

VITAMIN D DEFICIENCY DID NOT AUGMENT THE PROGRESSION OF HIGH-FRUCTOSE-INDUCED NONALCOHOLIC FATTY LIVER DISEASE IN RATS

VİTAMİN D EKSİKLİĞİ SIÇANLARDA ALKOLE BAĞLI OLMAYAN YAĞLI KARACİĞER HASTALIĞININ PROGRESYONUNU ARTTIRMADI

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ABSTRACT

Objective: Vitamin D has antioxidant, anti-inflammatory and antiglycation activities, and hepatoprotective potential. There is a relationship between vitamin D deficiency (VDD) and the severity of liver disorders. VDD has been proposed to contribute to the progression of nonalcoholic fatty liver disease (NAFLD). However, experimental results are not clear. Therefore, in this study, the effects of a VDD diet on high fructose (HFr) drinking-induced NAFLD was evaluated.

Material and Method: Male Wistar rats were divided into four groups as control, HFr, VDD+HFr, and VDD. Control and HFr groups were fed a control diet, and other groups with a VDD-diet for 12 weeks. HFr (30%; w/v; in drinking water) was given in the last 8 weeks. Insulin resistance (IR), serum lipids, hepatic triglyceride, lipid peroxide, protein carbonyl, advanced glycation end products (AGEs) and inflammation (TNF- α and myeloperoxidase) parameters, and histopathological changes were investigated.

Results: Increases in serum transaminases, hypertriglyceridemia, and IR were observed in HFr and VDD+HFr groups. Increased liver triglyceride, lipid and protein oxidation products, protein glycation and inflammation markers as well as microvesicular hepatic steatosis and hepatocyte ballooning were observed in both groups. Although IR and hepatic inflammation markers were higher in the VDD+HFr group, serum transaminases, hepatic triglyceride, lipid and protein oxidation products, and glycation indicators in the liver did not alter between the two groups. However, Nrf2 mRNA expression and superoxide dismutase and

ÖZET

Amaç: Vitamin D antioksidan, antiinflamatuvar ve antiglikasyon etkinliğe ve karaciğeri koruyucu potansiyele sahiptir. Vitamin D eksikliği/yetersizliği (VDD/VDI) ile karaciğer bozukluklarının ciddiyeti arasında bir ilişki bulunmaktadır. VDD'nin alkole bağlı olmayan yağlı karaciğer hastalığının (NAFLD) progresyonunda etkili olduğu bildirilmiştir. Fakat deneysel sonuçlar yeterli değildir. Bu nedenle, bu çalışmada VDD'nin yüksek fruktozlu (HFr) içme suyu uygulanarak oluşturulan NAFLD üzerine etkisi incelendi.

Gereç ve Yöntemler: Erkek Wistar sıçanlar kontrol, HFr, VDD+Fr ve VDD olmak üzere 4 gruba ayrıldı. Kontrol ve HFr grupları Vit D3 içeren, diğerleri ise Vit D₃ içermeyen yemle 12 hafta beslendiler. HFr (%30; w/v) içme suyu ile son 8 hafta uygulandı. İnsulin direnci (IR), serum lipitleri, hepatik trigliserit, lipit peroksit, protein karbonil, ileri glikasyon ürünleri (AGEs) ve inflamasyon (TNF- α ve miyeloperoksidaz) göstergeleri tayin edildi.

Bulgular: HFR ve VDD+HFr gruplarında serumda transaminazlar arttı, hipertrigliseridemi, ve insülin direnci oluştu. Her iki grupta da karaciğerde trigliserit düzeyleri, lipit ve protein oksidasyon ürünleri, protein glikasyon ve inflamasyon göstergeleri arttı, mikroveziküler steatoz ve hepatosit balonlaşması saptandı. IR ve inflamasyon göstergeleri VDD+HFr grubunda daha yüksek olmasına rağmen, iki grup arasında serum transaminazları, karaciğer trigliserit, lipit ve protein oksidasyon ürünleri ve glikasyon göstergeleri düzeylerinde bir farklılık bulunmadı. Ancak Nrf2 mRNA ekspresyonu ile süperoksit dismutaz ve glutatyon peroksidazın mRNA ekspresyonları ve aktivitelerinin VDD+HFr grubunda

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glutathione peroxidase mRNA expression and activities were significantly higher in the VDD+HFr group.

