Check for updates

OPEN ACCESS

EDITED BY Marcello Pinti, University of Modena and Reggio Emilia, Italy

REVIEWED BY Owen Kelly, Sam Houston State University, United States Zhelong Liu, Huazhong University of Science and Technology, China

*CORRESPONDENCE Lei Yang 🔀 xjyanglei_616@sina.com

[†]These authors have contributed equally to this work

SPECIALTY SECTION

This article was submitted to Endocrinology of Aging, a section of the journal Frontiers in Endocrinology

RECEIVED 28 November 2022 ACCEPTED 23 February 2023 PUBLISHED 08 March 2023

CITATION

Chen S, Yan S, Aiheti N, Kuribanjiang K, Yao X, Wang Q, Zhou T and Yang L (2023) A bi-directional Mendelian randomization study of sarcopenia-related traits and type 2 diabetes mellitus. *Front. Endocrinol.* 14:1109800. doi: 10.3389/fendo.2023.1109800

COPYRIGHT

© 2023 Chen, Yan, Aiheti, Kuribanjiang, Yao, Wang, Zhou and Yang. This is an openaccess article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

A bi-directional Mendelian randomization study of sarcopenia-related traits and type 2 diabetes mellitus

Simin Chen^{1†}, Shikang Yan^{1†}, Nuerbiyamu Aiheti¹, Kaidiriyan Kuribanjiang¹, Xuemei Yao¹, Qian Wang¹, Tao Zhou¹ and Lei Yang^{1,2*}

¹Department of Epidemiology and Health Statistics, School of Public Health, Xinjiang Medical University, Urumqi, Xinjiang, China, ²State Key Laboratory of Causes and Prevention of High Morbidity in Central Asia jointly established by the Ministry and the Province Urumqi, Xinjiang, China

Background: Previous studies have reported an association between sarcopenia and type 2 diabetes mellitus (T2DM), but causation was prone to confounding factors. A more robust research approach is urgently required to investigate the causal relationship between sarcopenia and T2DM.

Methods: The bi-directional two-sample MR study was carried out in two stages: Sarcopenia-related traits were investigated as exposure while T2DM was investigated as an outcome in the first step, whereas the second step was reversed. The GWAS summary data for hand-grip strength (n = 256,523), appendicular lean mass (ALM, n = 450,243), and walking pace (n = 459,915) were obtained from the UK Biobank. T2DM data were obtained from one of the biggest case-control studies on diabetes (DIAGRAM; n = 180,834 cases and 492,191 controls), which was published in 2022. The inverse-variance weighted (IVW) approach was used to obtain MR estimates, and various sensitivity analysis was also performed.

Results: Low hand-grip strength had a potential causal relationship with an increased incidence of T2DM (OR = 1.109; 95% CI, 1.008–1.222; p = 0.0350). T2DM risk was reduced by increasing ALM and walking pace: A 1 kg/m² increase in ALM decreased the risk of T2DM by 10.2% (OR = 0.898; 95% CI, 0.830–0.952; p < 0.001). A 1 m/s increase in walking pace decreased the risk of T2DM by 90.0% (OR = 0.100; 95% CI, 0.053–0.186; p < 0.001). The relationship was bidirectional, with T2DM as a causative factor of sarcopenia-related traits (p < 0.05) except for ALM ($\beta = 0.018$; 95% CI, -0.008 to -0.044; p = 0.168).

Conclusions: Hand-grip strength and T2DM had a potential bidirectional causal relationship, as did walking pace and T2DM. We suggest that sarcopenia and T2DM may mutually have a significant causal effect on each other.

KEYWORDS

type 2 diabetes mellitus, sarcopenia, mendelian randomization, hand-grip strength, appendicular lean mass, walking pace, aging

1 Introduction

With the worldwide population aging, the incidence of sarcopenia and type 2 diabetes mellitus (T2DM) is rapidly increasing, which is related to a worse standard of living and higher mortality (1, 2). They have contributed to a serious global health problem, putting a heavy medical and economic burden on society (3). Sarcopenia and TDM are both age-related diseases with the same underlying pathophysiological mechanism (4, 5). People with sarcopenia have impaired glucose tolerance, and elevated blood insulin levels, and are at increased risk of T2DM. T2DM is characterized by insulin resistance and muscular metabolism, which may function as an accelerator for sarcopenia. Based on observational research, significant lines of evidence quantitatively revealed a tight association between sarcopenia and T2DM. The prevalence of sarcopenia among the elderly with T2DM ranges from 15.7-29.3 percent (6, 7). A cohort study of 6895 adults with an average age of 52 years reported that people with a muscle mass index in the lowest third of the population were twice as likely to get T2DM as those in the control group (8). These results make it difficult for researchers to determine which condition causes the other, particularly when confounders such as age, gender, and lifestyle are present (5). Assessing the causal relationship between sarcopenia and T2DM may help in the development of new techniques for the prevention, diagnosis, and treatment of T2DM.

Mendelian randomization (MR) investigates the causal effects associated with exposure and outcome by using genetic variants as instrumental variables (IVs), which avoids confounding variables and reverse causation (9, 10). To the authors' knowledge, no MR between sarcopenia and T2DM has been studied. Moreover, no randomized controlled study (RCT) specifically evaluated the bidirectional relationship. Therefore, we used a bi-directional twosample MR analysis to investigate the links between sarcopenia (measured as hand-grip strength, ALM, and walking pace) and T2DM.

2 Materials and methods

2.1 Study design

In this study, we adopted a bi-directional MR study design, employing two-Sample MR methodologies and different GWAS summary level data sets to explain the causation and causative direction between the sarcopenia-related traits (hand-grip strength, appendicular lean mass, walking pace) and the risk of T2DM and glycemic traits (fasting insulin, fasting blood glucose, glycated hemoglobin, and two-hour blood glucose challenge) in a European population. This study consisted of two stages. In first, we investigated whether sarcopenia-related traits were causally related to T2DM and glycemic characteristics. In the second, we evaluated whether genetic T2DM and glycemic characteristics were linked to sarcopenia-related traits. (Figure 1B).

2.2 Data source

The open GWAS database established by the MRC Integrated Epidemiology Unit (IEU) (https://gwasmrcieu.ac.uk/) provided the majority of the summary-level data used in this investigation (11). The datasets are largely public and may be downloaded using R and Python packages for accessing the application programming interface. Table 1 shows the phenotypic and consortium details.

