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© 2024 Roumi, Mirzababaei, Abaj, Davaneghi, Aali and Mirzaei. This is an open-access 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. The interaction between polyphenol intake and genes (MC4R, Cav-1, and Cry1) related to body homeostasis and cardiometabolic risk factors in overweight and obese women: a cross-sectional study

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Background: Cardiovascular disease (CVD), which is an important global health challenge, is expanding. One of the main factors in the occurrence of CVD is a high genetic risk. The interaction between genetic risk in CVD and nutrition is debatable. Polyphenols are one of the important dietary components that may have a protective role in people who have a high genetic risk score (GRS) for cardiometabolic risk factors. This study, conducted in overweight and obese women, examines the interaction between polyphenol intake and specific genes (MC4r, Cav-1, and Cry1) related to maintaining body balance and their interaction with cardiometabolic risk factors.

Methods: This cross-sectional study included 391 women who were overweight or obese, aged 18 to 48 years, with a body mass index (BMI) between 25 and 40 kg/m². Body composition was measured using the InBody 770 scanner. Total dietary polyphenol intake (TDPI) was assessed with a validated 147-item food frequency questionnaire (FFQ), and polyphenol intakes were determined using the Phenol-Explorer database. Serum samples underwent biochemical tests. The Genetic Risk Score (GRS) was calculated based on the risk alleles of three genes: MC4r, Cav-1, and Cry1.

Results: The mean±standard deviation (SD) age and BMI of women were 36.67 (9.1) years and 30.98 (3.9) kg/m², respectively. The high GRS and high TDPI group had a significant negative interaction with fasting blood glucose (FBS) (p=0.01). Individuals who had a high GRS and a high phenolic acid intake were found to have a significant negative interaction with Triglyceride (p=0.04). Similarly, individuals with high GRS and a high intake of flavonoids had a significant negative interaction with TG (p<0.01) and a significant positive interaction with High-density lipoprotein (HDL) (p=0.01) in the adjusted model.

Conclusion: According to our findings, those with a high GRS may have a protective effect on cardiometabolic risk factors by consuming high amounts of polyphenols. Further studies will be necessary in the future to validate this association.

KEYWORDS

cardiometabolic risk factors, genetic risk score, homeostasis, obesity, polyphenols

Introduction

Cardiovascular disease (CVD), which is the leading cause of death and a global health challenge, is becoming more prevalent (1-3). It is projected that by 2030, the number of deaths from this disease will reach 23.6 million (4). The rate of CVD in women is reported to be 1 in every 3 women, with 45% of women over the age of 20 affected (5, 6). Cardiometabolic risk factors like obesity, high blood pressure, dyslipidemia, and inflammation play a role in the development of CVD (7, 8).

Genetic background plays an important role in CVD (9). CVD is greatly influenced by the genetic predisposition of individuals (10). Research has shown that having a genetic risk for developing cardiometabolic risk factors increases the likelihood of certain health problems, emphasizing the importance of genetic factors in understanding cardiometabolic diseases (11). A genetic risk score (GRS) is an estimate of an individual's genetic predisposition to a specific outcome, such as disease susceptibility. The combination of multiple genetic markers enables the prediction of disease risk based on an individual's genetic profile (12). Genes play a crucial role in maintaining body homeostasis by regulating various metabolic processes. The research suggests that gene expression is closely connected to metabolite homeostasis, influencing adaptations in response to environmental changes and influencing energy efficiency and product formation (13).

Genetic mutations, like the Melanocortin 4 receptor (MC4R) gene mutation, can lead to obesity (14). This gene is situated in the hypothalamus (15, 16) and its mutation may indirectly contribute to a higher risk of mortality from CVD (17). The MC4R gene (rs17782313) not only influences obesity but is also associated with other risk factors for CVD, including hypertension (HTN) (18). Inactivation of the MC4R gene has been shown to reduce blood pressure independently of obesity in previous studies (19).

CAV-1, also known as Caveolin-1, is a protein that has been related to different biological processes and diseases. Research has found that CAV-1 levels are increased in individuals with metabolic syndrome (20). Studies reported that CAV-1 might have a role in the impairment of endothelial function, which is a fundamental anomaly in the development of hypertension, atherosclerosis, and coronary artery disease (21).

Moreover, the presence of CAV-1 in the cells lining the blood vessels can be affected by various factors including green tea polyphenols. The presence of CAV-1 in individuals with metabolic syndrome, a disease linked to insulin resistance (IR), high blood glucose levels, hypertension, abnormal lipid levels, obesity, and increased WC, has been observed (20, 22).

The Cryptochrome 1 (Cry1) gene is a molecular clock gene that plays a role in generating circadian rhythms (23). Evidence suggests a potential association between CRY1 (rs10861688) polymorphism, obesity and related cardiovascular risk factors (24). Research has shown that CRY1 is associated with components of metabolic syndrome, such as hypertension and triglyceride (25) levels, as well as obesity and insulin resistance (IR) (26, 27). Furthermore, recent research has suggested that variations and genetic differences in the human genome, including different forms of the Cry1 gene, could have an effect on energy expenditure and body weight (28).

Nutrition is a factor that influences GRS on the incidence of cardiometabolic diseases (29). Polyphenols, chemical compounds present in plants such as fruits, vegetables, and tea, have been shown in studies to be effective in decreasing the risk factors associated with CVD (30, 31). Previous studies have discussed various types of polyphenols, including flavonoids, stilbenes, phenolic acids, and lignans (32). Numerous research studies have examined the potential benefits of polyphenols in preventing obesity, and there is evidence to suggest that plant polyphenols have the potential to be effective in this area (33). In a cohort study, researchers discovered that elevated levels of flavanones and lignans were correlated with adult body composition, including BMI and waist circumference (WC) (34). Furthermore, a separate study conducted on women showed that polyphenols were linked to decreased fasting blood sugar (FBS) and blood pressure levels. Moreover, there was a significant correlation between elevated levels of high-density lipoprotein (HDL) cholesterol (35, 36).

The GRS allows us to explore how various genes related to cardiometabolic diseases interact with dietary intake to influence cardiometabolic risk factors. In this study, we aim to examine how a high consumption of polyphenols affects cardiometabolic risk factors in individuals with a high GRS, to determine if consuming high levels of polyphenols is beneficial in improving these risk factors.

Method

Study population

In this cross-sectional study, 391 overweight or obese women, aged between 18 and 48 years and with a body mass index (BMI) between 25 and 40 kg/m² participated. These women were selected from people who visited 20 different health centers in Tehran using random sampling. Individuals with a prior medical history of cardiovascular or thyroid disease, malignancies, liver or kidney diseases, types of diabetes, acute or chronic diseases, pregnancy, lactation, or menopause, adherence to a specific diet or weight loss

supplements, consumption of glucose and lipid lowering drugs and blood pressure medications within the past year, and smoking were not included in the study. Before the study, all participants were required to sign a written informed consent form. The study protocol received ethics approval from the Human Ethics Committee of Tehran University of Medical Sciences, with the ethics number IR.TUMS. MEDICINE.REC.1402.636. The procedures were conducted in compliance with applicable guidelines and regulations.

Evaluation of dietary intake

The participants' nutritional status, including energy intake, macronutrients, and micronutrients, was assessed using the food frequency questionnaire (FFQ) consisting of 147 items. Previous studies have confirmed the validity of this questionnaire for the Iranian population (37). A trained nutritionist conducted interviews with women to complete this questionnaire. The data were then analyzed using version 7 of the NUTRITIONIST 4 software, after being converted to grams using household measure servings (38).

Evaluation of dietary polyphenol intake (DPI)

The Phenol-Explorer database (www.phenol explorer.eu/contents) was used to gather data on the overall polyphenol content in various foods (39). The total polyphenol content was determined either through the Folin Ciocalteu assay or by calculating the sum of four main subgroups, which include flavonoids, phenolic acids, stilbenes, lignans, and other polyphenols.

