

CARDIO-RENAL METABOLIC SYNDROME AND PRO-INFLAMMATORY FACTORS: THE DIFFERENTIAL EFFECTS OF DIETARY CARBOHYDRATE AND FAT

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Abstract

Background. We aimed to evaluate whether a high carbohydrate or a high fat diet differs in alteration of the inflammatory and metabolic risk factors in cardio-renal metabolic syndrome in rats.

Methods. Twelve male Wister rats were randomly divided into two groups: one received diet 1 standard pellet rat diet (D1) containing 10% fat, 50% carbohydrate, 25% protein and another group received diet 2 (D2) containing 59% fat, 30% carbohydrate and 11% protein for 16 weeks. Weight was recorded weekly. FSG and insulin levels were measured using an enzymatic spectrophotometric and a standard ELISA kit respectively. Inflammatory parameters including TGF- β , MCP-1, TNF- α , IL-1 β , IL-6 in the renal and cardiac tissues of rats were evaluated by ELISA technique.

Result. Food intake in D1 and D2 groups increased in the study period, however food intake in D2 group was significantly higher compared with D1 group. FSG, HOMA and TG concentrations in D2 group were significantly higher compared to D1 group. Moreover, TGF- β and MCP-1 concentrations in the renal tissues of D2 group and TNF- α in the cardiac tissues of D1 group were significantly higher compared with D1 group ($P < 0.05$). Positive associations between IL-1 β and TG and between HOMA, FSG with TGF- β and MCP-1 in the renal tissue of animals were also identified.

Key words: metabolic syndrome, diet, HOMA, insulin resistance.

INTRODUCTION

A cluster of interactive maladaptive cardiovascular and kidney disease risk factors including insulin resistance, obesity, dyslipidemia, hyperglycemia, hypertension, cardiac and renal failure due to over-nutrition and development of obesity contribute to what constitutes the Cardio-Renal Metabolic Syndrome (CRS) (1). According to numerous evidences, there are potent interactions between the heart and kidney disorders; renal dysfunction is an independent risk factor for the development of cardiovascular diseases (CVD) and

is associated with worsened outcome in patients with different kinds of CVDs (2). Moreover, cardiovascular diseases are the leading cause of mortality, consisting of 43.6% of all deaths in patients with end-stage renal disease (3). On the other hand, cardiac dysfunction, for instance post-myocardial infarction (MI), leads to a gradual decrease in renal function as reflected by an increase in creatinine levels. This interaction between the heart and kidney, where dysfunction of either of them leads to disorder of the other, accompanied by ingredients of metabolic syndrome is usually referred to as the CRS (4). The CRS is associated with early cardiac disease because of diastolic dysfunction, vascular, and renal disease and may lead to type 2 diabetes mellitus (T2DM) especially in overweight and obese individuals (5).

Obesity as a known risk factor for numerous health problems including metabolic syndrome, diabetes, hypertension and dyslipidemia (6) is associated with morbidity and mortality and reduces life expectancy in all age groups (7, 8). Obese individuals are at higher risk of chronic kidney disease (CKD) (9, 10) and obesity has been known as a major independent risk factor of focal glomerulosclerosis (11), podocyte lesions (12), end stage renal disease (13) and even renal cell and kidney cancers (14, 15). Moreover, obesity is a known risk factor for CVD worldwide and all of obesity-associated co-morbidities like hypertension, dyslipidemia, diabetes, or insulin resistance, and elevated levels of fibrinogen and C-reactive protein are associated with increased risk of CVD events (16, 17).

The complex heart-kidney bidirectional relationship in CRS involves numerous pro-inflammatory and metabolic mediators which, via bloodstream in the midst of the prevailing metabolic condition, reach target tissues and deliver the adverse effects of the disease (18). These pro-inflammatory mediators had a potent role in maladaptive remodelling of cardiac and renal tissues and are involved in tissue injuries; From the pathological point of view, several of these key inflammatory mediators play an important role in the pathogenesis of obesity related

