

L-ergothioneine and Metformin Alleviates Liver Injury in Experimental Type-2 Diabetic Rats via Reduction of Oxidative Stress, Inflammation, and Hypertriglyceridemia

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Abstract

Diabetic-induced liver toxicity is a serious complication that cause significant metabolic dysfunction. L-ergothioneine (L-egt) is a bioactive nutraceutical obtained from mushrooms and certain food products, with reported cytoprotective, antioxidant and anti-inflammatory properties and potential to improve efficacy of existing therapy. Thus, this study evaluates the effects of L-egt, and/or metformin, on diabetes-induced liver injury. Diabetes was induced in male Sprague-Dawley rats using 10% fructose for two weeks, followed by a single low dose streptozotocin (STZ, 40 mg/kg *i.p*) injection. After induction of diabetes, animals were treated either with de-ionized water (DW), L-egt (35 mg/kg bwt), metformin (500 mg/kg bwt), or a combination of L-egt and metformin orally for seven weeks. Body weight and glucose were monitored during the experiment. At the completion of experiment, blood samples were collected, and liver tissue was excised for biochemical analysis, enzyme-link immunosorbent assay (ELISA) of various liver function biomarkers, reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analysis of genes associated with inflammation, oxidative stress, and lipid metabolism, as well as histopathological evaluation. Diabetic animals showed liver hypertrophy, increased liver injury, hepatic triglycerides, oxidative stress, and hepatic inflammation. However, L-egt, and/or metformin, improved glycemic control, reduced liver injury, triglycerides, oxidative damage, inflammatory injury, and normalize liver histology by upregulating Nrf2/Sirt1, downregulate NF-kB/TGF-B1, and reduce SREBP1c/FAS expression. In conclusion, these results showed that combination of L-egt and metformin improve therapeutic efficacy than either treatment alone. Thus, L-egt can be used as an adjuvant to mitigate diabetes-induced liver complication.

Introduction

Type-2 diabetes (T2D), a chronic metabolic disorder of global prevalence, is drastically increasing in developing and industrialized countries due to excessive caloric intake, sedentary lifestyle, poor diagnosis, and disease management. The etiology and progression of T2D are multifaceted and frequently associated with obesity, hypertriglyceridemia, insulin resistance (IR), and compromised insulin secretion by the pancreatic islet with subsequent hyperglycemia (Lozano et al., 2016, Pandey et al., 2015). Liver complications are often seen in approximately 70% of people with diabetes and account for about 2–4% mortality in T2D patients (Hazlehurst et al., 2016, Zoppini et al., 2014). These complications (including non-alcoholic fatty liver disease, steatosis) results from hyperglycemia-induced oxidative injury and low-grade inflammation that mediate liver damage, including fibrosis and apoptosis. Also, IR increases *de novo* lipogenesis under the influence of SREBP-1c and alters lipid metabolism by upregulating FAS, thereby promoting the influx of fatty acid into the liver cells (Perry et al., 2015, Gehrke and Schattenberg, 2020, Arrese et al., 2019). Excessive accumulation and infiltration of fats into the hepatocytes contribute to liver toxicity by increasing production of reactive oxygen species (ROS), stimulate inflammatory cytokines, mitochondrial dysfunction, and endoplasmic reticular stress that mediate liver damage. Thus, therapeutic measures that can effectively inhibit oxidative injury, prevent inflammation, and reduce fatty infiltration into the hepatocytes may provide adequate cytoprotection

against liver damage. However, recent attention has been geared towards natural compounds with multiple bioactivities to provide adjuvant therapy to manage diabetic liver complications.

The beneficial role of bioactive compounds with antioxidants properties in the management of diabetic complications has been associated with their ability to increase antioxidant capacity and stimulate endogenous antioxidant enzymes and cytoprotective genes, activating various transcription factors such as Nrf2 and Sirt1 (Jadeja et al., 2016, Farghali et al., 2019). These compounds also exert anti-inflammatory activities by downregulating the Nuclear factor-kappa B signaling pathway to inhibit the production of cytokines, chemokines, and fibrosis in the liver (de Gregorio et al., 2020, McKinley and Willoughby, 2014). Furthermore, natural compounds also help reduce serum cholesterol and triglycerides by downregulating the expression of lipogenic proteins, thereby reducing the influx of triglyceride into hepatocytes (Xu et al., 2020, Romano et al., 2021). Thus, attention has been geared towards nutraceuticals that regulate glycemic control, upregulate the antioxidant pathway, inhibit cellular inflammation and improve lipid metabolism in the management of diabetic complications (Wang et al., 2018, Ma et al., 2019). Recently, the therapeutic benefits of mushrooms in the management of diabetes and associated complications have received significant attention owing to the presence of various bioactive compounds, including flavonoids and L-ergothioneine (Lo et al., 2020, Lindequist and Haertel, 2020, Azeem et al., 2021).

