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OPEN The association of dietary patterns with dietary inflammatory index, systemic inflammation, and insulin resistance, in apparently healthy individuals with obesity

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Inflammation is considered a key mechanism leading to obesity. Dietary patterns and certain food items influence inflammation. Few studies have investigated the contribution of major dietary patterns to biological measures of inflammation. Therefore, the present study aimed to examine the associations of different dietary patterns with dietary inflammatory index (DII), systemic inflammation, and insulin resistance (IR) in the apparently healthy obese. In this cross-sectional study, 151 abdominally obese subjects were recruited from the Northwest of Iran. Dietary intake, demographic data, anthropometric indices, and physical activity (PA) was assessed. DII scores were calculated based on a validated 168-item food frequency questionnaire (FFQ). Three dietary patterns were identified, using principal component analysis. Basal blood samples were collected to determine biochemical parameters. Linear regression test with adjusted beta estimates was applied for data analysis. Three dietary patterns were extracted as Healthy, Western, and Traditional. Body mass index (BMI) (p < 0.01) and fat mass (p < 0.001) were directly associated with the Western dietary pattern. Conversely, serum lipopolysaccharide-binding protein (LBP) (b = -0.1, p < 0.04) was negatively associated with Healthy dietary pattern, after controlling for confounders. The Traditional pattern was found to be inversely related to DII (b = -0.3, p < 0.001). The association was also reveresed between Traditional pattern and IR (Odds Ratio: 0.3 (95% Confidence Interval 0.1-0.9)). The results suggested that the Western dietary pattern was related to higher BMI and fat mass. In addition, the Healthy pattern was associated with decreased levels of LBP. Adherence to the Traditional dietary pattern was inversely related to DII as well as IR.

Abbreviations

Body mass index BMI CVD Cardiovascular disease DII Dietary inflammatory index

ELISA Enzyme-Linked Immunosorbent Assay

FFQ Food frequency questionnaire

HC Hip circumference

HDL-C High-density lipoprotein-cholesterol

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HOMA-IR Homeostasis model assessment of insulin resistance

IL18 Interleukin 18

IPAQ International Physical Activity Questionnaire

IR Insulin resistance

LBP Lipopolysaccharide-binding protein

LPS Lipopolysaccharide **MUFA** Monounsaturated fatty acid Metabolic Syndrome MetS NFκB Nuclear factor-κB Polyunsaturated fatty acid **PUFA** PA Physical activity SFA Saturated fatty acid TLR Toll-like receptor

TNF-α Tumor necrosis factor-alpha

WC Waist Circumference

Obesity is a major public health concern, affecting more than half a billion people in the world¹. It emerged as a threat in more wealthy countries nearly four decades ago². Globally, the pandemic of overweight and obesity in adults (i.e., those with body mass index (BMI) of \geq 25 kg/m²) has increased from 28.8 to 36.9% in men and from 29.8 to 38.0% in women during the years $1980-2013^3$. Over half of the Iranian adult population are overweight or obese. The prevalence of obesity and abdominal obesity increased from 23.1% and 47.9% in 1999 to 34.1% and 71.1% in 2011, respectively^{4,5}.

Risk factors associated with obesity are multiple and complex. The most closely related adverse health consequences include type 2 diabetes mellitus (T2DM), dyslipidemia, cardiovascular diseases (CVD), and insulin resistance (IR)⁶. There is a strong direct association between obesity and IR in nondiabetic subjects. Obesity also increases IR in diabetic subjects and may lead to enhanced acute-phase response⁷. Overweight and obesity are characterized by chronic low-grade inflammation that results in the secretion of pro-inflammatory factors engaged in the pathogenesis of IR⁸. In addition, an abundance of adipose mass can result in the secretion of inflammatory adipokines⁹.

Chemerin is a novel inflammatory adipokine that has a regulatory role in lipid and glucose homeostasis¹⁰. Moreover, there is a positive correlation between plasma concentration of chemerin and body mass index (BMI), triglycerides, and blood pressure¹¹. A body of evidence shows that chemerin concentration is increased in overweight and obese persons. These conditions exhibit a positive correlation with various aspects of metabolic syndrome.

