

# The effects of glucose and fructose on body weight and some biochemical parameters in rats

Esra Köseler, Gül Kızıltan, Perim Fatma Türker, Mendane Saka, Mehtap Akçil Ok, Didem Bacanlı, Tolga Reşat Aydos, Nilüfer Bayraktar, Handan Özdemir

Department of Nutrition and Dietetics, Faculty of Health Science, Baskent University, Ankara, Turkey-E-mail:koseler@baskent.edu.tr

**Summary.** *Objective:* Dietary fructose from added sugar as high fructose corn syrup may causes major risks in obesity, hyperlipidemia, cardiovascular diseases, hyperuricemia and fatty liver. The aim of this study was to investigate and compare the effects of high fructose and high glucose intake on body weight and some biochemical parameters in rats. *Subject and methods:* The study was conducted on adult, 32 Wistar albino male rats (300-350 g weeks) which fed with standard laboratory chow. In each group, 8 rats was selected randomly and which was be composed four groups. The rats in each group, in addition to standard meal, different amount of glucose and fructose containing solutions (10% and 30% glucose-fed group, 10% and 30% fructose-fed group) was given by oral gavage for 6 weeks. At baseline and after 6 weeks total cholesterol, VLDL-cholesterol, triglycerides, uric acid, AST and ALT as biochemical parameters and liver histopathological examination of rats were determined. Body weight of the rats was evaluated every week. *Results:* The 30% fructose group caused higher AST levels according to 10% glucose group, 30% glucose group and 10% fructose group. At the end of 6 weeks, the mean body weight in the fructose-fed groups was higher than the glucose-fed groups ( $p > 0.05$ ). No statistically significant difference between rat groups' portal inflammation rates were found and the moderate and severe ballooning were observed in 30% fructose rats ( $p < 0.05$ ). *Conclusions:* As a result, dietary fructose from added sugar as high fructose corn syrup may causes major metabolic disorders.

**Key words:** fructose, glucose, body weight, biochemical parameters, portal inflammation

## Introduction

Fructose, commonly known as fruit sugar, is also a major component of sweeteners such as table sugar, honey and high fructose corn syrup (HFCS) (1). Although fructose is a simple sugar that exists naturally in fruits and vegetables, the majority of dietary fructose comes from two sweeteners, sucrose and high-fructose corn syrup, which are commonly used in manufactured foods and beverages (2). Since the beginning of 20th century, fructose consumption has increased 4-fold by the introduction of HFCS (1). Especially, fructose consumption has increased as usage of HFCS in the Western diet. Based upon disappearance data, the annual per capita intake of HFCS from 1967 to 2006

increased from 0.03 to 58.2 lbs, whereas sucrose decreased from 98.5 to 62.3 lbs. Sucrose is a disaccharide and consists of 50% fructose and 50% glucose. The HFCS form used in soft drinks compose of 55% fructose, 42% glucose, and 3% oligosaccharides. Because of the higher fructose dose, soft drinks sweetened with HFCS would provide more fructose into the systemic circulation than soft drinks sweetened with sucrose. Furthermore, HFCS provides an immediate source of free fructose and glucose, whereas sucrose must first be broken down by sucrase (2,3). An increasing amount of fructose in the diet is suggested to play a causal role in the pathogenesis of the metabolic syndrome, insulin resistance, impaired glucose tolerance, type 2 diabetes, obesity, hyperlipidemia, cardiovascular diseases, hype-

uricemia and fatty liver (4). Fructose does not increase the satiety signals of blood glucose and insulin to the same extent as does sucrose or glucose. Short-term food intake is inversely related to the glycemic and insulin responses to sugars, and it has been proposed that fructose does not suppress gastric appetite hormone and reduced insulin and leptin signaling in the brain. High fructose causes an increase in the synthesis of non-esterified fatty acids production. Fructose is lipogenic and stimulates triglyceride synthesis. Acute oral or intravenous administration of fructose results in a rapid increase in serum levels of uric acid through accentuated degradation of purine nucleotides and increased purine synthesis. The aim of this study was determined the effect of different amounts of fructose and glucose in rats to body weight and some biochemical parameters.