2.2.1 Source of exposure data

The GWAS summary data for hand-grip strength were obtained from a meta-analysis of 256,523 people of European origin \geq 60 years of age and in 22 cohorts (12). Based on hand-grip strength using the European Working Group on Sarcopenia in Older People's (EWGSOP) definition of 30 kg for men and 20 kg for women, 46,596 (18.9%) of all participants reported muscular weakness. We collected GWAS summary data from the UK Biobank for appendicular lean mass (ALM) in 450,243 people and walking pace in 459,915 people who were included in the discovery cohort (13). The UK Biobank is an expansive biomedical database and a research resource (14). It includes in-depth information on the genetic makeup and health of about half a million people between 40 and 69 years of age who participated in the study in the UK. Genetic association studies (GWAS) on sarcopenia-related variables were undertaken in European populations using data from population-based cohorts.

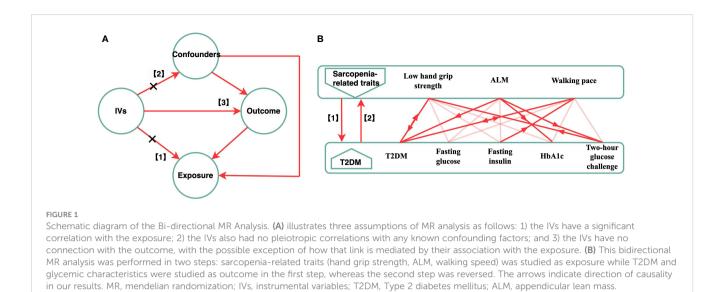
2.2.2 Source of outcome data

In this study, the instrument for the outcome (T2DM) was received from the DIAMANTE consortium and from the summary statistics of a GWAS meta-analysis. The GWAS meta-analysis comprised 22 GWAS that involved 180,834 patients with type 2 diabetes and 492,191 controls and five pedigree groups (51.1% of European ancestry). The GWAS database of European ancestors was released by Mahajan et al. in 2022 (15). Summary data for fasting glucose, fasting insulin, glycated hemoglobin (HbA1c), and 2-hour glucose challenge were collected from an online source (https:// gwas.mrcieu.ac.uk/) under the GWAS IDs ebi-a-GCST90002232 (fasting glucose, N = 200,622), ebi-a-GCST90002238 (fasting insulin, N =151), ieu-b-103 (HbA1C, N =46,368) and ebi-a-GCST000569 (two-hour glucose challenge, N = 15,234) (11).

2.3 IV selection and validation

Under the foundational premise of MR, specific SNPs for each exposure attribute were chosen as IVs. The IVs med three assumptions (Figure 1A). (1) The IVs had a significant relationship with the exposure. (2) The IVs had no pleiotropic correlations with any known confounding factors, and (3) had no association with the outcome, with the possible exception of how the link was mediated by their association with the exposure (16, 17). We obtained the IVs from the published GWAS using the clumping

Abbreviations: MR, Mendelian randomization; T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People; GWAS, genomewide association study; DIAGRAM, Diabetes Genetics Replication And Metaanalysis; IVW, inverse-variance weighted.



function in the Two-Sample MR package. Except for hand-grip strength, HbA1c, and two-hour glucose challenge ($p < 5 \times 10^{-6}$), all IV and exposure traits were significantly ($p < 5 \times 10^{-8}$) and independently [chain disequilibrium (LD) $r^2 < 0.01$) associated (18). Following the guidelines, we first chose several independent SNPs strongly related to each exposure variable before matching them in the outcome database. During the harmonization process, the harmonies data function from the two-sample MR package was used so the effect allele of each SNP could be matched with the corresponding allele of the exposure. To correct for multiple comparisons, the Bonferroni method was utilized (five outcomes). The associations with *p*-values < 0.003 (0.05/15) were regarded as statistically significant associations, and associations with *p*-values between 0.05 and 0.003 were considered as suggestive associations.

2.4 Analysis of horizontal pleiotropy and heterogeneity

Causal estimates and findings may be skewed due to the pleiotropy of SNPs in the IVW study (19). We used the

pleiotropy test function with the two-sample MR package to determine whether the SNPs were suitable IVs. If the pleiotropy was not significant (p > 0.05), then it was possible to use the IVs. We selected SNPs one by one on the Phenoscanner website (http://www.phenoscanner.medschl.cam.ac.uk/) if the *p*-value of the pleiotropy test was < 0.05 (20). Cochran's Q test was used to approximate heterogeneity among the chosen IVs. The fixed-effects model was used if the heterogeneity was negligible. Otherwise, the IVW with random effects approach was acceptable.

2.5 Analysis of Mendelian randomization

Mendelian randomization with genetic variations as the IVs allows investigation and assessment of the causal influence of an exposure on an outcome. First, we divided the per-allele log-odds ratio (or beta) of each variation in the outcome data by the logodds ratio (or beta) of the same variant in the exposure data to get the Wald ratio for each IV. After that, the inverse variance weighted (IVW) approach was used to arrive at an estimation of the connection between the exposures and the results. IVW

Phenotype	Consortium	Participants	Datatype	GWAS ID	Year of publication
Low hand grip strength (EWGSOP)	UKB	n = 48,596 cases and 207,927 controls	Binary	ebi-a-GCST90007526	2021
ALM	UKB	450,243	Continuous	ebi-a-GCST90000025	2020
Walking pace	UKB	459,915	Continuous	ukb-b-4711	2018
T2DM	DIAMANTE	n = 180,834 cases and 492,191 controls	Binary	NA	2022
Fasting glucose	MAGIC	200,622	Continuous	ebi-a-GCST90002232	2021
Fasting insulin	MAGIC	151,013	Continuous	ebi-a-GCST90002238	2021
HbA1c	MAGIC	46,368	Continuous	ieu-b-103	2010
Two-hour glucose challenge	MAGIC	15,234	Continuous	ebi-a-GCST000569	2010

TABLE 1 Phenotype source and description.

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

weights the Wald ratio of each SNP by its inverse variance and meta-analysis impact estimates using random or fixed effects. If the *p*-value of the Cochran Q test was < 0.05, we used the use of random-effect models; otherwise, we relied on fixed-effect models. To complement the outcome of IVW, MR-Egger and the weighted median approach were used (21).

2.6 Sensitivity analysis

To further guarantee the reliability of the MR causal impact estimate, we performed sensitivity analyses (22). To start, the MR-Egger approach was used to determine whether SNPs had horizontal pleiotropy (23). If the p-value of the intercept term was < 0.05, then the IVs may have pleiotropic effects. But there is no evidence of horizontal pleiotropy across the chosen IVs if the *p*-value of the intercept term is not < 0.05. Second, the standard MR testing outliers were identified with Mendelian Randomization Pleiotropy Residual Sum and Outlier (MR-PRESSO) test, which produced a robust estimate after outlier correction. A sensitivity analysis was performed to test for substantial distortions in the IVW causal estimate before and after MR-PRESSO adjustment. In this study, random-effects IVW was replicated by taking removing individual SNPs and then comparing the total analysis. The sensitivity of each SNP was inferred from how much the findings changed before and after it was removed

2.7 Reported results and software

MR findings were reported as odd ratios (ORs) with 95% confidence intervals (CIs) per standard deviation for dichotomous variables, and as beta values (β) with 95% CIs per standard deviation for continuous variables. Both were stated consistently as estimate values. A two-tailed *p*-value of 0.05 was the threshold of statistical significance. The statistical analysis was conducted with two-sample MR and MR-PRESSO packages of R version 4.0.0 (24).

