Value of Vitamin D Administration Versus Suppressing Gut Microbiota in Modifying Metabolic Associated Osteoarthritis in Rats

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Abstract

Background: Metabolic syndrome (MetS) is considered as common risk factors for knee osteoarthritis (OA).

Aim of Study: Comparing the effect of vit. D supplementation and gut microbiota suppression on the development of metabolic knee OA in rats is the aim of the study.

Material and Methods: Forty adult male rats were subdivided into control and high carbohydrate high fat (HCHF) fed rats that furtherly subdivided into: HCHF (positive control), HCHF-Vitamin D treated & HCHF-antibiotics treated. Body mass index (BMI), serum glucose, insulin, total cholesterol, triglycerides, HDL-C, LDL-C, inflammatory cytokines were estimated. Homeostatic model assessment for insulin resistance (HOMA-IR) was calculated. Synovial cytokine, tissue degenerative enzymes gene expression and knee histopathology were assessed.

Results: HCHF induced significant increase in BMI, glucose, insulin, HOMA-IR, serum and synovial cytokines and degenerative enzymes with dyslipidemia and knee joint damage compared with controls. Vit. D improved all parameters in comparison to HCHF fed group. Antibiotic administration improved glucose, insulin, HOMA-IR, dyslipidemia and serum cytokines but induced insignificant changes in BMI, synovial cytokines, degenerative enzymes and knee joint damage compared with HCHF fed group.

Conclusion: HCHF diet led to development of knee joint damage associated with changes in the metabolic profile in rats. Vit D supplementation dramatically improved knee joint damage, obesity insulin resistance, dyslipidemia and inflammation systemically and locally while controlling gut dysbiosis by antibiotics did not improve obesity, local inflammation and knee damage but improved the other parameters less than Vit D.

Key Words: Inflammation – Metabolic syndrome – Obesity – Oxidative stress.

Introduction

OSTEOARTHRITIS (OA) is a common chronic joint disorder and one of the causes of disability [1], with a multifactorial etiology including post-

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traumatic, ageing, genetic [2]. Metabolic OA is a newly defined phenotype associated with metabolic syndrome (MetS) [3]. MetS associates with a state of chronic low-grade inflammation, through elevated levels of cytokines, acute-phase inflammatory components as complements and C reactive protein (CRP) and other mediators has been implicated in the pathogenesis of OA [4].

It was reported that metabolic syndrome is accompanied with changes in gut microbiota [5]. All metazoan, from invertebrates to vertebrates. has intestinal microbiota that plays a key role in host metabolism, immune function, nutrition [6]. Gut microbial dysbiosis is considered one of the pivotal risk factors for obesity, atherosclerosis, cardiovascular diseases, type 2 diabetes (T2DM), and MetS [7]. The association of the gut microbiome with osteoarthritic pain and severity in the knee was recently suggested indicating that controlling microbiome is a possible therapeutic target for osteoarthritis-related knee pain [8]. The intestinal microbiota is strictly associated with the pathogenesis of numerous disorders and there is emerging evidence supports the hypothesis that the gut microbiome could be possible triggering factors for metabolic OA [9].

The main function of vitamin D is regulation of bone metabolism and calcium homeostasis via the nuclear receptors named vitamin D receptors (VDRs). Thus, inadequate vitamin D status is thought to affect the pathophysiology and progression of OA [10].

Furthermore, Vitamin D is believed to have effects on inflammation, alters cytokine synthesis and has positive effects on muscle strength [11].

Results from human studies had controversial findings as three randomized controlled trials

(RCTs) found an insignificant role of vitamin D supplementation on cartilage volume or pain in knee OA [12-14] while one RCT showed a significant decrease in OA pain, and an increase in knee function [15]. A small non-controlled study found a significant increase in muscle strength after two months of vitamin D supplementation to patients with knee OA [16]. In a recent trial, significantly improved structural and functional outcomes were found in patients with knee OA who maintained sufficient plasma vitamin D compared to those insufficient [17].

On basis of these data, we suppose that if we compare the effect of vit. D supplementation and gut microbiota suppression on the development of metabolic knee OA in diet-induced rat model of MetS; we could recommend new insights in controlling metabolic osteoarthritis. So, comparing the effect of vit. D supplementation and gut microbiota suppression on the development of metabolic knee OA in diet-induced rat model of MetS is the aim of the current study.