## Material and Methods

### *Experimental design*

This research conducted in Baskent University Production and Research Centre for Experimental Animal, Ankara, Turkey. This study was approved by Baskent University Ethical Committee for Experimental Research on Animals (Project no: DA14/14) and supported by Baskent University Research Fund.

Male rats were divided into four groups with each group comprising of eight animals. Male Wistar albino rats (32 weeks) weighing 300–350 g were randomly assigned to one of the four groups; 10% glucose-fed group, 30% glucose-fed group, 10% fructose-fed group and 30% fructose-fed group.

Group 1 n(8): Standart pellet+10% HFCS

Group 2 n(7): Standart pellet+30% HFCS

Group 3 n(8): Standart pellet+10% glucose solution

Group 4 n(7): Standart pellet+30% glucose solution

Sample size calculated on the basis of probability distribution of the measured values with a given significance level (e.g., 5%), medium effect size (e.g., 0.35) and the power of test (e.g., 85%). This analysis was performed using G\*Power 3.1.3 software program. Thus, the total sample size was obtained in 32 rats. All animals were housed in cages and subjected to a 12 h light-dark cycle at  $24 \pm 2$  °C and animals were

fed on a standard pellet diet and water ad libitum. The solutions have been prepared by feeding to rats, at four concentrations, 10 and 30 g/ 100 milliliter glucose; 10 and 30 g/ 100 milliliter fructose. Solutions to be administered by gavage were stored at 4°C and warmed to room temperature. The follow-up terminated at the end of 6 weeks.

### *Evaluation of Measurements*

At baseline and at the end of the 6 weeks, total cholesterol (TC), VLDL-cholesterol (VLDL-C), triglyceride (TG), uric acid (UA), alanine aminotransferase (ALT), aspartate transaminase (AST) measurements were sampled. For the experiment; the animals were starved overnight for 12 h before the blood collection process and approximately 1mL blood sample was collected from each rat by snipping the tail using heparin anti-coagulant under diethyl ether anaesthesia. Then, plasma was obtained from the blood using a centrifuge at 4 °C for 15 min. Serum total cholesterol, triglyceride and uric acid levels were assayed by enzymatic tests, using an Abbott® Architect C8000 Analyzer according to the manufacturers specifications. (Abbott Park, IL, USA). VLDL cholesterol was calculated from measurements obtained for triglyceride using the following formula:  $VLDL = \text{Triglyceride}/5$  (mg/dL). Serum ALT and AST levels were assayed by an UV test according to standardized method, using an Abbott® Architect C8000 Analyzer according to the manufacturers specifications. (Abbott Park, IL, USA). Body weight was measured weekly during the follow-up.

### *Liver histopathology*

Histopathologic examination was carried out at the end of 6 weeks. Steatohepatitis was evaluated using the grading and staging system of Brunt et al. (5). The grades were classified as grades 0–4, which were based on the percent of hepatocytes involved in the biopsy (0: none, 1: 10%, 2: 10–33%, 3: 33–66%, 4: 66%).

### *Statistical analysis*

The results were expressed as mean±SD or mean (95% CI). Paired t-tests were used to estimate the presence of changes in study parameters for each experiment group (e.g., Group 1: 10% HFCS; Group 2: 30% HFCS; Group 3: 10% glucose solution; Group