3 Results

3.1 Stage 1: Effect of genetically predicted sarcopenia-related traits on T2DM and glycemic traits

Screening of the IVs found SNP beta, SE, and *p*-values strongly related to exposure. We found the beta, SE, and p-value corresponding to the SNP in the outcome database. Supplementary Tables 1-3 has more information about IVs. For low hand-grip strength, ALM, and walking pace, we found 59, 690, and 57 LD-independent ($r^2 < 0.001$) IVs that reached genome-wide significance ($p < 5 \times 10^{-8}$), except low hand-grip strength ($p < 5 \times 10^{-6}$). The horizontal pleiotropy test findings revealed that all IVs lacked horizontal pleiotropy (p > 0.05). One IV groups were not heterogeneous (p > 0.05). The IVW random-effect model for IV

with heterogeneity was used; otherwise, the IVW fixed-effect model was used. The level pleiotropy and heterogeneity test results are shown in Table 2.

The findings of this MR study are mostly concerned with IVW analysis. Our findings reveal that sarcopenia-related features were all linked to T2DM. Low hand-grip strength had a potential causal relationship with an increased incidence of T2DM (OR = 1.109; 95% CI: 1.008–1.222; p = 0.035). T2DM risk was reduced by increasing ALM and walking pace. A 1 kg² increase in ALM decreased the risk of T2DM by 10.2% (OR = 0.898; 95% CI: 0.830-0.952; p < 0.001), but a 1 m/s increase in walking pace decreased the odds of T2DM by 90.0% (OR = 0.100; 95% CI: 0.053-0.186 p < 0.001). ALM also had a potential causal relationship with HbA1C (OR = -0.006; 95% CI: -0.029 to -0.017; p = 0.012) and 2-hour glucose post-challenge (OR = -0.292; 95% CI: -0.401 to -0.184; p = 0.001). Table 3 shows the findings of the thorough MR analysis. According to the intercept of the MR-Egger regression, there was no evidence of horizontal pleiotropy between the exposures and the outcomes (p > 0.05). As expected, IVW findings were consistent after MR-PRESSO adjustment. The sensitivity analysis findings are shown in Table 4.

3.2 Stage 2: Genetically predicted T2DM and glycemic traits influence sarcopenia-related traits

We obtained beta, standard error, and p-values of SNPs significantly associated with exposure using the instrumental variables (IV) screening. In the outcomes database, we discovered the beta value, standard error, and *p*-values that correlated with the SNPs. For fasting glucose, fasting insulin, and T2DM, we found 66, 38, and 187 linkage disequilibrium (LD)-independent ($r^2 < 0.001$) IVs that reached genome-wide significance level ($p < 5 \times 10^{-8}$). For HbA1c, 2-hour blood glucose challenge, we found 31 and 11 linkage disequilibrium (LD)-independent ($r^2 < 0.001$) IVs that reached genome-wide significance level ($p < 5 \times 10^{-6}$). Supplementary Tables 4-8 has detailed information about IVs. The horizontal pleiotropy test confirmed the of absence horizontal pleiotropy (p > 0.05). There were one IV groups that were not heterogeneous (p > 0.05). The IVW random-effect model was used if heterogeneity was present. Otherwise, the IVW fixed-effect model was used. The level pleiotropy and heterogeneity test results are shown in Table 5.

This MR analysis mostly refers to IVW. It was shown that T2DM was a potential pathogenic factor in sarcopenia-related characteristics (p < 0.05), except for ALM ($\beta = 0.018$; 95% CI: -0.008 to -0.044; p = 0.168). There was a potential positive correlation between fasting insulin and walking pace ($\beta = 0.062$; 95% CI: -0.0001 to -0.125; p = 0.050). There was no correlation between HbA1c and sarcopenia-related characteristics (p > 0.05). Low hand-grip strength (OR = 0.058; 95% CI: 0.005-0.111; p = 0.031) was the only sarcopenia-related characteristic causally linked with a 2-hour glucose test. A gene predictive of 2-hour glucose test results after the challenge would increase the of having a weak hand-grip strength. Table 6 shows the findings of the MR

F	0.1	Ne of Los	Heterogenei	ty test	Pleiotropy_test
Exposures	Outcomes	No.of Ivs	Cochran's Q	Р	Р
	T2DM	52	225.3713	<0.001	0.377
	Fasting glucose	56	108.4448	<0.001	0.244
Low hand grip strength (EWGSOP)	Fasting insulin	56	95.9703	< 0.001	0.661
	HbA1c	30	38.4647	0.112	0.675
	Two-hour glucose challenge	33	46.5341	0.047	0.305
	T2DM	552	3214.1050	< 0.001	0.289
	Fasting insulin	372	981.8712	< 0.001	0.424
ALM	Fasting glucose	617	1635.6020	<0.001	0.998
	HbA1c	362	474.1836	<0.001	0.903
	Two-hour glucose challenge	401	453.5987	0.033	0.886
	T2DM	56	496.6391	< 0.001	0.284
	Fasting glucose	56	112.1423	<0.001	0.962
Walking pace	Fasting insulin	56	125.2184	<0.001	0.268
	HbA1c	43	59.7183	0.037	0.745
	Two-hour glucose challenge	46	67.2943	0.017	0.473

TABLE 2 Association sarcopenia-related traits with T2DM and glycaemic traits using heterogeneity test and pleiotropy test.

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

TABLE 3 Association sarcopenia-related traits with T2DM and glycaemic traits using Mendelian randomization.