Conclusion: Our results show that VDD did not augmented HFr-induced hepatotoxicity and glycooxidative stress in the liver of rats.

Keywords: Vitamin D deficiency, high fructose diet, nonalcoholic liver disease, glycooxidative stress, antioxidant, inflammation daha yüksek olduğu bulundu.

Sonuç: Sonuçlarımız VDD'nin HFr ile oluşturulan karaciğer hasarı ve glikooksidatif streste bir değişiklik oluşturmadığını göstermektedir.

Anahtar Kelimeler: Vitamin D yetersizliği, yüksek fruktozlu diyet, alkolik olmayan karaciğer hastalığı, glikooksidatif stres, antioksidanlar, inflamasyon

INTRODUCTION

Nonalcoholic fatty liver disease (NAFLD) refers to an accumulation of fat in the liver due to causes other than alcohol. NAFLD is accepted as the hepatic manifestation of the metabolic syndrome. NAFLD starts with simple steatosis and it can progress to more serious conditions, such as steatohepatitis (NASH), fibrosis and cirrhosis. According to a two-hit hypothesis, the presence of steatosis (first hit) makes the liver susceptible to some factors such as oxidative stress, endotoxemia, inflammation and mitochondrial dysfunction as the second hit (1). Advanced glycation end products (AGEs) also play an important role as a contributing factor in the pathogenesis of NAFLD (2).

A high-fructose (HFr) diet affects glucose and lipid metabolism and causes various metabolic disorders such as insulin resistance (IR), hypertriglyceridemia and NAFLD (3). A HFr diet is one of the good dietary models of NA-FLD in animals (4). Increased lipogenesis, oxidative stress and inflammation, and decreased β -oxidation are strongly associated with HFr-induced NAFLD (5). Increases in non-enzymatic protein glycation result in the accumulation of advanced glycation end products (AGEs) in HFr rats (6). These products are known to affect structures and functions of proteins and cause further increased oxidative stress and inflammation by interacting with their receptors (2, 6). Therefore, AGEs were accepted to be effective in the progression of NAFLD (6).

There are two main forms of Vitamin D: Vit D_2 (ergocalciferol) and Vit D_3 (cholecalciferol). Vit D_2 is derived from the diet, but Vit D_3 is both derived from diet and synthesized from 7-dehydrocholesterol with the effect of ultraviolet light on the human skin. Dietary or synthesized Vit D are hydroxylated in the liver to 25-hydroxyvitamin D [25(OH) D]. It is subsequently hydroxylated in the kidney into 1,25-dihydroxyvitamin D [1,25(OH)₂ D] which is the biologically active form of Vit D (7).

 $1,25(OH)_2 D$ induces biological effects by binding to a vitamin D receptor (VDR) located in the nucleus of target cells. Its classical role is in calcium/phosphate homeostasis. Additionally, it has multiple functions and target organs and influences the expression of several genes and plays important roles in the regulation of cell proliferation and differentiation, immune function, oxidative stress, protein glycation, inflammation, and apoptosis (7-9). Therefore, Vit D has been reported to be effective in preventing many diseases such as diabetes, hypertension, cardiovascular and hepatic diseases and cancer (10-12).

Vit D deficiency (VDD) is guite common in the world. Sedentary lifestyle, inadequate sunlight exposure and the lack of Vit D in the food has resulted in VDD (13,14). Serum 25(OH)D levels are used as an indicator of Vit D stores. Although there are some differences in clinical guidelines, the Institute of Medicine (Washington; USA) defines VDD as a 25(OH)D level of <20 ng/mL and severe VDD as <10 ng/mL (15). VDD alone may induce IR, oxidative stress, inflammation, and mitochondrial dysfunction (8, 10, 14). However, epidemiological and laboratory evidence have shown that VDD may be associated with some common diseases such as tumors, bone diseases, immune disorders, cardiovascular diseases, diabetes mellitus, and chronic liver disorders (12-14). VDD has also been reported to be a secondary factor in the onset and progression of NAFLD, but a clear result has not been reached in clinical and experimental studies (7, 10, 11, 16, 17).