**Table 1.** Effect of fructose and glucose feeding on biochemical parameters and systolic blood pressure for 6 weeks

	10% Glucose		p <sup>1</sup>	30% Glucose		p <sup>2</sup>	10%Fructose		p <sup>3</sup>	30% Fructose		p <sup>4</sup>
	Baseline	After 6 Weeks		Baseline	After 6 Weeks		Baseline	After 6 Weeks		Baseline	After 6 Weeks	
Cholesterol	75.6±14.95	110.8±64.19	0.184	92.6±36.88	93.8±57.32	0.879	69.5±10.83	72.7±11.10	0.554	69.8±8.11	74.1±16.11	0.249
VLDL-C	14.0±3.52	37.2±37.43	0.177	17.9±9.25	20.3±15.49	0.418	16.0±5.14	17.1±4.67	0.596	14.0±2.94	16.9±5.55	0,242
Triglycerid	70.0±17.60	186.0±187.34	0.179	89.5±46.25	101.5±77.47	0.418	80.0±25.71	85.3±23.36	0.596	70.0±14.71	84.7±27.77	0.242
Uric asid	1.46±0.51	2.0±0.36	0.061	1.33±0.19	1.53±0.60	0.359	1.3±0.26	1.3±0.34	0.547	1.4±0.29	2.0±0.36	0.006*
ALT	74.8±64.19	65.4±20.05	0.620	54.2±17.81	53.5±10.23	0.907	65.6±16.68	74.6±31.05	0.544	69.6±24.64	89.0±19.44	0.015*
AST	124.8±71.36	177.7±50.6	0.010*	116.3±7.53	119.1±24.13	0.754	97.3±11.85	134.8±48.34	0.067	106.0±27.39	206.3±47.23	0.002*

p<sup>1-4</sup>: The significance test of differences between baseline and after six weeks values for each group.

VLDL-C: VLDL-Cholesterol

4: 30% glucose solution), In addition, the absolute changes (the difference between baseline values and after six weeks values) were tested between groups using one-way ANOVA. The distribution of changes was evaluated for normality assumptions using One Sample Kolmogorov–Smirnov test. The Fisher exact test was used for proportions. SPSS version 21.0 was used to analyze the recorded data. Significant values of  $p < 0.05$  were considered to be statistically significant.

## Results

The mean of plasma UA, TG, TC, VLDL-C, ALT and AST at baseline and after the six weeks were shown in Table 1. It was found that the significant differences in mean values of AST in 10% glucose-fed group ( $p=0.010$ ); uric acid, ALT and AST in 30% fructose fed group ( $p=0.011$ ,  $p=0.015$ ,  $p=0.002$ , respectively). The 30% fructose group caused the difference in AST levels according to 10% glucose group, 30% glucose group and 10% fructose group.

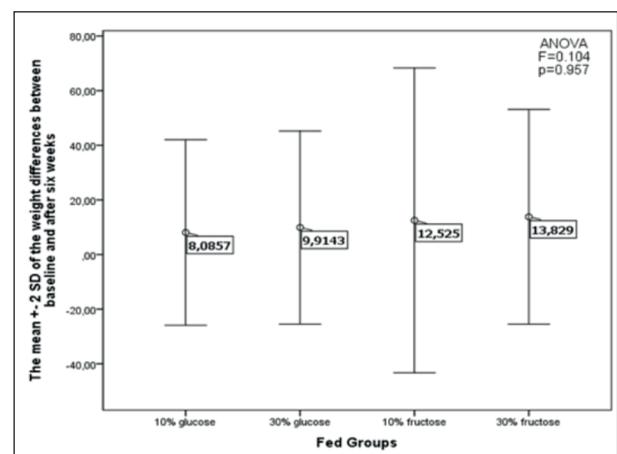
The difference of the initial and final body weight were shown in Figure 1. After a 6 week trial, the mean body weight in the fructose-fed groups was higher than the glucose-fed groups, but there were no significant differences in body weight gain among groups ( $p > 0.05$ ) (Figure 1).

The effect of fructose and glucose feeding on portal inflammation and hepatocyte ballooning in rats'

livers for 6 weeks were shown in Table 2. No statistically significant difference between rat groups' portal inflammation rates were found. Both 10% fructose and 30% fructose groups, 2 of the 8 rats were observed mild inflammation. There were statistically significant differences between the rat groups in terms of hepatocyte ballooning ( $p=0.025$ ). Mostly, the moderate and severe ballooning were observed in 30% fructose rats (Table 2).

