^{1,2}. Its occurrence also results from the progression and treatment of some diseases^{3,4}. Excessive daytime sleepiness has become the risk factor of motor vehicle crashes, work-related accidents and loss of productivity^{5,6}. In addition, these patients also commonly suffer from poor cognition, behavior and quality of life⁷.

Previous observational studies⁸⁻¹⁰ documented that excessive daytime sleepiness may be associated with the incidence of cardiometabolic disorders, stroke and osteoporosis, but these connections may be affected by potential confounding factors and reverse causality. Genetic factors contribute to the variation in daytime sleepiness because the heritability of daytime sleepiness is approximately between 0.37 and 0.48 in twin studies^{11,12}. Genome-wide association studies (GWASs) find that daytime sleepiness, cardiometabolic disorders and osteoporosis are highly polygenic traits¹³⁻²⁰.

In order to explore the causal roles of daytime sleepiness in cardiometabolic diseases and osteoporosis, Mendelian randomization (MR) study has been designed and conducted to establish the causal relationship between exposure phenotype and outcome phenotype¹⁹⁻²³. This two-sample MR study can increase the scope and statistical power of MR by using instrumental variables of daytime sleepiness^{20,24,25}, and aim to explore the causal effect of daytime sleepiness on the incidence of cardiometabolic diseases and osteoporosis.

Introduction

Excessive daytime sleepiness has the typical feature of chronic insufficient sleep and can cause several sleep disorders, including sleep apnea, nar-

Materials and Methods

Genetic Instrument for Daytime Sleepiness

The largest available GWAS reported the frequency of daytime sleepiness among 452,071 par-

Table I. Details of studies and datasets used for analyses.

	Traits	Samples size	Population	Consortium or cohort study (Link URL)
Exposure	Daytime sleepiness	452,071	European	UK Biobank
	Coronary artery disease	547,261	European	UK Biobank and CARDIoGRAMplusC4D (https://cvd.hugeamp.org/)
Cardiometabolic diseases	Heart failure	977,323	European	UK Biobank (http://www.broadcvdi.org/)
	Atrial fibrillation	587,446	Predominant European (Mixed)	Meta-analysis of more than 50 studies (http://www.broadcvdi.org/)
	Type 2 diabetes	898,130	European	DIAGRAM (http://diagram-consortium.org)
Stroke	Cerebral ischemia	401,937	European	UK Biobank (https://www.leelabsg.org/resources)
	Intracerebral hemorrhage	399,717	European	
Osteoporosis	forearm BMD	53236	European	GEFOS (http://www.gefos.org)
	femoral neck BMD	53236	European	
	lumbar spine BMD	53236	European	

Participants of European genetic ancestry (Table I). Daytime sleepiness was determined by the question: “How likely are you to doze off or fall asleep during the daytime when you don’t mean to? (e.g.: when working, reading or driving)”, and answer categories included “never” (n=347,285), “sometimes” (n=92,794), “often” (n=11,963) and “all of the time” (n=29). Four categories were analyzed as a continuous variable using a linear mixed regression model adjusted for age, sex, genotyping array, ten principal components (PCs) of ancestry and genetic relatedness matrix²⁶.

Initially, 37 SNPs were thought to have robust association with daytime sleepiness ($p < 5 \times 10^{-8}$, **Supplementary Table I**). Linkage disequilibrium (LD) between selected SNPs was calculated using European samples from the 1000 Genomes project, and three SNPs (rs34478464, rs7162082, rs189568347) were excluded due to high LD ($r^2 \geq 0.001$). Finally, 34 SNPs were used as instrumental variables of daytime sleepiness (**Supplementary Table II**).

Outcome Data Sources

Table I demonstrated the summary-level data for the genetic associations with the outcomes. Briefly, we used the GWAS summary data of cardiometabolic diseases including coronary artery disease (547,261 individuals) from UK Biobank and CARDIoGRAMplusC4D²⁷, heart failure (977,323 individuals) from UK Biobank²⁸, atrial fibrillation (587,446 individuals) from one large meta-analysis²⁹ and type 2 diabetes (898,130 individuals) from DIAGRAM³⁰. For the association with stroke, GWAS summary data reported cerebral ischemia (401,937 individuals) and intracerebral hemorrhage (399,717 individuals) from UK Biobank¹⁶. Another large meta-analysis³¹ reported

the genetic variants associated with FN-BMD, FA-BMD and LS-BMD among 53,236 individuals of European ancestry. BMD was measured at the trabecular structure of forearm (distal 1/3 of radius), femoral neck and lumbar spine (L1-4). BMD was measured by dual X-ray absorptiometry.

Most GWASs were adjusted for sex, BMI and genetic principal components. All participants were all from European descent except those with atrial fibrillation from predominantly European descent (mixed descents). Summary statistics for SNPs related to daytime sleepiness and corresponding statistics of outcomes were presented in **Supplementary Table II**.

Statistical Analysis

To determine MR estimates of daytime sleepiness on each outcome, we conducted the inverse variance weighted (IVW), weighted median and MR-Egger regression methods^{32,33}. MR-PRESSO was used to assess the presence of pleiotropy and the effect estimates were recalculated after excluding SNP outliers³⁴. The ethical approval for each study included in this investigation can be found in the original publications (including informed consent from each participant). The differences with $p < 0.05$ were considered statistically significant. All these analyses were conducted in R V.4.0.4 by using the R packages of ‘MendelianRandomization’³⁵, ‘TwoSampleMR’³⁶ and ‘MR-PRESSO’³⁷.

Results

Cardiometabolic Diseases

We evaluated the causal effect of daytime sleepiness on coronary artery disease, heart failure, atrial fibrillation and type 2 diabetes in this MR