Material and Methods

A total number of 40 adult male Wistar rats aged between 10 and 12 weeks (200 ±20g) were purchased from animal house of Faculty of Veterinary Medicine, Zagazig University, Egypt. All experimental procedures and protocols were following the guide for the care and use of laboratory animals (8th edition, National Academies Press) and have been reviewed and approved by Zagazig University institutional animal care unit committee (ZU-IACUC; Sharkia; Egypt) with approval number: ZU-IACUC/3/F/115/2019. Before starting the experiment, all animals subjected to 14 days for house acclimatization under conditions of controlled temperature (24-26°C), humidity (50-60%) and 12 hrs light-dark cycle. They were fed on a standard diet with free access to water.

The animals were randomly divided into 2 main groups: Control group (n=10): Were fed standard rat chow which consisted of 25.8% protein, 62.8% carbohydrate, and 11.4% fat, and tap water and high carbohydrate high fat (HCHF) group (n=30): The rats fed high carbohydrate high fat (HCHF) diet for 12 weeks, (1kg) contained 375g of sweetened condensed milk, 200g of ghee, 175g of fructose, 155g of powdered rat food, 25g of Hubble Mendel and Wakeman salt mixture and 50mL of water. The drinking water was supplemented with 25% fructose [18]. HCHF animals were subdivided into three subgroups, 10 rats each: HCHF (positive control): Rats were fed HCHF diet, HCHF+D: Rats

were fed HCHF diet with vit D3 supplementation and HCHF+AB: Rats were fed HCHF diet receiving broad spectrum antibiotics in drinking water. The experiment lasts for 12 weeks after acclimatization of the rats to the experimental conditions. Both food formulae were obtained from Faculty of Agriculture, Zagazig University, Egypt.

Vitamin D3 supplementation:

Oral vitamin D3 (vidrop) (cholecalciferol) (Medical union pharmaceuticals, Abu-Sultan, Ismailia, Egypt) was diluted in corn oil and then given orally by gavage daily (500IU/Kg/day) for a total duration of 12 weeks [19]. At the same time, rats in groups; control, HCHF, I HCHF+AB were given oral corn oil by gavage.

Broad-spectrum antibiotics regimen:

Rats received antibiotics in drinking water composed of Vancomycin + metronidazole 0.5-1.0g/L each for a total duration of 12 weeks [20,21].

Blood sample collection:

The rats were sacrificed after 12 hours of fasting under anesthesia (chloral hydrate) inhalation. Blood samples were obtained by exsanguination at the time of scarification, collected and allowed to clot for 2 hours at room temperature before centrifugation. Sera were stored at -20 °C until analysis. Repeated freezing and thawing was avoided.

Determination of metabolic parameters:

Body mass index (BMI) was calculated for each animal at the end of the experiment by the equation: BMI (gm/cm²)=Body weight/length² (nose to anus length) this index can be used as an indicator of obesity where the cutoff value of obesity BMI is more than 0.68gm/cm² [22]. The sera were examined for level of insulin by ELISA kits (Bio Basic INC, NY, USA) and glucose by enzymatic colorimetric assay using commercial kits (Cayman chemicals, Michigan, USA). The homeostatic model assessment for insulin resistance (HOMA-IR) was calculated from the formula: $[HOMA-IR = Insulin (mIU/L) \times glucose (mg/dL)/$ 405 [23]. For lipid profile assessment, total cholestero l (TC), HDL-cholesterol, LDL-cholesterol and triglycerides (TG) using enzymatic colorimetric assays using commercial kits (Calgary Lab Service, Calgary, Canada) were examined.

Determination of inflammatory cytokines:

By using ELISA kits; serum and synovial fluid pro inflammatory cytokines; interleukin-6 (IL-6) (Ray Biotech Inc. USA), USA) & tumor necrosis factor α (TNF- α) (Koma Biotech. Inc. Korea) and anti-inflammatory cytokine interleukin-10 (IL-10)

(DRG International Inc., were determined as the recommendations of the manufacturers.

Real time-Polymerase chain reaction (RT-PCR) measurement of tissue degenerative enzymes gene expression:

Right knee joints and subchondral bones were dissected, rinsed in ice-cold saline and were frozen at -80°C for subsequent biochemical measurements of myeloperoxidase (MPO), matrix metall-oproteinase2 (MMP2), and tissue inhibitor metall-oproteinase2 (TIMP2) by RT-PCR technique [24].

