POTENTIAL ROLE OF HAEMATOCOCUS PLUVIALIS AGAINST DIABETES INDUCED OXIDATIVE STRESS AND INFLAMMATION IN RATS

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ABSTRACT

Objective: The aim of this study is to investigate the impact of Haematococcus pluvialis extract against oxidative stress and inflammatory cytokines induced by hyperglycemia in diabetic rats.

Methods: Oxidative stress; lipid peroxide (as presented by Malondialdehyde; MDA) and nitric oxide (NO), beside total antioxidant capacity, enzymatic and non-enzymatic antioxidants including reduced glutathione, glutathione peroxidase, and glutathione reductase were evaluated. The inflammatory cytokines; tumor necrosis factor-alpha and interleukin-1 beta were also investigated in rats’ serum. Several analyses including expression of antioxidant enzyme related genes, reactive oxygen species (ROS) formation and DNA adducts were performed.

Results: The results showed that diabetes mellitus induced-rats exhibited increase in oxidative stress biomarkers and inflammatory cytokines, lower expression levels of the antioxidant enzyme genes; superoxide dismutase and glutathione S-transferase than those in control rats. In addition, diabetic rats exhibited significantly higher levels of ROS generation and 8-hydroxy-2'-deoxyguanosine (8-OHdG) formation. In contrary, supplementation of diabetic rats with H. pluvialis extract improved the negative effect of the hyperglycemia on antioxidant enzymes, the gene expression of antioxidant enzymes, and ROS generation as well as 8-OHdG formation.

Conclusion: H. pluvialis extract decreased the oxidative stress, enhanced antioxidant status and inflammatory cytokines induced by hyperglycemia in diabetic rats. The effect of H. pluvialis extract involved in the increase of expression levels of antioxidant enzyme genes; decreased the levels of ROS generation and 8-OHdG formation which may be attributed to the presence of astaxanthin in H. pluvialis extract.

Keywords: Haematococcus pluvialis, Hyperglycemia, Diabetes mellitus, Oxidative stress, Inflammatory cytokines, DNA adducts.
MATERIALS AND METHODS

Chemicals
All chemicals in this study are of analytical grade, products of Sigma, Merck and Aldrich. STZ was purchased from Sigma-Aldrich, India. All kits were the products of Biosystems (Alocbendas, Madrid, Spain), Sigma Chemical Company (St. Louis, MO, USA), and Biodiagnostic Company (Cairo, Egypt). TRizol reagent was bought from Invitrogen (Germany). The reverse transcription and polymerase chain reaction (PCR) kits were obtained from Stratagene (USA).

Materials
Cultivation of H. pluvialis
H. pluvialis (strain No. CCAP 34/7) was isolated by spreading 0.1 ml of water samples collected from Nile River phytoplankton using BG11 media for algal isolation [14] into petri dishes containing 1.5% agar for solidification. Then, single colonies of algae were re-cultivated in the specified liquid media as nonaxenic batch cultures (50 ml) at 25±2°C and 24 hr with continuous white fluorescent lamp intensity ≈2500 Lux. Cultivation was carried out on an open pond with a capacity of 70 L containing 55 L of growth media. After cultivation, the biomass was initially separated from the water by gravitational settling, then further concentrated by centrifugation [15] and dried at 40°C.

Ethanolic extract preparation of H. pluvialis
About 100 g of H. pluvialis powder was soaked in ethanol (80%) and shook on shaker (Heidolph UNIMAX 2010) for 48 hr at 150 rpm. The extract was filtered using a Buchner funnel and Whatman No. 4 filter paper, and the algal residue was re-extracted with the addition of fresh ethanol for another two times. Combined filtrates were concentrated using Rotaty evaporator (Heidolph-Germany) at 40°C under vacuum to dryness. The dry resulting extract was stored at −20°C in a freeze and dried at 40°C.