## Discussion

When we analyzed the difference of body weight during a 6 week treatment, the mean body weight in



**Figure 1.** Effect of fructose and glucose feeding on body weight in rats for 6 weeks

**Table 2.** Effect of fructose and glucose feeding on portal inflammation and hepatocyte ballooning in rats' livers for 6 weeks

	10% Glucose		30% Glucose		10% Fructose		30% Fructose		
	n	%	n	%	n	%	n	%	
Portal Inflammation									
No	7	26.9	7	26.9	6	23.1	6	23.1	p=0.886
Mild	1	25.0	-	-	2	50.0	1	25.0	
Hepatocyte Ballooning									
No	3	25.0	5	41.7	-	-	4	33.3	p=0.021*
Yes	5	27.8	2	11.1	8	44.4	3	16.7	

the fructose-fed groups has shown higher than the glucose-fed groups ( $p>0.05$ ) (Figure 1). This study was suggested that the weight gain by fructose feeding as previous studies. Over the past several years, the reasons for the increase in obesity prevalence have shown that increased the sugar added to food and it has taken the place of the sucrose to HFCS by researchers (6-8). HFCS caused an increase in body weight greater than sucrose in both male and female rats. This increase in body weight was accompanied by an increase in fat accumulation and circulating levels of TG (9).

In recent studies having drawn attention to fructose has emphasized the absence of satiety such as other sugars. Plasma glucose and insulin levels effected the state of satiety after food consumption. Although fructose does not contribute to the feeling of fullness, has the same energy load with the blood sugar glucose. Therefore, as long as the amount of glucose decreases and the amount of fructose increases, the feeling of fullness occurs later and it is consist of more eating behavior (10,11). The excessive consumption of HFCS may contribute to the incidence of obesity by reducing insulin and leptin levels (12). The intake of HFCS would not lead to insulin or leptin-induced satiety. Because fructose leads to increased plasma free fatty acids, leptin, adiponectin, abdominal adipose tissue and impaired insulin sensitivity (13,14). The recent studies demonstrate that compared to pure glucose, chronic fructose feeding does not suppress the appetite hormone ghrelin and does not provide enough insulin and leptin secretion (15,16). In a study which was analyzed the long-term effects of HFCS on body weight, the rats with access to HFCS gained significantly more

body weight than sucrose groups (9). Fructose (or sucrose) administration to humans and rats also induces attributes of liver diseases and may have a role in the pathogenesis of fatty liver diseases (17,18). The fatty liver disease includes a broad spectrum of manifestations of fatty liver, ranging from steatosis alone, steatosis with inflammation, steatosis with hepatocyte injury, or steatosis with sinusoidal fibrosis in relation to the progress of the pathological state (19,20). In this study we investigated whether fructose could play a role metabolic disorders in liver.

Administration of high doses fructose can also cause elevation of portal inflammation rates hepatocyte ballooning. The moderate and severe ballooning were observed in most 30% fructose rats. But there were no statistically significant difference between rat groups' portal inflammation rates were found. If only both 10% fructose and 30% fructose groups, 2 of the 8 rats were observed mild inflammation. Ackerman et al., demonstrated that implementation of fructose to rats results in hepatic steatosis with a 198% increase in hepatic triglycerides and an 89% increase in hepatic cholesterol concentration and Davail et al., evidenced high fructose diets also develop fatty liver (21,22)

## Conclusion

As a conclusion, dietary fructose from added sugar as high fructose corn syrup may causes major risks in obesity, fatty liver disease, insulin resistance, hyperlipidemia, impaired glucose tolerance, Type 2 diabetes, cardiovascular diseases, hyperuricemia, gout and meta-

bolic syndrome. So, the healthy preference of fructose source in diets is fruit and the amount of safe dietary intake of fructose may accept as 10% of total energy.