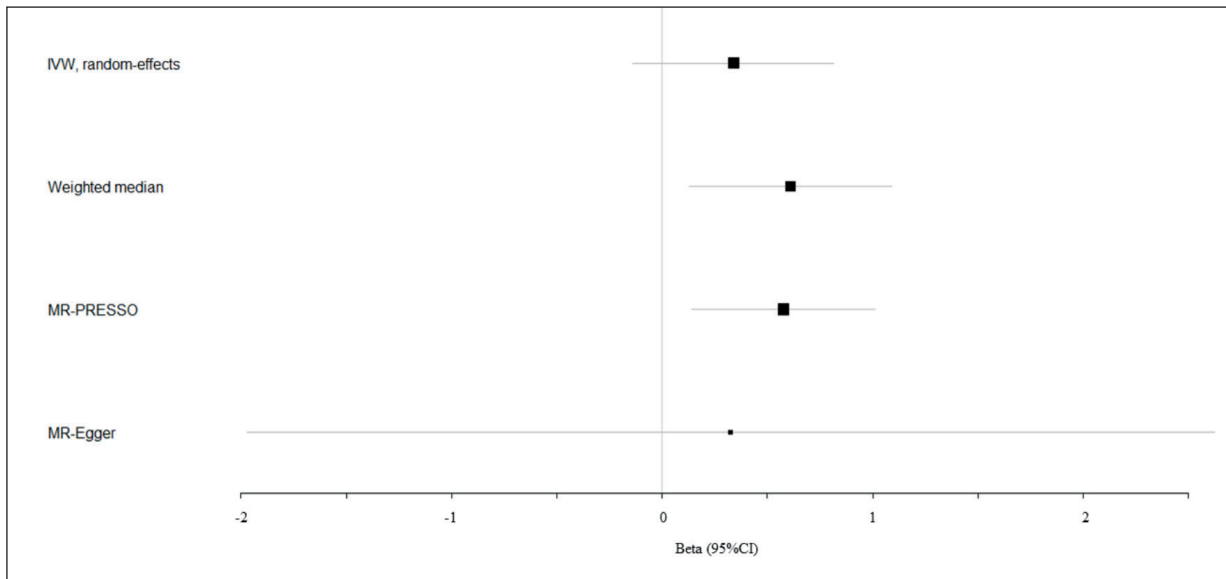


Figure 1. Beta (95% CIs) for causal association between daytime sleepiness and coronary artery disease through multiple analyses.

study (Table II). According to the weighted-median analysis, genetically excessive daytime sleepiness played a significant causal role in increased the risk of coronary artery disease (beta-estimate: 0.610, 95% CI: 0.128 to 1.093, SE: 0.246, p -value=0.013), but it was not supported by IVW analysis (beta-estimate: 0.339, 95% CI: -0.137 to 0.816, SE: 0.243, p -value=0.163, Figure 1).

In addition, the weighted-median analysis suggested some evidence to support the causal effect of excessive daytime sleepiness on increased risk of type 2 diabetes (beta-estimate: 0.614, 95% CI: 0.009 to 1.219, SE: 0.309, p -value=0.047), which was not confirmed by the IVW analysis (beta-estimate: 0, 95% CI: -0.794 to 0.794, SE: 0.405, p -value=1, Figure 2). IVW analyses showed that

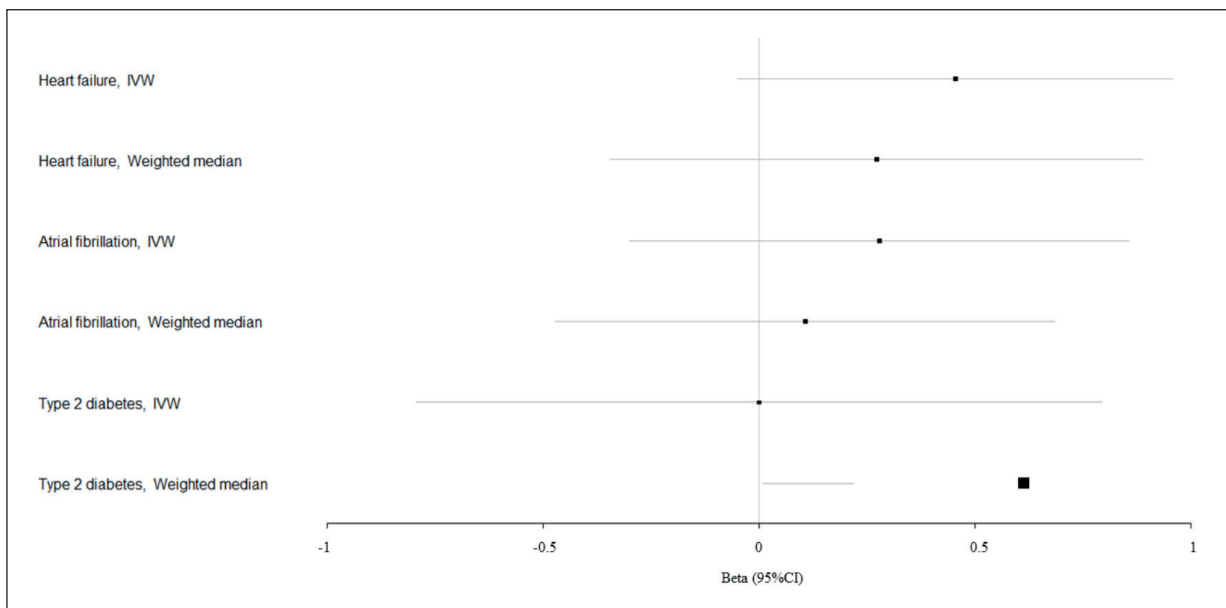


Figure 2. Beta (95% CIs) for causal influence of daytime sleepiness on heart failure, atrial fibrillation and type 2 diabetes through multiple analyses.

daytime sleepiness demonstrated no significant causal impact on heart failure (beta-estimate: 0.455, 95% CI: -0.049 to 0.958, SE: 0.257, p -value=0.077) or atrial fibrillation (beta-estimate: 0.279, 95% CI: -0.301 to 0.858, SE: 0.296, p -value=0.439), which were both confirmed by the weighted-median analyses (Figure 2).

Stroke

Excessive daytime sleepiness was often reported in patients who experienced stroke³⁸. Our IVW analyses found no causal roles of daytime sleepiness in cerebral ischemia (beta-estimate: -0.241, 95% CI: -1.587 to 1.105, SE: 0.687, p -value=0.726) or intracerebral hemorrhage (beta-estimate: 1.988, 95% CI: -4.727 to 0.751, SE: 1.397, p -value=0.115, Figure 2), which were both confirmed by the weighted-median and MR-Egger analyses (Table II).

Osteoporosis

Previous studies³⁹ demonstrated that daytime sleepiness was a risk factor of low BMD. Here, we explored the causal effect of daytime sleepiness on FA-BMD, FN-BMD and LS-BMD (Table II). According to IVW analysis, daytime sleepiness showed no causal influence on FA-BMD (beta-estimate: 0.058, 95% CI: -0.824 to 0.939, SE: 0.450, p -value=0.898), FN-BMD (beta-estimate: 0.238, 95% CI: -0.145 to 0.621, SE: 0.195, p -value=0.173) or LS-BMD (beta-estimate: -0.222, 95% CI: -0.835 to 0.390, SE: 0.312, p -value=0.173, Figure 3). These results were confirmed by the weighted-median and MR-Egger analyses (Table II).

Evaluation of Assumptions and Sensitivity Analyses

We found little evidence of directional pleiotropy for all models (MR-Egger intercept p -value>0.05, Table II). Significant heterogeneity remained for coronary artery disease, atrial fibrillation, and type 2 diabetes. Thus, MR-PRESSO method was conducted to identify 2 outliers (rs1846644, rs11078398) for coronary artery disease, 2 outliers (rs7607363, rs843372) for atrial fibrillation, 3 outliers (rs12140153, rs4665972, rs11078398) for type 2 diabetes and 1 outlier (rs11078398) for LS-BMD (Table III).