Evenosuros	Outcomes	No.of	MR Egger		Weighted median		IVW(random-effect model)	
Exposures	Outcomes	lvs	Estimate (95% Cl)	Р	Estimate (95% Cl)	Р	Estimate (95% Cl)	Р
	T2DM	52	1.285(0.917,1.802)	0.151	1.090(1.006,1.180)	0.032	1.109(1.008,1.222)	0.035
	Fasting glucose	56	-0.030 (-0.102,0.042)	0.413	0.001(-0.020,0.021)	0.945	0.011(-0.007,0.030)	0.224
Low hand grip strength (EWGSOP)	Fasting insulin	56	-0.004 (-0.081,0.073)	0.914	0.030(0.003,0.050)	0.027	0.012(-0.007,0.032)	0.213
()	HbA1c	30	-0.020 (-0.213,0.172)	0.836	0.012(-0.037,0.060)	0.633	0.020(-0.017,0.058)	0.226
	Two-hour glucose challenge	33	-0.319 (-1.280,0.642)	0.520	0.227(-0.034.0.489)	0.077	0.181(-0.021,0.383)	0.080
	T2DM	552	0.823(0.704,0.964)	0.016	0.843(0.797,0.893)	< 0.001	0.889(0.830,0.952)	< 0.001
	Fasting glucose	617	0.004(-0.03,0.03)	0.800	-0.005(-0.02,0.01)	0.526	0.004(-0.009,0.017)	0.550
ALM	Fasting insulin	372	-0.041 (-0.110,0.032)	0.263	-0.019 (-0.051,0.0004)	0.054	-0.011 (-0.040,0.010)	0.264
	HbA1c	362	-0.003 (-0.059,0.053)	0.921	-0.019 (-0.052,0.015)	0.266	-0.006 (-0.029,0.017)	0.012
	Two-hour glucose challenge	401	-0.311(-0.591,- 0.031)	0.030	-0.292(-0.459,- 0.125)	<0.001	-0.292(-0.401,- 0.184)	<0.001
	T2DM	56	0.403(0.030,5.440)	0.496	0.152(0.102,0.226)	< 0.001	0.100(0.053,0.186)	< 0.001
Walking pace	Fasting glucose	56	0.004(-0.387,0.379)	0.989	0.008(-0.087,0.103)	0.864	-0.005 (-0.090,0.080)	0.906

(Continued)

TABLE 3 Continued

Exposures	Outcomes	No.of Ivs	MR Egger		Weighted median		IVW(random-effect model)	
	Outcomes		Estimate (95% Cl)	Р	Estimate (95% Cl)	Р	Estimate (95% Cl)	Р
	Fasting insulin	56	0.193(-0.268,0.655)	0.415	-0.026 (-0.144,0.092)	0.641	-0.063 (-0.166,0.039)	0.224
	HbA1c	43	-0.020 (-0.708,0.668)	0.955	-0.119 (-0.313,0.075)	0.230	-0.128 (-0.285,0.030)	0.113
	Two-hour glucose challenge	46	-1.157 (-5.000,2.682)	0.558	0.402(-0.618,1.423)	0.440	0.227(-0.613,1.066)	0.597

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

TABLE 4 Association sarcopenia-related traits with T2DM and glycaemic traits using MR-PRESSO and MR Egger.

		MR-PRESSO					MR Egger		
Exposures	Outcomes	No.of Ivs	P IVS OUTUERS		lvs outliers	No.of Ivs	Intercept	Р	
	T2DM	50	1.111 (1.032,1.190)	0.007	rs10952289,rs34415150, rs8061064,rs823130		-0.008	0.322	
	Fasting glucose	59	0.013 (-0.007,0.031)	0.239	NA		0.003	0.234	
Low hand grip strength	Fasting insulin	58	0.013 (-0.007,0.031)	0.244	rs7740188	59	0.001	0.569	
(EWGSOP)	HbA1c	36	0.021 (-0.022,0.049)	0.335	NA	36	0.002	0.669	
	Two-hour glucose challenge	35	0.192 (-0.001,0.380)	0.060	NA		0.024	0.257	
	T2DM	511	0.872 (0.831,0.900)	<0.001	rs10225945,rs11260035,rs11605297,rs12074850,rs12150907, rs1260326,rs12724708,rs12909863,rs13391980,rs17205463, rs173135,rs2071450,rs2112617,rs2142331,rs2194411, rs2236406,rs2287821,rs2289629,rs28379706,rs28592876, rs28678024,rs2871960,rs34522021,rs3764002,rs40270, rs4274112,rs4752689,rs4788218,rs4985445,rs55872725, rs6142059,rs610694,rs61919240,rs62106258,rs62370472, rs6860245,rs6931421,rs6977416,rs7020491,rs7144307, rs73052033,rs73125634,rs7418410,rs7563362,rs7610055, rs7701233,rs7768973,rs7816345,rs7826059,rs9388490, s951366,rs9634212,rs9647379,rs9669278	565	0.002	0.257	
ALM	Fasting glucose	596	0.001 (-0.009,0.010)	0.907	rs12188208,rs12188208,rs17205463,rs17681189,rs2025609, rs2142331,rs2237485,rs2871960,rs34517439,rs40270, rs4752829,rs4852257,rs488621,rs57513571,rs7107356, rs7328187,rs73384223,rs7448554,rs9375188,rs9391254, rs9669278	617	6.60E-05	0.842	
	Fasting insulin	377	-0.021 (-0.042,0.001)	0.042	rs11098677,rs11098677,rs12724708,rs13391980,rs2142331, rs2237485,rs2287821,rs310796,rs36012032,rs40270, rs4865956,rs7229520,rs73006226,rs7328187,rs7816345	392	0.001	0.323	
	HbA1c	359	0.003 (-0.017,0.023)	0.747	rs12074850,rs2923411,rs3184504	362	-0.0002	0.786	
	Two-hour glucose challenge	398	-0.252 (-0.351,- 0.152)	<0.001	rs1260326,rs17205463,rs7610055	401	0.0004	0.896	
Walking pace	T2DM	52	0.133 (0.091,0.200)	<0.001	rs113825410,rs11732213,rs1592,rs4643373,rs62048402	57	-0.013	0.274	

(Continued)

TABLE 4 Continued

	Outcomes			MR Egger				
Exposures		No.of Ivs	Estimate (95%Cl)	Р	lvs outliers	No.of Ivs	Intercept	Р
	Fasting glucose	54	0.031 (-0.040,0.110)	0.414	rs113825410,rs7896518	56	-2.08E-06	0.999
	Fasting insulin 54 -0.071 (-0.162,0.021)		0.147	rs11039324,rs28519617	56	-0.003	0.303	
	HbA1c	43	-0.153 (-0.301,- 0.002)	0.053	NA	43	-0.001	0.821
	Two-hour glucose challenge	46	0.231 (-0.613,1.071)	0.600	NA	46	0.013	0.473

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

analysis. No horizontal pleiotropy (p > 0.05) was found in the MR-Egger regression of exposures and outcomes, as shown by the intercept. As expected, the IVW findings were consistent after MR-PRESSO adjustment. The sensitivity analysis findings are shown in Table 7.

4 Discussion

The bi-directional two-sample MR analysis in this study was performed using summary-level data. We examined the potential causative connections between sarcopenia-related characteristics (hand-grip strength, ALM, and walking pace) and T2DM and glycemic features. Major findings were: (1) Sarcopenia-related variables predicted genetically were all causally associated with T2DM; (2) There was a potential causal relationship between genetically predicted T2DM and sarcopenia-related characteristics but not expected ALM in the other direction. (3) The results showed there was potential bi-directional causation between sarcopenia and T2DM to a greater extent.