## References

- Echtay KS. Mitochondrial uncoupling proteins—what is their physiological role? *Free Radic Biol Med* 2007; 43: 1351–71.
- Horvath TL, Diano S, Barnstable C. Mitochondrial uncoupling protein 2 in the central nervous system: neuro-modulator and neuroprotector. *Biochem Pharmacol* 2003; 65: 1917–21.
- Vincent AM, Olzmann JA, Brownlee M, Sivitz WI, Russell JW. Uncoupling proteins prevent glucose-induced neuronal oxidative stress and programmed cell death. *Diabetes* 2004; 53:726–34.
- Jiffri EH. Association of the UCP2 45-bp insertion/deletion polymorphism with diabetes type 2 and obesity in Saudi population, *The Egyptian Journal of Medical Human Genetics* 2012; 13: 257–262.
- Yanovski JA, Diament AL, Sovik KN, Nguen TT, Li H, Sebring NG, et al. Associations between uncoupling protein 2, body composition, and resting energy expenditure in lean and obese African American, white, and Asian children. *Am J Clin Nutr* 2000; 71: 1405–20.
- Kovacs P, Ma L, Hanson RL, Franks P, Stumvoll M, Bogardus C, et al. Genetic variation in UCP2 (uncoupling protein-2) is associated with energy metabolism in Pima Indians. *Diabetologia* 2005; 48(11): 2292–5.
- Liu YJ, Liu PY, Long J, Lu Y, Elze L, Recker RR, et al. Linkage and association analyses of the UCP3 gene with obesity phenotypes in Caucasian families. *Physiol Genom* 2005; 22(2): 197–203.
- Ochoa MC, Santos JL, Azcona C, Moreno-Aliaga MJ, MartinezGonzalez MA, Martinez JA, et al. Association between obesity and insulin resistance with UCP2–UCP3 gene variants in Spanish children and adolescents. *Mol Genet Metab* 2007; 92(4): 351–8.
- Otaegui D, Saenz A, Ruiz-Martinez J, Olaskoaga J, Lopez de Munain A. UCP2 and mitochondrial haplogroups as a multiple sclerosis risk factor. *Mult Scler* 2007; 13(4): 454–8.
- Vogler S, Goedde R, Mitterski B, Gold R, Kroner A, Koczan D, et al. Association of a common polymorphism in the promoter of UCP2 with susceptibility to multiple sclerosis. *J Mol Med* 2005; 83(10): 806–11.
- Rudofsky G-Jr, Schroedter A, Schlotterer A, Voron'ko OE, Schlimme M, Tafel J, et al. Functional polymorphisms of UCP2 and UCP3 are associated with a reduced prevalence of diabetic neuropathy in patients with type 1 diabetes. *Diabetes Care* 2006; 29(1): 89–94.
- Yamasaki H, Sasaki H, Ogawa K, Shono T, Tamura S, Doi A, et al. Uncoupling protein 2 promoter polymorphism 866G/A affects peripheral nerve dysfunction in Japanese type 2 diabetic patients. *Diabetes Care* 2006; 29(4): 888–94.
- Humphries SE, Cooper JA, Talmud PJ, Miller GJ. Candidate gene genotypes, along with conventional risk factor assessment, improve estimation of coronary heart disease risk in healthy UK men. *Clin Chem* 2007; 53(1): 8–16.