After excluding these SNP outliers, genetical-ly excessive daytime sleepiness was confirmed to have a causal effect on increased risk of coronary artery disease (beta-estimate: 0.576, 95% CI: 0.140 to 1.011, SE: 0.222, p -value=0.010, Figure 1 and

Table III). In addition, the MR association between daytime sleepiness with other outcomes were not changed after excluding the SNP outlier variants (Table III).

Discussion

Overall, our two-sample MR study confirms that excessive daytime sleepiness is strongly and causally associated with increased risk of coronary artery disease and may result in the increased risk of type 2 diabetes. However, we found no causal effect of daytime sleepiness on heart failure, atrial fibrillation, cerebral ischemia, intracerebral hemorrhage and three sites of BMD (i.e., FA-BMD, FN-BMD and LS-BMD).

Many studies^{8,40,41} reported the associations between daytime sleepiness and cardiovascular diseases, such as coronary heart disease and stroke, but yielded conflicting results. Effective treatment of daytime sleepiness significantly improves the sleepiness symptoms and the quality of life⁴²⁻⁴⁴. If there are causal relationships between daytime sleepiness and cardiovascular diseases, the assessment, diagnosis, and treatment of excessive daytime sleepiness is of critical significance for the primary prevention of cardiovascular diseases.

These inconsistent results may be subject to potential confounding factors and reverse causality, which can be effectively prevented in this two-sample MR study^{19,32}. Large-scale patient population were involved in our study, including the GWAS meta-analyses of daytime sleepiness (n=452,071) and coronary artery disease (n=547,261 individuals). Our research results revealed excessive daytime sleepiness displayed a robustly causal role in the increased risk of coronary artery disease, which was further confirmed by the PRESSO test (beta-estimate: 0.576, 95% CI: 0.140 to 1.011, SE: 0.222, p -value=0.010).

The positive MR association between daytime sleepiness and coronary artery disease is revealed in this study, and its mechanisms are attributed to systemic inflammation and metabolic syndrome⁴⁵. Inflammatory cytokines contribute to sleepiness and fatigue, and they are presumed to be protective responses and promote recovery from illness⁴⁶. During the process of atherosclerosis in patients with coronary artery disease, multiple mediators of inflammation induce immune cells to infiltrate the arterial intima^{47,48}. The systemic inflammation in patients with daytime sleepiness participates in the pathogenesis of metabolic disorders, such as obe-

Table II. Mendelian randomization estimates of daytime sleepiness on outcomes.

Variables	IVW							Weighted median				MR-Egger					Pleiotropy <i>p</i> value			
	Estimate	SE	95% CI	<i>p</i> -value	Q value	I ²	Heterogeneity <i>p</i> -value	Estimate	SE	95% CI	<i>p</i> -value	Estimate	SE	95% CI	<i>p</i> -value	Intercept		SE	95% CI	
Cardiometabolic disease																				
Coronary artery disease	0.339	0.243	-0.137,0.816	0.163	86.326	62.90%	0.000	0.610	0.246	0.128,1.093	0.013	0.326	1.171	-1.970,2.622	0.781	0.000	0.009	-0.017,0.017	0.991	
Heart failure	0.455	0.257	-0.049,0.958	0.077	52.484	37.10%	0.017	0.273	0.315	-0.345,0.890	0.387	1.892	1.192	-0.443,4.228	0.112	-0.011	0.009	-0.028,0.006	0.217	
Atrial fibrillation	0.279	0.296	-0.301,0.858	0.346	78.777	59.40%	0.000	0.107	0.296	-0.473,0.686	0.718	0.308	1.418	-2.472,3.088	0.828	0.000	0.010	-0.021,0.020	0.983	
Type 2 diabetes	0.000	0.405	-0.794,0.794	1.000	142.509	77.50%	0.000	0.614	0.309	0.009,1.219	0.047	2.067	1.885	-1.629,5.762	0.273	-0.015	0.014	-0.043,0.012	0.262	
Stroke																				
Cerebral ischemia	-0.241	0.687	-1.587,1.105	0.726	30.963	0.00%	0.519	-1.230	0.975	-3.140,0.680	0.207	-5.113	3.202	-11.389,1.163	0.110	0.037	0.024	-0.009,0.083	0.119	
Intracerebral hemorrhage	-1.988	1.397	-4.727,0.751	0.115	30.548	0.00%	0.540	-2.996	2.017	-6.950,0.958	0.137	2.082	6.511	-10.680,14.845	0.749	-0.031	0.048	-0.125,0.063	0.522	
Osteoporosis																				
FA-BMD	0.058	0.450	-0.824,0.939	0.898	40.536	18.60%	0.172	0.620	0.592	-0.540,1.780	0.295	2.941	2.060	-1.096,6.978	0.153	-0.022	0.015	-0.052,0.008	0.152	
FN-BMD	0.238	0.195	-0.145,0.621	0.223	33.198	0.60%	0.458	0.330	0.287	-0.232,0.892	0.250	-0.082	0.920	-1.885,1.722	0.929	0.002	0.007	-0.011,0.016	0.722	
LS-BMD	-0.222	0.312	-0.835,0.390	0.477	62.324	47.10%	0.002	-0.101	0.356	-0.798,0.596	0.777	0.133	1.470	-2.749,3.014	0.928	-0.003	0.011	-0.024,0.019	0.805	