EXPERIMENT

Animals
About 50 male Wistar albino rats (180-200 g) procured from Central Animal House, National Research Centre (NRC) were used. Animals were acclimatized to the laboratory conditions at room temperature before the experimentation. Animals were kept under standard conditions of a 12 hr light/dark cycle with food and water in plastic cages with soft bedding. All the experiments were carried out between 9.00 and 15.00 hr. The protocol was approved by the NRC Ethics Committee Guidelines (approval no: 0111457) for the use and care of animals.

Drug and treatment schedule
Induction of diabetes model: For the evaluation of streptozocin (STZ) diabetic effect, type 2 diabetes was induced by intraperitoneal injection of a single dose of STZ (45 mg/kg.b.wt.) dissolved in 0.01 M citrate buffer immediately before use [17]. After STZ injection, rats had free access to food, water and were given 5% glucose solution to drink overnight to encounter hypoglycemic shock [18]. Rats were checked daily for the presence of glycosuria. Rats were considered to be diabetic if glycosuria was present for 3 consecutive days [19]. Three days after STZ injection, fasting blood samples were obtained and blood sugar was determined (≥300 mg/dl). The antioxidant glibenclamide (daonil) reference drug was orally administrated at a dose of 10 mg/kg b.wt. daily for 30 days [20].

Design
A total of 50 rats were divided randomly into 5 groups (10 rats each), as follows:
Group 1: Considered as normal, healthy control rats.
Group 2: Considered as normal rats treated with H. pluvialis ethanolic extract (150 mg/kg b.wt.) [21].
Group 3: Considered as diabetic group.
Group 4: Considered as diabetic rats orally administered with H. pluvialis ethanolic extract (150 mg/kg b.wt.).
Group 5: Considered as diabetic rats orally administered antidiabetic glibenclamide reference drug (10 mg/kg b.wt.) daily for 30 days.

Collection of blood, organs, and tissue samples
Rats were fasted overnight (12-14 hrs), anesthetized by diethyl ether and blood collected by puncture of the sublingual vein in clean and dry test tube, left 10 minutes to clot and centrifuged at 3000 rpm for serum. The separated serum was used for biochemical analysis of total antioxidant capacity (TAC), TNF-alpha (TNF-α), and IL-1 beta (IL-1β). Liver was removed immediately, homogenized in 5-10 volumes of appropriate medium using electrical homogenizer, centrifuged at 3000 rpm for 15 minutes, the supernatants (10%) were collected and placed in Eppendorf tubes then stored at −80°C and quantified for the determination of oxidative stress markers; MDA, NO as well as nonenzymatic antioxidant; reduced glutathione (GSH), antioxidant enzyme glutathione peroxidase (GPx) and glutathione reductase (GR) spectrophotometrically.

Biochemical determinations

GPx activity
GPx activity was assayed according to the method of Paglia and Valentine [22], using NADPH-coupled reduction of GSSG cata-lyzed by GR which can be measured at 340 nm.

GSH
Brain GSH was measured colorimetrically according to the method of Beutler et al. [23]. This method is based on determination of the relatively stable yellow color when 5,5’-dithiobis-2-nitrobenzoic acid (DTNB) is added to sulfhydryl compounds which can be measured at 503 nm.

Lipid peroxide (MDA)
MDA was measured by the thiobarbituric reactive species assay, which measures the production of MDA that reacts with thiobarbituric acid [24].

Nitric oxide (NO)
NO level was assayed in liver tissue by the spectrophotometric method according to Berkels et al. [25]. NO level was assayed by the spectrophotometric method. Promega’s griess reagent system is based on the chemical reaction between sulfanilamide and N-1-naphthylethylenediamine dihydrochloride under acidic condition (phosphoric acid) to give colored azo-compound which can be measured at 520–550 nm.

TAC
TAC was assayed according to the method of Koracevic et al. [26]. The method is based on determination of the ability to eliminate added hydrogen peroxide. The remaining H 2O2 is determined colorimetrically by an enzymatic reaction converting 3,5-dichloro-2-hydroxybenzene sulfonate to a colored product that is measured at 532 nm.