Total RNA was extracted from joint tissue samples using SV Total RNA isolation system (Promega, Madison, WI, USA) which provides fast and simple technique for isolation of RNA.SV mean (spin or vacuum) purification protocol. The extracted RNA was reverse transcribed into cDNA and amplified by PCR using RT-PCR kit (Stratagene, USA). The relative gene expression ratio is calculated from the real-time PCR using the $2-\Delta\Delta$ Ct method and glyceraldehyde-3-phosphate dehydrogenase (GAP-DH) as reference gene. The sets of primers used were listed in Table (1).

Table (1): Primer sequences.

Gene	Forward	Reverse
MMP2 TIMP2	5'-TATAGAACTCCAACCACAC-3' 5'-CGCTACCAACGGTGGAAGRC-3'	5'-CCGTAGAAGGTCCTCGTC-3' 5'-CAAGCTCCCCCTCTTTTTCA-3'
MPO	5'-ACCTACCCAGTACCGATCC-3'	5'-AACTCTCCAGCTGGCAAAAA-3'
GAPDH	5'-ATGACTCTACCCACGGCAAG-3'	5'-CTGGAAGATGGTGATGGGTT-3'

Myeloperoxidase = MPO.
Matrix metalloproteinase2 = MMP2.

 $\label{eq:total_continuous_continuous} Tissue\ inhibitor\ metalloproteinase 2 = TIMP2.$ $Glyceraldehyde-3-phosphate\ dehydrogenase = GAPDH$

Knee joint histopathology: The left knee was harvested by cutting the femur and tibia/fibula 2cm above and below the joint line, respectively. Muscles were removed and joints were fixed kept in 10% buffered formalin and 1% HNO3 for 1 day or more till they become soft, rinsed with runny water then put in alcohol series (different concentrations of alcohol) in order to remove all water from tissues before putting in paraffin, and were kept in xylene paraffin. Serial sagittal plane sections of 10 mutickness were obtained, stained with hematoxylin and eosin (H & E) for histopathological examination. All histological sections were scored using a Modified Mankin Histology Scoring System [25].

Statistical analysis: The results are presented as descriptive statistics (mean \pm standard deviation). Statistical analysis was performed using the Statistical Package for Social Science (SPSS) version 25 (SPSS, Inc., IBM Company, Chicago, IL, USA). The normal distribution of data from each group was confirmed using the Kolmogorov-Smirnov normality test. Since the test indicated that variables followed normal distribution, comparisons among the experimental groups were analyzed by oneway analysis of variance (ANOVA) followed by least significance differences (LSD) test to evaluate statistical difference between two groups. p-value <0.05 was considered to be statistically significant.

Results

Metabolic parameters: There was significant increase in insulin, glucose, HOMA-IR, in HCHF

and HCHF+AB group in comparison to control one (p<0.001) with insignificant change in these parameters in HCHF+D when compared with control group (p=0.75, 0.54, 0.173 respectively) while there was significant increase in BMI, TC, TG, LDL-C with decrease in HDL-C in all groups in comparison to control one (p<0.001). Both Vit D and Antibiotic administration induced significant decrease in all parameters (p<0.001) and significant increase in HDL-C (p<0.001, p=0.002 respectively) in comparison to HCHF group except BMI which showed insignificant change in HCHF+AB group compared with HCHF group (p=0.119). There was significant increase in insulin, glucose, HOMA-IR (p<0.001), BMI (p=0.016), TC (p=0.007), TG (p=0.002), LDL-C (p=0.001) and significant decrease in HDL-C (p=0.025) in HCHF+AB when compared with HCHF+D (Table 2).

Serum and synovial inflammatory cytokines: There was significant increase in serum and synovial levels IL-6 and TNF-α with decrease in IL-10 in all groups in comparison to control one (p<0.001). There was significant decrease in serum and synovial levels IL-6 and TNF-α with significant increase in IL-10 in HCHF+D group in comparison to HCHF one (p<0.001). Antibiotic administration induced the same effect like Vit D on the serum cytokines (p<0.001) while produced insignificant changes in synovial IL-6, TNF-α and IL-10 in comparison to HCHF group (p=0.064, 0.089, 0.112respectively). There was significant increase in serum and synovial levels IL-6 and TNF- α with decrease in IL-10 in HCHF+AB group in comparison to HCHF+D group (p<0.001) (Table 3).