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CRS and inhibition of their expression is a target of therapeutic approaches to mimic the adverse effects of the CRS; among them, transforming growth factor (TGF)- β is expressed at high concentrations in both the heart and kidneys and is a potent activator of fibroblasts, known to induce myo-fibroblastic activation and increased collagen deposition and wound contraction (19). TGF- β is a key mediator of fibrosis in myocardial injury (20) and has a key role in glomerular and tubule-interstitial pathobiology in chronic kidney diseases, mediating apoptosis and epithelial-to-mesenchymal trans-differentiation (EMT) in chronic progressive renal disease (21). Therapeutic strategies to inhibit TGF- β expression are increasingly investigated with the aim of developing novel renal and cardiac events. It has been shown that abrogation of TGF- β signalling using neutralizing antibodies or oral pharmacological inhibitors has promising results in animal models of cardiac remodelling and renal failure (22, 23). Moreover, TGF- β and monocyte chemoattractant protein (MCP)-1 act in a close relationship with each other in the pathogenesis of renal and cardiac events; MCP-1 is a potent inducer of TGF- β production by infarcted heart and damaged kidneys and enhances the fibrogenic potential of mature macrophages by inducing TGF- β and stimulating collagen synthesis, while TGF- β exerts potent pro-fibrotic actions by inducing myo-fibroblast activation and stimulating the synthesis of various extracellular matrix proteins (24). MCP-1 plays a key role in the pathophysiological processes of cardiovascular and renal biology. As a key attractant for mononuclear cells it is involved in the development and progression of cardiovascular diseases like atherosclerosis, restenosis and thrombosis and its production is up-regulated in patients with congestive heart failure and dilated cardiomyopathy (25). Moreover, MCP-1 has a major role in the pathology of numerous renal diseases and the role of MCP-1 has been revealed in whole cascade of the monocyte-associated inflammation from release of monocytes from the bone marrow, building a chemokine gradient for adhesion and infiltration, stimulation, and differentiation of monocytes and macrophages in the kidney (26). Alongside with the pro-inflammatory role of TGF- β and MCP-1 in the cardio-renal injury, it closely acts in parallel with several other pro-inflammatory cytokines like tumor necrosis factor (TNF)- α , interleukin (IL)-6 and IL-1 β in the cardio-renal inflammation linking these injuries with metabolic syndrome. Their pro-inflammatory roles linking them to obesity and cardiovascular or renal disorders have been implicated in previous studies (27-29).

Diet and dietary ingredients are one of the main inducers of metabolic syndrome and its adverse

effects. By induction of obesity and feeding high fat or high carbohydrate diets or a combination of these two diets are the most common types of diets for inducing metabolic syndrome in animals such as Wistar rats as the most common rodent strains used (30). Many researchers have employed different types of high-fat diets that vary between 20% and 60% of total energy (31). High fat diets have been extensively used to induce metabolic syndrome in experimental animals (32, 33). Studies have also indicated that high-fat diet is effective in inducing hyperglycemia, insulin resistance and dyslipidemia (34). Studies evaluating the role of high carbohydrate diet on inducing metabolic syndrome are scarce while most of the studies used the combination of these two diets for induction of metabolic syndrome and the animals experienced abdominal obesity, hypertension, dyslipidemia and impaired glucose tolerance (35, 36).

To our review of literature, there was no study evaluating the role of numerous potent inflammatory factors in relation with CRS in rats or to evaluate the differential effects of fat-enriched or carbohydrate-enriched diets on the metabolic or pro-inflammatory ingredients of the cardio-renal metabolic syndrome in animals. Therefore, the current experimental project aimed to evaluate these two hypotheses.

ANIMALS AND METHODS

Animals and diets

Male Wistar rats (250-300 g, n=12) were purchased from the Pasteur Institute, Animal Care Center (Karaj, Iran). Each cage included six rats with 12 hours dark/light under standard situation as temperature-controlled environment ($25\pm 2^\circ\text{C}$) with *ad libitum* availability to food and water. The study protocol was approved by the veterinary Ethics Committee of Tabriz University of Medical Sciences (TBZMED.REC.1394.747). After one week of acclimatization and feeding a standard laboratory chow diet, rats were randomly divided into two groups, one group received diet 1 or standard pellet rat diet (D1) containing 10% fat, 50% carbohydrate, 25% protein and another group received diet 2 (D2) containing 59% fat, 30% carbohydrate and 11% protein (Table 1). Each group was fed for 16 weeks. Moreover, all rats were weighed weekly by digital scale (PAND industries, px3000, 5kg \pm 1g). The study procedure and experimental protocol have been published before. Here, the procedure is explained briefly (37-39).