In observational studies, hand-grip strength was shown to have a bidirectional relationship with T2DM. A previous English longitudinal study found that for each unit gain in hand-grip strength, T2DM risk was reduced by 2% (HR = 0.98; 95% CI: 0.96-0.99) (25). In another cohort study, T2DM was an independent predictor of weak hand-grip strength (26). A TSMR

	Accesietien	T2DM and alw		ithe environmentio	unlated tunits	union la stava e		I m la i a hua muu ha ah
I ADLE D	Association	IZDM and gly	caemic traits w	ith sarcopenia	-related traits	using neteroge	eneity test and	pleiotropy test.

Fundation	Outcomes	No.of Ivs	Heterogeneity 1	test	Pleiotropy_test
Exposures	Outcomes	INO.OI IVS	Cochran's Q	Р	Р
	Low hand grip strength (EWGSOP)	180	367.2484	< 0.001	0.287
T2DM	ALM	181	5109.0300	< 0.001	0.157
	Walking pace	163	395.5066	< 0.001	0.139
	Low hand grip strength (EWGSOP)	64	106.3683	< 0.001	0.886
Fasting glucose	ALM	64	1120.4190	<0.001	0.648
	Walking pace	40	113.3818	<0.001	0.141
	Low hand grip strength (EWGSOP)	38	80.3508	< 0.001	0.691
Fasting insulin	ALM	38	2377.4080	<0.001	0.326
	Walking pace	37	164.1926	<0.001	0.523
	Low hand grip strength (EWGSOP)	28	43.4848	0.023	0.061
HbA1c	ALM	29	368.1774	< 0.001	0.073
	Walking pace	29	60.4644	<0.001	0.970
	Low hand grip strength (EWGSOP)	7	8.6807	0.192	0.086
Two-hour glucose challenge	ALM	7	288.4325	< 0.001	0.406
	Walking pace	7	25.4576	<0.001	0.528

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

Evenosuros	Quitcomes	No.of	MR Egger		Weighted me	dian	IVW(random-effect model)	
Exposures	Outcomes	lvs	OR(95%CI)	Р	Estimate(95% Cl)	Р	Estimate(95% Cl)	Р
	Low hand grip strength (EWGSOP)	180	1.073 (1.008,1.142)	0.030	1.043(1.004,1.082)	0.028	1.043(1.012.1.069)	0.005
T2DM	ALM	181	-0.020 (-0.080,0.039)	0.504	0.009 (-0.003,0.021)	0.117	0.018 (-0.008,0.044)	0.168
	Walking pace	182	0.002 (-0.013,0.017)	0.799	-0.002 (-0.008,0.004)	0.504	-0.013(-0.020,- 0.006)	< 0.001
	Low hand grip strength (EWGSOP)	64	-0.033 (-0.251,0.18)4	0.764	-0.102 (-0.255,0.052)	0.194	-0.017 (-0.136,0.101)	0.769
Fasting glucose	ALM	64	0.073 (-0.109,0.254)	0.436	0.062(0.017,0.106)	0.006	0.037 (-0.062,0.136)	0.463
	Walking pace	40	-0.040 (-0.101,0.031)	0.290	-0.022 (-0.051,0.019)	0.386	0.011 (-0.013,0.045)	0.609
	Low hand grip strength (EWGSOP)	38	0.026 (-0.728,0.781)	0.946	-0.141 (-0.421,0.139)	0.324	-0.119 (-0.367,0.128)	0.345
Fasting insulin	ALM	38	0.773 (-0.248,1.794)	0.147	0.243(0.089,0.395)	< 0.001	0.283 (-0.058.0.625)	0.104
	Walking pace	37	0.0002 (-0.199.0.199)	0.998	0.044 (-0.009,0.098)	0.106	0.062 (-0.0001,0.125)	0.050
	Low hand grip strength (EWGSOP)	29	-0.324 (-0.651,0.003)	0.063	-0.076 (-0.247,0.095)	0.385	0.046 (-0.109,0.201)	0.385
HbA1c	ALM	29	0.141 (-0.084,0.366)	0.230	0.064(0.015,0.113)	0.010	-0.051 (-0.154,0.052)	0.329
	Walking pace	29	-0.011 (-0.076,0.054)	0.743	-0.013 (-0.046,0.020)	0.453	-0.012 (-0.040,0.016)	0.394
	Low hand grip strength (EWGSOP)	7	0.298 (0.071,0.524)	0.049	0.084(0.013,0.154)	0.020	0.058(0.005,0.122)	0.031
Two-hour glucose challenge	ALM	7	0.110 (-0.298,0.517)	0.620	0.007 (-0.014,0.029)	0.514	-0.073 (-0.167,0.021)	0.126
	Walking pace	7	0.034 (-0.050,0.119)	0.458	-0.002 (-0.015,0.011)	0.739	0.006 (-0.013,0.025)	0.517

TABLE 6 Association T2DM and glycaemic traits with sarcopenia-related traits using Mendelian randomization.

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

investigation was conducted to evaluate the causal relationship and discovered a bi-directional causal relationship between hand-grip strength and T2DM (27). Our results were consistent with these previous studies and provided evidence of a potential bi-directional causality between hand-grip strength and T2DM. In a long-term retrospective analysis of 159 older women, researchers found that patients with low ALM had a 3.81 higher incidence of IFG/T2DM than those with normal ALM (OR = 3.81; 95% CI: 1.09-9.80) (28). Our MR findings follow prior research that ALM decreased the risk of T2DM. But in the other direction, T2DM was not causally associated with ALM. Notably, the major glucose metabolic reservoir of T2DM was in muscle mass, suggesting that our results only partially demonstrate a causal relationship between ALM and diabetes. More studies with better MR methods and data

are needed in the future to validate the causal relationship between ALM and diabetes. Prospective studies have shown that a slower walking pace was associated with an increased risk of T2DM. Men and women who walked slowly were shown to have an increased risk of incident T2DM (29). A Baltimore Longitudinal Study of Aging found that T2DM was a significant predictor of slower gait speed in the elderly (30). Our MR findings, in contrast to those of the traditional studies, included a strong bi-directional causative connection between walking pace and T2DM.

Many previous studies had demonstrated the positive association between sarcopenia and T2DM, and the interaction between the two conditions seems to tilt more toward a bidirectional causation correlation owing to the complexity and proximity of their relationship. Landin et al. described sarcopenia

TABLE 7 Association T2DM and glycaemic traits with sarcopenia-related traits using MR-PRESSO and MR Egger.