L-ergothioneine (L-egt), an adaptive antioxidant obtained from mushroom and some meat products (e.g., kidney and liver), has been shown to exert antioxidant and anti-inflammatory activities as well as exhibit adaptive cytoprotective function by accumulating at the site of injury to protect against tissue damage (Halliwell et al., 2018, Ey et al., 2007, Salama and Omar, 2021, Ko et al., 2021). L-egt protects against hyperglycemia-induced cell senescence, prevents embryo malformations in diabetic pregnant rats, and enhances the therapeutic efficacies in-vitro and in-vivo (Song et al., 2017, D'Onofrio et al., 2016, Guijarro et al., 2002). Furthermore, administration of L-egt rich foods ameliorated dimethylnitrosamine-induced liver fibrosis and oxidative stress in mice (Tang et al., 2016) and has been shown in the treatment of non-alcoholic fatty liver disease (NAFLD) to downregulate SREBP1c and FAS expression, thereby inhibiting hepatic lipogenesis and lipid accumulation (Carbonero et al., 2019, Jeong and Park, 2020). Also, the hepatoprotective effect of biguanides, e.g., Metformin, may result from its potency to regulate glycemic index by reducing hepatic gluconeogenesis and increased glucose uptake (Rena et al., 2017). However, metformin is associated with side effects, and patients still present with liver complications despite glycemic control, suggesting that metformin alone does not confer overall effective treatment. Therefore, this study evaluated the role of L-egt, with or without metformin, on liver injury in a rat model of type-2 diabetes.

Material And Methods

Drugs and chemicals

Pure L-egt was obtained from Tetrahedron limited, Paris, France. QPCR iTAQ SYBR Green and cDNA synthesis kits were purchased from Lasec (Cape Town, South Africa). Primers were synthesized by Inqaba Biotec (Pretoria, South Africa). Metformin was obtained from a local pharmacy (Pharmed, South Africa). All other chemicals, reagents, and equipment were procured from standard commercial suppliers and high analytical grades.

Experimental animals and ethical approval

Thirty-six (36) male Sprague-Dawley rats (175 ± 20 g) were obtained from the Biomedical Research Unit, Westville Campus, University of KwaZulu-Natal (UKZN), South-Africa and were housed in a room with standard laboratory conditions (12 hours light-dark cycles; temperature $23 \pm 1^\circ\text{C}$, 40–60% humidity). The animals were allowed access to rat feed and water *ad libitum* for an acclimatization period of one week before the experiment. All animal and experimental procedures were approved by the Animal Research Ethics Committee (AREC) of the University of KwaZulu-Natal, Durban, South Africa (Ethic number: AREC/006/019D).

Experimental design

After acclimatization, the animals were randomly divided into two major groups: the non-diabetic ($n = 10$) and the diabetic ($n = 20$) groups. All animals in the diabetic group were treated with fructose and streptozotocin (STZ) to induce type-2 diabetes using the established model described by (Wilson and Islam, 2012). Briefly, the animals were supplied 10% fructose in drinking water *ad-libitum* for two weeks to induce IR and later injected (*i.p.*) 40mg/kg bwt STZ freshly prepared in 0.1M citrate buffer. The animals in the non-diabetic group were injected with the same volume of 0.1M citrate buffer. Animals with non-fasting blood glucose levels of $> 16.7\text{mmol/L}$ after one-week post-STZ injection were confirmed diabetic (Srinivasan et al., 2005) and included in the study. After successful diabetes induction, the non-diabetic animals were subdivided into two groups, while the diabetic animals were subdivided into four groups as follows.

Group 1 (NC): non-diabetic plus de-ionized water (negative control).

Group-2 (NE): non-diabetic plus L-egt

Group-3 (DC): diabetic plus de-ionized water (positive control)

Group-4 (DE): diabetic plus L-egt

Goup-5 (DM): diabetic plus metformin

Group-6 (DEM): diabetic plus l-egt plus Metformin.