In this study, we aimed to examine whether a diet deficient in VitD₃ has a progressive effect on HFr-induced NAFLD in rats. For this reason, parameters affecting the progression of NAFLD such as IR, hepatic prooxidant and antioxidant parameters, protein glycation and inflammation markers together with histopathological changes were evaluated in the liver.

MATERIAL AND METHOD

Chemicals

Fructose (Fr) and other chemicals were purchased from Sigma-Aldrich (Saint-Louise, MI, USA).

Animals

Male Wistar rats (140-160 g) were provided from the Aziz Sancar Experimental Medical Research Institute of Istanbul University. The animals were supplied with food and water *ad libitum*. They were kept in polypropylene cages (three to four per cage) at 22°C, with 12-h light and 12-h darkness without shielding from ultraviolet B radiation (290-320 mn).

Diets and experimental design

Rats were randomly assigned into four groups as control (n=6), HFr (n=7), HFr+VDD (n=8) and VDD (n=8) groups. Control and HFr groups were fed on PicoLab Rodent diet 5053 (2300 IU Vit D3/kg added); however, VDD+HFr and VDD groups were fed on Modified LabDiet 5053 (Vit D3 not added) for 12 weeks. These diets were supplied from LabDiet (St. Louis-Missouri, USA). Fructose (30%, w/v) was added to drinking water for the last 8 weeks to HFr and VDD+HFr groups.

Samples

The animals were fasted overnight and anesthetized with ketamine (35 mg/kg, i.p., Pfizer, USA) and xylazine HCl (15 mg/kg, i.p., Bioveta, Czech Republic). Blood was collected into dry tubes by cardiac puncture and centrifuged at 1,500xg for 10 min to get serum.

Liver tissues were removed, washed with ice-cold 0.9% NaCl and homogenized in ice-cold PBS (phosphate-buffered saline; 0.01M, pH:7.4). Liver homogenates were centrifuged at 600 g for 10 min at 4°C to obtained the postnuclear fraction (PNF). Serum and postnuclear fraction of liver were stored at -80°C until they were analyzed. The liver index was calculated as liver weight /body weightx100.

Determinations in serum

Serum fasting glucose, triglyceride (TG), calcium and inorganic phosphorus levels, and alanine aminotransferase (ALT) and aspartate aminotransferase (AST) activities were measured using a Cobas Integra 800 autoanalyzer (Roche Diagnostics, Mannheim, Germany). Serum $25(OH)D_3$ and insulin levels were determined by ELISA kits (Abbkine, Wuhan-China). Homeostasis model assessment (HOMA) index was used to evaluate insulin resistance (IR) and calculated using the formula: fasting insulin concentration (pmol/L)xfasting glucose concentration (mmol/L)/135. High HOMA scores indicate IR (low insulin sensitivity).

Determinations in the liver

TG, hydroxyproline (Hyp), tumor necrosis factor-alfa (TNF- α) levels and myeloperoxidase (MPO) activity

TG levels were determined in hepatic lipid extracts by using kits provided by Biolabo Biochemistry and Coagulation (Maizy, France). Liver tissues were homogenized in 10 volumes of PBS with a glass homogenizer on ice. Tissue homogenates were sonicated, and then centrifuged at 5,000 g for 5 min to get the supernatants. Hyp (Bioassay Technology Laboratory, Shanghai, China) and TNF- α (USCN, Wuhan, China) levels were determined in these supernatants by using ELISA kits.

To determine MPO activity, liver tissue was homogenized in 0.5% hexadecyltrimethylamine bromide in 50 mM phosphate buffer, pH 6. The homogenates were

sonicated, freeze-thawed three times and centrifuged at 10,000xg. MPO activity was measured in the supernatant samples (18).