Measurement of anthropometric indicators

Participants' height was measured using a Seca stadiometer with an accuracy of 0.1 cm, and their weight was measured with a Seca digital scale (Hamburg, Germany) with an accuracy of 0.1 kg. To measure these two indicators, the participants must be without shoes and in the lightest clothes. BMI was calculated from the ratio of weight (kg) to the square of height (m²). In addition, to measure abdominal obesity, WC in the smallest circumference and hip circumference (HC) in the largest circumference were measured with an accuracy of 0.1 cm (40). Waist-to-hip ratio (WHR) was also calculated.

Assessment of body composition

The InBody 770 Scanner, a multi-frequency bioelectrical impedance analyzer, was used to measure body composition parameters such as the amount and proportion of visceral fat level (VFL) and obesity degree (3). The measurements were taken in the morning while participants were in a fasted state and wearing light clothing. Participants were instructed to refrain from exercising, carrying electrical devices, and to urinate before the analysis to ensure accuracy. Following the manufacturer's instructions, participants stood on the scale barefoot and held the machine's handles for 20 s, after which the results were printed (41).

To assess the levels of biochemical factors (such as glucose and lipids) in the participants, blood samples were collected after a period of fasting and the serum was separated using a centrifuge. The serum was then divided into smaller portions and stored at -80°C until it could be analyzed. All blood parameters were measured in the Bionanotechnology Laboratory of the Endocrine and Metabolism Research Institute of Tehran University of Medical Sciences and analyzed using an exclusive assay based on the instructions provided by the manufacturer. All calculations were performed using a package from Randox Laboratories (Hitachi 902). The GPO-PAP method was employed to determine the levels of TG, while enzymatic and clearance endpoint assays were utilized to measure the total cholesterol (TC) and HDL cholesterol, respectively, in this research (42). Alanine aminotransferase (4) and aspartate aminotransferase (AST) were measured via standard protocols.

Measurement of genetic risk score (GRS)

The DNA was obtained from whole blood samples through salting out techniques (43). The quality and quantity of the extracted DNA were evaluated using 1% agarose gel and the Nanodrop 8000 Spectrophotometer, respectively. TaqMan Open Array was used to genotype single nucleotide polymorphisms (SNPs), including CAV-1 (rs3807992), Cry1 (rs2287161), and MC4R (rs17782313) (44). These SNPs have been associated with obesity-related traits in previous studies (45–47). The GRS was computed by summing up the scores of the three SNPs, which were coded as 0, 1, or 2 based on their association with higher BMI. The unweighted GRS ranges from 0 to 6, with higher scores indicating a greater genetic predisposition to high BMI (48).

Measurement of blood pressure

Systolic blood pressure (SBP) and diastolic blood pressure (DBP) were measured using standard sphygmomanometer and cuff through auscultation. After each subject had sat for at least 5 min, two consecutive blood pressure measurements were taken. Systolic blood pressure (SBP) and diastolic blood pressure (DBP) were measured with a standard mercury sphygmomanometer using the first and fifth Korotkoff sounds, to within 2 mmHg. If the difference between the two systolic or diastolic blood pressures was more than 5 mmHg, an additional measurement was performed.

Assessment of other variables

Trained nutritionists filled out a demographic questionnaire that consisted of information regarding job, education, marital, and economic status. The present study measured PA as a confounding variable using the International Physical Activity Questionnaire (IPAQ). The data obtained from this questionnaire was measured on a scale of metabolic hours per week (MET. h week⁻¹) (49).

Statistical analysis

In the present study, cardiometabolic risk factors were determined based on biochemical, anthropometric and body composition criteria. The statistical analyses were performed using the IBM SPSS Statistics 23 software, with a significance level of less than 0.05. The normality of the distribution of the quantitative data of the study was performed by the Kolmogorov-Smirnov test. In this study, qualitative variables (Marriage, Education, Job and Economic Status) were described as numbers/percentages and quantitative variables (demographic variables, anthropometric measurements, body composition, blood parameters and blood pressure) were described as mean ± standard deviation (SD). The characteristics of the study participants among the tertile of GRS were compared with ANOVA and the characteristics of the study participants among the total dietary polyphenols index (TDPI) were compared with the independent t-test. To eliminate any confounding outcomes, ANCOVA was utilized. Both the crude and adjusted models employed a generalized linear model (GLM) to evaluate the interactions between metabolic factors and GRS, phenolic acid, lignans, flavonoids, and polyphenol. The outcomes were adjusted for age, energy intake, and PA.

Results

Study population characteristic

Our study was conducted on 391 overweight or obese women. The mean (\pm SD) age, weight, BMI and WC of participants were 36.67 (9.1) years, 80.28 (11.05) kg, 30.98 (3.9) kg/m² and 99.16 (9.42) cm, respectively. Most of the participants were married (70.8%) and had no academic education (51%). They were also in a Moderate economic situation (45.5%).

Characteristics of the study participants among tertile of GRS

The baseline characteristics of the study participants were presented in Table 1, categorized based on tertiles of their GRS. According to the table, in crude model there were a significant difference in mean values among the GRS tertiles for weight (p=0.03), height (p=0.03), WC (p=0.03), and WHR (p=0.03). There was also a marginally significant difference for TG (p=0.06) among the GRS tertiles. After adjusting for confounding factors such as age, PA, and energy intake, the VFL (p=0.03), SBP (p=0.01), FBS (p=0.02), and LDL (p=0.01) became significantly different among the GRS tertiles. Additionally, there was a significant difference in height (p=0.005), WC (p=0.04), and WHR (p=0.02), and a marginally significant difference for weight (p=0.06) and TC (p=0.08) among the GRS tertiles.

Characteristics of the study participants among intake of TDPI

In Table 2, the characteristics of the participants are compared based on receiving low and high TDPI. The results showed that in the

crude model, there is a significant difference between the two groups (low and high TDPI) in terms of job (p=0.02), However in the adjusted model (age, PA, and energy intake), in addition to job (p=0.01), the participants also had significant differences in terms of FBS (p<0.001), TC (p=0.01), and LDL (p<0.001). Also a marginal difference for WC (p=0.08) and WHR (p=0.09) between the low intake and high intake groups of TDPI.

The interaction between GRS, TDPI, stilbenes, phenolic acid, lignans, flavonoids and polyphenol on cardiometabolic risk factors

The findings on the interaction between GRS, TDPI, and stilbenes on cardiometabolic risk factors are presented in Table 3. The crude and adjusted models showed a significant negative interaction between high GRS and high intake TDPI with FBS (crude: 95%CI = -20.39, -3.28, p < 0.001; adjusted: 95%CI = -19.95, -1.8, p = 0.01).

The findings also showed that the significant negative interaction between moderate GRS and high intake TDPI with BMI (95%CI=-5.05, -1.03, p < 0.001), WC (95%CI=-12.53, -2.33, p < 0.001), VFL (95%CI=-3.88, -0.33, p = 0.02), FBS (95%CI=-20.39, -3.28, p < 0.001) and LDL (95% CI=-28.75, 0.028, p = 0.05) in the crude model. After adjusting for age, IPAC and total energy intake in model 1, the interaction between moderate GRS and high intake TDPI with BMI (95% CI=-4.55, -0.46, p = 0.01), WC (95% CI=-12.51, -2.38, p < 0.001), VFL (95% CI=-3.85, -0.26, p = 0.02) and FBS (95% CI=-13.38, -1.63, p = 0.01) remained negative. The interaction between TDPI and GRS on BMI, WC, VFL and FBS is shown in Figure 1.

Furthermore, there was no significant interaction found between moderate/high GRS and stilbenes with cardiometabolic risk factors in both the crude and adjusted models.

The interaction between GRS, phenolic acid and lignans on cardiometabolic risk factors were presented in Table 4. In the crude model, a significant positive interaction was observed between high GRS and high intake phenolic acid on HDL (95%CI=0.16, 19.23, p=0.04) But in adjusted model, this interaction was not reported. However, a significant negative interaction was observed between high GRS and high intake phenolic acid on TG (95%CI=-115.66, -2.42, p=0.04). Interaction between Phenolic acid and GRS on TG is shown in Figure 2.