Groups 1 and 3 were administered de-ionized water (1ml/100g), groups 2 and 4 were administered L-egt (35mg/kg bwt), group 5 was administered metformin (500mg/kg bwt), while group 6 was administered a combination of L-egt and metformin. The dosage of L-egt used in this study was based on previous in-

vivo studies using this nutraceutical (Tang et al., 2018, Williamson et al., 2020). All treatments were done daily by oral gavage and lasted for seven weeks.

Blood and tissue collection

After the seven-week treatment period, all animals were sacrificed by decapitation and blood was immediately collected into a serum vacutainer EDTA bottle and allowed to stand for 30 mins. The blood was then centrifuged at 3000rpm for 10mins at 4⁰C to obtain serum. The serum samples obtained were stored in the bio-freezer (Snijers Scientific, Holland) at -80⁰C until used for biochemical analysis.

Afterward, incisions were made along the linea alba of the anterior abdominal wall to excise the liver. This organ weighed, rinsed with cold normal saline, and snap-frozen in liquid nitrogen before been stored in the bio-freezer at -80⁰C until used for analysis. Liver tissue was fixed in 10% neutral-buffered formalin for histological assessment.

Analysis of Bodyweight and liver index

Body weight was monitored using a sensitive electronic weighing scale (Metler, Greifensee, Switzerland). The liver index (use to assess liver hypertrophy) was calculated as the ratio of harvested liver weight to the body weight and expressed in percentage i.e.

Liver index = Liver weight × 100

Body weight

Preparation of liver homogenates

The liver was thawed and homogenized in 10% phosphate buffer (0.1M, pH7.4, 1:9 w/v). The homogenates were centrifuged a 600g for 10min to remove cell debris. The supernatant was subsequently centrifuged at 10,000g for 20mins to obtain the cytosolic fraction, which was used immediately for biochemical analyses.

Biochemical analysis

The serum concentration of liver enzymes (AST, ALT, and ALP) and triglyceride were analyzed at an accredited pathology laboratory (Global Clinic and Viral laboratories, Amazimtoti, South Africa). The concentration of triglyceride was also measured in the liver homogenate using an automatic biochemical analyzer. Blood glucose was measured at the end of the experiment using a glucometer (Accu-Chek Performa, USA). Serum insulin levels were measured by ELISA kits (Mercodia kit), and used to evaluate homeostasis model assessment of insulin resistance (HOMA-IR), which was calculated as previously described by (Matthews et al., 1985) using the following formula:

HOMA-IR = fasting serum insulin (mU/L) X fasting blood glucose (mg/dl).

Analysis of lipid peroxidation (MDA), antioxidant enzymes (SOD and CAT), and reduced glutathione level (GSH)

The liver homogenates were used to measure the concentration of malondialdehyde (MDA), superoxide dismutase (SOD), catalase (CAT), and reduced glutathione (GSH) by spectrophotometric assay. MDA, a marker of lipid peroxidation, was evaluated by measuring the content of thiobarbituric acid (TBA) reactive product in the liver homogenates using the method of Mkhwanazi et al. (Mkhwanazi et al., 2014) and expressed as nmol MDA per milligram protein. GSH level and SOD activity were assessed using the method of Ellman and Marklund, respectively (Ellman, 1959, Marklund, 1985), while CAT activity was assessed using the protocol of Aebi (Aebi, 1984).

Analysis of inflammatory biomarkers

The concentration of cytokine- tumor necrosis factor- α (TNF- α); chemokine, monocyte chemotactic protein-1 (MCP-1) and fibrotic cytokine, tumor growth factor- β (TGF- β) were quantified in the liver homogenates using Elabscience commercial ELISA kits according to the manufacturer's protocol.