- Yasuno K, Ando S, Misumi S, Makino S, Kulski JK, Muratake T, et al. Synergistic association of mitochondrial uncoupling protein (UCP) genes with schizophrenia. *Am J Med Genet B Neuropsychiatr Genet* 2007; 144(2): 250–3.
- Park D, Han CZ, Elliott MR, Kinchen JM, et al. Continued clearance of apoptotic cells critically depends on the phagocyte UCP2 protein. *Nature* 2011; 477: 220–224.
- Jimeno RE, Dembski M, Weng X, Deng N, et al. Cloning and characterization of an uncoupling protein homolog: a potential molecular mediator of human thermogenesis. *Diabetes* 1997; 46: 900–906.
- Zhang CY, Baffy G, Perret P, Krauss S, et al. Uncoupling protein-2 negatively regulates insulin secretion and is a major link between obesity, beta cell dysfunction, and type 2 diabetes. *Cell* 2001; 105: 745–755.
- Yang W, Lu J, Weng J, Jia W, et al. Prevalence of diabetes among men and women in China. *N. Engl. J. Med.* 2010; 362: 1090–1101.
- Dalgaard LT. UCP2 mRNA expression is dependent on glucose metabolism in pancreatic islets. *Biochem. Biophys. Res. Commun.* 2012; 417: 495–500.
- Chan CB, De Leo D, Joseph JW, McQuaid TS, et al. Increased uncoupling protein-2 levels in beta-cells are associated with impaired glucose-stimulated insulin secretion: mechanism of action. *Diabetes* 2001; 50: 1302–1310.
- Sun LL, Jiang BG, Li WT, Zou JJ, et al. MicroRNA-15a positively regulates insulin synthesis by inhibiting uncoupling protein-2 expression. *Diabetes Res. Clin. Pract.* 2011; 91: 94–100.
- Krempler F, Esterbauer H, Weitgasser R, Ebenbichler C, et al. A functional polymorphism in the promoter of UCP2 enhances obesity risk but reduces type 2 diabetes risk in obese middle-aged humans. *Diabetes* 2002; 51: 3331–3335.
- Zheng YM, Xiang KS, Zhang R and Jia WP. Association between Ala55Val variant in the uncoupling protein 2 gene and glucose stimulated insulin secretion in type 2 diabetic Chinese. *Chin. J. Endocrinol. Metab.* 1999; 15: 199–202.
- Xiu LL, Weng JP, Sui Y, Wang J, et al. Common variants in 3-adrenergic-receptor and uncoupling protein-2 genes are associated with type 2 diabetes and obesity. *Zhonghua Yi Xue Za Zhi* 2004; 84: 375–379.
- Shen XJ, Zhu DL, Tong GY and Hu Y. Association of -866G/A polymorphism in uncoupling protein 2 gene of patients with type 2 diabetes in Nanjing. *Chin. J. Practical Internal Med.* 2007; 27: 670–673.
- Gu GY, Zheng SX, Liu DM and Chen LM. Association of functional polymorphism in the promoter of uncoupling protein 2 (UCP2) gene with type 2 diabetes. *Chin. J. Diabetes* 2007; 15: 411–412.
- Li JN, He L, Ye F and Dong CP. Association of uncoupling