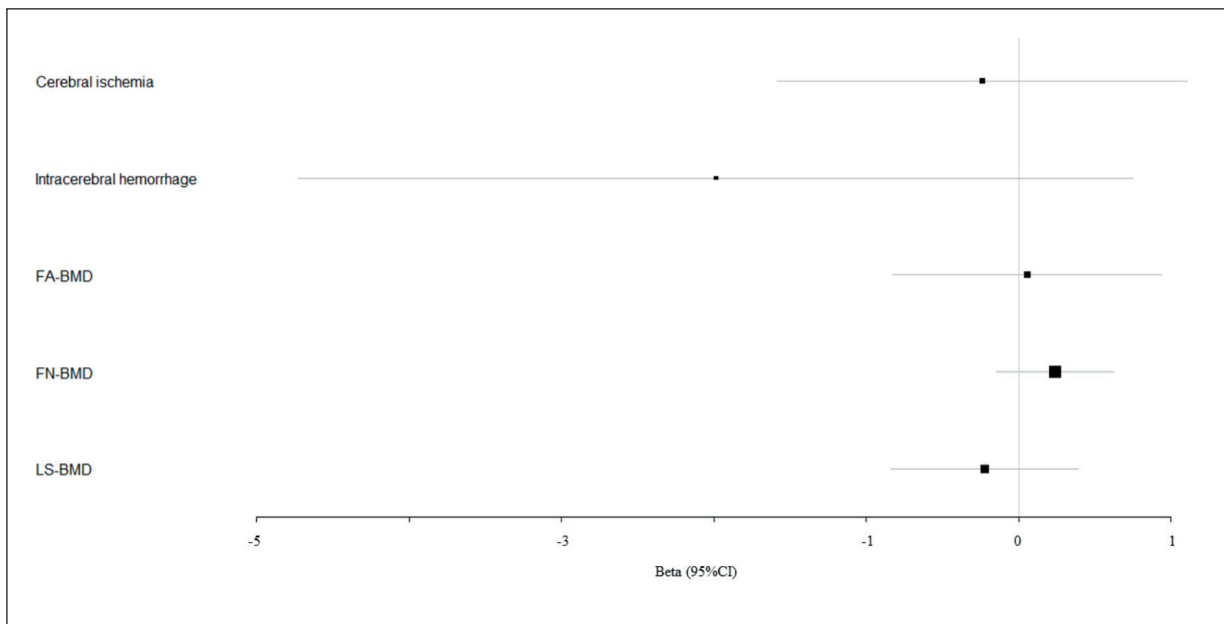


Figure 3. Beta (95% CIs) for causal influence of daytime sleepiness on cerebral ischemia, intracerebral hemorrhage, FA-BMD, FN-BMD and LS-BMD through the IVW analyses.

sity and diabetes⁴⁹. Metabolic syndrome is associated with compromised insulin sensitivity, raised blood pressure and increased low-density lipoprotein cholesterol levels, which are widely accepted as the risk factors of coronary artery disease⁵⁰⁻⁵³. In addition, we also find the potential causal association between excessive daytime sleepiness and increased risk of type 2 diabetes, which is regulated by reduced insulin sensitivity⁵¹.

Osteoporosis leads to low bone mass and microstructure deterioration. These patients have increased risk of fracture⁵⁴⁻⁵⁷. Excessive daytime sleepiness is common in the elderly, which suggests some association between daytime sleepiness and BMD^{9,58}. In one observational study, daytime napping duration ≥ 30 min in postmenopausal women (n=4,962) and ≥ 60 min in premenopausal women (n=1,548) were found to increase the risk of osteopenia and osteoporosis⁹. However, our MR study found no causal association between daytime

sleepiness and three sites of BMD (i.e., FA-BMD, FN-BMD and LS-BMD), which was confirmed by multiple MR analyses and sensitivity analyses. The positive association found in observational studies may result from confounding factors and reverse causality. In addition, we find limited evidence of associations between daytime sleepiness and heart failure, atrial fibrillation, cerebral ischemia and intracerebral hemorrhage.

Limitations

We should consider several limitations. Firstly, all included participants are of predominantly European people, and we do not know whether our findings are applicable to other populations. Secondly, excessive daytime sleepiness shows the MR association with increased risk of coronary artery disease, but the related mechanisms

Table III. Mendelian randomization estimates between daytime sleepiness and outcomes after excluding outliers detected by MR-PRESSO.

Outcomes	Estimate	SE	95% CI	p-value
Coronary artery disease excluding 2 outliers (rs1846644, rs11078398)	0.576	0.222	0.140,1.011	0.010
Atrial fibrillation excluding 2 outliers (rs7607363, rs843372)	0.261	0.247	-0.224,0.746	0.291
Type 2 diabetes excluding 3 outliers (rs12140153, rs4665972, rs11078398)	0.297	0.254	-0.199,0.794	0.241

are not clear. Thirdly, MR association between daytime sleepiness and type 2 diabetes is not robust, and more populations should be included to confirm it.

Conclusions

This two-sample MR study supports that excessive daytime sleepiness causally results in increased risk of coronary artery disease, which may help prevent it.

Conflict of Interest

The authors declare that they have no conflict of interest.

Acknowledgments

The authors acknowledged the GENetic Factors for OSTeoporosis Consortium, the UK Biobank and DIAGRAM consortium for contributing the data used in this work.

Funding

None.

Data Availability

Data supporting the findings of this study were available within the paper.

References

- Ohayon MM. From wakefulness to excessive sleepiness: what we know and still need to know. *Sleep Med Rev* 2008; 12: 129-141.
- Slater G, Steier J. Excessive daytime sleepiness in sleep disorders. *J Thorac Dis* 2012; 4: 608-616.
- Bixler EO, Vgontzas AN, Lin HM, Calhoun SL, Vela-Bueno A, Kales A. Excessive daytime sleepiness in a general population sample: the role of sleep apnea, age, obesity, diabetes, and depression. *J Clin Endocr Metab* 2005; 90: 4510-4515.