Tissue degenerative enzyme gene expression:

There was significant increase in tissue MPO and MMP2 with decrease in TIMP2 in all groups in comparison to control animals (p < 0.001). There was significant decrease in tissue MPO and MMP2 with significant increase in TIMP2 in HCHF+D group in comparison to HCHF one (p<0.001).

Antibiotic administration produced insignificant changes in tissue MPO, MMP2 and TIMP2 in comparison to HCHF group (p=0.132, 0.86, 0.173respectively). There was significant increase in tissue MPO and MMP2 with decrease in TIMP2 in HCHF+AB group in comparison to HCHF+D group (p < 0.001) (Table 4).

Table (2): Effect of vit D3 supplementation and antibiotic administration to HCHF fed rats on BMI. Glucose metabolism and lipid profile parameters.

	Control	HCHF	HCHF+D	HCHF+AB
BMI (gm/cm ²) Insulin (mIU/L) Glucose (mg/dL) HOMA-IR TC (mg/dL) TG (mg/dL) HDL-C (mg/dL) LDL-C (mg/dL)	0.41±0.036	0.76±0.027a	$0.71\pm0.022\mathbf{a'b}$	0.74±0.035 a'c
	18.8±2.5	36.1±2.3a	$20.9\pm1.8\mathbf{b}$	30.1±3.3 a,b,c
	96.8±6.4	136.9±5.3a	$102.3\pm6.4\mathbf{b}$	118.1±5.04a'b'c
	4.6±0.4	12.5±2.3a	$5.4\pm0.7\mathbf{b}$	8.1±0.9a'b'c
	107.2±7.3	157.1±7.9	$119.3\pm5.03\mathbf{a'b}$	128.3±7.3 a'b'c
	66.8±10.4	124.6±7.1a	$86.8\pm11.7\mathbf{a'b}$	102.5±11.3 a,b,c
	54.8±9.1	26.7±6.7a	$43.2\pm4.5\mathbf{a,b}$	36.3±5.03 a'b'c
	42.9±4.9	112.5±7.5a	$71.8\pm6.9\mathbf{a'b}$	82.7±7.1 a,b,c

n=10 in each group.

Data are represented as mean \pm SD.

BMI: Body mass index.

HDL-C: High-density lipoprotein-cholesterol.

HCHF: High carbohydrate high fat.

HCHF+ AB: High carbohydrate high fat+ antibiotic. HCHF+D: High carbohydrate high fat + vit D.

HOMA-IR: Homeostasis model assessment-insulin resistance.

LDL-C: Low-density lipoprotein-cholesterol.

TC: Total cholesterol.

TG: Triglyceride. Significance (*p*<0.05):

- (a) Significant when compared with control group.
- (b) Significant when compared with HCHF fed group.
- (c) Significant when compared with HCHF+D group.

Table (3): Effect of vit D3 supplementation and antibiotic administration to HCHF fed rats on serum and synovial fluid inflammatory cytokines (IL-6, TNF- α & IL-10) levels.

	Control	HCHF	HCHF+D	HCHF+AB
Serum: IL-6 (pg/ml) TNF-α (pg/ml) IL-10 (pg/ml)	10.8±2.9 15.3±3.6 54.1±5.01	43.9±4.4a 64.2±6.6a 17.9±5.07a	20.3±3.4 a 'b 30.5±5.6 a 'b 40±6.03 a 'b	33.6±4.4a'b'c 42.8±4.9a'b'c 30.2±3.7a'b'c
Synovial fluid: IL-6 (pg/ml) TNF-α (pg/ml) IL-10 (pg/ml)	596±55 91.6±6.4 928 ±27.8	$2074.5 \pm 109 \mathbf{a}$ $240.3 \pm 11.5 \mathbf{a}$ $280 \pm 11.7 \mathbf{a}$	970.7±87 a ' b 137.9±6.8 a ' b 735±34.8 a ' b	1994±119 a,c 232.5±13.4 a °c 298±18.3 a °c

⁻ n = (10) in each group Data are represented as

mean ± standard deviation. IL-6: Inerlukin-6.

IL-10: Inerlukin-10.

TNF-α: Tumor necrotic factor-α.