					MR-PRESSO	MR Egger			
Exposures	Outcomes	No.of Ivs	Estimate (95%Cl)	Р	lvs outliers	No.of Ivs	Intercept	Р	
	Low hand grip strength (EWGSOP)	182	1.031 (1.012,1.060)	0.002	rs1117610,rs601945,rs9379084	185	-0.002	0.411	
T2DM	ALM	118	0.014 (0.004,0.025)	0.007	rs1007090,rs10097617,rs10419627,rs11063069,rs1117610, rs115505614,rs11699802,rs11708067,rs11709077,rs11759026, rs11856307,rs12325539,rs1260326,rs12910361,rs12920022, rs13022337,rs13385171,rs13389219,rs1359790,rs1426371, rs1517037,rs1562396,rs1665901,rs1705263,rs17744783, rs2023681,rs2277536,rs2383205,rs2820441,rs2925979, rs2972144,rs3094682,rs34341,rs348330,rs35777422, rs35895680,rs3768321,rs3786900,rs3798519,rs459193, rs4686471,rs4709746,rs474513,rs4804833,rs490689, rs4925109,rs4977213,rs505922,rs55653563,rs55872725, rs58432198,rs58642235,rs601945,rs62107261,rs62271373, rs6937438,rs72802358,rs739846,rs7633675,rs7667864, rs76895963,rs7732130,rs77464186,rs8097210,rs8192675, rs878521,rs9368222,rs9379084	186	0.003	0.132	
	Walking pace	159	-0.007 (-0.012,- 0.002)	0.010	rs13330951,rs34990153,rs601945, rs9873519		-0.001	0.160	
	Low hand grip strength (EWGSOP)	66	-0.021 (-0.142,0.092)	0.695	NA		5.89E-05	0.981	
Fasting glucose	ALM	44	-0.010 (-0.071,0.041)	0.528	rs10838693,rs11603349,rs11708067,rs1260326,rs12888855, rs157512,rs16851397,rs16913693,rs17265513,rs174583, rs1820176,rs189548,rs2238435,rs3842753,rs6113722, rs6489811,rs7163757,rs77981966,rs78132593,rs878521		-0.001	0.648	
	Walking pace	42	-0.002 (-0.032,0.028)	0.885	rs10974438,rs11603349,rs4862423,rs6662924	42	0.001	0.240	
	Low hand grip strength (EWGSOP)	36	-0.133 (-0.341,0.081)	0.238	rs1351394,rs6905288	68	-0.002	0.691	
Fasting insulin	ALM	38	0.279 (-0.061,0.632)	0.112	NA	38	-0.009	0.326	
	Walking pace	32	0.080 (0.031,0.122)	0.003	rs10865959,rs11708067,rs1351394,rs3775380,rs5017305	37	0.001	0.523	
	Low hand grip strength (EWGSOP)	30	0.022 (-0.121,0.150)	0.822	NA	30	0.009	0.051	
HbA1c	ALM	19	0.021 (-0.033,0.232)	0.502	rs12819124,rs16926246,rs17533945,rs1800562,rs2723517, rs4844390,rs552976,rs7355559,rs7805661,rs837763	29	-0.006	0.052	
	Walking pace	28	-0.002 (-0.031,0.030)	0.884	rs837763	29	0.0003	0.722	
Two-hour	Low hand grip strength (EWGSOP)	7	0.061 (-0.005,0.122)	0.122	NA	7	-0.029	0.086	
glucose challenge	ALM	7	-0.070 (-0.171,0.022)	0.177	NA	7	-0.022	0.406	
	Walking pace	6	-0.005 (-0.010,0.003)	0.28	rs10423928	7	-0.003	0.528	

T2DM, Type 2 diabetes mellitus; ALM, appendicular lean mass; HbA1c, glycated hemoglobin; EWGSOP, European Working Group on Sarcopenia in Older People.

10.3389/fendo.2023.1109800

and T2DM as two sides of the same coin when discussing the correlation between the two conditions (31). Through the use of transcriptome analysis, Huang et al. discovered 15 shared genes associated with both sarcopenia and T2DM, in addition to multiple shared pathways between the two disorders (32). A meta-analysis of community-dwelling Asian populations found that T2DM patients had a considerably greater incidence of sarcopenia than nondiabetics (33). In another recent meta-analysis of observational studies, T2DM was related to an elevated risk of sarcopenia (34). Our bidirectional MR study further complements previous studies and provides evidence for a potential causal relationship between sarcopenia and T2DM. Sarcopenia and TDM are both age-related diseases with the same underlying pathophysiological mechanism (4, 5, 35). T2DM is often associated with insulin resistance, increased inflammation, accumulation of advanced glycation end products, and elevated oxidative stress (36). These pathways would result in activated inflammation, mitochondrial and vascular dysfunction, and protein metabolism deficiencies, all of which would be harmful to muscle health, including muscle mass, muscular strength, and muscle quality (32). Because of reduced insulin sensitivity, the anabolic activity of insulin in skeletal muscle may be gradually lost in T2DM. Furthermore, poor insulin action may result in increased protein breakdown and reduced protein synthesis, resulting in a loss of muscle mass and strength (37, 38). As a result, T2DM patients are at a higher risk of sarcopenia (39, 40). On the other hand, older people may be at a higher risk for developing T2DM due to the effects of sarcopenia. Sarcopenia is a condition defined by decreased muscular mass, low muscle strength, or poor athletic performance (41), and may contribute to impaired glucose clearance, increased inflammation, and reduced metabolic rate and physical activity. Skeletal muscle is essential for glucose clearance, accounting for more than 80% of postprandial glucose uptake. As a result, sarcopenia alters glucose disposal by lowering muscle mass and increasing localized inflammation, which may contribute to the development and progression of T2DM (42, 43).

Our research has several advantages. First, this was the first bidirectional MR study to investigate the causal association between three sarcopenia-related traits (EWGSOP) and T2DM, especially evaluating the causal relationship between walking pace and T2DM, which had been neglected in previous studies. Second, the database for this study was the most recent and reliable. The GWAS summary data for low hand-grip strength was defined by EWGSOP; T2DM data originated from one of the largest diabetes case-control studies, which was published in 2022. Third, different MR analysis methods were used to confirm the correctness and validity of the results, and sensitivity analyses such as MR-PRESSO were performed to obtain consistent estimations of the causal impact sizes of MR. Finally, we investigated the link between walking pace and diabetes, which had been neglected in earlier research. Still, the study's weaknesses should be mentioned. To begin, this study was carried out using summary-level data, which restricted our capacity to carry out subgroup analyses, such as those regarding low hand-grip strength and ALM according to gender. Besides, we applied ALM (appendicular lean mass) rather than ASM (appendicular skeletal muscle mass) to evaluate muscle mass, which may be inaccurate due to bias from other non-fat soft tissue components such as lungs, kidneys, and other internal organs. In addition, our research population was exclusive of European descent, so our results may not apply to other racial or ethnic groups. At last, residual bias cannot be avoided, as it is a recognized shortcoming of the MR technique, even when the pleiotropy test and MR-PRESSO procedures are used to prevent confounding by pleiotropy.