GR activity
The activity of GR was determined spectrophotometrically where the decrease in absorbance at 340 nm was red according to the method of Zanetti [27].

Inflammatory cytokines; TNF-α and IL-1β
Serum inflammatory cytokines were performed by ELISA; a sandwich enzyme immunoassay.

Gene expression analysis

Extraction of total RNA and complementary DNA (cDNA) synthesis
Liver tissues of male rats were used to extract the total RNA using TRIzol® Reagent (Invitrogen, Germany) Kit. The isolation method was carried out according to the manufacturer’s instructions of the above Kit. To synthesize the cDNA isolated RNA from liver tissues was reverse transcribed into cDNA [28,29].

**Quantitative real time-PCR (qRT-PCR)**

A StepOne RT-PCR System (Applied Biosystem, USA) was used to assess the copy of the cDNA of male rats to detect the expression values of the tested genes. To perform the PCR reaction, a volume of 25 μl of reaction mixtures was prepared containing 12.5 μl of SYBR® green (TaKaRa, Biotech. Co. Ltd.), 0.5 μl of 0.2 mM forward and reverse primers, 6.5 μl DNA-RNA free water, and 2.5 μl of the synthesized cDNA. The sequences of specific primer of the genes used are listed in Table 1. The relative quantification of the target genes to the reference (β-Actin) was determined using the 2−ΔΔCT method.

**Determination of intracellular ROS formation**

Intracellular ROS generation was measured in pancreatic tissue by a flow cytometer with an oxidation-sensitive dichlorodihydrofluorescein diacetate fluorescent probe, after single-cell suspensions were made [32].

**Determination of 8-OHdG by high-performance liquid chromatography (HPLC)**

DNA was extracted from rat pancreatic tissue by homogenization in buffer containing 1% sodium dodecyl sulfate, 10 mM Tris, 1 mM EDTA (pH 7.4) and an overnight incubation in 0.5 mg/ml proteinase K at 55°C. Homogenates were incubated with RNase (0.1 mg/ml) at 50°C for 55°C. Homogenates were incubated with RNase (0.1 mg/ml) at 50°C for 55°C. Homogenates were incubated with RNase (0.1 mg/ml) at 50°C for 55°C. Homogenates were incubated with RNase (0.1 mg/ml) at 50°C for 55°C.

- DNA was then digested and the adduct 8-OHdG was measured with HPLC equipped with a Coui Array system (Model 5600). Analytes were detected on two coulometric array modules, each containing four electrochemical sensors attached in series which allowed identification targets based on reduction potential. The UV detection was set to 260 nm. The HPLC was controlled and the data acquired and analyzed using CouiArray software. The mobile phase was composed of 50 mM sodium acetate 5% methanol at pH 5.2. Electrochemical detector potentials for 8-OHdG and 2-deoxyguanosine (2-dG) were 120/230/280/420/600/750/840/900 mV and the flow rate was 1 ml/minutes [33].

**Statistical analysis**

Data were analyzed by one-way analysis of variance using the Statistical Package for the Social Sciences program, version 11 followed by least significant difference to compare significance between groups. In addition, co-state computer program was applied, where unshared letters are significant at p<0.05.

**RESULTS AND DISCUSSION**

**Effect of the H. pluvialis extract on oxidative stress biomarkers in different therapeutic groups**

Table 2 declared insignificant change in oxidative stress biomarkers, MDA and NO levels in normal rats treated with *H. pluvialis* ethanolic extract while, significant increase in MDA and NO levels in STZ-induced rats with percentages 373.68% and 158.48%, respectively. The treatment of hyperglycemic rats with ethanolic extract of *H. pluvialis* showed marked amelioration in MDA and NO levels with percentages of improvement 294.74% and 127.14%, respectively, compared to standard drug which recorded improvement percentages reached to 315.78% and 129.24%, respectively.