Omentin, a 38–40 kDa adipokine, was observed from a cDNA in visceral omental adipose tissue¹². Serum omentin levels may play a key role in the pathogenesis of IR and diabetes¹³. Omentin can increase insulin sensitivity and stimulate glucose transport and Akt phosphorylation in human adipocytes¹⁴. Furthermore, some studies have demonstrated that circulating omentin was most notably decreased in obesity¹⁵.

Lipopolysaccharide binding protein (LBP), as an acute-phase protein, is a 50 kDa polypeptide that is commonly synthesized in the liver¹⁶. Serum LBP level is a marker of coronary artery disease¹⁷. It is reported to be associated with obesity and related disorders in apparently healthy people¹⁸. Lipopolysaccharide stimulates the release of several cytokines that are key inducers of IR which is a putative factor for triggering metabolic disturbances¹⁹. A recent study indicated that increased metabolic concentrations of plasma LPS are sufficient molecular mechanisms for the development of IR, obesity, and T2DM²⁰.

Diet is an important predictor of circulating levels of inflammatory markers²¹. Diets rich in pro-inflammatory constituents such as saturated fatty acids (SFAs) and trans fatty acids have consistently been associated with proliferation²² and oxidative stress that can promote inflammation²³. By contrast, polyunsaturated fatty acids (PUFAs), monounsaturated fatty acids (MUFAs), and fiber have been shown to attenuate the inflammatory cascade²⁴. Recently, a survey on the inflammatory potential of diet and its influence on obesity and chronic diseases has received special attention²⁵.

The dietary inflammatory index (DII) is a novel scoring algorithm that provides an estimate of the inflammatory potential of the overall diet based on the inflammatory properties of dietary constituents²⁶. Food pattern analysis is a way to investigate the relationship between diet and risk of chronic diseases²⁷. Currently, few studies have studied the association between dietary patterns and DII. Dietary patterns with more protein, specifically animal protein may also aggravate glucose metabolism, leading to the development of IR²⁸. Moreover, a body of evidence shows that certain dietary patterns have also been associated with the markers of inflammation²⁹. A cross-sectional study of the Hispanic elderly living in Massachusetts reported lower concentrations of CRP with higher fruit and vegetable consumption³⁰. To our knowledge, there exists no study that evaluates the association of different dietary patterns with DII, systemic inflammation, and IR in apparently healthy obese individuals in the Middle East. Therefore, the present study was aimed to identify the dietary patterns of apparently healthy obese and to determine their association with DII as well as measures of systemic inflammation and IR in Iran.

Materials and methods

Sampling. The present cross-sectional study was conducted from June to November 2015 in Tabriz, Iran. Five-hundred volunteers were randomly invited from the general population. Of these, 151 apparently healthy obese (BMI≥30 kg/m²) people (aged 18–60 years) were consecutively enrolled, based on defined eligibility criteria³1,32. Exclusion criteria included having any chronic disease such as hypertension, diabetes, CVD, hepatic disorders, renal diseaseor gastrointestinal disease; surgery during the past year; any current cancer, infectious

disease, or severe mental illness. Moreover, those on a diet or who misreported dietary intake (< 800 kcal/day or > 4200 kcal/day) or were taking any medications that could influence lipid and glucose metabolism, corticosteroids, contraceptives, antibiotics, anti-obesity drugs, anti-inflammatory drugs, beta-blockers, anti-coagulants or any diettary supplements including multivitamins, multiminerals, antioxidants, herbal supplements, and omega 3 or omega 6 fatty acids in the last two months were also excluded. The study was approved by the ethics committee of Tabriz University of Medical Sciences, Tabriz, Iran. Informed consent was obtained from all the atudy individuals. The whole research was performed in compliance with the Declaration of Helsinki (Ethical code: TBZMED.REC.1395.1169).

Measurements. Height, weight, WC, and hip circumference (HC) were measured, while the participant wore clothes without shoes. Body mass index (BMI) was estimated as weight in kilograms divided by the square of height in meters. WC was measured at the midpoint between the lower ribs and the iliac crest and HC was measured at the level of the maximum extension of the buttocks. WHR was calculated as WC divided by HC³³. Physical activity (PA) data was extracted from long-form International Physical Activity Questionnaire (IPAQ)³⁴. PA was expressed as metabolic equivalents (Mets)-h per week.