- Fernandez-Mendoza J, Vgontzas AN, Kritikou I, Calhoun SL, Liao D, Bixler EO. Natural history of excessive daytime sleepiness: role of obesity, weight loss, depression, and sleep propensity. *Sleep* 2015; 38: 351-360.
- Ferrie JE, Kumari M, Salo P, Singh-Manoux A, Kivimäki M., Sleep epidemiology--a rapidly growing field. *Int J Epidemiol* 2011; 40: 1431-1437.
- Lloberes P, Levy G, Descals C, Sampol G, Roca A, Sagales T, de la Calzada MD. Self-reported sleepiness while driving as a risk factor for traffic accidents in patients with obstructive sleep apnoea syndrome and in non-apnoeic snorers. *Resp Med* 2000; 94: 971-976.
- Ohayon MM, Vecchierini MF. Daytime sleepiness and cognitive impairment in the elderly population. *Arch Intern Med* 2002; 162: 201-208.
- Gangwisch JE, Rexrode K, Forman JP, Mukamal K, Malaspina D, Feskanich D. Daytime sleepiness and risk of coronary heart disease and stroke: results from the Nurses' Health Study II. *Sleep Med* 2014; 15: 782-788.
- Wang K, Wu Y, Yang Y, Chen J, Zhang D, Hu Y, Liu Z, Xu J, Shen Q, Zhang N, Mao X, Liu C. The associations of bedtime, nocturnal, and daytime sleep duration with bone mineral density in pre- and post-menopausal women. *Endocrine* 2015; 49: 538-548.
- Martynowicz H, Jodkowska A, Skomro R, Gać P, Brylka A, Bładowski M, Wojakowska A, Mazur G, Poręba R. The estimation of excessive daytime sleepiness in post-stroke patients - a polysomnographic study. *Resp Physiol Neurobi* 2019; 267: 1-5.
- Desai AV, Cherkas LF, Spector TD, Williams AJ. Genetic influences in self-reported symptoms of obstructive sleep apnoea and restless legs: a twin study. *Twin Res* 2004; 7: 589-595.
- Watson NF, Goldberg J, Arguelles L, Buchwald D. Genetic and environmental influences on insomnia, daytime sleepiness, and obesity in twins. *Sleep* 2006; 29: 645-649.
- Lane JM, Liang J, Vlasac I, Anderson SG, Bechtold DA, Bowden J, Emsley R, Gill S, Little MA, Luik AI, Loudon A, Scheer FA, Purcell SM, Kyle SD, Lawlor DA, Zhu X, Redline S, Ray DW, Rutter MK, Saxena R. Genome-wide association analyses of sleep disturbance traits identify new loci and high-light shared genetics with neuropsychiatric and metabolic traits. *Nat Genet* 2017; 49: 274-281.
- Warrington NM, Beaumont RN, Horikoshi M, Day FR, Helgeland, Laurin C, Bacelis J, Peng S, Hao K, Feenstra B, Wood AR, Mahajan A, Tyrrell J, Robertson NR, Rayner NW, Qiao Z, Moen GH, Vaudel M, Marsit CJ, Chen J, Nodzinski M, Schnurr TM, Zafarmand MH, Bradfield JP, Grarup N, Kooijman MN, Li-Gao R, Geller F, Ahluwalia TS, Paternoster L, Rueedi R, Huikari V, Hottenga JJ, Lyytikäinen LP, Cavadino A, Metrustry S, Cousminer DL, Wu Y, Thiering E, Wang CA, Have CT, Vilor-Tejedor N, Joshi PK, Painter JN, Ntalla I, Mähre R, Pitkänen N, van Leeuwen EM, Joro R, Lagou V, Richmond RC, Espinosa A, Barton SJ, Inskip HM, Holloway JW, Santa-Marina L, Estivill X, Ang W, Marsh JA, Reichetzeder C, Marullo L, Hoche B, Lunetta KL, Murabito JM, Relton CL, Kogevinas M, Chatzi L, Allard C, Bouchard L, Hivert MF, Zhang G, Muglia LJ, Heikkinen J, Morgen CS, van Kampen AHC, van Schaik BDC, Mentch FD, Langenberg C, Luan J, Scott RA, Zhao JH, Hemani G, Ring SM, Bennett AJ, Gaulton KJ, Fernandez-Tajes J, van Zuydam NR, Medina-Gomez C, de Haan HG, Rosendaal FR, Kutalik Z, Marques-Vidal P, Das S, Willemsen G, Mbarek H, Müller-Nurasyid M, Standl M, Appel EVR, Fonvig CE, Trier C, van Beijsterveldt CEM, Murcia M, Bustamante M, Bonas-Guarch S, Hougaard DM, Mercader JM, Linneberg A, Schraut KE, Lind PA, Medland SE, Shields BM, Knight BA, Chai JF, Panoutsopoulou K, Bartels M, Sánchez F, Stokholm J, Torrents D, Vinding RK, Willems SM, Atalay