Significance (p < 0.05):

- (a) Significant when compared with control group.
- (b) Significant when compared with HCHF fed group.
- (c) Significant when compared with HCHF+D group.

Table (4): Effect of vit D3 supplementation and antibiotic administration to HCHF fed rats on knee joint tissue degenerative enzyme gene expressions with (GAPDH) as reference gene.

	Control	HCHF	HCHF+D	HCHF+AB
MPO/GADPH MMP2/GADPH TIMP2/GADPH	0.1±0.01 1.7±0.1 26.3±2.2	1.1±0.08 a 10.2±1.6 a 13.4±2.6 a	0.5±0.06 a 'b 4.3±0.6 a 'b 21.7±2.03 a 'b	1±0.08 a 'c 9.2±1.5 a ,c 14.9±2.7 a ,c

⁻ n = (10) in each group Data are represented as mean ± standard deviation.

MPO: Myeloperoxidase.

MMP2: Matrix metalloproteinase2.

TIMP2: Tissue inhibitor metalloproteinase2.

Significance (p < 0.05):

- (a) Significant when compared with control group.
- (b) Significant when compared with HCHF fed group.
- (c) Significant when compared with HCHF+D group.

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Knee joint damage:

There was significant increase in Modified Mankin scores in all groups in comparison to control animals (p<0.001). There was significant decrease Modified Mankin in HCHF+D group in comparison to HCHF one (p<0.001). Antibiotic

administration produced insignificant changes Modified Mankin scores in comparison to HCHF group (p=0.07). There was significant increase in Modified Mankin scores in HCHF+AB group in comparison to HCHF+D group (p<0.001) (Fig. 1).

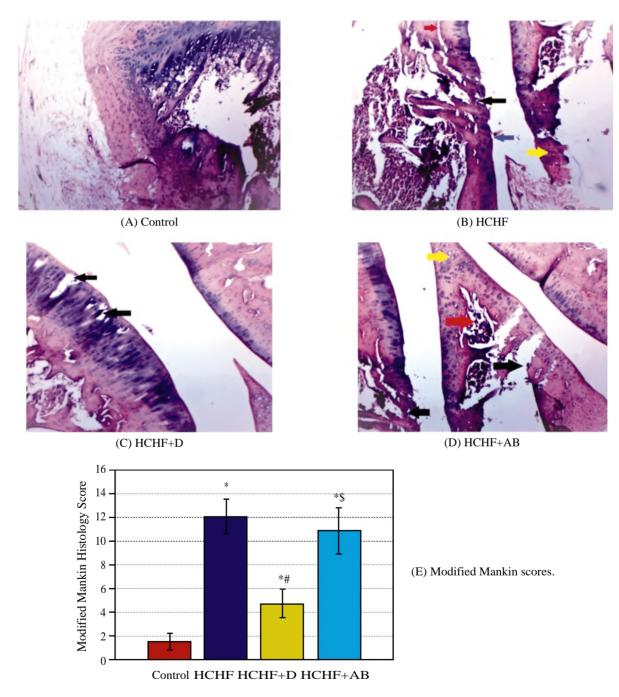


Fig. (1): Histology images taken at x200, showing the normal features of the articular cartilage of the control group and the joint damage features in the other groups in the form of weak staining of chondrocytes (yellow arrow), fibrillation and fissuring of the cartilage (black arrow), vascularization into the articular cartilage (red arrow), and osteophyte formation (blue arrow). HCHF, high carbohydrate high fat; HCHF + AB, high carbohydrate high fat + antibiotic; HCHF+D, high carbohydrate high fat + vit D. Modified Mankin scores: n = (10) in each group. Data are represented as mean \pm standard deviation. Significant when compared with HCHF fed group, (\$) significant when compared with HCHF+D group.

Discussion

In the present study, adult male rats on the HCHF diet showed increased BMI compared with normal chew fed rats and this obesity was associated with dyslipidemia, elevated insulin and glucose, levels with reduction in insulin sensitivity evidenced by increased HOMA-IR. These results come in agreement with Wong et al. [18] who demonstrated that MetS is successfully established in rats induced by the HCHF diet after 12 weeks. Other reviews discussed the various animal models of MetS [26,27].