Furthermore, there was no significant interaction found between moderate/high GRS and lignans with cardiometabolic risk factors in both the crude and adjusted models.

The interaction between GRS, flavonoids and polyphenol on cardiometabolic risk factors were presented in Table 5. The crude and adjusted models showed a significant negative interaction between high GRS and high intake flavonoids with TG (crude: 95%CI = -137.58, -31.2, p < 0.001; adjusted: 95%CI = -135.52, -22.69, p < 0.001). Also, a significant positive interaction was observed between high GRS and high intake flavonoids with HDL (95%CI = 2.47, 22.65, p = 0.01) in the adjusted model.

The findings also showed that the significant negative interaction between moderate GRS and high intake flavonoids with BMI

TABLE 1 Characteristics of the study participants among tertile of genetic risk score (GRS).

Variables				GRS		
		Low risk <3 (n = 164)	Moderate risk (3&4) (n = 97)	High risk ≥5 (<i>n</i> = 130)	<i>p</i> -value	<i>p</i> -value*
Demographic vari	ables					
Age (years)		36.02±8.78	36.53±8.15	36.08±8.61	0.91	0.89
PA (MET-minutes/	/week)	1052.27 ± 1116.01	1353.96±2742.28	1389.51 ± 2576.15	0.61	0.58
Anthropometric n	neasurements					
Weight (kg)		79.31 ± 9.79	76.9903 ± 10.29312	82.47±9.33	0.03	0.06
Height (cm)		162.46 ± 5.41	160.26 ± 5.93	161.41 ± 4.45	0.03	0.005
BMI (kg/m²)		29.98 ± 3.31	30.09 ± 3.5	31.44 ± 3.32	0.16	0.22
WC (cm)		97.39 ± 8.62	96.4 ± 8.56	101.42 ± 9.03	0.03	0.04
WHR		0.92 ± 0.05	0.92 ± 0.04	0.95 ± 0.05	0.03	0.02
Body composition	l					
VFL (cm ²)		14.9±3.15	15±3.07	16.16±2.98	0.19	0.03
OD (%)		139.37±15.47	139.91±16.32	146.25 ± 15.54	0.16	0.22
Blood pressure						
SBP (mmHg)		111.24 ± 12.06	111.26±15.38	113.08 ± 14.57	0.83	0.01
DBP (mmHg)		77.14 ± 10.08	77.44±9.51	77.58 ± 11.04	0.97	0.16
Blood parameters						
FBS (mg/dL)		87.32±9.2	86.46±10.32	88.7±9.82	0.57	0.02
TC (mg/dL)		186.41±33.3	182.95 ± 37.7	179.83±35.46	0.68	0.08
TG (mg/dL)		120.83 ± 57.88	108.35 ± 53.01	136.91±75.67	0.06	0.11
HDL (mg/dL)		47.35±9.69	47.59±11.62	46.87±11.6	0.95	0.98
LDL (mg/dL)		96.96±21.67	95.12±25.51	90.45 ± 25.82	0.51	0.01
AST (IU/L)		17.51±6.9	17.97±8.08	17.91±8.56	0.92	0.79
ALT (IU/L)		18.16±14.02	18.77±13.31	20.25±13.68	0.8	0.53
Qualitative variab	le N (%)					
Manifestation	Single	29 (49.2)	25 (42.4)	5 (8.5)	0.16	0.10
Marriage status	Married	69 (35.9)	96 (50)	27 (14.1)	0.16	0.18
P.1	Non academic	49 (76.8)	62 (88.4)	22 (34.9)	6.2	0.00
Education levels Academic		49 (42.2)	57 (49.1)	10 (8.6)	- 0.3	0.32
T . I.	Unemployed	52 (34.7)	77 (51.3)	21 (14)	0.16	0.10
Job Employed		45 (46.9)	40 (41.7)	11 (11.5)	0.16	0.18
	Poor	19 (34.5)	27 (49.1)	9 (16.4)		
Economic status	Moderate	46 (39.3)	56 (47.9)	15 (12.8)	0.77	0.77
	Good	28 (43.8)	30 (46.9)	6 (9.4)		

BMI, Body mass index; WC, waist circumference; WHR, waist height ratio; FBS, fasting blood sugar; Chol, Cholesterol; TG, Triglyceride; HDL, High density lipoprotein; LDL, Low density lipoprotein; SBP, Systolic Blood Pressure; DBP, Diastolic Blood Pressure; OD, Obesity degree; VFL, Visceral fat level; TC, total cholesterol.

Quantitative variable: Mean \pm SD (Standard deviation), Qualitative variable: N (%) Number (Percentage).

 $p\text{-values}\,{<}\,0.05$ are in bold.

p-value calculated by analysis of variance (ANOVA).

*p-value was found by ANCOVA, and adjusted for age, IPAC and total energy intake.

(95%CI=-5.44, -1.49, p < 0.001), WC (95%CI=-12.48, -2.39, p < 0.001), VFL (95%CI=-3.96, -0.46, p = 0.01) and OD (95%CI=-24.62, -1.47, p = 0.02) in the crude model. After adjusting for age, IPAC, and total energy intake in model 1, the interaction between moderate GRS and high intake flavonoids with BMI (95%CI=-4.59, -0.52, p = 0.01), WC (95%CI=-10.79, -0.56,

p = 0.03) and OD (95%CI = -22.67, 0.27, p = 0.05) remained negative. Interaction between Flavonoids and GRS on BMI, WC, OD, TG and HDL is shown in Figure 3.

Furthermore, there was no significant interaction found between moderate/high GRS and polyphenol with cardiometabolic risk factors in both the crude and adjusted models.

TABLE 2 Characteristics of the study participants among intake of total dietary polyphenols index (TDPI).

Quantitative va	ariables		TDPI			
		Low intake (<i>n</i> = 196)	High intake (<i>n</i> = 195)	<i>p</i> -value	<i>p</i> -value *	
Demographic variab	les					
Age (years)		35.44 ± 8.58	36.77±8.42	0.23	0.31	
PA(MET-minutes/we	ek)	1086.64 ± 2159.97	1292.74±2061.70	0.44	0.21	
Anthropometric mea	asurements					
BMI (kg/m ²)		30.6 ± 4.06	30.29±3.4	0.53	0.43	
Weight (kg)		80.51 ± 11.04	79.73±10.56	0.48	0.21	
Height (cm)		160.82 ± 6.31	161.25±5.31	0.47	0.35	
WC (cm)		98.17±9.06	97.32±9.24	0.48	0.08	
NHR		0.93 ± 0.046	0.92 ± 0.05	0.45	0.09	
Body composition						
/FL (cm ²)		15.32±3.3	15.06±3.3	0.55	0.45	
OD (%)		142.21 ± 18.98	140.88 ± 15.84	0.56	0.43	
Blood parameters						
FBS (mg/dL)		88.19 ± 11.18	86.77 ± 8.4	0.27	<0.001	
TC (mg/dL)		186.86 ± 40.29	182.19 ± 32.73	0.33	0.01	
TG (mg/dL)		116.29 ± 56.32	118.61 ± 60.03	0.76	0.31	
HDL (mg/dL)		46.63 ± 11.28	47.39 ± 10.35	0.59	0.73	
LDL (mg/dL)		94.68 ± 24.53	95.55 ± 23.98	0.78	<0.001	
AST (IU/L)		17.97 ± 6.37	17.68 ± 8.11	0.76	0.81	
ALT (IU/L)		18.68 ± 10.33	19.38 ± 14.93	0.68	0.48	
Qualitative variable						
Marriage status	Single	58 (52.7)	52 (47.3)	0.29	0.36	
Walliage status	Married	138 (49.1)	143 (50.9)	0.29	0.50	
Education	Nonacademic	97 (49.5)	107 (55.4)	0.26	0.14	
Laucation	Academic	99 (50.5)	86 (44.6)	0.20	0.14	
Job	Unemployed	105 (46.3)	122 (53.7)	0.02	0.01	
100	Employed	90 (57)	68 (43)	0.02	0.01	
	Poor	42 (47.7)	46 (52.3)			
Economic status	Moderate	86 (47.3)	96 (52.7)	0.18	0.17	
	Good	62 (57.9)	45 (42.1)			

BMI, Body mass index; WC, waist circumference; WHR, waist height ratio; FBS, fasting blood sugar; Chol, Cholesterol; TG, Triglyceride; HDL, High density lipoprotein; LDL, Low-density lipoprotein; OD, Obesity degree; VFL, Visceral fat level; TC, total cholesterol.