- M, Chawes BL, Kovacs P, Prokopenko I, Tuke MA, Yaghoobkar H, Ruth KS, Jones SE, Loh PR, Murray A, Weedon MN, Tönjes A, Stumvoll M, Michaelsen KF, Eloranta AM, Lakka TA, van Duijn CM, Kiess W, Körner A, Niinikoski H, Pahkala K, Raitakari OT, Jacobsson B, Zeggini E, Dedoussis GV, Teo YY, Saw SM, Montgomery GW, Campbell H, Wilson JF, Vrijkotte TGM, Vrijheid M, de Geus E, Hayes MG, Kadarmideen HN, Holm JC, Beilin LJ, Pennell CE, Heinrich J, Adair LS, Borja JB, Mohlke KL, Eriksson JG, Widén EE, Hattersley AT, Spector TD, Kähönen M, Viikari JS, Lehtimäki T, Boomsma DI, Sebert S, Vollenweider P, Sørensen TIA, Bisgaard H, Bønnelykke K, Murray JC, Melbye M, Nohr EA, Mook-Kanamori DO, Rivadeneira F, Hofman A, Felix JF, Jaddoe VVW, Hansen T, Pisinger C, Vaag AA, Pedersen O, Uitterlinden AG, Järvelin MR, Power C, Hyppönen E, Scholtens DM, Lowe WL, Jr, Davey Smith G, Timpson NJ, Morris AP, Wareham NJ, Hakonarson H, Grant SFA, Frayling TM, Lawlor DA, Njølstad PR, Johansson S, Ong KK, McCarthy MI, Perry JRB, Evans DM, Freathy RM. Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors, *Nat Genet* 2019; 51: 804-814.
- 15) Iotchkova V, Huang J, Morris JA, Jain D, Barbieri C, Walter K, Min JL, Chen L, Astle W, Cocca M, Deelen P, Elding H, Farmaki AE, Franklin CS, Franberg M, Gaunt TR, Hofman A, Jiang T, Kleber ME, Lachance G, Luan J, Malerba G, Matchan A, Mead D, Memari Y, Ntalla I, Panoutsopoulou K, Pazoki R, Perry JRB, Rivadeneira F, Sabater-Lleal M, Sennblad B, Shin SY, Southam L, Traglia M, van Dijk F, van Leeuwen EM, Zaza G, Zhang W, Amin N, Butterworth A, Chambers JC, Dedoussis G, Dehghan A, Franco OH, Franke L, Frontini M, Gambaro G, Gasparini P, Hamsten A, Issacs A, Kooner JS, Kooperberg C, Langenberg C, Marz W, Scott RA, Swertz MA, Toniolo D, Uitterlinden AG, van Duijn CM, Watkins H, Zeggini E, Maurano MT, Timpson NJ, Reiner AP, Auer PL, Soranzo N. Discovery and refinement of genetic loci associated with cardiometabolic risk using dense imputation maps. *Nat Genet* 2016; 48: 1303-1312.
 - 16) Zhou W, Nielsen JB, Fritsche LG, Dey R, Gabrielsen ME, Wolford BN, LeFaive J, VandeHaar P, Gagliano SA, Gifford A, Bastarache LA, Wei WQ, Denny JC, Lin M, Hveem K, Kang HM, Abecasis GR, Willer CJ, Lee S. Efficiently controlling for case-control imbalance and sample relatedness in large-scale genetic association studies. *Nat Genet* 2018; 50: 1335-1341.
 - 17) Trajanoska K, Rivadeneira F. The genetic architecture of osteoporosis and fracture risk. *Bone* 2019; 126: 2-10.
 - 18) Yang TL, Shen H, Liu A, Dong SS, Zhang L, Deng FY, Zhao Q, Deng HW. A road map for understanding molecular and genetic determinants of osteoporosis. *Nat Rev Endocrinol*. 2020; 16: 91-103.
 - 19) He B, Yin L, Zhang M, Lyu Q, Quan Z, Ou Y. Causal Effect of Blood Pressure on Bone Mineral Density and Fracture: A Mendelian Randomization Study. *Front Endocrinol* 2021; 2: 716681.
 - 20) He B, Lyu Q, Yin L, Zhang M, Quan Z, Ou Y. Depression and Osteoporosis: A Mendelian Randomization Study. *Calcif Tissue Int*. 2021; 109: 675-684.
 - 21) Burgess S, Dudbridge F, Thompson SG. Combining information on multiple instrumental variables in Mendelian randomization: comparison of allele score and summarized data methods. *Stat Med* 2016; 35: 1880-1906.
 - 22) Dalbeth N, Topleless R, Flynn T, Cadzow M, Bolland MJ, Merriman TR. Mendelian randomization analysis to examine for a causal effect of urate on bone mineral density. *J Bone Miner Res* 2015; 30: 985-991.
 - 23) He B, Xia L, Zhao J, Yin L, Zhang M, Quan Z, Ou Y, Huang W. Causal Effect of Serum Magnesium on Osteoporosis and Cardiometabolic Diseases. *Front Nutr* 2021; 8: 738000.
 - 24) Pierce BL, Burgess S. Efficient design for Mendelian randomization studies: subsample and 2-sample instrumental variable estimators. *Am J Epidemiol* 2013; 178: 1177-1184.
 - 25) Burgess S, Scott RA, Timpson NJ, Davey Smith G, Thompson SG. Using published data in Mendelian randomization: a blueprint for efficient identification of causal risk factors. *Eur J Epidemiol* 2015; 30: 543-552.
 - 26) Wang H, Lane JM, Jones SE, Dashti HS, Ollila HM, Wood AR, van Hees VT, Brumpton B, Winsvold BS, Kantojärvi K, Palviainen T, Cade BE, Sofer T, Song Y, Patel K, Anderson SG, Bechtold DA, Bowden J, Emsley R, Kyle SD, Little MA, Loudon AS, Scheer F, Purcell SM, Richmond RC, Spiegelhalter K, Tyrrell J, Zhu X, Hublin C, Kaprio JA, Kristiansson K, Sulkava S, Paunio T, Hveem K, Nielsen JB, Willer CJ, Zwart JA, Strand LB, Frayling TM, Ray D, Lawlor DA, Rutter MK, Weedon MN, Redline S, Saxena R. Genome-wide association analysis of self-reported daytime sleepiness identifies 42 loci that suggest biological subtypes. *Nat Commun* 2019; 10: 3503.
 - 27) van der Harst P, Verweij N. Identification of 64 Novel Genetic Loci Provides an Expanded View on the Genetic Architecture of Coronary Artery Disease. *Circ Res* 2018; 122: 433-443.
 - 28) Shah S, Henry A, Roselli C, Lin H, Sveinbjörnsson G, Fatemifar G, Hedman Å K, Wilk JB, Morley MP, Chaffin MD, Helgadottir A, Verweij N, Dehghan A, Almgren P, Andersson C, Aragam KG, Ärnlöv J, Backman JD, Biggs ML, Bloom HL, Brandimarto J, Brown MR, Buckbinder L, Carey DJ, Chasman DI, Chen X, Chen X, Chung J, Chutkova W, Cook JP, Delgado GE, Denaxas S, Doney AS, Dörr M, Dudley SC, Dunn ME, Engström G, Esko T, Felix SB, Finan C, Ford I, Ghanbari M, Ghasemi S, Giedraitis V, Giulianini F, Gottdiener JS, Gross S, Guðbjartsson DF, Gutmann R, Haggerty CM, van der Harst P, Hyde CL, Ingelsson E, Jukema JW, Kavousi M, Khaw KT, Kleber ME, Kober L, Koekemoer A, Langenberg C, Lind L, Lindgren CM, London B, Lotta LA, Lovering RC, Luan J, Magnusson P, Mahajan A, Margulies KB, März W, Melander O, Mordi IR, Morgan T, Morris AD, Morris AP, Morrison AC, Nagle MW, Nelson CP, Niessner A, Niiranen T, O'Donoghue ML, Owens AT, Palmer CNA, Parry HM, Perola M, Portilla-Fernandez E, Psaty BM, Rice KM, Ridker PM, Romaine SPR, Rotter JI, Salo P, Salomaa V, van Setten J, Shalaby AA, Smelser DT, Smith NL, Stender S, Stott DJ, Svensson P, Tammesoo ML, Taylor KD, Teder-Laving M, Teumer A, Thorgeirsson G, Thorsteinsdóttir U, Torp-Pedersen C, Trompet S,