Laboratory assays. Blood samples were drawn after an overnight fast at the beginning of the study. The serum was centrifuged at 3000 rpm for 15 min. Lipid profile and blood sugar were calculated immediately, using the commercial kits (Pars Azmoon, Tehran, Iran) and a Selectra 2 auto-analyzer (Vital Scientific, Spankeren, the Netherlands). Inter-assay and intra-assay coefficients of variation (CVs) were < 5% for all assays. Serum samples were stored at – 80 °C, until analysis. Sandwich Enzyme-Linked Immunosorbent Assay (9) was used to measure serum insulin, LBP, chemerin, and omentin concentrations (Monobind Inc., Lake Forest, CA, USA). Other parameters were measured according to the manufacturer's instructions (Bioassay Technology Laboratory, Shanghai Korean Biotech, Shanghai City, China). The intra-assay and inter-assay CVs were < 8% and 10% for chemerin, omentin, and LBP and < 8% and 9.8% for insulin, respectively. The homeostasis model assessment of IR (HOMA-IR) was calculated, using fasting plasma glucose and fasting insulin values, according to the formula.

Assessment of dietary intake. The food intake of the indivuals during last year was assessed, using a validated semi-quantitative food frequency questionnaire (FFQ)³⁴. The FFQ consisted of 168 food items and the participants reported their frequency of intake of each food item on a daily, weekly, monthly, or yearly basis³⁵. Then, daily grams of food intake were estimated from food intake data. The *Nutritionist IV* software (Axxya Systems, Stafford, TX, 1994), modified for Iranian foods was used to calculate the energy and nutrient contents of foods³⁶. Foods from FFQ were categorised into 29 food groups, on the basis of nutrient profile or culinary usage (Appendix 1).

Food patterns, consisting of 29 food groups were analyzed. The principal component analysis was used to identify explanatory factors³⁷. Three dietary patterns were recognized, according to eigen value (>1), scree plot, factor interpretability, and the variance explained (>5%). Factors were rotated with varimax rotation to develop the interpretability of the factors and minimize the correlation between them. Factor loadings are equivalent to the correlation between food items and factors. Higher loadings represent a higher shared variance with the factor. Factor loadings of >0.20 indicated the foods that were significantly related to the identified factor^{37,38}. After recognizing dietary patterns, the participants were assigned, based on their patterns factor score. Factor scores were divided into quartiles in terms of their distribution in each stratum, implying increased intake from tertile 1 to tertile 3.

The DII scores were calculated according to FFQ questionnaire²⁶. Briefly, dietary data for each participant were first linked to a regionally representative global database that provided a robust estimate of means and standard deviations for each of the food parameters considered (i.e. foods, nutrients, and other food components). The 'standard mean' was subtracted from the actual food parameter value and divided by its standard deviation. This z-score was then converted to a percentile (so as to minimize the effect of outliers or right-skewing, a common occurrence with dietary data). This value was then converted to a proportion (values 0–1) and centered by multiplying by 2 and subtracting one. This value was then multiplied by the respective inflammatory effect score of the food parameters (derived from a literature review and scoring of 1943 'qualified' articles) to obtain the subject's food parameter-specific DII score. All of the food parameter-specific DII scores were then summed to create the overall DII score for each subject in the study.

Sample size estimation. Five persons per group are recommended for factor analysis³⁹; hence, 145 people were adequate to be recruited based on 29 food groups in the present study. However, 150 apparently healthy obese were recruited for this purpose.

Statistical analysis. Statistical Package for the Social Sciences (SPSS) software (Version 25, SPSS, Inc., Chicago, IL, USA) was applied to perform data analysis. Twenty-one cases of study participants were excluded because of the missing data; the final analysis included 150 participants. The Kolmogorov- Smirnov test was used to check the normality of data distribution. The characteristics of the participants are expressed as means (standard deviation) for continuous variables and percentages for categorical variables. We used multiple linear regression analysis to find out the association of dietary patterns with systemic inflammatory markers and DII. All models were adjusted for sex, weight, and energy intake (continuous). In these models, scores of the dietary pattern were the independent variable (grouped as tertile of the dietary pattern) and HOMA-IR was dependent variable (IR+ vs. IR-). These models were adjusted for variables found to be significantly associated with IR such as age, sex, energy intake, physical activity, and level of education.