TDPI Low intake < 2060.65 (mg/day), TDPI High intake \geq 2060.65(mg/day).

p-values < 0.05 are in bold.

Quantitative variable: Mean \pm SD (Standard deviation).

Qualitative variable N (%): N (%) Number (Percentage).

p-value calculated by analysis of independent sample *t*-test.

p-value *Adjusted for age, IPAC, and total energy intake calculated by analysis of covariance (ANCOVA).

Discussion

The purpose of this cross-sectional study was to investigate the relationship between polyphenol consumption and genes (MC4r, Cav-1, and Cry1) and cardiometabolic risk factors in overweight and obese Iranian women.

The findings of our study revealed a significant negative interaction between high GRS and high intake TDPI with FBS in both crude and adjusted models. Also, a significant negative interaction was observed between high GRS and high intake of phenolic acid on TG in the adjusted model. Moreover, a significant negative interaction was observed between high GRS and high intake flavonoids with TG, and a significant positive interaction was observed between high GRS and high intake flavonoids with HDL in the adjusted model. Furthermore, a significant negative interaction between moderate GRS and high intake TDPI on BMI, WC, VLF and FBS levels. According to our results, high intake of TDPI is associated with significant interaction with decreased levels of BMI, WC, VLF, and FBS in participants at moderate risk of GRS. In addition, in this study reported a significant negative interaction between moderate GRS and flavonoid on BMI, WC and OD levels.

Variables	GRS				TDPI						Stil	benes			
		Low			High	intake			Low	High intake					
		intake < 2060.65		Crude			Adjust		intake < 0.49		Crude			Adjust	
		(mg/ day)	β	95% Cl	р	β	95% Cl	р	(mg/ day)	β	95% Cl	p		95% Cl	p
Anthropometri	ic factors														
	Low risk	Reference			Refe	rence			Reference			Refe	rence		
BMI (kg/m²)	Moderate risk	_	-3.04	-5.05 to -1.03	<0.001	-2.50	-4.55 to -0.46	0.01	_	0.48	-1.49 to 2.47	0.62	0.12	-1.89 to 2.14	0.9
	High risk	-	-1.81	-4.96 to 1.34	0.26	-1.79	-5.01 to 1.42	0.27	-	-1.16	-4.13 to 1.8	0.44	-0.43	-3.47 to 2.59	0.77
	Low risk	Reference	Reference					Reference		1	Refe	rence			
WC (cm)	Moderate risk	_	-7.43	-12.53 to -2.33	<0.001	-7.45	-12.51 to -2.38	<0.001	-	3.27	-1.73 to 8.29	0.2	2.68	-2.36 to 7.73	0.29
	High risk	_	-5.29	-13.12 to 2.53	0.18	-5.59	-13.32 to 2.14	0.15	-	-2.35	-9.8 to 5.09	0.53	-1.21	-8.72 to 6.29	0.75
	Low risk	Reference		1	Refe	rence			Reference			Refe	rence	1	
WHR	Moderate risk	_	-0.01	-0.048 to 0.008	0.17	-0.02	-0.052 to 0.004	0.09	-	0.02	0.001- 0.056	0.06	0.02	-0.005 to 0.05	0.1
	High risk	-	-0.02	-0.056 to 0.031	0.56	<0.001	-0.051 to 0.035	0.71	-	-0.01	-0.051 to 0.03	0.61	<0.001	-0.05 to 0.32	0.68
Body composit	ion	1							1						
	low risk	Reference		1	Refe	rence			Reference			Refe	rence	1	1
VFL (cm ²)	Moderate risk	-	-2.11	-3.88 to -0.33	0.02	-2.06	-3.85 to -0.26	0.02	-	0.89	-1.66 to 1.83	0.92	-0.23	-2.02 to 1.55	0.79
	High risk	-	-1.02	-3.76 to 1.71	0.46	-0.82	-3.58 to 1.93	0.55	-	-0.96	-3.56 to 1.64	0.47	-0.31	-2.98 to 2.34	0.81
	Low risk	Reference			Refe	rence			Reference			Refe	rence		
OD (%)	Moderate risk	_	-6.70	-18.47 to 5.06	0.26	-6.77	-18.37 to 4.82	0.25	-	2.17	-9.32 to 13.67	0.71	-2.32	-13.78 to 9.13	0.69
	High risk	_	-8.52	-26.69 to 9.65	0.35	-9.56	-27.36 to 8.23	0.29	_	-7.73	-24.85 to 9.39	0.37	-4.72	-21.77 to 12.32	0.58
Biochemical va	riables														
-	Low risk	Reference	e Reference F									Refe	rence		
FBS (mg/dL)	Moderate risk	_	-9.16	-14.74 to -3.58	<0.001	-7.51	-13.38 to -1.63	0.01	_	3.06	-2.41 to 8.53	0.27	3.48	-2.29 to 9.17	0.24
	High risk	_	-11.83	-20.39 to -3.28	<0.001	-10.88	-19.95 to -1.8	0.01	_	2.58	-5.68 to 10.85	0.54	4.09	-4.66 to 12.84	0.36

TABLE 3 The interaction between GRS, TDPI and stilbenes on cardiometabolic risk factors.

(Continued)

TABLE 3 (Continued)

Variables	GRS			1	[DPI						Stil	benes	5		
		Low			High	intake			Low	High intake					
		intake < 2060.65		Crude			Adjust		intake < 0.49		Crude			Adjust	
		(mg/ day)	β	95% Cl	р	β	95% Cl	p	(mg/ day)		95% Cl	p		95% Cl	p
	Low risk	Reference			Refe	rence			Reference			Refe	erence		
TC (mg/dL)	Moderate risk	-	-25.02	-46.51 to -3.54	0.22	-18.77	-40.94 to 3.39	0.09	-	1.51	-19.24 to 22.28	0.88	-3.86	-25.18 to 17.44	0.72
	High risk	-	-21.92	-54.89 to 11.04	0.19	-18.74	-52.97 to 15.49	0.28	-	3.88	-27.47 to 35.24	0.8	17.08	-15.44 to 49.62	0.3
TG (mg/dL)	Low risk	Reference			Refe	rence			Reference			Refe	erence		1
	Moderate risk	-	-7.63	-44.58 to 29.23	0.68	0.45	-38.6 to 39.52	0.98	-	18.50	-16.65 to 53.67	0.3	14.71	-22.49 to 51.91	0.43
	High risk	-	-23.11	-81.29 to 35.06	0.43	-21.68	-83.49 to 40.12	0.49	-	4.36	-49.41 to 58.13	0.87	14.59	-42.95 to 72.15	0.61
HDL (mg/dL) Low risk		Reference	Reference									Refe	erence		
	Moderate risk	-	-1.42	-8.12 to 5.26	0.67	1.23	-5.8 to 8.26	0.73	-	0.30	-5.97 to 6.58	0.92	0.51	-6.12 to 7.14	0.88
	High risk	_	2.73	-7.53 to 13	0.6	1.09	-9.76 to 11.95	0.84	-	2.71	-6.7 to 12.26	0.56	4.77	-5.35 to 14.89	0.35
LDL (mg/dL)	Low risk	Reference			Refe	rence			Reference			Refe	erence		
	Moderate risk	_	-14.36	-28.75 to 0.028	0.05	-6.08	-21.04 to 8.87	0.42	-	6.25	-7.41 to 19.91	0.37	7.05	-7.12 to 21.24	0.33
	High risk	-	-20.03	-42.11 to 2.049	0.07	-21.10	-44.2 to 1.99	0.07	-	6.39	-14.24 to 27.03	0.54	12.57	-9.08 to 34.22	0.25
AST (IU/L)	Low risk	Reference			Refe	rence			Reference			Refe	erence		
	Moderate risk	_	-0.50	-5.05 to 4.038	0.82	0.83	-3.91 to 5.59	0.72	-	0.52	-3.8 to 4.86	0.81	1.94	-2.58 to 6.47	0.4
	High risk	-	-4.80	-11.78 to 2.16	0.17	-3.90	-11.25 to 3.43	0.29	-	-3.69	-10.51 to 2.57	0.23	-3.76	-10.68 to 3.14	0.28
ALT (IU/L)	Low risk	Reference			Refe	rence			Reference			Refe	erence		
	Moderate risk	-	-3.25	-11.27 to 4.76	0.42	-1.39	-9.92 to 7.14	0.74	-	0.64	-7.01 to 8.29	0.86	3.00	-5.14 to 11.14	0.47
	High risk	_	-5.93	-18.23 to 6.37	0.34	-4.02	-17.2 to 9.15	0.54	-	-2.71	-14.27 to 8.85	0.64	-0.96	-13.39 to 11.46	0.87