- Tyl B, Uitterlinden AG, Veluchamy A, Völker U, Voors AA, Wang X, Wareham NJ, Waterworth D, Weeke PE, Weiss R, Wiggins KL, Xing H, Yerges-Armstrong LM, Yu B, Zannad F, Zhao JH, Hemingway H, Samani NJ, McMurray JJV, Yang J, Visscher PM, Newton-Cheh C, Malarstig A, Holm H, Lubitz SA, Sattar N, Holmes MV, Cappola TP, Asselbergs FW, Hingorani AD, Kuchenbaecker K, Ellinor PT, Lang CC, Stefansson K, Smith JG, Vasani RS, Swerdlow DI, Lumbers RT. Genome-wide association and Mendelian randomisation analysis provide insights into the pathogenesis of heart failure. *Nat Commun* 2020; 11: 163.
- 29) Roselli C, Chaffin MD, Weng LC, Aeschbacher S, Ahlberg G, Albert CM, Almgren P, Alonso A, Anderson CD, Aragam KG, Arking DE, Barnard J, Bartz TM, Benjamin EJ, Bihlmeyer NA, Bis JC, Bloom HL, Boerwinkle E, Bottinger EB, Brody JA, Calkins H, Campbell A, Cappola TP, Carlquist J, Chasman DI, Chen LY, Chen YI, Choi EK, Choi SH, Christophersen IE, Chung MK, Cole JW, Conen D, Cook J, Crijs HJ, Cutler MJ, Damrauer SM, Daniels BR, Darbar D, Delgado G, Denny JC, Dichgans M, Dörr M, Dudink EA, Dudley SC, Esa N, Esko T, Eskola M, Fatkin D, Felix SB, Ford I, Franco OH, Geelhoed B, Grewal RP, Gudnason V, Guo X, Gupta N, Gustafsson S, Gutmann R, Hamsten A, Harris TB, Hayward C, Heckbert SR, Hernesniemi J, Hocking LJ, Hofman A, Horimoto A, Huang J, Huang PL, Huffman J, Ingelsson E, Ipek EG, Ito K, Jimenez-Conde J, Johnson R, Jukema JW, Kääh S, Kähönen M, Kamatani Y, Kane JP, Kastrati A, Kathiresan S, Katschnig-Winter P, Kavousi M, Kessler T, Kietselaer BL, Kirchhof P, Kleber ME, Knight S, Krieger JE, Kubo M, Launer LJ, Laurikka J, Lehtimäki T, Leineweber K, Lemaitre RN, Li M, Lim HE, Lin HJ, Lin H, Lind L, Lindgren CM, Lokki ML, London B, Loos RJF, Low SK, Lu Y, Lyytikäinen LP, Macfarlane PW, Magnusson PK, Mahajan A, Malik R, Mansur AJ, Marcus GM, Margolin L, Margulies KB, März W, McManus DD, Melander O, Mohanty S, Montgomery JA, Morley MP, Morris AP, Müller-Nurasyid M, Natale A, Nazarian S, Neumann B, Newton-Cheh C, Nie-meijer MN, Nikus K, Nilsson P, Noordam R, Oellers H, Olesen MS, Orho-Melander M, Padmanabhan S, Pak HN, Paré G, Pedersen NL, Pera J, Pereira A, Porteous D, Psaty BM, Pulit SL, Pullinger CR, Rader DJ, Refsgaard L, Ribasés M, Ridker PM, Rienstra M, Risch L, Roden DM, Rosand J, Rosenberg MA, Rost N, Rotter JI, Saba S, Sandhu RK, Schnabel RB, Schramm K, Schunkert H, Schurman C, Scott SA, Seppälä I, Shaffer C, Shah S, Shalaby AA, Shim J, Shoemaker MB, Siland JE, Sinisalo J, Sinner MF, Slowik A, Smith AV, Smith BH, Smith JG, Smith JD, Smith NL, Soliman EZ, Sotoodehnia N, Stricker BH, Sun A, Sun H, Svendsen JH, Tanaka T, Tanriverdi K, Taylor KD, Teder-Laving M, Teumer A, Thériault S, Trompet S, Tucker NR, Tveit A, Uitterlinden AG, Van Der Harst P, Van Gelder IC, Van Wagener DR, Verweij N, Vlachopoulou E, Völker U, Wang B, Weeke PE, Weijs B, Weiss R, Weiss S, Wells QS, Wiggins KL, Wong JA, Woo D, Worrall BB, Yang PS, Yao J, Yoneda ZT, Zeller T, Zeng L, Lubitz SA, Lunetta KL, Ellinor PT. Multi-ethnic genome-wide association study for atrial fibrillation. *Nat Genet* 2018; 50: 1225-1233.
- 30) Mahajan A, Taliun D, Thurner M, Robertson NR, Torres JM, Rayner NW, Payne AJ, Steinthorsdóttir V, Scott RA, Grarup N, Cook JP, Schmidt EM, Wuttke M, Sarnowski C, Mägi R, Nano J, Gieger C, Trompet S, Lecoeur C, Preuss MH, Prins BP, Guo X, Bielak LF, Below JE, Bowden DW, Chambers JC, Kim YJ, Ng MCY, Petty LE, Sim X, Zhang W, Bennett AJ, Bork-Jensen J, Brummett CM, Canouil M, Ec Kardt KU, Fischer K, Kardia SLR, Kronenberg F, Läll K, Liu CT, Locke AE, Luan J, Ntalla I, Nylander V, Schönherr S, Schurmann C, Yengo L, Bottinger EP, Brandslund I, Christensen C, Dedousis G, Florez JC, Ford I, Franco OH, Frayling TM, Giedraitis V, Hackinger S, Hattersley AT, Herder C, Ikram MA, Ingelsson M, Jørgensen ME, Jørgensen T, Kriebel J, Kuusisto J, Ligthart S, Lindgren CM, Linneberg A, Lyssenko V, Mamakou V, Meitinger T, Mohlke KL, Morris AD, Nadkarni G, Pankow JS, Peters A, Sattar N, Stančáková A, Strauch K, Taylor KD, Thorand B, Thorleifsson G, Thorsteinsdóttir U, Tuomilehto J, Witte DR, Dupuis J, Peyser PA, Zeigini E, Loos RJF, Froguel P, Ingelsson E, Lind L, Groop L, Laakso M, Collins FS, Jukema JW, Palmer CNA, Grallert H, Metspalu A, Dehghan A, Köttgen A, Abecasis GR, Meigs JB, Rotter JI, Marchini J, Pedersen O, Hansen T, Langenberg C, Wareham NJ, Stefansson K, Gloyn AL, Morris AP, Boehnke M, McCarthy MI. Fine-mapping type 2 diabetes loci to single-variant resolution using high-density imputation and islet-specific epigenome maps. *Nat Genet* 2018; 50: 1505-1513.
- 31) Zheng HF, Forgetta V, Hsu YH, Estrada K, Rosello-Diez A, Leo PJ, Dahia CL, Park-Min KH, Tobias JH, Kooperberg C, Kleinman A, Styrkarsdóttir U, Liu CT, Uggla C, Evans DS, Nielson CM, Walter K, Pettersson-Kymmer U, McCarthy S, Eriksson J, Kwan T, Jhamai M, Trajanoska K, Memari Y, Min J, Huang J, Danecek P, Wilmot B, Li R, Chou WC, Mokry LE, Moayyeri A, Claussnitzer M, Cheng CH, Cheung W, Medina-Gómez C, Ge B, Chen SH, Choi K, Oei L, Fraser J, Kraaij R, Hibbs MA, Gregson CL, Paquette D, Hofman A, Wibom C, Tranah GJ, Marshall M, Gardiner BB, Cremin K, Auer P, Hsu L, Ring S, Tung JY, Thorleifsson G, Enneman AW, van Schoor NM, de Groot LC, van der Velde N, Melin B, Kemp JP, Christiansen C, Sayers A, Zhou Y, Calderari S, van Rooij J, Carlson C, Peters U, Berlivet S, Dostie J, Uitterlinden AG, Williams SR, Farber C, Grinberg D, LaCroix AZ, Haessler J, Chasman DI, Giulianini F, Rose LM, Ridker PM, Eisman JA, Nguyen TV, Center JR, Noguez X, Garcia-Giralto N, Launer LL, Gudnason V, Mellström D, Vandenput L, Amin N, van Duijn CM, Karlsson MK, Ljunggren Ö, Svensson O, Hallmans G, Rousseau F, Giroux S, Bussière J, Arp PP, Koromani F, Prince RL, Lewis JR, Langdahl BL, Hermann AP, Jensen JE, Kaptoge S, Khaw KT, Reeve J, Formosa MM, Xuereb-Anastasi A, Åkeson K, McGuigan FE, Garg S, Olmos JM, Zarrabaitia MT, Riancho JA, Ralston SH, Alonso N, Jiang X, Goltzman D, Pastinen T, Grundberg E, Gauguier D, Orwoll ES, Karasik D, Davey-Smith G, Smith AV, Siggeirsdóttir K, Harris TB, Zillikens MC, van Meurs JB, Thorsteinsdóttir U, Maurano MT, Timpson NJ, Soranzo N, Durbin R, Wilson SG, Ntzani EE, Brown MA, Stefansson K, Hinds DA, Spector T, Cupples