Preparation of blood, heart and kidney tissue samples

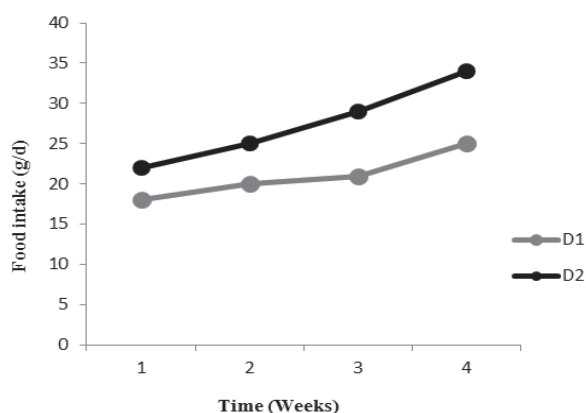
Table 1. Composition of high carbohydrate diet (D1) and high fat diet (D2)

D1 diet		D2 diet	
Ingredient	Percentage	Ingredient	Percentage
Protein	25 %	Butter	28 %
Fat	10%	Chow diet (D1)	28%
Carbohydrate	50 %	Sugar	14 %
Ash	10 %	Yolk egg	19 %
-		White egg	11 %
Energy	100	Total	100

After an overnight fasting, the rats were anesthetized with Ketamin (6.6 mg/kg) and Xylazine (0.3mg/kg) intraperitoneally. Blood samples were obtained from cardiac puncture and centrifuged at 10000 g at 4°C or 20 min; sera were separated and stored in an ultra-low temperature freezer (Jal-Tajhiz Production, Iran) at -80 °C until assaying. Finally, after rats were sacrificed by decapitation, the left nephrectomy specimens and heart samples were homogenized in phosphate buffered saline (PBS) and centrifuged at 10000 g at 4°C for 20 min, and clear supernatants were collected for biochemical assays.

ELISA

Total protein concentration in the tissue supernatants was measured by protein assay kit (Pars

**Figure 1.** Increase of food intake across the groups during the study period. D1: high carbohydrate diet; D2: high fat diet.**Table 2.** Weight, glycemic factors and serum lipids in treatment groups

Parameters	D1 group	D2 group	P-value †
Baseline weight (kg)	0.220±0.004	0.217±0.006	0.718
Terminal weight (kg)	0.308±0.014	0.428±0.007	0.001
FSG (mmol/L)	7.33±1.32	10.23±0.71	0.001
Insulin (pmol/L)	87.92±24.79	64.93±2.22	0.470
HOMA-IR	0.64 ± 0.13	1.21±0.36	0.005
TG (mg/dL)	41.60 ± 5.01	54.02±10.51	0.03
TC (mg/dL)	74.02 ± 3.15	73.38 ±7.68	0.94
LDL-C (mg/dL)	27.82 ± 5.88	23.96± 5.81	0.32
HDL-C (mg/dL)	30.08 ±7.54	42.50 ± 7.73	0.17
AIP	2.49 ± 1.12	0.77 ± 0.23	0.17

D1, high carbohydrate diet, D2, High fat diet, FSG, Fasting serum glucose, HOMA-IR, Homeostatic Model Assessment for Insulin Resistance. Data are presented as means ± SD. †p value indicated intra-group differences compared by independent samples t-test. P < 0.05 was considered as statistically significant.

Azmun, Tehran, Karaj); accordingly, insulin, TNF- α , MCP-1, IL-1 β , IL-6 and TGF- β concentrations in the supernatants were also determined using routine ELISA kits (Hangzhou Eastbiopharm, Zhejiang, China). Serum lipids were measured using the Autoanalyser (Alcyon 300 USA). Atherogenic index was measured with formula (40): Atherogenic index = ((Total Cholesterol-HDL))/HDL. Homeostatic Model Assessment for Insulin Resistance (HOMA-IR) was calculated using the formula: fasting insulin (microU/L) x fasting glucose (nmol/L)/22.5.

Statistical analysis

All statistical analyses were performed using SPSS software, version 16. Kolmogorov–Smirnov test was performed for normality of the distribution of variables. Data are expressed as the mean ± SD. The data were analyzed using independent sample t-test for comparisons between two groups. P < 0.05 was considered as statistically significant.