In conclusion, sarcopenia and T2DM may mutually have a potential causal influence on each other. Our study suggests that sarcopenia-related traits (hand-grip strength, ALM, walking pace) may benefit T2DM. Instead, T2DM may be linked to decreased hand-grip strength and walking pace. This study provides solid evidence that hand-grip strength, ALM and walking pace are possible predictors of T2DM in middle-aged and elderly people.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Materials. Further inquiries can be directed to the corresponding author.

Author contributions

Study design: LY, SC. Manuscript writing: SC, SY. Statistical analysis and data interpretation: SY, NA, KK. Critical revision of the manuscript: LY, XY. Literature Search: SC, SY, NA, KK, QW, TZ. All authors contributed to the article and approved the submitted version.

Funding

This work was supported by the the 14th Five-Year Plan Key Discipline of Public Health and Preventive Medicine in Xiniiang Uvgur Autonomous Region, State Key Laboratory Project of Xinjiang Medical University (SKL-HIDCA-2020-ER4) and National Key Research and Development Program of China (No.2017YFC0907200; 2017YFC0907203).

Acknowledgments

Thanks to everyone who assisted with this project. And we would like to express our gratitude to all the GWAS consortia and the UKB for supplying us with the sarcopenia-related traits and T2D and glycemic traits GWAS summary data. We also thank International Science Editing (http://www.internationalscienceediting.com) for editing this manuscript.

Conflict of interest

We declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated

References

1. Petermann-Rocha F, Balntzi V, Gray SR, Lara J, Ho FK, Pell JP, et al. Global prevalence of sarcopenia and severe sarcopenia: A systematic review and meta-analysis. J Cachexia Sarcopenia Muscle (2022) 13(1):86–99. doi: 10.1002/jcsm.12783

2. Cho NH, Shaw JE, Karuranga S, Huang Y, Fernandes J, Ohlrogge AW, et al. Idf diabetes atlas: Global estimates of diabetes prevalence for 2017 and projections for 2045. *Diabetes Res Clin Pract* (2018) 138:271–81. doi: 10.1016/j.diabres.2018.02.023

3. Norman K, Otten L. Financial impact of sarcopenia or low muscle mass - a short review. *Clin Nutr* (2019) 38(4):1489–95. doi: 10.1016/j.clnu.2018.09.026

 Sherwani SI, Khan HA, Ekhzaimy A, Masood A, Sakharkar MK. Significance of Hba1c test in diagnosis and prognosis of diabetic patients. *biomark Insights* (2016) 11:95–104. doi: 10.4137/bmi.S38440

5. Shou J, Chen PJ, Xiao WH. Mechanism of increased risk of insulin resistance in aging skeletal muscle. *Diabetol Metab Syndr* (2020) 12(1):14. doi: 10.1186/s13098-020-0523-x

6. Liccini AP, Malmstrom TK. Frailty and sarcopenia as predictors of adverse health outcomes in persons with diabetes mellitus. *J Am Med Directors Assoc* (2016) 17 (9):846–51. doi: 10.1016/j.jamda.2016.07.007

7. Fung FY, Koh YLE, Malhotra R, Ostbye T, Lee PY, Ghazali SS, et al. Prevalence of and factors associated with sarcopenia among multi-ethnic ambulatory older asians with type 2 diabetes mellitus in a primary care setting. *BMC Geriatrics* (2019) 19(1):122. doi: 10.1186/s12877-019-1137-8

8. Son JW, Lee SS, Kim SR, Yoo SJ, Cha BY, Son HY, et al. Low muscle mass and risk of type 2 diabetes in middle-aged and older adults: Findings from the koges. *Diabetologia* (2017) 60(5):865–72. doi: 10.1007/s00125-016-4196-9

9. Davey Smith G, Hemani G. Mendelian randomization: Genetic anchors for causal inference in epidemiological studies. *Hum Mol Genet* (2014) 23:R89–98. doi: 10.1093/hmg/ddu328

10. Burgess S, Small DS, Thompson SG. A review of instrumental variable estimators for mendelian randomization. *Stat Methods Med Res* (2017) 26(5):2333–55. doi: 10.1177/0962280215597579

11. Hemani G, Zhengn J, Elsworth B, Wade KH, Haberland V, Baird D, et al. The Mr-base platform supports systematic causal inference across the human phenome. *Elife* (2018) 7:e34408. doi: 10.7554/eLife.34408

12. Jones G, Trajanoska K, Santanasto AJ, Stringa N, Kuo CL, Atkins JL, et al. Genome-wide meta-analysis of muscle weakness identifies 15 susceptibility loci in older men and women. *Nat Commun* (2021) 12(1):654. doi: 10.1038/s41467-021-20918-w

13. Pei YF, Liu YZ, Yang XL, Zhang H, Feng GJ, Wei XT, et al. The genetic architecture of appendicular lean mass characterized by association analysis in the uk biobank study. *Commun Biol* (2020) 3(1):608. doi: 10.1038/s42003-020-01334-0

14. Sudlow C, Gallacher J, Allen N, Beral V, Burton P, Danesh J, et al. Uk biobank: An open access resource for identifying the causes of a wide range of complex diseases of middle and old age. *PloS Med* (2015) 12(3):e1001779. doi: 10.1371/journal.pmed.1001779

15. Mahajan A, Spracklen CN, Zhang WH, Ng MCY, Petty LE, Kitajima H, et al. Multiancestry genetic study of type 2 diabetes highlights the power of diverse populations for discovery and translation. *Nat Genet* (2022) 54(5):560. doi: 10.1038/s41588-022-01058-3

16. Emdin CA, Khera AV, Kathiresan S. Mendelian randomization. Jama-Journal Am Med Assoc (2017) 318(19):1925–6. doi: 10.1001/jama.2017.17219

17. Zhao Q, Wang J, Hemani G, Bowden J, Small DS. Statistical inference in twosample summary-data mendelian randomization using robust adjusted profile score. *Ann Stat* (2020) 48(3):1742–69. doi: 10.1214/19-aos1866

18. Burgess S, Butterworth A, Thompson SG. Mendelian randomization analysis with multiple genetic variants using summarized data. *Genet Epidemiol* (2013) 37 (7):658–65. doi: 10.1002/gepi.21758

19. Hemani G, Bowden J, Smith GD. Evaluating the potential role of pleiotropy in mendelian randomization studies. *Hum Mol Genet* (2018) 27(R2):R195–208. doi: 10.1093/hmg/ddy163

organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2023. 1109800/full#supplementary-material