Hepatic reactive oxygen species (ROS), lipid peroxide and protein oxidation products

ROS generation was determined by the method described by Wang and Joseph (19). After excitation at 485 nm, the fluorescence emission of 2',7'-dichlorofluorescein at 538 nm was recorded using a microplate fluorometer and luminometer (Fluoroskan Ascent FL, Thermo Scientific Inc, USA).

Thiobarbituric acid reactive substances (TBARS) and diene conjugate (DC) levels were measured to assess hepatic lipid peroxidation. TBARS levels were determined according to Buege and Aust and calculated from absorption at 532 nm by using the molar extinction coefficient of $1.56 \times 10^5 M^{-1}$ (20). DC levels were determined spectrophotometrically at 233 nm. For this assay, liver lipids were extracted in chloroform/methanol (2:1; v/v) mixture, and then the extracted lipids were dissolved in cyclohexane. DC levels were calculated using a molar extinction coefficient of $2.52 \times 10^4 M^{-1}$ cm⁻¹ (20).

PC levels were evaluated according to the method of Reznick and Packer, which is based on the measurement of protein hydrazones (21). Results were calculated from the maximum absorbance (360 nm) using a molar extinction coefficient of 22,000 M^{-1} cm⁻¹.

Protein glycation products in liver and serum

Hepatic advanced oxidation products of protein (AOPP) levels were measured spectrophotometrically at 340 nm (22). AGEs levels were determined spectrofluometrically in the liver and serum. For this reason, liver homogenates or serum were diluted with PBS (pH 7.4) and the fluorescence intensity ($\lambda_{emission}$: 440 nm; $\lambda_{excitation}$: 350 nm) was determined (23). N^e-(carboxymethyl)lysine (CML) levels were measured using ELISA kits (Abbkine, Wuhan, China).

Determination of antioxidant system parameters

Hepatic ferric reducing anti-oxidant power (FRAP) levels were determined spectrophotometrically to evaluate antioxidant power in liver. Reducing a ferric-tripyridyltriazine complex to the ferrous form by the antioxidants in liver homogenates is monitored by measuring the change in absorption at 593 nm (24). Hepatic glutathione (GSH) levels were determined by using 5,5-dithiobis-(2-nitrobenzoate) at 412 nm (25).

Superoxide dismutase (SOD) activity was determined according to Mylorie et al., and calculated using a standard curve prepared by bovine SOD (26). Glutathione peroxidase (GSH-Px) activity was measured using cumene hydroperoxide as a substrate (27). The reaction was followed spectrophotometrically (340 nm) at 37°C, and activity was calculated using the extinction coefficient of NADPH (6.22x10³ M⁻¹cm⁻¹). Catalase (CAT) activity was measured spectrophotometrically at 240 nm by using hydrogen peroxide (H_2O_2) as substrate (28). One unit of CAT was considered as the activity of enzyme needed to degrade 1 μ mol H₂O₂ per min at 25°C.

Determination of mRNA expressions of nuclear erythroid factor 2-related factor (Nrf2) superoxide dismutase (SOD), glutathione peroxidase (GSH-Px) and catalase (CAT)

In order to obtain mRNA expressions of Nrf2, SOD, GSH-Px and CAT in liver, the tissue was homogenized using a handheld homogenizer (SCILOGEX D160, USA). Total RNA was isolated using an RNA isolation kit (NucleoSpin RNA Kit, #740955, Macherey-Nagel, Germany). cDNA was synthesized with a SCRIPT cDNA Synthesis Kit (Jena Bioscience, GmbH, Jena, Germany) by using 5 ng RNA. Nrf2 (F: 5'-GTGGATCT-GTCAGCTACTCCC-3'; R: 5'-CTGGGAATATCCAGGG-CAAGC-3'), SOD (F: 5'-GGTCCAGCGGATGAAGAG-3'; 5'-GGACACATTGGCCACACC-3'), GSH-Px (F: R: 5'-CGACATCGAACCCGATATAGA-3'; R: 5'-ATGCCT-TAGGGGTTGCTAGG-3'), CAT (F: 5'-AGGTGCTTTTG-GATACTTTGAGG-3'; R: 5'-CGACTGTGGAGAATC-GGACGG-3') primers, and the housekeeping gene GAPDH (F: 5'-CAGGGCTGCCTTCTCTGTG-3'; R: 5'-AACTTGCCGTGGGTAGAGTC-3') were purchased from LGC Biosearch Technologies (Denmark). Quantitative real-time polymerase chain reaction (qPCR) was performed using qPCR Green Master with UNG (Jena Bioscience, GmbH, Jena, Germany) in a real-time PCR system (Biorad CFX Connect, California, USA). The expression levels of mRNAs were quantified using the 2 -AACt method.