RESULTS

Changes in weight, serum lipids, glycemic status and food intake

Baseline weight in two groups was identical whereas after dietary treatment, rats in D2 group achieved significantly more weight compared with rats in D1 group (P < 0.001). D2 group also led to the significantly higher concentrations of FSG, insulin resistance and TG concentrations compared with D1 group (P < 0.05). Rats in D1 and D2 groups experienced higher food intake during the study period while the food intake in D2 group was significantly higher compared with rats in D1 group (Figure 1, P < 0.05).

Changes in TNF- α , TGF- β , MCP-1, IL-6 and IL-1 β concentrations in the cardiac and renal tissues of treatment groups

The comparison of inflammatory factors including TNF- α , TGF- β , MCP-1, IL-6 and IL-1 β in the renal and cardiac tissues of D1 and D2 groups is presented in Table 3. In the cardiac tissue, only the TNF- α concentrations in D1 group were significantly higher compared with D2 group whereas, in the renal tissue, TGF- β and MCP-1 concentrations in D2 group were significantly higher compared with D1 group (P <0.05). No significant difference between other inflammatory parameters of cardiac or renal tissues between groups was identified. Also, we assessed the association between pro-inflammatory parameters and markers of metabolic syndrome in renal tissues (Table 4). Renal IL-6 concentrations were in positive relationship with serum TG (r=0.57, P= 0.009), while, renal TNF- α and MCP-1 were in negative association with HOMA-IR and serum FSG concentrations.

DISCUSSION

In the current study, we evaluated the differential effects of dietary carbohydrate or fat enriched diet on pro-inflammatory factors in the cardiac and renal tissues of male Wistar rats with cardio-renal metabolic syndrome. According to our findings, high fat diet induced more pronounced features of cardio-renal metabolic syndrome in rats. Moreover, there was an association between inflammatory parameters of the renal tissue with markers

of glycemic status and lipids while no association between these parameters in the cardiac tissue was found.

The most common feature of the cardio-renal metabolic syndrome is insulin resistance, metabolic dyslipidemia alongside with cardiac and renal complications (41). In the current study, we used dietary approaches to induce the cardio-renal metabolic syndrome and a high fat diet revealed the biggest effect on renal and cardiac tissue inflammation and insulin resistance. Numerous previous studies revealed the major potential of a high fat diet in inducing metabolic syndrome and its related metabolic disorders while very scarce experimental studies used high carbohydrate diet and most of them used the combination of high fat and high carbohydrate dietary regimens (30). In the study by Castro *et al.* the high fat diet for 13 weeks led to a marked increase in the blood pressure, heart rate and major elevations in visceral lipid store in Wistar rats and the researchers suggested that a high fat diet could be the best nutritional intervention procedure in inducing metabolic syndrome (42). Other studies also revealed that high-fat diet is effective in promoting hyperglycemia, insulin resistance and dyslipidemia either independently or concurrently (43).

The potential effects of a high fat diet in increasing the pro-inflammatory parameters including TGF- β and MCP-1 in a renal tissue of rats in the current study was also similar to the findings of previous studies highlighting the

Table 3. Pro-inflammatory factors in cardiac and renal tissues of treatment groups

Parameters	Cardiac Tissues			Renal Tissues		
	D1 group	D2 group	P-value †	D1 group	D2 group	P-value †
TNF- α (ng/mg Protein)	8.28 ±1.04	5.96±0.66	0.003	0.08±0.02	0.11±0.06	0.25
TGF- β (ng/mg Protein)	1.12± 0.39	1.11±0.3	0.39	0.08±0.04	0.14±0.05	0.027
MCP-1 (ng/mg Protein)	0.62±0.15	0.68±0.07	0.12	0.05±0.22	0.09±0.02	0.05
IL-6 (ng/mg Protein)	0.53± 0.12	0.42±0.34	0.98	0.06±0.02	0.07±0.03	0.35
IL-1 β (ng/mg Protein)	0.43 ±0.11	0.55±0.15	0.43	0.59±0.21	0.55±0.17	0.69

D1, high carbohydrate diet, D2, High fat diet, Data are presented as means ± SD. †p value indicated intra-group differences compared by independent samples t-test. P < 0.05 was considered as statistically significant.