**Effect of the *H. pluvialis* extract on TAC, enzymatic, nonenzymatic antioxidant in different therapeutic groups**

Regarding to TAC, enzymatic and nonenzymatic antioxidant (Table 3), significant change was recorded in normal untreated rats while a significant increase in GPx (17.36%), was detected as compared to normal untreated rats. On the other hand, STZ-induced diabetic rats showed a significant reduction in TAC, GSH levels, GPx and GR activities with percentages 65.31, 44.52, 53.15 and 67.46%, respectively, as compared to normal control rats. Marked amelioration in all detected parameters percentages 52.19, 52.43, 37.37 and 61.23%, respectively, for TAC, GSH levels, GPx and GR activities.

**Table 1: Primer sequences used for qPCR**

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer sequence (5’−3’)</th>
<th>References</th>
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<tbody>
<tr>
<td>SOD-1</td>
<td>F: TAGCAGGACAGCAGATGAC</td>
<td>Sad1 et al. [30]</td>
</tr>
<tr>
<td></td>
<td>R: GCAGAAGGCCAGGCTGAAC</td>
<td></td>
</tr>
<tr>
<td>GST-Mu</td>
<td>F: AGAAGGAAAAGGATCTTC</td>
<td>El-Baz et al. [31]</td>
</tr>
<tr>
<td></td>
<td>R: GGGGGATGATTAGGATT</td>
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</tr>
<tr>
<td>β-actin</td>
<td>F: GTG GCC GCC TCT AGG CAC CAA</td>
<td>El-Baz et al. [31]</td>
</tr>
<tr>
<td></td>
<td>R: CTC TTT GTA CAC GAT TCT</td>
<td></td>
</tr>
</tbody>
</table>

F: Forward primer; R: Reverse primer; PISK: Phosphati-dylinositol-3 kinase, Akt: Serine/threonine protein kinase B, qPCR: Quantitative-polymerase chain reaction

<table>
<thead>
<tr>
<th>Groups</th>
<th>Biomarkers</th>
<th>Control</th>
<th>Control+<em>H. pluvialis</em></th>
<th>Hyperglycemia</th>
<th>Hyperglycemia+<em>H. pluvialis</em></th>
<th>Hyperglycemia+standard drug</th>
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<td></td>
<td>MDA (μmol/l)</td>
<td>0.19±0.01 a</td>
<td>0.17±0.03 a</td>
<td>0.9±0.01 b</td>
<td>0.34±0.05 c</td>
<td>0.30±0.04 c</td>
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<tr>
<td></td>
<td>% change</td>
<td>-</td>
<td>10.53</td>
<td>373.68</td>
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<tr>
<td></td>
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<td>-</td>
<td>-</td>
<td>294.74</td>
<td>315.78</td>
<td></td>
</tr>
<tr>
<td></td>
<td>NO (μmol/l)</td>
<td>6.19±0.60 a</td>
<td>5.22±0.95 a</td>
<td>16.00±1.22 b</td>
<td>8.13±0.52 a</td>
<td>8.00±0.92 a</td>
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<td>158.48</td>
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<td>29.24</td>
</tr>
<tr>
<td></td>
<td>% of improvement</td>
<td>-</td>
<td>-</td>
<td>127.14</td>
<td>129.24</td>
<td></td>
</tr>
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</table>

Data are expressed as mean±SD. Statistical analysis is carried out using SPSS computer program (one-way ANOVA) coupled with co-state computer program, where unshared letters are significant at p<0.05. SD: Standard Deviation, ANOVA: Analysis of variance, *H. pluvialis*: Haematococcus pluvialis, NO: Nitric oxide. Calculations:

% Change to control = \frac{\text{Mean of control} − \text{Mean of treated}}{\text{Mean of control}} \times 100