B, Standard Error; GRS, Genetic risk score; BMI, Body mass index; WC, waist circumference; WHR, waist height ratio; FBS, fasting blood sugar; TG, Triglyceride; LDL, Low density lipoprotein; HDL, High density lipoprotein; OD Obesity degree; VFL, Visceral fat level; TC, total cholesterol. TDPI High intake ≥ 2060.65 (mg/day), Stilbenes High intake ≥ 0.49(mg/day).

 $p\text{-values}\,{<}\,0.05$ are in bold.

Adjust = adjusted for potential confounding factors including (age, IPAC and total energy intake).

Low Risk: 0,1,2 Risk alleles, Moderate Risk: 3,4 Risk allele, High Risk: 5,6 Risk allele.



Interaction between TDPI and GRS on (A) BMI, (B) WC, (C) VFL, (D) FBS. The interaction between low and high intake of TDPI and GRS. Data shown are mean \pm standard error of the mean. BMI, Body mass index; WC, Waist circumference; VFL, Visceral fat level; FBS, Fasting Blood Sugar; GRS, genetic risk score; TDPI, total dietary polyphenol intake. Adjust = adjusted for potential confounding factors including (age, IPAC and total energy intake). The asterisk (*) represents the *p*-value of the statistical test. The asterisk means that the *p*-value is less than 0.05.

Variables	GRS			Phen	olic a	cid					Lig	nans			
		Low			High	intake			Low			High	intake		
		intake <		Crude			Adjust		intake <		Crude			Adjust	
		55.95 (mg/ day)	β	95% Cl	р	β	95% Cl	р	0.0065 (mg/ day)	β	95% Cl	р	β	95% Cl	p
Anthropometr	ic factors														
	Low risk	Reference		1	Refe	erence	1		Reference		1	Refe	rence	1	
BMI (kg/m²)	Moderate risk	-	-0.83	-2.83 to 1.16	0.41	-0.60	-2.62 to 1.41	0.55	-	-0.30	-2.29 to 1.69	0.76	0.48	-1.54 to 2.52	0.63
	High Risk	_	-0.59	-3.56 to 2.37	0.69	-1.26	-4.29 to 1.77	0.41	_	-1.08	-4.04 to 1.88	0.47	-0.60	-3.65 to 2.44	0.69
	Low Risk	Reference			Refe	erence			Reference			Refe	rence		
WC (cm)	Moderate Risk	_	0.33	-4.74 to 5.4	0.89	-0.06	-5.15 to 5.01	0.97	_	-0.69	-5.76 to 4.38	0.78	1.05	-4.03 to 6.15	0.68
	High Risk	-	2.09	-5.38 to 9.56	0.58	<0.001	-7.55 to 7.53	0.99	_	-3.52	-11 to 3.95	0.35	-2.20	-9.8 to 5.39	0.56
	Low Risk	Reference			Refe	erence			Reference			Refe	rence		
WHR	Moderate Risk	_	0.01	-0.01 to 0.04	0.35	0.01	-0.01 to 0.03	0.43	_	<0.001	-0.03 to 0.02	0.67	-2.24	-0.02 to 0.02	0.99
	High Risk	_	0.03	-0.009 to 0.072	0.12	0.02	-0.01 to 0.06	0.21	_	-0.01	-0.05 to 0.02	0.52	<0.001	-0.04 to 0.03	0.86
Body composit	tion			1	1										
	Low risk	Reference			Refe	erence			Reference			Refe	rence		
VFL (cm ²)	Moderate risk	-	-0.66	-2.42 to 1.09	0.45	-0.66	-2.44 to 1.12	0.46	_	0.57	-1.17 to 2.33	0.51	0.92	-0.86 to 2.72	0.31
	High risk	_	0.07	-2.51 to 2.67	0.95	-0.24	-2.89 to 2.41	0.85	_	-0.83	-3.42 to 1.76	0.39	0.03	-2.64 to 2.7	0.98
	Low risk	Reference			Refe	erence			Reference			Refe	rence		
OD (%)	Moderate risk	_	2.99	-8.57 to 14.56	0.61	0.25	-11.18 to 11.69	0.96	_	3.79	-7.58 to 15.54	0.5	5.08	-6.4 to 16.57	0.38
	High Risk	_	0.27	-16.82 to 17.37	0.97	-4.73	-21.75 to 12.28	0.58	_	-7.17	-24.29 to 9.94	0.41	-5.25	-22.42 to 11.9	0.54
Biochemical va	riables				1				1						
	Low risk	Reference			Refe	erence			Reference			Refe	rence		
FBS (mg/dL)	Moderate risk	_	0.45	-5.03 to 5.94	0.87	0.30	-5.45 to 6.05	0.91	_	1.19	-4.34 to 6.72	0.67	1.03	-4.78 to 6.84	0.72
	High risk	_	-2.71	-10.97 to 5.55	0.52	-3.39	-12.21 to 5.41	0.45	_	0.37	-7.96 to 8.7	0.93	0.41	-8.7 to 9.54	0.92
	Low risk	Reference			Refe	erence			Reference			Refe	rence		
TC (mg/dL)	Moderate risk	_	0.65	-20.11 to 21.42	0.95	-3.75	-24.96 to 17.45	0.72	-	16.88	-3.92 to 37.68	0.11	12.40	-8.99 to 33.81	0.25
	T3	_	-11.81	-43.07 to	0.45	-21.04	-53.54 to 11.45	0.2	-	4.59	-26.73 to	0.77	14.60	-18.97 to	0.39

TABLE 4 The interaction between GRS, phenolic acid and Lignans on cardiometabolic risk factors.