The results of the current study revealed that a 12-week MetS induction period led to obesity, hyperglycemia, insulin resistance and altered lipid metabolism that associated with increased serum and synovial pro inflammatory and decreased antiinflammatory markers with development of histopathological OA-related changes in knee joints. These findings are supported by results of other researches that investigated the role of MetS in developing joint ostearthritis [28] or specifically knee osteoarthritis [29] in a rat model. There is a strong evidence that inflammation plays a fundamental role in osteoarthritis pathogenesis, by affecting chondrocyte catabolic responses that can be triggered within the joint either locally from tissue damage and stress responses or systemically from obesity-related inflammation that might enhance these local responses [30]. Many studies have demonstrated that adipose tissue-derived inflammatory mediators, hyperglycemia, insulin resistance, or dyslipidemia could affect the metabolism of the joint as for other organs [31,32].

TNF-a and IL-6 seem to be the main pro inflammatory cytokines and are critical mediator of the disturbed metabolism and enhanced catabolism of cartilage destruction in OA [33]. Moreover, Collins et al. [34] identified dysregulation of proinflammatory markers in serum and synovial fluid that may be involved in accelerated OA progression in rat model of MetS. IL-10 is an anti-inflammatory cytokine that has cartilage protective effect in the course of OA as it inhibits TNF a synthesis, increases proteoglycan synthesis, inhibits chondrocytes apoptosis, decrease metalloproteinases synthesis and secretion, and decreases level of PGE2. [35].

The present study showed remarkable increase in MPO and MMP2 along with TIMP2 decrease in HCHF rats that may be attributed to certain imbalance between these enzymes as MPO has been found to disrupt the physiological balance of MMPs and functions of MMP inhibitors besides the negative correlation between MPO and both TIMP2 and IL-10 reflecting a high rate of turnover in articular cartilage [36]. It could be explained by that TNF-a stimulates chondrocytes to release several proteolytic enzymes, and can drive the catabolic process in OA leading to cartilage degradation [33].

Vit D administration in the current study led to improved obesity, hyperglycemia, insulin resistance, altered lipid metabolism, disturbed serum and synovial inflammatory markers and the structural changes in knee joints produced by HCHF diet. These results come in agree with the results of Mostafa et al. [37] who demonstrated that vitamin D supplementation ameliorated hyperinsulinemia, insulin resistance, obesity as well as dyslipidemia but not hyperglycemia in rat model of MetS. In addition, vit.D administration reduced weight and food intake, inflammation and several oxidative stress markers in adipose tissue of high-fat diet fed rats [38]. Human and experimental data indicate a role for vitamin D deficiency in the development of type-2 diabetes [39], Rheumatoid Arthritis [40] and osteoarthritis [41].

In contrary, a human study demonstrated positive relationship of Vit D levels with HOMA-IR and inverse correlation with HDL but didn't found association between Vit D levels and other metabolic syndrome components [42]. Other human study founded no significant relationships between vitamin D3 levels and MetS parameters as triglycerides, LDL, HDL, and total cholesterol [43]. Several human studies have examined the relationship between vit. D supplementation and serum levels cytokines. Barker et al. [44] showed a positive relationship with IL-10 while Peterson and Heffernan [45] reported a significant inverse relationship with serum TNF-a but no relationship with IL-6, or IL-10. Another study founded no significant changes in serum levels of IL-10, IL-6, or TNFa [46].

Suppressing gut microbiota led to improved hyperglycemia, insulin resistance, altered lipid metabolism, disturbed serum but not synovial inflammatory markers while failed to modify obesity or the structural changes in knee joints produced by HCHF diet. Vit. D showed superior results when compared with controlling gut dysbiosis. Many studies suggested that a high-fat diet can lead to gut microbiota dysbiosis that increased gut permeability and metabolic endotoxemia. This in turn contributes to low-grade inflammation and insulin resistance, obesity and diabetes [47].

In our study, the antibiotics were used to deplete the gut microbiota of rats as vancomycin, which is a Gram+-targeting and metronidazole encompasses various protozoans and most anaerobic bacteria to assess the effect microbiota depletion on MetS associated OA. Although many studies examined phenotypic differences after antibiotics treatment, some studies have assessed how regimens affect commensal populations at other sites [21].

Our results supported by findings of Rios et al., who postulated that controlling gut microbiota dysbiosis by prebiotic fiber supplementation improved insulin resistance, dyslipidemia observed in this rat model of obesity. In contract, prebiotic fiber supplementation completely prevented knee joint damage that could not be achieved by using antibiotics in our study [29].