Protein levels

Protein levels were determined spectrophotometrically using bicinchoninic acid (29).

Histopathologic examination

Livers were fixed in 10% buffered formalin, embedded in paraffin, sectioned and stained with hematoxylin and eosin (H&E) for histologic examinations. Masson's trichrome (MTC) staining was also performed to show reticulin fibers of fibrotic areas. Steatosis, liver damage and fibrosis scores were made according to the protocol proposed by Goodman, as previously described in detail in our previous study (30, 31).

Statistical analysis

Statistical analysis was evaluated by using the Statistical Package for The Social Sciences program (21.0; SPSS Inc., Chicago, IL, USA) program. All variables were expressed as mean±standard deviation (SD). Data distributions and test of normality were investigated by Kolmogorov-Smirnov test. One-way ANOVA test (post-hoc Tukey's test) was used to assess the parameters with normal distribution. Homogeneity of variances was evaluated with Levene test. Kruskal-Wallis test (post-hoc Mann Whitney-U test) was used to compare the parameters without normal distribution. In all cases, a difference was considered significant when p<0.05.

RESULTS

Final body weight did not change, but liver weight and liver index were elevated in HFr and VDD+HFr groups as compared to control group. Serum 25(OH)D₃ levels did not alter in the HFr group, but it decreased (47.3%) in the VDD+HFr group. Calcium and inorganic phosphorus levels remained unchanged in both groups as compared to the control (Table 1).

Significant increases in glucose, TG and HOMA levels, ALT and AST activities were detected in the HFr and VDD+HFr groups as compared to the control. However, insulin levels remained unchanged in both groups. Glucose and HOMA levels were higher in the VDD+HFr group than the HFr group (Table 2).

levels of 25(OH)D3, calcium, and phosphorus in high fructose (HFr)-treated rats (Mean±SD)					
	Control (n=6)	VDD (n=8)	HFr (n=7)	VDD+HFr (n=8)	
Body weight (g)	299.5±16.1	315.5±29.8	312.1±31.5	336±16.5	
Liver weight (g)	8.25±0.93	7.55±0.70	10.3±1.46 ^{a2}	10.9±0.83 ^{a1}	
Liver index (%)	2.75±0.21	2.41±0.36	3.28±0.27 ^{a2}	3.22±0.21 ^{a3}	
25(OH)D₃ (ng/mL)	28.4±3.09	15.7±1.07 ^{a2}	25.3±5.88	15.0±1.51 ^{a2,b1}	
Calcium (mmol/L)	2.42±0.16	2.50±0.18	2.36±0.12	2.39±0.15	
Phosphorus (mmol/L)	2.55±0.18	2.58±0.26	2.30±0.16	2.40±0.29	

Table 1: The effect of vitamin D deficiency (VDD) on body weight, liver weight, and liver index* values and serum

^{a1}p<0.001, ^{a2}p<0.01, ^{a3}p<0.05 as compared to control; ^{b1}p<0.001 HFr vs VDD+HFr *Liver index= Liver weightx100/body weight

(in)-treated rats. (mean±3D)				
	Control (n=6)	VDD (n=8)	HFr (n=7)	VDD+HFr (n=8)
Glucose (mmol/L)	7.53±1.44	9.76±1.04 ^{a3}	12.2±1.09 ^{a1}	14.3±1.67 ^{a1,b3}
Insulin (mmol/L)	29.2±4.37	29.3±2.98	24.5±2.01	27.9±5.29
HOMA-IR	1.61±0.33	2.10±0.21 ^{a2}	2.21±0.23 ^{a2}	2.96±0.68 ^{a2,b3}
TG (mmol/L)	0.48±0.09	1.13±0.26 ^{a2}	0.79±0.18 ^{a2}	0.71±0.05ª3
ALT (U/L)	43.3±13.1	53.6±8.53	58.0±4.86 ^{a3}	67.9±10.3 ^{a1}
AST (U/L)	108.0±10.2	114.6±19.0	139.6±22.1 ^{a2}	148.8±27.1 ^{a2}