20. Staley JR, Blackshaw J, Kamat MA, Ellis S, Surendran P, Sun BB, et al. Phenoscanner: A database of human genotype-phenotype associations. *Bioinformatics* (2016) 32(20):3207–9. doi: 10.1093/bioinformatics/btw373

21. Bowden J, Smith GD, Haycock PC, Burgess S. Consistent estimation in mendelian randomization with some invalid instruments using a weighted median estimator. *Genet Epidemiol* (2016) 40(4):304–14. doi: 10.1002/gepi.21965

22. Burgess S, Bowden J, Fall T, Ingelsson E, Thompson SG. Sensitivity analyses for robust causal inference from mendelian randomization analyses with multiple genetic variants. *Epidemiology* (2017) 28(1):30–42. doi: 10.1097/ede.00000000000559

23. Bowden J, Smith GD, Burgess S. Mendelian randomization with invalid instruments: Effect estimation and bias detection through egger regression. *Int J Epidemiol* (2015) 44(2):512–25. doi: 10.1093/ije/dyv080

24. Yavorska OO, Burgess S. Mendelianrandomization: An r package for performing mendelian randomization analyses using summarized data. *Int J Epidemiol* (2017) 46(6):1734–9. doi: 10.1093/ije/dyx034

25. Cuthbertson DJ, Bell JA, Ng SY, Kemp GJ, Kivimaki M, Hamer M. Dynapenic obesity and the risk of incident type 2 diabetes: The English longitudinal study of ageing. *Diabetic Med* (2016) 33(8):1052–9. doi: 10.1111/dme.12991

26. Lin Y-L, Chen S-Y, Lai Y-H, Wang C-H, Kuo C-H, Liou H-H, et al. Angiotensin ii receptor blockade is associated with preserved muscle strength in chronic hemodialysis patients. *BMC Nephrol* (2019) 20(1):54. doi: 10.1186/s12882-019-1223-3

27. Yeung CHC, Yeung SLA, Fong SSM, Schooling CM. Lean mass, grip strength and risk of type 2 diabetes: A bi-directional mendelian randomisation study. *Diabetologia* (2019) 62(5):789–99. doi: 10.1007/s00125-019-4826-0

28. Buscemi C, Ferro Y, Pujia R, Mazza E, Boragina G, Sciacqua A, et al. Sarcopenia and appendicular muscle mass as predictors of impaired fasting Glucose/Type 2 diabetes in elderly women. *Nutrients* (2021) 13(6):1909. doi: 10.3390/nu13061909

29. Boonpor J, Ho FK, Gray SR, Celis-Morales CA. Association of self-reported walking pace with type 2 diabetes incidence in the uk biobank prospective cohort study. *Mayo Clinic Proc* (2022) 97(9):1631–40. doi: 10.1016/j.mayocp.2022.02.028

30. Gonzalez-Freire M, Moaddel R, Sun K, Fabbri E, Zhang P, Khadeer M, et al. Targeted metabolomics shows low plasma lysophosphatidylcholine 18:2 predicts greater decline of gait speed in older adults: The Baltimore longitudinal study of aging. *Journals Gerontol Ser a-Biological Sci Med Sci* (2019) 74(1):62–7. doi: 10.1093/gerona/gly100

31. Landi F, Onder G, Bernabei R. Sarcopenia and diabetes: Two sides of the same coin. J Am Med Directors Assoc (2013) 14(8):540–1. doi: 10.1016/j.jamda.2013.05.004

32. Huang S, Xiang C, Song Y. Identification of the shared gene signatures and pathways between sarcopenia and type 2 diabetes mellitus. *PloS One* (2022) 17(3): e0265221. doi: 10.1371/journal.pone.0265221

33. Chung SM, Moon JS, Cheol M. Prevalence of sarcopenia and its association with diabetes: A meta-analysis of community-dwelling Asian population. *Front Med* (2021) 8:681232. doi: 10.3389/fmed.2021.681232

34. Qiao YS, Chai YH, Gong HJ, Zhuldyz Z, Stehouwer CDA, Zhou JB, et al. The association between diabetes mellitus and risk of sarcopenia: Accumulated evidences from observational studies. *Front Endocrinol* (2021) 12:782391. doi: 10.3389/ fendo.2021.782391

35. Kim JW, Kim R, Choi H, Lee SJ, Bae GU. Understanding of sarcopenia: From definition to therapeutic strategies. *Arch Pharmacal Res* (2021) 44(9-10):876–89. doi: 10.1007/s12272-021-01349-z

36. Galicia-Garcia U, Benito-Vicente A, Jebari S, Larrea-Sebal A, Siddiqi H, Uribe KB, et al. Pathophysiology of type 2 diabetes mellitus. *Int J Mol Sci* (2020) 21(17):6275. doi: 10.3390/ijms21176275

37. Wang T, Feng X, Zhou J, Gong H, Xia S, Wei Q, et al. Type 2 diabetes mellitus is associated with increased risks of sarcopenia and pre-sarcopenia in Chinese elderly. *Sci Rep* (2016) 6:38937. doi: 10.1038/srep38937

38. Perez-Sousa MA, del Pozo-Cruz J, Cano-Gutierrez CA, Izquierdo M, Ramirez-Velez R. High prevalence of probable sarcopenia in a representative sample from Colombia: Implications for geriatrics in Latin America. *J Am Med Directors Assoc* (2021) 22(4):859. doi: 10.1016/j.jamda.2020.10.021

39. Wiedmer P, Jung T, Castro JP, Pomatto LCD, Sun PY, Davies KJA, et al. Sarcopenia - molecular mechanisms and open questions. *Ageing Res Rev* (2021) 65:101200. doi: 10.1016/j.arr.2020.101200

40. Nowotny K, Jung T, Hohn A, Weber D, Grune T. Advanced glycation end products and oxidative stress in type 2 diabetes mellitus. *Biomolecules* (2015) 5(1):194–222. doi: 10.3390/biom5010194

41. Fielding RA, Vellas B, Evans WJ, Bhasin S, Morley JE, Newman AB, et al. Sarcopenia: An undiagnosed condition in older adults. current consensus definition: Prevalence, etiology, and consequences. international working group on sarcopenia. J Am Med Directors Assoc (2011) 12(4):249-56. doi: 10.1016/ j.jamda.2011.01.003

42. Wu H, Ballantyne CM. Skeletal muscle inflammation and insulin resistance in obesity. J Clin Invest (2017) 127(1):43–54. doi: 10.1172/jci88880

43. Bano G, Trevisan C, Carraro S, Solmi M, Luchini C, Stubbs B, et al. Inflammation and sarcopenia: A systematic review and meta-analysis. *Maturitas* (2017) 96:10–5. doi: 10.1016/j.maturitas.2016.11.006