% of improvement = \frac{\text{Mean of disease} − \text{Mean of treated}}{\text{Mean of control}} \times 100
The results of this study demonstrated that *H. pluvialis* can promote the glycemic regulation, reduce oxidative stress, attenuate antioxidant biomarkers, improve inflammatory cytokines of diabetic rats and consequently improve the immune reaction against hyperglycemia induced oxidative stress. In a good agreement with the present results, several studies clearly indicated significant increase in oxidative stress biomarkers while low level of antioxidant biomarkers in type 2 DM [9,34-36]. They explained this perturbations in antioxidant system on the basis of, high blood sugar levels elicited high levels of ROS which caused depletion of antioxidant pool, GSH, GPx, and GR in liver rats indicates the damage of the second line of antioxidant defense system. This probably further exacerbates oxidative damage via adverse effect on critical GSH-related processes [37]. Reduced antioxidant status as a result of increased ROS production in experimental STZ induced rats has been reported previously by Ceriello [38] who declared that, hyperglycemia causes impairment of the antioxidant defense system as well as low antioxidant concentration that may lead to oxidative stress. In addition, the drastic depletion of liver GSH in this study may be due to high cytotoxic effect of H$_2$O$_2$ in endothelial cells as a result of inhibition of GR [37].

**Effect of the *H. pluvialis* extract on the inflammatory cytokines in different therapeutic groups**

TNF-α and IL-1β exhibited insignificant change in normal rats treated with ethanolic extract of *H. pluvialis* as compared to untreated rats (Table 4). A significant increase in TNF-α and IL-1β levels (50.62 and 42.23%) were recorded in diabetic rats as compared to normal rats. Treatment of diabetic rats with *H. pluvialis* extract showed noticeable improvement in TNF-α (47.24%) and IL-1β (44.29%) levels comparing to standard which revealed 40.43% and 49.92%, respectively.

The current results demonstrated elevated levels of TNF-α and IL-1β in diabetic rats. In accordance with the present finding El-Baz *et al.* [9] declared that, chronic hyperglycemia initiated an inflammatory state where cytokines levels are elevated which may result in the destruction of the pancreatic β-cells and malfunction of the endocrine pancreas. TNF-α is an inflammatory adipocytokine that has a major effect in the progress of insulin resistance [39]. Furthermore, IL-1β has an inverse relation with low insulin level [39]. The present results are in comconitant with the results of Al-Dahr and [Jiffri [40] who found an adverse effect as mean±standard error of mean. Followed by different superscripts are significantly different (p<0.05)

![Fig. 1: Expression levels of superoxide dismutase-1 gene in liver tissues of diabetes mellitus-induced rats treated with *Haematococcus pluvialis* extract. Data are presented as mean±standard error of mean.](image)

<table>
<thead>
<tr>
<th>Groups</th>
<th>Biomarkers</th>
<th>Control</th>
<th>Control+<em>H. pluvialis</em></th>
<th>Hyperglycemia</th>
<th>Hyperglycemia+<em>H. pluvialis</em></th>
<th>Hyperglycemia+Standard drug</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>TAC (mmol/mg protein)</td>
<td>15.54±1.21$^a$</td>
<td>16.90±0.91$^a$</td>
<td>5.39±0.25$^b$</td>
<td>13.50±1.51$^b$</td>
<td>10.50±1.21$^b$</td>
</tr>
<tr>
<td></td>
<td>% Change</td>
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<td>8.04</td>
<td>65.31</td>
<td>13.13</td>
<td>32.43</td>
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<td>-</td>
<td>-</td>
<td>-</td>
<td>52.19</td>
<td>32.88</td>
</tr>
<tr>
<td></td>
<td>GSH (μmol/mg protein)</td>
<td>36.05±2.13$^a$</td>
<td>37.00±3.45$^a$</td>
<td>20.00±1.00$^b$</td>
<td>38.90±2.23$^b$</td>
<td>28.99±2.29$^b$</td>
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<td>52.43</td>
<td>24.93</td>
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<td>GPX (μmol/mg protein)</td>
<td>1.90±0.01$^a$</td>
<td>2.23±0.01$^b$</td>
<td>0.89±0.05$^b$</td>
<td>1.60±0.1$^b$</td>
<td>1.20±0.09$^b$</td>
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<td>53.16</td>
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<td>37.37</td>
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<tr>
<td></td>
<td>GR (μmol/mg protein)</td>
<td>13.00±2.20$^b$</td>
<td>13.65±1.00$^b$</td>
<td>4.23±0.55$^b$</td>
<td>12.19±0.56$^b$</td>
<td>9.10±0.20$^b$</td>
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<tr>
<td></td>
<td>% change</td>
<td>-</td>
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<td>67.46</td>
<td>6.23</td>
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<tr>
<td></td>
<td>% of improvement</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>61.23</td>
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Data are expressed as mean±SD. Statistical analysis is carried out using SPSS computer program (one-way ANOVA) coupled with co-state computer program, where unshared letters are significant at p≤0.05. SD: Standard deviation, ANOVA: Analysis of variance, *H. pluvialis*: *Haematococcus pluvialis*, TAC: Total antioxidant capacity, GSH: Reduced glutathione, GPx: Glutathione peroxidase, GR: Glutathione reductase