- protein 2 -866G/A polymorphism with type 2 diabetes in northern Chinese. *J. Fourth Mil. Med. Univ.* 2008; 29: 163-166.
28. Wang XX, Xian TZ, Wang SL and Sun XM. Correlation between -866G/A variation in the promoter region of uncoupling protein-2 gene and the risk of type 2 diabetes in population from Beijing. *CRTER* 2009; 13: 4754-4758.
29. Liu L, Guan YF, Li Z and Sun W. UCP-2 gene promoter -866G/A polymorphism related to the development of type 2 diabetes mellitus in Chinese. *Medicine & Philosophy (Clinical Decision Making Forum Edition)* 2009; 30: 50-52.
30. She YM. SUR1 and UCP2 Gene Polymorphism with Type 2 Diabetes and the Impact on Nateglinide Effectiveness. Master's thesis, CSU, Changsha. 2009.
31. Yang M, Huang Q, Wu J, Yin JY, et al. Effects of UCP2 -866G/A and ADRB3 Trp64Arg on rosiglitazone response in Chinese patients with Type 2 diabetes. *Br. J. Clin. Pharmacol.* 2009; 68: 14-22.
32. Hu ZQ, Ma GQ, Ma CH, Liu J. An analysis of association of UCP-2 A55V polymorphism with overweight, obesity and type 2 diabetes in Dongxiang of Gansu people. *Chin. J. Diabetes* 2010; 18: 115-117.
33. Qin LJ, Wen J, Qu YL, Huang QY. Lack of association of functional UCP2 -866G/A and Ala55Val polymorphisms and type 2 diabetes in the Chinese population based on a case-control study and a meta-analysis. *Genetics and Molecular Research* 2013; 12 (3): 3324-3334.
34. Souza BM, Assmann TS, Kliemann LM, Gross JL, Canani LH, et al. The role of uncoupling protein 2 (UCP2) on the development of type 2 diabetes mellitus and its chronic complications. *Arq Bras Endocrinol Metabol* 2011; 55: 239-248.
35. Bulut F, Erol D, Elyas H, Do an H, Ozdemir FA, Keskin L. Protein tyrosine phosphatase non-receptor 22 gene C1858T polymorphism in patients with coexistent type 2 diabetes and hashimoto's thyroiditis. *Balkan Medical Journal* 2014; 31: 37-42.
36. Elhadd TA, Al-Amoudi AA, Alzahrani AS. Epidemiology, clinical and complications profile of diabetes in Saudi Arabia: a review. *Ann Saudi Med* 2007; 27(4): 241-50.
37. Tayeb MT. Association of the UCP2 866G/A polymorphism with type 2 diabetes and obesity in Saudi population. *Egypt J Med Hum Genet* 2009; 10(2): 228-36.
38. Chan CB, MacDonald PE, Saleh MC, Johns DC, Marban E, Wheeler MB. Overexpression of uncoupling protein 2 inhibits glucose-stimulated insulin secretion from rat islets. *Diabetes* 1999; 48(7): 1482-6.
39. Esterbauer H, Schneitler C, Oberkofler H, Ebenbichler C, Paulweber B, Sandhofer F, et al. A common polymorphism in the promoter of UCP2 is associated with decreased risk of obesity in middle-aged humans. *Nat Genet* 2001; 28: 178-83.
40. Yu X, Jacobs DR Jr, Schreiner PJ, Gross MD, Steffes MW, et al. The uncoupling protein 2 Ala55Val polymorphism is associated with diabetes mellitus: the CARDIA study. *Clin Chem* 2005; 51: 1451-1456.
41. Bulotta A, Ludovico O, Coco A, Di Paola R, Quattrone A, et al. The common -866G/A polymorphism in the promoter region of the UCP-2 gene is associated with reduced risk of type 2 diabetes in Caucasians from Italy. *J Clin Endocrinol Metab* 2005; 90: 1176-1180.
42. Crispim D, Fagundes NJ, Dos Santos KG, Rheinheimer J, Bouc, As AP, et al. Polymorphisms of the UCP2 gene are associated with proliferative diabetic retinopathy in patients with diabetes mellitus. *Clin Endocrinol (Oxf)* 2010; 72: 612-619.
43. Lindholm E, Klannemark M, Agardh E, Groop L, Agardh CD. Putative role of polymorphisms in UCP1-3 genes for diabetic nephropathy. *J Diabetes Complications* 2004; 18: 103-107.
44. D'Adamo M, Perego L, Cardellini M, Marini MA, Frontoni S, et al. The 2866A/A genotype in the promoter of the human uncoupling protein 2 gene is associated with insulin resistance and increased risk of type 2 diabetes. *Diabetes* 2004; 53: 1905-1910.
45. Wang H, Chu W, Lu T, Hasstedt S, Kern P, Elbein S. Uncoupling protein-2 polymorphisms in type 2 diabetes, obesity, and insulin secretion. *Am J Physiol Endocrinol Metab.* 2004; 286:1-7.
46. Oberkofler H, Iglseider B, Klein K, Unger J, Haltmayer M, Krempler F, et al. Associations of the UCP2 gene locus with asymptomatic carotid atherosclerosis in middle-aged women. *Arterioscler Thromb Vasc Biol.* 2005; 25: 604-610.

Correspondence:

Esra Kösele

Department of Nutrition and Dietetics,

Faculty of Health Science,

Baskent University Ankara Turkey

E-mail: koseler@baskent.edu.tr