	Healthy				Traditional				Western			
	<47.46	47.47-72.93	>72.93	2 _p	<9.87	9.88-14.64	>14.65	2 _p	<5.20	5.21-12.02	>12.03	2 _p
Age (years)	37 ± 8.5	36 ± 8.0	34±8.7	0.20	37 ± 7.5	38 ± 9.1	37 ± 8.8	0.79	39 ± 8.5	37 ± 8.5	34 ± 7.7	0.01
Sex				< 0.001				0.22				0.53
Male	16 (32%)	24 (49%)	34 (69.4%)		24 (49%)	29 (59.2%)	74 (72%)		22 (44.9%)	28 (56%)	24 (49%)	
Female	34 (68%)	25 (51%)	15 (30.6%)		25 (51%)	20 (40.8%)	29 (28%)		27 (55.1%)	22 (44%)	25 (51%)	
BMI (kg/m²)	31.3 ± 3.7	32.0 ± 5.0	31.8 ± 3.7	0.68	32.0 ± 4.5	31.5 ± 3.6	31.6±4.4	0.86	31.9 ± 4.1	30.4 ± 3.8	32.9 ± 4.3	0.01
Fat Mass	28.3 ± 7.9	28.6 ± 10.3	27.2 ± 7.2	0.68	28.0 ± 9.1	27.5 ± 8.1	28.6 ± 8.4	0.81	28.3 ± 8.3	25.4 ± 8.3	30.5 ± 8.4	0.001
PA (MET-h/d)				0.29				0.22				0.10
< 600	2 (4.2%)	8 (18.2%)	6 (12.5%)		2 (4.3%)	8 (17.8%)	6 (12.2%)		4 ((8.9%)	2 (4.3%)	10 (20.4%)	
600-3000	22 (45.8%)	19 (43.2%)	19 (39.6%)		18 (39.1%)	19 (42.2%)	23 (46.9%)		18 (40%)	24 (52.2%)	18 (36.7%)	
> 3000	24 (50%)	17 (38.6%)	23 (47.9%)		26 (56.5%)	18 (40.0%)	20 (40.8%)		23 (51.1%)	20 (43.5%)	21 (42.9%)	
Education				0.99				0.46				0.40
Middle school	13 (26%)	12 (24.5%)	12 (24.5%)		13 (26.5%)	15 (30.6%)	9 (18%)		9 (18.1%)	16 (32%)	12 (24.5%)	
High school	23 (46%)	23 (46.9%)	23 (46.9%)		20 (40.8%)	21 (42.9%)	28 (56%)		26 (53.1%)	23 (46%)	120 (40.8%)	
University & Higher education	14 (28%)	14 (28.6%)	14 (28.6%)		16 (32.7%)	13 (26.5%)	13 (26%)		14 (28.6%)	11 (22%)	17 (24.7%)	
Fat% (%E)	27.9 ± 9.2	27.0 ± 10.9	24.8 ± 8.1	0.28	25.9 ± 8.0	26.2 ± 10.6	227.3 ± 9.5	0.79	25.0 ± 10.54	26.2 ± 7.7	28.5 ± 9.5	0.22
CHO% (%E)	59.0 ± 9.1	59.1 ± 10.3	62.2 ± 9.2	0.45	61.1 ± 8.4	61.1 ± 10.5	59.2 ± 9.3	0.75	63.3 ± 9.2	59.1 ± 8.1	57.2 ± 9.4	0.02
Pro% (%E)	14.2 ± 3.3	13.7 ± 2.8	14.8 ± 3.9	0.40	14.6 ± 3.2	14.3 ± 3.8	13.9 ± 3.2	0.67	13.4 ± 3.1	14.5 ± 2.9	15 ± 4.1	0.09
Energy intake (kcal/day)	2399 ± 803	3030±966	3562 ± 677	0.001	2389 ± 734	2999±915	3591 ± 79	0.001	2453±775	3107 ± 939	3410 ± 876	0.001

Table 1. Descriptive characteristics of the study population and their association with tertile of dietary patterns. *%E* percent of energy. Data are presented as means Sd for continuous variables and n (%) foe categorical variables. ^aANOVA for quantitative variables. ^bChi-square for qualitative variables.