Table 2: The effect of vitamin D deficiency (VDD) on some biochemical parameters in serum of high fructose (HFr)-treated rats. (Mean±SD)

^{a1}p< 0.001, ^{a2}p< 0.01, ^{a3}p< 0.05 as compared to control; ^{b3}p<0.05 HFr vs VDD+HFr

In hematoxylin and eosin (H&E) and Masson's trichrome (MTC) staining of liver sections, the HFr group exhibited marked microvesicular steatosis and hepatocyte ballooning without fibrotic changes. Some alleviations in histopathological findings were obtained in the VDD+HFr group (Figure 1). Steatosis score decreased significantly in the VDD+HFr (1.63±0.52) group as compared to HFr (3.00±0.00). However, the ballooning score did not change significantly in the VDD+HFr (1.63±0.52) as compared to HFr (2.00±0.00) group.

Hepatic TG and TNF- α levels and MPO activity were elevated, but Hyp levels remained unchanged in the HFr and



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Figure 1: The effect of vitamin D deficiency (VDD) on histopathological changes in liver tissues of high fructose (HFr)-treated rats (H&E and MTC, x100)

VDD+HFr groups as compared to the control. TNF- α levels and MPO activity, but not TG and Hyp levels were higher in the VDD+HFr group than the HFr group (Figure 2).



Figure 2. The effect of vitamin D deficiency (VDD) on hepatic triglyceride (TG), hydroxyproline (Hyp), tumor necrosis factor-alpha (TNF- α) levels and myeloperoxidase (MPO) activities in high fructose (HFr)-treated rats (Mean±SD). ^{a1}p< 0.001, ^{a2}p< 0.01, ^{a3}p< 0.05 as compared to control; ^{b1}p<0.001, ^{b3}p<0.05 HFr vs VDD+HFr

Hepatic ROS, TBARS, DC and PC levels were elevated in both groups. However, the TBARS levels were higher in the VDD+HFr group than the HFr group (Figure 3). Hepatic AGEs and AOPP together with serum AGE and CML levels were increased in the HFr and VDD+HFr groups as compared to the control (Table 3).

There were no changes in FRAP and GSH levels in the HFr and VDD+HFr groups. Hepatic SOD and GSH-Px activities diminished significantly, but CAT activity did not alter in the HFr group as compare to the control. The VD-D+HFR and control groups did not differ in terms of antioxidant enzymes activities. When the HFr and VDD+HFr groups were compared, increases in SOD (30.8%), GSH-Px(16.6%) and CAT (20.7%) activities were increased in the VDD+HFr group. However, increases in GSH-Px and CAT activities were not significant (Table 4).

mRNA expressions of Nrf2, SOD, GSH-Px, and CAT were decreased in the HFr group. In the VDD+HFr group,



Figure 3. The effect of vitamin D deficiency (VDD) on hepatic reactive oxygen species (ROS), thiobarbituric acid reactive substances (TBARS), diene conjugates (DC) and protein carbonyl (PC) levels in high fructose (HFr)-treated rats (Mean±SD). ^{a1}p<0.001, ^{a2}p<0.01 as compared to control; ^{b3}p<0.05 HFr vs VDD+HFr

mRNA expression of SOD increased significantly as compared to the control, and Nrf2 and GSH-Px mRNA expressions returned to control levels. However, Nrf2, SOD and GSH-Px expressions were significantly higher than the HFr group. mRNA expression of CAT did not alter among the groups (Figure 4).