(Continued)

TABLE 4 (Continued)

Variables	GRS			Phen	olic a	cid					Lig	nans			
		Low			High	intake			Low			High	intake		
		intake <		Crude			Adjust		intake <		Crude		Adjust		
		55.95 (mg/ day)	β	95% Cl	р	β	95% Cl	p	0.0065 (mg/ day)	β	95% Cl	p	β	95% Cl	р
TG (mg/dL)	Low Risk	Reference			Refe	erence			Reference			Refe	rence		
	Moderate risk	_	31.83	-2.74 to 66.41	0.07	28.60	-7.62 to 64.82	0.12	_	1.96	-33.53 to 37.45	0.91	-9.24	-46.61 to 28.13	0.62
	High risk	-	-39.54	-92.26 - 13.17	0.14	-59.04	-115.66 to -2.42	0.04	-	16.72	-37.22 to 70.67	0.54	28.58	-30.54 to 87.7	0.34
HDL (mg/	Low risk	Reference			Refe	erence			Reference			Refe	rence		
dL)	Moderate risk	-	3.10	-3.23 to 9.44	0.33	2.87	-3.77 to 9.52	0.39	_	-0.13	-6.56 to 6.3	0.96	2.13	-4.6 to 8.87	0.53
	High risk	-	9.70	0.16– 19.23	0.04	7.89	-2.3 to 18.08	0.12	_	1.85	-7.83 to 11.54	0.7	7.02	-3.55 to 17.59	0.19
LDL (mg/dL)	Low risk	Reference			Refe	erence			Reference			Refe	rence	1	<u> </u>
	Moderate risk	_	6.33	-7.48 to 20.14	0.36	9.11	-5.01 to 23.25	0.2	_	9.69	-4.22 to 23.61	0.17	11.26	-3.14 to 25.66	0.12
	High risk	-	-7.33	-28.12 to 13.45	0.48	-13.89	-35.54 to 7.76	0.2	_	0.19	-20.76 to 21.16	0.98	3.34	-19.24 to 25.9	0.77
AST (IU/L)	Low risk	Reference			Refe	erence			Reference			Refe	rence		
	Moderate risk	-	-2.93	-7.25 to 1.37	0.18	-2.68	-7.19 to 1.83	0.24	-	-2.01	-6.39 to 2.36	0.36	-0.54	-5.13 to 4.04	0.81
	High risk	-	-2.31	-8.8 to 4.17	0.48	-1.86	-8.78 to 5.05	0.59	_	-1.35	-7.95 to 5.23	0.68	2.03	-5.16 to 9.23	0.58
ALT (IU/L)	Low risk	Reference			Refe	erence			Reference			Refe	rence		
	Moderate risk	-	-6.83	-14.38 to 0.71	0.07	-5.45	-13.49 to 2.59	0.18	-	-7.23	-14.9 to 0.43	0.06	-4.59	-12.79 to 3.59	0.27
_	High Risk	_	-0.28	-11.64 to 11.07	0.96	-0.12	-12.45 to 12.2	0.98	_	-3.70	-15.26 to 7.84	0.52	1.02	-11.83 to 13.87	0.87

B, Standard Error; GRS, Genetic risk score; BMI, Body mass index; WC, waist circumference; WHR, waist height ratio; FBS, fasting blood sugar; TG, Triglyceride; LDL, Low-density lipoprotein; HDL, High-density lipoprotein; OD, Obesity degree; VFL, Visceral fat level; TC, total cholesterol.

Phenolic acid High intake \geq 55.95(mg/day), Lignans High intake \geq 0.0065(mg/day).

p-values < 0.05 are in bold.

Adjust = adjusted for potential confounding factors including (age, IPAC and total energy intake).

Low Risk: 0,1,2 Risk alleles, Moderate Risk: 3,4 Risk allele, High Risk: 5,6 Risk allele.

However, there was no significant interaction found between moderate/high GRS and high intake stilbenes, polyphenols, and lignans with cardiometabolic risk factors in both the crude and adjusted models. In a study, there was no significant association total polyphenol and stilbenes with FBS, TG, HDL in participants. Also, there was no association between lignans with cholesterol, TG, HDL, DBP (50). The another study after adjusted for age, BMI, physical activity, and total energy intake, there was no association between Cry1 genotypes with cholesterol, TG, HDL, LDL, SBP, DBP (51). There was no association between Cav-1 rs3807992 genotypes with FBS, insulin, TC, TG (52).

Polyphenols are effective in cardiovascular health due to their antioxidant, blood sugar control, anti-inflammatory, and lipid profile control effects (25, 53, 54). In a study, it was discovered that a better diet quality, which might include polyphenol-rich foods, was significantly linked to a reduction in cardiometabolic risk factors (55). Furthermore, a cohort study spanning 10 years and involving more than 450,000 participants across 10 European countries revealed that



higher diet quality index (DQI) scores were linked to a reduced risk of CVD mortality and its associated risk factors, such as dyslipidemia and hyperglycemia (56).

GRS play a key role in comprehending various aspects of body homeostasis, especially with respect to conditions such as type 2 diabetes (T2DM) and cardiometabolic risk factors (9, 57). It seems that the effects of MC4R, Cav-1 and Cry1 genes on body composition and metabolic parameters may depend on the quality of the diet (58). Polyphenols have antioxidant and anti-inflammatory properties that can improve the quality of food. Based on the evidence, these compounds have the ability to influence the interaction and interactions between diet, genes, and metabolic parameters (59, 60). Polyphenols and their various types play a role in this interaction by either activating or deactivating genes that are associated with obesity. The precise molecular and cellular mechanism behind this process is still not entirely comprehended (36). Previous research has shown that gene-diet interactions have an impact on metabolic factors, and our study's findings are consistent with this. A cross-sectional study found a relationship between the dietary inflammatory index (DII) and the rs17782313 mutation, which influences body composition (61). Hianza et al. conducted a study on individuals with a genetic predisposition to obesity and discovered that adopting a healthy diet led to a decrease in risk factors associated with CVD (62).

A study conducted by Aali et al. has significantly validated the findings of our study. Both studies have found a negative correlation between the consumption of polyphenols and its various forms with indicators of body composition (such as WC, WHR, WHtR) and metabolic parameters such as glycemia (FBG, HOMA-IR) and lipids (CHOL, TG). Also, Aali's study has reported a positive correlation with HDL cholesterol. However, our study did not find any significant correlation between the consumption of lignans and stilbenes with body composition and CVD risk factors, Aali's study did report such a correlation, which is the only point of difference between the two studies (50).

Numerous studies have confirmed that GRS increases the risk of CVD (9), the exact way in which food compounds like polyphenols

can mitigate this effect remains unclear. The statistical power of the analysis can be affected by factors such as sample size, food components used, genetic variations, and gender, which can cause discrepancies in study findings. Evidence shows that polyphenols and its types reduce TG accumulation, increase lipolysis, decrease lipogenesis, and increase energy consumption, which may be an acceptable reason to justify the negative interaction between the consumption of polyphenols and GRS on body composition and some of the risk factors of CVD (63). Polyphenols play a crucial role in regulating blood sugar levels and improving the body's ability to respond to insulin by decreasing the production of specific hormones (64). Moreover, these compounds have the ability to control the process of lipolysis by activating hormone-sensitive lipase (65). Polyphenols enhance the body's antioxidant system, reduce fat oxidation, and enhance the activity of antioxidant enzymes (66, 67). Therefore, it is anticipated that enhancing the consumption of a diet rich in polyphenols and their various forms will result in enhanced body composition and a reduction in certain risk factors associated with CVD (68). These results are confirmed by our study.

The present study is one of the first studies to examine the interaction of polyphenol intake with GRS on metabolic parameters, which is one of the strengths of this study. Some limitations of this research include the cross-sectional design of the study, the lack of investigation into causal interactions, the use of memory-based tools such as FFQ, and the inability to generalize the results to men's gender.

Conclusion

Our findings indicate that individuals who consume high GRS and polyphenols have a significant negative effect on FBS and TG levels, as well as a significant positive effect on HDL. Therefore, a high intake of polyphenols in individuals with high GRS may have a protective effect on cardiometabolic risk. This finding indicates that the interaction of dietary components, such as polyphenols, with genetic risk factors over cardiometabolic risk factors is of great

TABLE 5 The interaction between GRS, flavonoids and polyphenol on cardiometabolic risk factors.