Ethical approval. Dr. James R. Hébert owns a controlling interest in Connecting Health Innovations LLC (CHI), a company that has licensed the right to his invention of the dietary inflammatory index (DII) from the University of South Carolina in order to develop computer and smartphone applications for patient counseling and dietary intervention in clinical settings. The subject matter of this paper will not have any direct bearing on that work, nor has that activity exerted any influence on this project. Dr. Nitin Shivappa is an employee of CHI.

Results

Three major patterns were identified, using factor analysis (Appendix 2): the Healthy dietary pattern (higher intakes of yellow vegetables including carrots, as row orboiled, nuts, unsweetened drinks, tomato, grains (including White breads (lavash, baguette), whole breads (sangak, barbari, taftoon), noodles, pasta, rice, biscuit, barley, bulgur, cornflakes, olive, low-fat dairy, eggs, fish, legumes, potato, and dry fruits); the Western dietary pattern (higher intakes of mayonnaise, high-fat dairy, soft drinks, butters, fruit juices, French fries, pizza, sweets, processed meat, and pickles); and the Traditional dietary pattern (higher intakes of poultry, organ meats, fruits (whole, juice, or puree), green vegetables (including green leafy vegetables such as lettuce, spinach, as row or boiled), other vegetables (including Cabbage, cauliflower, Brussels sprouts, kale, cucumbers, mixed vegetable, eggplant, celery, green peas, green beans, green peppers, corn, turnips, squash, mushrooms, onions, and garlic), red meats, and hydrogenated fat. Together, these factors explained 30% of the variance.

Characteristics of the study participants across tertile categories of food patterns scores are presented in Table 1. Compared with participants in the lower tertile, those in the upper tertile of the Healthy and Traditional dietary patterns had lower energy intake. Unlike females, male patients presented higher adherence to the upper tertile of the Healthy pattern. Those in the upper tertile of the Western dietary pattern had higher BMI, fat mass, energy intake, and lower carbohydrate intake and had a lower age.

The results of multiple linear regression models are shown in Table 2. The Healthy pattern score was inversely related to plasma concentration of LBP, after controlling for potential confounders. The Traditional pattern score was negatively associated with DII when controlled for confounders including sex, weight, and energy intake.

The results of logistic regression analysis indicated that the highest tertile of the Traditional dietary pattern was associated with approximately 70% decreased odds of IR, as indicated by HOMA-IR (OR: 0.32, 95% CI 0.11–0.93) (Table 3). The Western or Healthy dietary pattern showed no significant association with the odds of IR in the study population.

Discussion

In the present study, three major dietary patterns were identified, as follows: Healthy, Western, and Traditional. Of these, the Healthy pattern was inversely associated with serum LBP concentration. Also, a positive association between the Western diet and BMI as well as fat mass was found. The Traditional pattern was inversely associated with IR and DII.

	Healthy		Traditional		Western	
Inflammatory markers and DII score	β	p	β	p	β	p
Omentin (ng/mL)	0.10	0.28	-0.03	0.78	0.05	0.54
Chemerin (ng/mL)	0.001	0.84	-0.001	0.99	-0.15	0.09
Insulin (mg/dL)	0.18	0.48	-0.13	0.19	-0.11	0.20
LBP (µg/mL)	-0.18	0.04	0.05	0.62	0.12	0.16
DII score	-0.13	0.15	-0.37	0.001	-0.11	0.19

Table 2. Linear regression analysis for the association between dietary patterns and circulating concentrations of inflammatory markers and DII. The model was adjusted for energy intake, weight, and sex. *LBP* lipopolysaccharide binding protein, *DII* dietary inflammatory index.

Tertiles of dietary patterns	Unadjusted model OR; 95% CI	Adjusted model OR; 95% CI					
Healthy							
1	Ref	Ref					
2	1.5 (0.6–3.4)	2.1 (0.8–5.3)					
3	1.5 (0.7–3.5)	2.3 (0.8-6.5)					
Traditional							
1	Ref	Ref					
2	1 (0.4–2.3)	0.6 (0.2–1.7)					
3	0.7 (0.3–1.6)	0.3 (0.1-0.9)					
Western							
1	Ref	Ref					
2	0.9 (0.4-2)	0.9 (0.3–2.3)					
3	0.9 (0.4-2.1)	0.5 (0.1–1.3)					

Table 3. Adjusted logistic regression model describing the association between various dietary patterns and insulin resistance in the study population. Adjusted for age, sex, energy intake, physical activity, and education.