<table>
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<tr>
<th>Groups</th>
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<th>Hyperglycemia+standard drug</th>
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<tr>
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<td>TNF-α (ng/l)</td>
<td>26.12±3.10$^a$</td>
<td>27.00±3.52$^b$</td>
<td>28.78±1.20$^b$</td>
<td>3.37</td>
<td>10.18</td>
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<tr>
<td></td>
<td>25.00±1.92$^b$</td>
<td>39.34±3.49$^a$</td>
<td>50.62</td>
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<td>-</td>
<td>4.29</td>
<td>50.62</td>
<td>3.37</td>
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<td></td>
<td>% of improvement</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>47.24</td>
<td>40.43</td>
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<tr>
<td></td>
<td>IL-1β (Pg/l)</td>
<td>3370.70±78.10$^a$</td>
<td>3389.00±98.45$^a$</td>
<td>4794.00±560.60$^a$</td>
<td>331.00±57.89$^a$</td>
<td>311.13±100.70$^a$</td>
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<tr>
<td></td>
<td>% change</td>
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<td>0.5</td>
<td>42.23</td>
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<td>% of improvement</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>44.29</td>
<td>49.92</td>
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Data are expressed as mean±SD. Statistical analysis is carried out using SPSS computer program (one-way ANOVA) coupled with co-state computer program, where unshared letters are significant at p≤0.05. SD: Standard deviation, ANOVA: Analysis of variance, *H. pluvialis*: *Haematococcus pluvialis*, TNF-α: Tumor necrosis factors-α, IL-1β: Interleukin-1 beta

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**Table 3: Effect of *H. pluvialis* extract on TAC, enzymatic and nonenzymatic antioxidant in different therapeutic groups**

**Table 4: Effect of *H. pluvialis* extract on proinflammatory TNF-α and IL-1β in different therapeutic groups**
increment in TNF-α level in all diabetic patients. In addition, Mojtaba et al. [39] found elevation in IL-1β level in type II diabetic patients which was higher than the nondiabetic counterparts. The authors backed this elevation in IL-1β to β-cell dysfunction.

**Effect of the *Haematococcus pluvialis* extract on the expression of genes encoding antioxidant enzymes**

The results of the gene expression analysis using quantitative real-time RT-PCR are summarized in Figs. 1 and 2. The genes encoding superoxide dismutase (SOD-1) and glutathione S-transferase (GST)-Mu were determined in liver tissues of DM rats before or after treatment with *Haematococcus pluvialis* extract. The results found that, DM-rats revealed very lower expression levels of SOD-1 and GST-Mu genes than those in control rats, where the differences were highly significant (p≤0.05). In contrary, treatment of male DM-rats with *Haematococcus pluvialis* extract increased the expression levels of SOD-1 and GST-Mu genes compared with those in DM-rats without treatment, where the differences were also highly significant (p≤0.05). In addition, treatment of male DM-rats with the reference drug for diabetes, namely glibenclamide, increased significantly the expression levels of SOD-1 and GST-Mu genes compared with those in DM-rats without treatment.