- LA, Ohlsson C, Greenwood CM, Jackson RD, Rowe DW, Loomis CA, Evans DM, Ackert-Bicknell CL, Joyner AL, Duncan EL, Kiel DP, Rivadeneira F, Richards JB. Whole-genome sequencing identifies EN1 as a determinant of bone density and fracture. *Nature* 2015; 526: 112-117.
- 32) He B, Zhao J, Zhang M, Yin L, Quan Z, Ou Y, Huang W. Causal roles of circulating adiponectin in osteoporosis and cancers. *Bone* 2022; 155: 116266.
 - 33) Zhao J, Zhang M, Quan Z, Deng L, Li Y, He B. Systematic Influence of Circulating Bilirubin Levels on Osteoporosis. *Front Endocrinol* 2021; 12: 719920.
 - 34) Burgess S, Thompson SG. Multivariable Mendelian randomization: the use of pleiotropic genetic variants to estimate causal effects. *Am J Epidemiol* 2015; 181: 251-260.
 - 35) Yavorska OO, Burgess S. MendelianRandomization: an R package for performing Mendelian randomization analyses using summarized data. *Int J Epidemiol* 2017; 46: 1734-1739.
 - 36) Hemani G, Zheng J, Elsworth B, Wade KH, Haberland V, Baird D, Laurin C, Burgess S, Bowden J, Langdon R, Tan VY, Yarmolinsky J, Shihab HA, Timpson NJ, Evans DM, Relton C, Martin RM, Davey Smith G, Gaunt TR, Haycock PC. The MR-Base platform supports systematic causal inference across the human phenome. *eLife* 2018; 7: e34408.
 - 37) Verbanck M, Chen CY, Neale B, Do R. Detection of widespread horizontal pleiotropy in causal relationships inferred from Mendelian randomization between complex traits and diseases. *Nat Genet* 2018; 50: 693-698.
 - 38) Suh M, Choi-Kwon S, Kim JS. Sleep Disturbances at 3 Months after Cerebral Infarction. *Eur Neurol* 2016; 75: 75-81.
 - 39) Saetung S, Reutrakul S, Chailurkit LO, Rajatanavin R, Ongphiphadhanakul B, Nimitphong H. The Association between Daytime Napping Characteristics and Bone Mineral Density in Elderly Thai Women without Osteoporosis. *Sci Rep* 2018; 8: 10016.
 - 40) Wang L, Liu Q, Heizhati M, Yao X, Luo Q, Li N. Association between Excessive Daytime Sleepiness and Risk of Cardiovascular Disease and All-Cause Mortality: A Systematic Review and Meta-Analysis of Longitudinal Cohort Studies. *J Am Med Dir Assoc* 2020; 21:1979-1985.
 - 41) Endeshaw Y, Rice TB, Schwartz AV, Stone KL, Manini TM, Satterfield S, Cummings S, Harris T, Pahor M. Snoring, daytime sleepiness, and incident cardiovascular disease in the health, aging, and body composition study. *Sleep* 2013; 36: 1737-1745.
 - 42) Lin MT, Lin HH, Lee PL, Weng PH, Lee CC, Lai TC, Liu W, Chen CL. Beneficial effect of continuous positive airway pressure on lipid profiles in obstructive sleep apnea: a meta-analysis. *Sleep Breath* 2015; 19: 809-817.
 - 43) Vitarelli A, Terzano C, Saponara M, Gaudio C, Mangieri E, Capotosto L, Pergolini M, D'Orazio S, Continanza G, Cimino E. Assessment of Right Ventricular Function in Obstructive Sleep Apnea Syndrome and Effects of Continuous Positive Airway Pressure Therapy: A Pilot Study. *Can J Cardiol* 2015; 31: 823-831.
 - 44) Fu Y, Xia Y, Yi H, Xu H, Guan J, Yin S. Meta-analysis of all-cause and cardiovascular mortality in obstructive sleep apnea with or without continuous positive airway pressure treatment. *Sleep Breath* 2017; 21: 181-189.
 - 45) Lee CH, Ng WY, Hau W, Ho HH, Tai BC, Chan MY, Richards AM, Tan HC. Excessive daytime sleepiness is associated with longer culprit lesion and adverse outcomes in patients with coronary artery disease. *J Clin Sleep Med* 2013; 9: 1267-1272.
 - 46) Vgontzas AN, Chrousos GP. Sleep, the hypothalamic-pituitary-adrenal axis, and cytokines: multiple interactions and disturbances in sleep disorders. *Endocrin Metab Clin* 2002; 31: 15-36.
 - 47) Libby P. Inflammation and cardiovascular disease mechanisms. *Am J Clin Nutr* 2006; 83: 456s-460s.
 - 48) Libby P. Role of inflammation in atherosclerosis associated with rheumatoid arthritis. *Am J Med* 2008; 121: S21-31.
 - 49) Wisse BE. The inflammatory syndrome: the role of adipose tissue cytokines in metabolic disorders linked to obesity. *J Am Soc Nephrol* 2004; 15: 2792-2800.
 - 50) Spiegel K, Leproult R, Van Cauter E. Impact of sleep debt on metabolic and endocrine function. *Lancet* 1999; 354: 1435-1439.
 - 51) Spiegel K, Tasali E, Penev P, Van Cauter E. Brief communication: Sleep curtailment in healthy young men is associated with decreased leptin levels, elevated ghrelin levels, and increased hunger and appetite. *Ann Intern Med* 2004; 141 846-850.
 - 52) Tochikubo O, Ikeda A, Miyajima E, Ishii M. Effects of insufficient sleep on blood pressure monitored by a new multibiomedical recorder. *Hypertension* 1996; 27: 1318-1324.
 - 53) Kerkhofs M, Boudjeltia KZ, Stenuit P, Brohée D, Cauchie P, Vanhaeverbeek M. Sleep restriction increases blood neutrophils, total cholesterol and low density lipoprotein cholesterol in postmenopausal women: A preliminary study. *Maturitas* 2007; 56: 212-215.
 - 54) Ensrud KE, Crandall CJ. Osteoporosis. *Ann Intern Med* 2017; 167: Itc17-itc32.
 - 55) Kanis JA, Cooper C, Rizzoli R, Reginster JY. European guidance for the diagnosis and management of osteoporosis in postmenopausal women. *Osteoporosis Int* 2019; 30: 3-44.
 - 56) He B, Zhao J, Zhang M, Quan Z. Zoledronic acid and fracture risk: a meta-analysis of 12 randomized controlled trials. *Eur Rev Med Pharmacol Sci*. 2021; 25:1564-1573.
 - 57) He B, Zhao J, Zhang M, Jiang G, Tang K, Quan Z. Effect of Surgical Timing on the Refracture Rate after Percutaneous Vertebroplasty: A Retrospective Analysis of at Least 4-Year Follow-Up. *Biomed Res Int* 2021; 2021: 5503022.
 - 58) Chen G, Chen L, Wen J, Yao J, Li L, Lin L, Tang K, Huang H, Liang J, Lin W, Chen H, Li M, Gong X, Peng S, Lu J, Bi Y, Ning G. Associations between sleep duration, daytime nap duration, and osteoporosis vary by sex, menopause, and sleep quality. *J Clin Endocr Metab* 2014; 99: 2869-2877.