The present work showed that LBP was the only biomarker to appear inversely associated with a Healthy dietary pattern; i.e. serum LBP was lower in those with Healthy dietary pattern. The finding for LBP is consistent with the notion that Healthy diets afford protective effects from diverse perspectives. One plausible mechanism is through a TLR4-dependent pathway, implicated in the launch of the transcription factor, nuclear factor- κ B (NF κ B)⁴⁰. This signaling cascade results in the enhanced release of pro-inflammatory cytokines, including tumor necrosis factor-alpha (TNF- α) or IL-6, thus promoting systemic inflammation⁴¹. Endotoxin levels have indicated a direct correlation with IR and chemerin and an inverse association with omentin⁴². The negative correlation between serum LBP and chemerin was also observed in our previous study²⁵.

Other studies reported a link between a high-fat diet especially saturated fatty acids and the release of endotoxins in the plasma of mice and humans^{21–43}. In this regard, López-Moreno et al. ⁴³ showed that consumption of diets with highly-saturated fatty acids (HSFAs) increases the intestinal absorption of LPS which, in turn, increases postprandial endotoxemia levels and the postprandial inflammatory response. In fact, chronic consumption of a high-fat diet especially rich in SFAs as well as obesity could increase endotoxemia and low-grade inflammation due to the repeated endotoxin absorption from the gut through the digestion of lipids which, in turn, could raise the risk of IR and atherosclerosis⁴⁴. It was also reported that fruits intake and a fiber-rich meal reduce postprandial LPS levels, as compared with a high-fat, especially saturated, meal in human. Moreover, it was shown that the n-3 fatty acid-rich test meal reduces postprandial serum endotoxin, compared to the meal rich in saturated fatty acid⁴³. However, in the present study, those with the Healthy dietary pattern had a lower intake of fat (i.e. saturated and trans fatty acids as well as dairy products) and subsequently, lower levels of serum LBP. No other correlation was found between the Healthy dietary pattern and other biomarkers.

Our findings also confirm a contrary association between the Traditional pattern and DII. It was also identified that those in the highest tertile of the Traditional diet had approximately 70% lower HOMA-IR scores, compared to the individuals in the lower tertile. The Traditional diet mainly included poultry, fruit, green leafy vegetables, red meat, organ meat, and hydrogenated fat which taken together could be responsible for the lower IR and DII in the study population. Higher consumption of fruit and vegetables in the Traditional diet could be considered as the key factor causing the inverse association between the Traditional dietary pattern, DII, and IR. The main ingredients of fruit and vegetables are dietary fiber, vitamin E, folate, and magnesium which are independently associated with reduced metabolic risks⁴⁴. Reduced insulin demand may be another protective mechanism associated with higher intakes of these foods. In general, due to physical properties and viscous structures of fibers, such foods are primarily non-digestible carbohydrates and absorbed more slowly; thus, they have relatively low glycemic indices.

Regarding the consumption of meat, a cross-sectional analysis of data from 3690 diabetes-free female participants in the Nurses' Health study found that a greater total, unprocessed, and processed red meat intakes were associated with higher inflammatiry biomarkers. Substitution of a serving of total red meat intake with alternative protein food consumed in a combination of poultry, fish, legumes, and nuts was associated with a healthier biomarkers profile of inflammation⁴⁵. However, a further cross-sectional study demonstrated that the relationship between red meat intake and inflammatory biomarkers was no longer observed after adjustment for BMI⁴⁶. It was found that a pro-inflammatory diet, defined as a higher DII score, is cheifly rich in some inflammatory ingredients such as SFA, and comparatively, poor in anti-inflammatory parameters including fibers, MUFA, and PUFA²⁵.

Our previous work indicated that participants with higher DII score had lower intake of polyunsaturated and monounsaturated fats and fiber as well as higher saturated fats²⁵. Similarly, Shivappa et al. reported a negative association between lower DII score and consumption of healthy foods and nutrients⁴⁷. Shivappa et al. ⁴⁷ also illustrated that a diet rich in pro-inflammatory food parameters such as saturated fatty acids and moderately poor in anti-inflammatory food ingredients such as fruit and vegetables can increase inflammatory biomarkers, as evidenced by the increased levels of IL-6 and homocysteine. In line with our research, another study suggested that a food pattern rich in fiber was indirectly associated with hyperinsulinemia in women, but not in men⁴⁸. In the Health Professionals Follow-up Study, a prudent dietary pattern full of fruits, vegetables, whole grains, and poultry was inversely associated with insulin levels⁴⁹, which is consistent with our results. In a UK cross-sectional study, the Isle of Ely Study, a dietary pattern poor in fruits and vegetables and rich in processed meats and French fries was associated with previously diagnosed diabetes⁵⁰.