**Effect of *Haematococcus pluvialis* extract on the ROS generation as oxidative stress indicator**

The impact of *Haematococcus pluvialis* extract as preventive effect on the intracellular ROS generation is summarized in Fig. 3. The results indicated that the generation of ROS in DM-rats was significantly higher than those in control rats. However, the generation of ROS in DM-rats treated with *Haematococcus pluvialis* extract decreased significantly the ROS generation compared with those in DM-rats without treatment. Moreover, treatment of male DM-rats with glibenclamide decreased significantly the ROS generation compared with those in DM-rats without treatment.

**Effect of *Haematococcus pluvialis* extract on the apoptosis rate**

The determination of the 8-OHdG generation in pancreatic tissues of male DM-rats' genome before or after *Haematococcus pluvialis* extract treatment as an indicator for oxidative stress induced DNA damage is summarized in Fig. 4.

The present results found that the ratio of 8-OHdG/2-dG generation increased significantly in DM-rats in comparison to that of the control group. In contrast, the ratio of 8-OHdG/2-dG generation decreased significantly following *Haematococcus pluvialis* extract treatment compared with those in DM-rats. Furthermore, treatment of male DM-rats with glibenclamide decreased significantly the ratio of 8-OHdG/2-dG generation compared with those in DM-rats without treatment.

A numerous studies have been suggested that hyperglycemia increased the formation of ROS and reduced the antioxidant enzyme activity through glycation of these proteins (binding the protein molecules with the sugar molecules) [38]. In agreement with these findings, the present results found that DM-rats exhibited lower expression levels of the antioxidant enzyme genes, SOD-1 and GST-Mu, than those in control rats. In an explanation for these results, the low expression of the genes encoding antioxidant enzymes in the diabetic animal tissues is attributed to the attack by the ROS molecules to the diabetic tissues [41,42]. On the other hand, the current results revealed that, diabetic rats exhibited significantly higher levels of ROS generation and 8-OHdG formation. In the same line, several studies reported that increase the levels of glucose in diabetic rats was coincided with increase the levels of 8-OHdG [5,43].

It has been found that the increase of the antioxidants levels is acting to delay or prevent the disorders of the pancreatic β-cells in diabetic rats which caused by increase the glucose concentrations [43,44]. Therefore, one of the potentially therapeutic tools in diabetes treatment is using antioxidant supplementation which could improve the control of this disease. This study showed that treatment of diabetic rats with *Haematococcus pluvialis* extract increased the levels of the TAC, antioxidant biomarkers, antioxidant enzyme genes and decreased the levels of oxidative stress biomarkers, inflammatory cytokines, ROS generation.
and 8-OHdG formation compared with those in diabetic rats without *H. pluvialis* extract treatment.

The results of the current work were in the same line with Ojo et al. [45] and El-Baz et al. [9] who found the amelioration in TNF-α IL-1β levels post-treatment of STZ-induced rats with *H. pluvialis* may be attributed to the fact that, antioxidants have the ability to suppress inflammatory markers like TNF-α and IL-1β. Furthermore, Uchiyama et al. [46] reported that treatment of diabetic mice with the active ingredient of *H. pluvialis* extracts namely ASTA improved the tolerance against diabetes which coincided with increase the levels of the antioxidants and decrease the ROS formation. They also found that the improving action of ASTA in diabetic mice was associated with decrease the levels of glucose concentration as the main reason in diabetes toxicity. Moreover, they found at 120 minutes, ASTA treatment decreased the levels of glucose (p<0.001) and increased the levels of insulin levels (p<0.0001) in diabetic mice compared with diabetic mice without ASTA treatment. Moreover, treatment of diabetic mice with ASTA improved the lesions in diabetic tissues.

**CONCLUSION**

The findings of this study suggested that *H. pluvialis* extract supplementation was able to decline the oxidative stress, enhanced antioxidant status and improved inflammatory cytokines induced by hyperglycemia in diabetic rats. The therapeutic action of *H. pluvialis* also involved in the increase of expression levels of antioxidant enzyme genes, decrease the levels of ROS generation and 8-OHdG formation which may be attributed to the presence of ASTA in *H. pluvialis* extract.

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