Figure 4. The effect of vitamin D deficiency (VDD) on hepatic mRNA expressions of nuclear erythroid factor 2-related factor (Nrf2) superoxide dismutase (SOD), glutathione peroxidase (GSH-Px) and catalase (CAT) in high fructose (HFr)-treated rats (Mean \pm SD). ^{a1}p<0.001, ^{a2}p<0.01 as compared to control; ^{b1}p<0.001, ^{b3}p<0.05 HFr vs VDD+HFr

Table 3: The effect of vitamin D deficiency (VDD) on serum advanced glycation end products (AGE) andcarboxymethyllysine (CML) levels as well as hepatic advanced oxidized protein products (AOPP) and AGE in highfructose (HFr)-treated rats (Mean±SD)

	Control (n=6)	VDD (n=8)	HFr (n=7)	VDD+HFr (n=8)
Serum AGE (RFU)	186.3±15.4	185.1±17.3	250.6±29.8°1	260.5 ± 17.6^{a1}
Hepatic AGE (RFU)	351.3±18.1	330.5±20.7	549.5 ± 22.8^{a2}	589.5 ± 82.4^{a2}
Hepatic AOPP (nmol/mg protein)	22.8±0.96	24.5±1.89	28.7 ± 3.02^{a1}	30.6±2.46 ^{a1}
Serum CML (µg/L)	12.3±1.75	13.7±4.35	17.34±3.00 ^{a3}	16.9±1.20ª3

 $^{\rm a1}p<0.001,\,^{\rm a2}p<0.01,\,^{\rm a3}p<0.05$ as compared to control

Table 4: The effect of vitamin D deficiency (VDD) on hepatic ferric reducing antioxidant power (FRAP), and glutathione (GSH) levels as well as hepatic superoxide dismutase (SOD), glutathione peroxidase (GSHPx), and catalase (CAT) activities in high fructose (HFr)-treated rats (Mean±SD)

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	Control (n=6)	VDD (n=8)	HFr (n=7)	VDD+HFr (n=8)	
FRAP ((nmol/mg protein)	80.3±11.9	76.9±15.8	65.6±12.4	77.5±11.8	
GSH (nmol/mg protein)	23.9±3.02	24.0±6.30	19.4±4.02	17.3±4.16	
SOD (U/mg protein)	20.1±2.89	18.7±1.43	16.4±1.78°3	21.47 ± 2.98^{b2}	
GSH-Px (nmol/min/mg protein)	520.9±64.1	433.0±71.9	383.7±32.5ª2	447.4±60.5	
CAT (µmol/min/mg protein)	352.6±47.7	366.3±76.2	314.4±36.4	379.4±44.7	

 $^{\rm a2}p{<}\,0.01,\,^{\rm a3}p{<}\,0.05$ as compared to control, $^{\rm b2}p{<}0.01$ HFr vs VDD+HFr

In the VDD group, there were significant decreases in serum 25(OH)D₃ and TG levels and significant increases in glucose and HOMA values as compared to the controls. In addition, hepatic TG and TNF- α levels and MPO activity were also increased. There were no changes in oxidative stress and protein glycation parameters and histopathological findings remained unchanged.

DISCUSSION

Several dietary models such as high fat, HFr, westernand methionine choline deficiency are used to produce NAFLD/NASH (4). Some investigators have reported that the HFr diet resulted in increased oxidative stress, increases in cytokines and AGEs levels in serum and liver (32-36). This diet caused hepatic lesions such as fatty liver, hepatocyte ballooning and lobular inflammation (32-36). Therefore, in this study, the role of the VDD diet in the pathogenesis of NAFLD was investigated by examining hepatic glycooxidative stress, inflammation and histopathology together with IR in a HFr-induced NAFLD model.

In the current study, rats received 30% fructose containing drinking water for eight weeks as previously reported (35, 36). Prooxidant status was evaluated by determining ROS, oxidation products of lipids (TBARS and DC) and proteins (PC and AOPP). AGEs are heterogenous products, and fluorescent AGEs and non-fluorescent CML levels were determined as protein glycation indicators (6). AOPP was reported to have some homologies with AGEs and to reflect protein glycooxidation (37). Determinations of TNF- α levels and MPO activity were used to examine hepatic inflammation. According to our results, the HFr diet caused IR, hypertriglyceridemia, increases in glycooxidative stress and inflammation together with histopathological changes (marked microvesicular steatosis, hepatocyte ballooning and no fibrosis) in rats. These findings indicate that the hepatic lesions were produced successfully in the HFr-induced NAFLD model.