Table 4. Correlation matrix between metabolic syndrome markers and inflammatory factors in the renal tissue

Parameter	r (†P)	LDL	HDL	TG	TC	FSG	HOMA-IR
IL-6	r	0.005	-0.12	0.57	0.28	-0.18	0.09
	P	0.98	0.6	0.009	0.22	0.42	0.81
IL-1 β	r	-0.17	-0.22	0.28	-0.02	-0.4	-0.1
	P	0.46	0.34	0.22	0.93	0.07	0.67
TNF- α	r	-0.12	-0.24	0.04	0.04	0.58	0.506
	P	0.59	0.32	0.85	0.83	0.008	0.023
TGF- β	r	-0.23	-0.31	0.03	0.01	0.39	0.27
	P	0.32	0.18	0.87	0.94	0.08	0.25
MCP-1	r	-0.19	-0.03	0.01	-0.06	0.58	0.45
	P	0.41	0.91	0.96	0.78	0.007	0.04

IL, interleukin; TNF- α , tumor necrosis factor α ; TGF- β , transforming growth factor β ; MCP-1 monocyte chemoattractant protein 1. FSG, Fasting serum glucose, HOMA-IR, Homeostatic Model Assessment for Insulin Resistance. † P values obtained by Pearson correlation analysis. P < 0.05 was considered as statistically significant.

pathogenic role of MCP-1 in high-fat diet-induced renal damage; in the study by Declèves *et al.* (44) a high-fat diet led to renal damage and hypertrophy and increased MCP-1 concentrations in the renal tissue and urine of C57BL/6J mice; thereby, the authors suggested that the enzyme adenosine monophosphate-activated protein kinase (AMPK) was the most important mediator of this high fat diet induced damages. They also revealed that the early increase in glomerular MCP-1 contributes to the subsequent recruitment of macrophages that would then contribute to an enhanced release of pro-inflammatory TNF- α , and the most renal profibrotic factor, TGF- β . TGF- β is a potent activator of renal damage and injury and it has been suggested that hyperlipidemia induced by high fat diet is the most important activator of TGF-beta1/Smad signalling pathway and eventually renal damage (45). TGF- β and MCP-1 are potentially linked with obesity, metabolic syndrome and diabetes mellitus; it has been suggested that TGF- β has a common target molecule for the development of CVD, renal insufficiency and metabolic syndrome (46).

In the current study, high carbohydrate diet increased TNF- α concentrations in the cardiac tissue of rats. TNF- α is a cytokine produced mainly by macrophages, has also been found in vascular smooth muscle cells and elevation of its concentrations had been reported in various cardiac diseases, especially in congestive heart failure (CHF); in the study by Doyama K *et al.* (47) the elevated TNF- α concentrations have been reported in the cardiac tissue of patients with heart failure. In a similar study, the effect of long term high-carbohydrate or high-fat diet on adiposity, glucose tolerance, and secretion of TNF- α and MCP-1 by adipose tissue and liver in Swiss mice has been evaluated. According to their findings, a high carbohydrate diet but no high fat diet was able to increase the tissue TNF- α concentrations (48).

In another similar work by Ferreira *et al.*, 30-weeks supplementation with carbohydrate - enriched diet in male Wistar rats significantly increased TNF- α and MCP-1 concentrations in the heart tissue of rats (49). The underlying mechanism linking carbohydrate intake and inflammation is related to stimulated NADPH oxidase in poly-morphonuclear leukocytes and mononuclear cells and increased the generation of reactive oxygen species (ROS) due to reduced antioxidant capacity (50). Moreover, intravenous glucose administration increases concentrations of the inflammatory markers IL-6, IL-18, and tumor necrosis factor α which can be prevented by simultaneous infusion of the anti-oxidant glutathione (51, 52).

In conclusion, in the current work, we showed

potential effects of high fat diet on inducing metabolic abnormalities related to cardio-renal syndrome including hyperglycemia, high insulin resistance, and higher inflammatory parameters including TGF- β and MCP-1 in renal tissue of rats. Although, high carbohydrate diet was also capable of inducing higher TNF- α in the cardiac tissue, but because of a more extent metabolic and inflammatory response after a high fat diet, it can be suggested as a dietary intervention tool for inducing the cardio-renal metabolic syndrome in animal models.

Conflict of interest

The authors declare that they have no conflict of interest.

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