Controlling gut dysbiosis by antibiotic in the current study did not improve obesity that supported by results of other studies used antibiotics in human [48] and mice [49]. This obesity could explain unimproved knee arthritis in spite of improved insulin resistance, dyslipidemia and systemic inflammatory cytokines in antibiotic treated rats in the current study.

In conclusion, HCHF diet led to development of knee joint damage associated with changes in the metabolic profile in rats. Vit D supplementation dramatically improved knee joint damage, obesity insulin resistance, dyslipidemia and inflammation systemically and locally while controlling gut dysbiosis by antibiotics did not improve obesity, local inflammation and knee damage but improved the other parameters less than Vit D. Further investigation of controlling gut microbiota by other like prebiotic fibers or probiotic supplements on metabolic OA is recommended.

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Conflicts of interest:

All authors have no conflicts of interest.

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قيمة فيتامين (د) التكميلي مقابل قمع ميكروبات الأمعاء في تعديل إلتهاب المفاصل المرتبطة بالأيض في الجرذان

الخلفية: تعتبر متلازمة التمثيل الغذائي من عوامل الخطر الشائعة التي تؤدى إلى إلتهاب المفاصل. لذلك صممت هذه الدراسة لمقارنة تأثير مكلات فيتامين (د) بتأثير قمع ميكروبات الآمعاء الدقيقة على تطور إلتهاب المفاصل المرتبط بالآيض للركبة في الجرذان.

مواد وطرق البحث: شملت الدراسة أربعين جرذان الذكور البالغة والتي تم تصنيفها إلى أربعة مجموعات متساوية كل منها عشرة جرذان:

- مجموعة ضابطة والتي تغذت على الغذاء المعتاد للجرذان.
- مجموعة حصلت على تغذية عالية الكربو هيدرات عالية الدهون.
- مجموعة حصلت على تغذية عالية الكربوهيدرات عالية الدهون بالإضافة إلى فيتا مين (د).
- مجموعة حصلت على تغذية عالية الكربوهيدرات عالية الدهون بالإضافة إلى المضادات الحبوية.

تم تقدير مؤشر كتلة الجسم وتحليل أمصال الدم لكل من قيمة الجلوكوز، الإنسو لين، الدهون الثلاثية، الكوليسترول الكلى، البروتين الدهنى عالى الكثافة، البروتين الدهنى منخفض الكثافة، السيتو كينات الإلتهابية. كما تم حساب تقييم نموذج مقاومة الإنسولين وكذلك قياس السيتوكينات الإلتهابية في ذلال الركبة. كما تم تقييم التعبير الجينى للانزيمات التنكسية وكذلك تم عمل دراسة ميكروسكوبية للركبة.

النتائج: أثبتت النتائج أن تغذية الجرذان بتغذية عالية الكربوهيدرات عالية الدهون قد أحدثت زيادة ذات دلالة إحصائية لكل من مؤشر كتلة الجسم، الجلو كوز، الإنسولين، مقاومة الإنسولين، السيتو كينات المصلية والزلالية والتعبير الجينى للإنزيمات التنكسية بالإضافة إلى اضطراب دهون الدم وتلف مفصل الركبة بينما أدى فيتامين (د) إلى تحسين ذى دلالة إحصائية للتغيرات التى أحدثتها التغذية عالية الكربوهيدرات عالية الدهون ولكن المضادات الحيوية أدت إلى تحسن ذى دلالة إحصائية لكل من مستويات الجلوكوز، الإنسولين، مقاومة الإنسولين وخلل الدهون والسيتو كينات المصلية فقط.

الخلاصة: خلصت هذه الدراسة إلى أن الحمية الغذائية عالية الكربوهيدرات عالية الدهون أدت إلى تطور تلف مفصل الركبة المرتبط بالتغيرات في التمثيل الغذائي في الجرذان وأن مكملات فيتامين (د) حسنت بشكل كبير تلف مفصل الركبة ومقاومة الإنسولين والسمنة واضطراب دهون الدم والإلتهابات المصلية والزلالية بينما السيطرة على ميكروبات الأمعاء بالمضادات الحيوية لم تحسن السمنة أو الآلتهاب الموضعي للركبة أوتلفها واكنها حسنت العوامل الأخرى واكن بنسبة أقل من فيتامين (د).