Rt-PCR Analysis of SREBP1c, FAS, Nrf2, Sirt1, NF-kB and TGF- β 1 mRNA expression

The relative mRNA expression of SREBP1c, FAS, Nrf2, Sirt1, NF-kB, and TGF- β 1 was quantified in the liver homogenates and analyzed using a light cycler. Total RNA was isolated in the liver using TRIzol reagent (40mg of tissue/mL, Trizol reagent). The isolated RNA quantity was determined by measuring absorbance at 260/280nm using nanodrop ND-1000 spectrophotometer (Thermo scientific, Johannesburg, South Africa). The total RNA was converted into cDNA using iScript cDNA synthesis kit, Life Science research (Biorad, South Africa) following manufacturers' instruction. The complete reaction mixture was incubated on SimpliAmp™ thermal cycler, Applied Biosystems (Thermo Fischer Scientific), using the following reaction condition; priming 5mins at 25⁰C, reverse transcription 20mins at 46⁰C, and RT inactivation 1min at 95⁰C. Real-time polymerase chain reaction (RT-PCR) was done using iTaq Universal SYBR Green supermix (Biorad, CA, USA) as fluorescent dye on a light cycler 96 RT-PCR system (Roche, Mannheim, Germany). RT-qPCR was performed in a 10 μ L reaction volume containing 5 μ L SYBR Green Master Mix, 1 μ L of each primer, 1 μ L of nuclease-free water, and 2 μ L of cDNA template. The primer sequences used are provided in Table-1. The purity and specificity of amplified PCR products were verified by melting curves generated at the end of each PCR. Relative mRNA expression was calculated using the $2^{-\Delta\Delta ct}$ method (Livak and Schmittgen, 2001) and normalized in relation to the endogenous control expression, GAPDH. The primers sets were homology searched using an NCBI BLAST search to ensure that they were specific.

Table-1: Oligonucleotide sequence and accession number

Genes	Primer sequence	Genbank Accession number
GAPDH	F: 5'-TGATGACATCAAGAAGGTGGTGGAG-3' R: 5'-TCCTTGGAGGCCATGTAGGCCAT-3'	XM_017593963.1
Nrf ₂	F: 5'-CAGCATGATGGACTTGAATTG - 3' R: 5'-GCAAGCGACTCATGGTCATC - 3'	NM_031789.2
Srt1	F: 5'-CCCAGATCCTCAAGCCATGTTC - 3' R: 5'-TGTGTGTGTGTTTTTCCCCC - 3'	NM_001372090.1
TGF-β1	F: 5'-GGGCTACCATGCCAACTTCTG - 3' R: 5'-GAGGGCAAGGACCTTGCTGTA-3'	NM_021578.2
NF-kB	F: 5'-ACGATCTGTTTCCCCTCATCT-3' R: 5'-TGCTTCTCTCCCCAGGAATA-3'	NM_199267.2
FAS	F: 5'-TGTGGGGTGGAAATCATCGG-3' R: 5'-CATTGCTCCTTTGGGGTTGC-3'	NM_012820.1
SREBP-1c	F: 5'- GGAGCCATGGATTGCACATT-3' R: 5'- AGGAAGGCTTCCAGAGAGGA-3'	NM_001276708.1

Histopathological analysis of Liver

Liver specimens were embedded in paraffin wax after dehydration in a graded series of ethanol and cleared in xylene. Serial sections were done using a rotary microtome; liver slices of 5-µm thick were fixed on a slide and stained with hematoxylin & eosin (H&E). The stained sections were visualized and captured using a nanozoomer S360 digital slide scanner (Hamamatsu Photonics, Japan) and nanozoomer digital pathology version 2.8 software for analysis by a pathologist.

Statistical analysis

Data were reported as mean ± SEM. GraphPad Prism Software version 7 (San Diego, CA) was used for statistical analysis. The differences between means were analyzed using a one-way analysis of variance (ANOVA) followed by Tukey's multiple comparison test to determine the difference between groups. Statistical significance between groups was considered at $P < 0.05$.

Results

Effect body weight, liver hypertrophy, blood glucose, HOMA-IR, and serum TG in T2DM-rats

As presented in table 2, the DC group had a significant reduction ($p < 0.001$) in body weight vs. NC. Co-administration of L-egt with metformin in diabetic rats caused a significant increase (DE: $p < 0.05$) in body weight vs. DC. There was a significant increase ($p < 0.001$) in the liver hypertrophy of DC rats vs. NC rats. However, L-egt, with or without metformin to diabetic rats significantly reduced (DE: $p < 0.05$; DEM: $p < 0.01$) liver hypertrophy vs. DC. There was a significant increase ($p < 0.001$) in blood glucose level in the DC group vs NC group. Co-administration of L-egt with metformin significantly reduced ($p < 0.01$) blood glucose level vs. DC. There was a significant increase in HOMA-IR ($p < 0.01$) in the DC group vs NC group. However, combination of L-egt with metformin significantly reduced HOMA-IR ($p < 0.05$) vs. DC. Serum TG increased significantly ($p < 0.01$) in the DC rats vs. NC rats, while L-egt alone, and its combination with metformin, significantly reduced (DE: $p < 0.05$; DEM: $p < 0.01$) serum TG vs. DC rats.