Variables	GRS				Flavonoid	S					Polyp	henol			
		Low		Higl	h intake \geq	81.55(mg/	'day)		Low		High int	ake ≥62	2.59(mg	/day)	
		intake < 81.55		Crude			Adjust		intake < 62.59		Crude			Adjust	
		(mg/ day)	β	95% CI	p	β	95% CI	p	(mg/day)	β	95% CI	p	β	95% CI	р
Anthropometri	c factors														
	Low risk	Reference			Refe	rence			Reference			Referei	nce		
BMI (kg/m ²)	Moderate risk	-	-3.46	-5.44 to -1.49	<0.001	-2.56	-4.59 to -0.52	0.01	-	0.26	-1.7 to 2.23	0.79	0.43	-1.56 to 2.44	0.66
	High risk	-	-0.91	-3.84 to 2.02	0.54	-0.27	-3.28 to 2.73	0.85	-	-0.66	-3.66 to 2.34	0.66	0.62	-2.45 to 3.69	0.69
	Low risk	Reference			Refe	rence			Reference			Referen	nce		
WC (cm)	Moderate risk	-	-7.43	-12.48 to -2.39	<0.001	-5.67	-10.79 to -0.56	0.03	-	0.82	-4.17 to 5.81	0.74	2.53	-2.48 to 7.54	0.32
	High risk	-	-0.82	-8.22 to 6.58	0.82	0.65	-6.78 to 8.08	0.86	-	-3.08	-10.67 to 4.5	0.42	1.60	-6.04 to 9.26	0.68
	Low risk	Reference		Reference								Referen	nce		
WHR	Moderate risk	-	-0.01	-0.043 to 0.013	0.29	<0.001	-0.033 to 0.023	0.71	-	0.00	-0.02 to 0.03	0.85	0.01	-0.014 to 0.041	0.34
	High risk	_	0.02	-0.016 to 0.066	0.23	0.03	-0.007 to 0.075	0.1	-	<0.001	-0.051 to 0.032	0.65	0.01	-0.023 to 0.061	0.37
Body composit	tion					1									
	Low risk	Reference			Refe	rence			Reference			Referei	nce		
VFL (cm ²)	Moderate risk	-	-2.21	-3.96 to -0.46	0.01	-1.61	-3.4 to 0.17	0.07	-	0.10	-1.63 to 1.83	0.91	0.87	-0.87 to 2.63	0.32
	High risk	-	0.57	-2 to 3.15	0.66	1.44	-1.15 to 4.05	0.27	-	-0.00	-2.65 to 2.64	0.99	1.67	-1.01 to 4.36	0.22
	Low risk	Reference	e Reference Reference Re						Referen	nce					
OD (%)	Moderate risk	-	-13.04	-24.62 to -1.47	0.02	-11.20	-22.67 to 0.27	0.05	-	-4.03	-15.45 to 7.38	0.48	0.85	-10.48 to 12.18	0.88
	High risk	-	-3.69	-20.75 to 13.36	0.67	-1.25	-17.99 to 15.47	0.88	-	-8.65	-26.06 to 8.76	0.33	-0.90	-18.24 to 16.43	0.91
Biochemical va	ariables	·	·			·					·				
	Low risk	Reference			Refe	rence			Reference			Referen	nce		
FBS (mg/dL)	Moderate risk	-	-0.85	-6.43 to 4.72	0.76	0.07	-5.82 to 5.97	0.98	-	-0.09	-5.53 to 5.34	0.97	0.86	-4.82 to 6.56	0.76
	High risk	-	-3.58	-11.92 to 4.75	0.4	-2.92	-11.71 to 5.86	0.51	-	-0.76	-9.2 to 7.68	0.86	-0.12	-9.15 to 8.91	0.97

(Continued)

Variables	GRS			l de la companya de l	Flavonoid	s			Polyphenol							
		Low		Hig	h intake \geq	81.55(mg	/day)		Low		High int	ake ≥6	2.59(mg	/day)		
		intake < 81.55		Crude			Adjust		intake < 62.59		Crude			Adjust		
		(mg/ day)	β	95% CI	p	β	95% CI	р	(mg/day)	β	95% CI	p	β	95% CI	p	
	Low risk	Reference			Refe	erence			Reference			Refere	nce			
TC (mg/dL)	Moderate risk	_	-13.03	-34.14 to 8.08	0.22	-7.61	-29.5 to 14.26	0.49	-	-3.11	-23.63 to 17.39	0.76	2.69	-18.31 to 23.71	0.8	
	High risk	-	-16.48	-48.02 to 15.06	0.3	-4.17	-36.79 to 28.43	0.8	-	-10.82	-42.67 to 21.02	0.5	-11.42	-44.78 to 21.93	0.	
TG (mg/dL)	Low risk	Reference			Refe	rence			Reference			Refere	nce			
	Moderate risk	-	-5.96	-41.06 to 29.12	0.73	2.96	-34.27 to 40.21	0.87	-	4.04	-30.93 to 39.01	0.82	9.17	-27.45 to 45.79	0.6	
	High risk	-	-84.30	-137.58 to -31.02	<0.001	-79.11	-135.52 to -22.69	<0.001	-	8.23	-47.21 to 63.68	0.77	15.75	-44.36 to 75.86	0.0	
HDL (mg/dL)	Low risk	Reference	Reference								·	Refere	nce			
	Moderate risk	-	1.75	-4.72 to 8.22	0.59	3.94	-2.82 to 10.71	0.25	-	-4.03	-10.35 to 2.28	0.21	-2.17	-8.8 to 4.44	0.5	
	High risk	-	8.39	-1.28 to 18.06	0.08	12.56	2.47-22.65	0.01	-	-7.74	-17.56 to 2.06	0.12	-7.31	-17.82 to 3.19	0.1	
LDL (mg/dL)	Low risk	Reference			Refe	rence			Reference			Refere	nce			
	Moderate risk	-	-2.68	-16.77 to 11.39	0.7	3.94	-10.74 to 18.63	0.59	-	0.95	-10.35 to 2.28	0.21	3.46	-8.8 to 4.44	0.5	
	High Risk	-	-11.13	-32.18 to 9.91	0.3	-4.83	-26.72 to 17.06	0.66	-	-16.77	-17.56 to 2.06	0.12	-22.00	-17.82 to 3.19	0.1	
AST (IU/L)	Low risk	Reference			Refe	erence			Reference			Refere	nce			
	Moderate risk	-	-2.14	-6.57 to 2.28	0.34	-1.86	-6.51 to 2.79	0.43	-	0.70	-3.61 to 5.03	0.74	1.15	-3.36 to 5.67	0.6	
	High risk	-	-0.78	-5.83 to 7.4	0.81	2.35	-4.58 to 9.29	0.5	-	-0.79	-7.5 to 5.9	0.81	1.25	-5.91 to 8.42	0.7	
ALT (IU/L)	Low risk	Reference			Refe	erence			Reference	Reference						
	Moderate risk	-	-7.02	-14.74 to 0.69	0.07	-5.31	-13.58 to 2.96	0.2	-	0.83	-6.77 to 8.43	0.83	2.34	-5.72 to 10.42	0.5	
:	High risk	-	3.75	-7.78 to 15.28	0.52	6.04	-6.28 to 18.37	0.33	-	-5.77	-17.58 to 6.02	0.33	-2.46	-15.27 to 10.34	0.3	

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B, Standard Error; GRS, Genetic risk score; BMI, Body mass index; WC, waist circumference; WHR, waist height ratio; FBS, fasting blood sugar; TG, Triglyceride; LDL, Low density lipoprotein; HDL, High density lipoprotein; OD, Obesity degree; VFL, Visceral fat level. Flavonoids High intake \geq 81.55(mg/day), Polyphenol High intake \geq 62.59(mg/day).

p-values < 0.05 are in bold.

Adjust = adjusted for potential confounding factors including (age, IPAC and total energy intake).

Low Risk: 0,1,2 Risk allele, Moderate Risk: 3,4 Risk allele, High Risk: 5,6 Risk allele.



importance. Further research is necessary in the future to validate this association.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Ethics statement

The studies involving humans were approved by the authors and the Ethics Committee of Tehran University of Medical Sciences (IR. TUMS.MEDICINE.REC.1402.636) have received approval for the method and protocol of the current study. The studies were conducted in accordance with the local legislation and institutional requirements. The participants provided their written informed consent to participate in this study.