A direct association between the Western diet and BMI (including fat mass) is in line with previous researches. Both cross-sectional⁵¹ and prospective studies⁵² have reported the parallel findings. A "meats" dietary pattern including "meat", rich in processed and red meats, fish, poultry, eggs, fats and oils, and condiments was correlated with higher BMI in a group of Hawaiian women⁵¹. A direct association between Western dietary patterns and obesity was also confirmed by Slattery et al.⁵³. In an 8-year prospective study among 50,000 adult women, Schulz et al.⁵⁴ demonstrated that Western dietary pattern was attributed to increased weight gain and higher intakes of red and processed meats and sweets. Our results are similar to this study showing an association of Western patterns with higher BMI and fat mass. It is noteworthy that their Western pattern was reflected in diets high in red and processed meats, refined grains, sweets and desserts, and potatoes which was like to ours. Joung et al.⁵⁵ reported that diets containing processed and red meats, oils or fats including (corn oil, margarin, butter, sesame oil, and soybean oil), and sugar may be correlated with obesity in Korean adults.

Levels of plasma leptin reflect the amount of fat stored in adipose tissue. Leptin has a key role in energy homeostasis. Circulating leptin increases by 40% after acute overfeeding, whereas fasting is correlated with significantly decreased leptin secretion⁵⁶. Also, Western dietary pattern principally characterized by highly saturated fat was found to be directly associated with higher leptin concentration⁵⁶. It is noteworthy to mention that our Western pattern was similar to those of Western or unhealthy dietary patterns in different studies^{49–58}.

The present study contains several strengths. It used a validated FFQ. Moreover, to our knowledge, this is the first to identify the association of dietary patterns of the obese with DII and inflammatory parameters. The DII has previously been validated in the Iranian population⁵⁹. This tool is standardized for food intake from various populations around the world²⁵⁻⁶⁰; therefore, scores in Iran can be compared with those obtained in other populations. It should be noted, however, that the DII was calculeted with data from 32 of the 45 food variables, thus 13 parameters are missing in our study. In fact, a total of 32 variables (including energy, carbohydrate, protein, total fat, fiber, cholesterol, saturated fat, mono-unsaturated fat, poly unsaturated fat, omega-3, omega-6, niacin, thiamin, riboflavin, vitamin B12, vitamin B6, iron, magnesium, selenium, zinc, vitamin A, vitamin C, vitamin D, vitamin E, folic acid, beta carotene, garlic, ginger, onion, turmeric, saffron, pepper) were available from the 45-item FFQ which could be used to compute DII. An additional limitation should be noted; because of the cross-sectional nature of our study, it is not possible to infer anything about the temporal component of a putative cause-effect relationship. In addition, measurement accuracy errors inherent in the use of FFQs for dietary assessment may exist^{61,62}.

Conclusion

The results suggested that Western dietary patterns, including higher intakes of red meat and sweets, was related to greater BMI and fat mass. In addition, the Healthy pattern with lower intake of SFA and trans fatty acids led to a significant reduction in the proinflammatory markers i.e. LBP. Adherence to Traditional dietary pattern rich in fruit and vegetables was inversely related to DII as well as IR. Altogether, it seems that dietary patterns can affect systemic inflammation, either directly or indirectly. Further studies with larger sample size are warranted to investigate the associations of diet-induced inflammation with altered levels of adipokines as well as IR markers.

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Author contributions

S.M. and M.S.A. conceived the study design and wrote the study protocol. S.M. was involved with data collection. M.A.J., N.S.H., M.S.A., and F.V. analyzed and interpreted the data. J.H., M.S.A., S.M. and V.E.A. were involved in drafting the article or revising it critically for content. All authors gave final approval of the version to be published.

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Competing interests

The authors declare no competing interests.

Additional information

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