Nrf2 is a transcription factor that is the main regulator of the antioxidant system. It regulates redox homeostasis by increasing the expression of many antioxidant enzymes such as SOD, GSH-Px, and CAT and plays a key role in combating oxidative stress (12, 14). The antioxidant potential of $1,25(OH)_2D_3$ is related to its direct antioxidant potential and the upregulation of Nrf2 and its target antioxidant genes (8, 12, 14, 38). It has a suppressive effect against oxidative stress in dietary NAFLD models due to these properties (39-41).

On the other hand, VDD was reported to have an additional effect in the experimental dietary model of NA-FLD as a second hit (42-45). However, some contradictory results are also available (46, 47). VDD diet exacerbated high fat diet (HFD)- (42, 43) or HFD+HFr (44)-induced metabolic changes and progressed NAFLD into NASH by increasing IR, inflammation and lipogenesis and decreasing β -oxidation. Similarly, long term feeding of VDD plus HFr-diet was reported to cause progression of NA-FLD into NASH by increasing hepatic steatosis, inflammatory lesions and interstitial fibrosis (45). Contrarily, VDD alleviated HFD-induced overweight, hyperinsulinemia, and hepatic steatosis through promoting fatty acid β -oxidation (46). It has also been reported that there was no difference between HFr and VDD+HFr groups in terms of inflammation, steatosis and expressions of genes related to lipogenesis and β -oxidation (47). However, there is not enough information about how VDD affects glycooxidative stress and antioxidant system parameters in dietary NAFLD models. For this reason, this subject was focused on in our study.

Serum 25(OH)D levels were reported as 27-30 ng/ml in normal mice fed on standard rodent chow containing Vit D_3 (1,500 IU/kg). In the same study, it was reported that the administration of a diet containing low doses of Vit D_3 (50 and 250 IU/kg, respectively) resulted in serum 25(OH)D levels compatible with severe VDD and VDD, respectively, as early as four weeks, and these levels were maintained for 3-4 months (48). Therefore, in our study, HFr drinking was started four weeks after VDD diet application and continued for eight weeks and then serum Vit (OH)D levels were found between 10-20 ng/ml in rats fed on a diet to which Vit D_3 is not added (48).

Some authors have reported that VDD diet alone resulted in moderate steatosis, inflammation, IR together with upregulation of genes related to lipogenesis and inflammation in the liver of rodents (43, 47). However, some investigators have found that there were no changes in metabolic parameters and liver damage markers due to a VDD diet (46). There is limited knowledge about prooxidant-antioxidant status due to VDD (49, 50). In this study, the VDD diet alone was observed not to alter hepatic prooxidant-antioxidant balance, mRNA expressions of Nrf2 and antioxidant enzymes, protein glycation parameters and histopathological findings despite increased inflammation, liver TG levels and IR.

In this study, when VDD and HFr diets were combined, the VDD diet did not augment HFr-induced oxidative stress, protein glycation, and hepatic damage. However, IR and inflammation parameters were higher in the VDD+HFr group than HFr. There was no difference in GSH and FRAP levels and CAT activity between the VDD+HFr and HFr groups. Although SOD and GSH-Px activities decreased in the HFr group, these activities returned to normal in the VDD+HFr group. This situation may be related to increases in gene expressions of these enzymes. Indeed, interestingly, Nrf2, SOD and GSH-Px mRNA expressions were significantly higher in the VD-D+HFr group than the HFr group. These changes in the antioxidant system may provide a protection against an additional increase in prooxidant status and HFr-induced liver damage in the VDD+HFr group.

CONCLUSION

Our results indicate that VDD augmented HFr-induced insulin resistance and hepatic inflammation, and upregulated Nrf2 and antioxidant enzymes while exhibiting no change in glycooxidative stress and histopathological changes. These results suggest that more studies are necessary to elucidate the effect of VDD on NAFLD progression.

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