Author contributions

ZR: Writing – original draft, Writing – review & editing. AM: Investigation, Validation, Writing – review & editing. FA: Data curation, Formal analysis, Methodology, Software, Writing – review & editing. SD: Conceptualization, Writing – original draft. YA: Methodology, Writing – review & editing. KM: Funding acquisition, Visualization, Writing – review & editing.

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Conflict of interest

The authors 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.

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Glossary

ALT Alamine aminotransferase AST Asparate aminotransferase CVD Cardiovascular diseases BMI Body mass indec WC Waist circumference DPI Dietary polyphenols intake FBS Fasting blood glacose FRG Fasting plasma glacose MC4R Melanocortin 4 receptor Cry1 Cryptochronne 1 gene CAV1 Carcolin-1 FFQ Food frequency questionnaire DQ1 diet quality index PA Physical activity questionnaire DQ1 diet quality index PA Physical activity questionnaire-short form HC Hp circumference WHR Waits-to-hip ratio VFL Visceral fat level OD Obesity degree GOD-PAP Glicoce oxidase-phenol 4-aminoantityrine peroxidase GPOPAP Glicoce oxidase-phenol 4-aminoantityrine peroxidase TG Trighyeeride HDL Low-density lipoprotein LDL Low-density lipoprotein SIAP Systolic blood pressure GRS Genetic risk score TC Total cholesterol TDPI Total deletary polyphenols index SIM		
CVDCardiovascular diseasesBM1Body mass indexWCWaist circumfrenceDP1Detary polyhenols intakeFBSFasting blood glucoseFBGFasting blood glucoseGK4RMelancortin 4 receptorCry1Cryptochrome 1 geneCAV1Careolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activity questionnaire -ahort formIFQInternational physical activity questionnaire -ahort formIFQHep circumferenceWHRWaist to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase phenol 4-aminoantipyrine peroxidaseIFQTrigbycroteinIDLInder dust phoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal chelestrolTDPITotal dietary polyhenols indexSIAMSoft kan massFFMFat-free mass	ALT	Alanine aminotransferase
BMI Body mass index WC Waist circumference DPI Dietary polyphenols intake FBS Fasting blood glucose FBG Fasting plasma glucose MC4R Melancortin 4 receptor CAR Melancortin 4 receptor CAVI Cryptochrome 1 gene CAVI Caveolin-1 FFQ Food frequency questionmaire DQI diet quality index PA Physical activity IPAQ International physical activity questionnaire short form IIC Ip circumference WHR Waist ch-bip ratio VFL Vasceral fat level OD Obesity degree GOD-PAP Glucose oxidase-phenol 4-aminoantipyrine peroxidase TG Triglyceride IDL Low-density lipoprotein LDL Low-density lipoprotein LDL Low-density lipoprotein SBP Systoic blood pressure GRS Genetic risk score TC Total dictary polyphenols index SIM <td>AST</td> <td>Aspartate aminotransferase</td>	AST	Aspartate aminotransferase
WCWalst circumferenceDPIDietary polyphenols intakeFISFasting blood glucoseFFGFasting plasma glucoseMCRMelanocortin 4 receptorCry1Crytochrome 1 geneCAV1Caveolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLViscar fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinDRSystolic blood pressureDBPDiabicic blood pressureCRGenet risk scoreTCTotal dietary polyphenols indexSIMSoft lean massFPASoft lean mass	CVD	Cardiovascular diseases
DPIDietary polyphenols intakeFBSFasting blasma glucoseFPGFasting plasma glucoseMC4RMelanocortin 4 receptorCry1Cryptochrome 1 geneCAV1Caweolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activityIFAQInternational physical activity questionnaire-short formHCHp circuniferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSPPDiatolic blood pressureGRSGenetic risk scoreTCTotal cholesterolFIMSof lean massFEMSof lean mass	BMI	Body mass index
PBS Fasting blood glucose FPG Fasting plasma glucose MC4R Melanocortin 4 receptor Cry1 Cryptochrome 1 gene CAV1 Caveolin-1 FFQ Food frequency questionnaire DQI diet quality index PA Physical activity IPAQ International physical activity questionnaire-short form HC Hp circumference WHR Wais-to-hip ratio VFL Visceral fat level OD Obesity degree GOD-PAP Glucase oxidase-phenol 4-aminoantipyrine peroxidase GPOPAP Glycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidase TG Triglyceride HDL High-density lipoprotein LDL Low-density lipoprotein SBP Systolic blood pressure GRS Genetic risk score TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass	WC	Waist circumference
PFGFasting plasma glucoseMC4RMelanocortin 4 receptorCry1Cryptochrome 1 geneCAV1Caveolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWais-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseTGTrigbyceridHDLHigh-density lipoproteinLDLLow-density lipoproteinBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal cholesterolFMMSofi lean massFFMFat-free mass	DPI	Dietary polyphenols intake
McARMelancortin 4 receptorCry1Cryptochrome 1 geneCAV1Caveolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTrigbycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinBPDiastolic blood pressureGRSGenetic risk scoreTCTotal dietary polyphenols indexSLMSoft lean massFFMFat. free mass	FBS	Fasting blood glucose
CrylCryptochrome l geneCAV1Caveolin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	FPG	Fasting plasma glucose
CAV1Cavedin-1FFQFood frequency questionnaireDQ1diet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	MC4R	Melanocortin 4 receptor
FFQFood frequency questionnaireDQIdiet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexFMSoft lean mass	Cryl	Cryptochrome 1 gene
DQIdiet quality indexPAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	CAV1	Caveolin-1
PAPhysical activityIPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	FFQ	Food frequency questionnaire
IPAQInternational physical activity questionnaire-short formHCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	DQI	diet quality index
HCHp circumferenceWHRWaist-to-hip ratioVFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	РА	Physical activity
WHRWaist-to-hip ratioWFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	IPAQ	International physical activity questionnaire-short form
VFLVisceral fat levelODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFMFat-free mass	НС	Hp circumference
ODObesity degreeGOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFMFat-free mass	WHR	Waist-to-hip ratio
GOD-PAPGlucose oxidase-phenol 4-aminoantipyrine peroxidaseGPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFarfree mass	VFL	Visceral fat level
GPOPAPGlycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidaseTGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	OD	Obesity degree
TGTriglycerideHDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	GOD-PAP	Glucose oxidase-phenol 4-aminoantipyrine peroxidase
HDLHigh-density lipoproteinLDLLow-density lipoproteinSBPSystolic blood pressureDBPDiastolic blood pressureGRSGenetic risk scoreTCTotal cholesterolTDPITotal dietary polyphenols indexSLMSoft lean massFFMFat-free mass	GPOPAP	Glycerol-3-phosphate oxidase-phenol 4-aminoantipyrine peroxidase
LDL Low-density lipoprotein SBP Systolic blood pressure DBP Diastolic blood pressure GRS Genetic risk score TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	TG	Triglyceride
SBP Systolic blood pressure DBP Diastolic blood pressure GRS Genetic risk score TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	HDL	High-density lipoprotein
DBP Diastolic blood pressure GRS Genetic risk score TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	LDL	Low-density lipoprotein
GRS Genetic risk score TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	SBP	Systolic blood pressure
TC Total cholesterol TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	DBP	Diastolic blood pressure
TDPI Total dietary polyphenols index SLM Soft lean mass FFM Fat-free mass	GRS	Genetic risk score
SLM Soft lean mass FFM Fat-free mass	TC	Total cholesterol
FFM Fat-free mass	TDPI	Total dietary polyphenols index
	SLM	Soft lean mass
SMM Skeletal muscle mass	FFM	Fat-free mass
	SMM	Skeletal muscle mass
DII Dietary inflammatory index	DII	Dietary inflammatory index