Table-2: Effect of L-egt with or without metformin for seven weeks on body weight, liver index, blood glucose and HOMA-IR in T2DM-rats. ** $p < 0.01$, *** $p < 0.001$ vs NC; # $p < 0.05$, ## $p < 0.01$, vs DC. (n = 5). NC = Nondiabetic control; NE = nondiabetic plus L-egt; DC = diabetic control; DE = diabetic plus L-egt; DM = diabetic plus metformin and DEM = diabetic plus L-egt plus metformin. (supplemental data)

Indices	NC	NE	DC	DE	DM	DEM
Body weight (g)	431.20 ± 16.68	424.20 ± 15.85	277.80 ± 20.68***	321.40 ± 8.44	338.60 ± 15.79	358.60 ± 12.97#
Liver index (%)	3.11 ± 0.08	3.17 ± 0.08	4.28 ± 0.24***	3.49 ± 0.23#	3.35 ± 0.08##	3.28 ± 0.17##
Blood glucose (mg/dl)	80.64 ± 7.64	73.08 ± 8.57	257.34 ± 28.15***	214.44 ± 23.02	175.90 ± 20.87#	143.54 ± 12.96##
HOMA-IR	2.51 ± 0.42	2.39 ± 0.47	5.04 ± 0.46**	3.94 ± 0.45	3.77 ± 0.45	3.07 ± 0.37#
TG (mmol/L)	1.47 ± 0.17	0.97 ± 0.19	2.88 ± 0.30**	1.78 ± 0.13#	1.84 ± 0.17#	1.65 ± 0.16##

Effect on liver injury in T2DM rats.

The effects of L-egt treatment with or without metformin for seven weeks on biomarkers of liver injury (ALP, AST, and ALT) and liver triglyceride in type-2 diabetic rats were presented in table-3. In the DC group, liver injury biomarkers increased significantly (ALP: $p < 0.05$; AST: $p < 0.001$ and ALT: $p < 0.001$) vs. the NC group. However, L-egt, with or without metformin, significantly reduced (DE: $p < 0.05$; DEM: $p < 0.01$) the concentration of liver biomarkers vs. DC. Liver triglyceride significantly ($p < 0.01$) increase in the DC group vs. NC group while L-egt, with or without metformin cause a significant decrease ($p < 0.05$) in hepatic triglyceride level vs. DC.

Table 3

Effect of L-egt with or without metformin on liver enzymes, and triglycerides in T2DM-rats. ***p < 0.001, **p < 0.01 vs NC and ###p < 0.001, ##p < 0.01, #p < 0.05 vs DC (n = 5). NC = Nondiabetic control; NE = nondiabetic plus L-egt; DC = diabetic control; DE = diabetic plus L-egt; DM = diabetic plus metformin and DEM = diabetic plus L-egt plus metformin. (supplemental data).

Indices	NC	NE	DC	DE	DM	DEM
ALP (U/L)	129.60 ± 15.58	124.20 ± 24.00	239.60 ± 29.04**	147.40 ± 16.33#	141.60 ± 10.26#	132.60 ± 11.61##
AST (U/L)	87.40 ± 11.21	89.20 ± 14.18	184.20 ± 14.51***	127.20 ± 9.88#	126.00 ± 9.82#	97.40 ± 15.22###
ALT (U/L)	87.20 ± 9.28	86.80 ± 14.97	172.04 ± 10.35***	122.40 ± 11.41#	122.20 ± 7.31#	105.80 ± 11.82##
TG (mmol//g liver)	1.17 ± 0.23	0.89 ± 0.19	3.29 ± 0.35**	1.84 ± 0.36#	1.85 ± 0.33#	1.80 ± 0.28#

Effect on lipid peroxidation and antioxidant enzymes

As presented in Fig. 1a-d, liver MDA concentration in the DC group was significantly increased (p < 0.001) vs. the NC group. Administration of L-egt alone and its co-treatment with metformin to diabetic rats caused a significant decrease (DC: p < 0.05; DEM: p < 0.01) in liver MDA concentration vs. the DC group. There was a significant decrease (P < 0.01) in SOD, GSH, and CAT in the DC rats vs. NC. However, L-egt, with or without metformin, significantly increase SOD, CAT (DE: p < 0.05; DEM: p < 0.01) and GSH (p < 0.05) in the diabetic rat vs. DC.

Fig-1c: GSH levels in L-egt, with or without metformin, treated T2DM rats

Figure 1a-d

Effect of L-egt, with or without metformin on Lipid peroxidation, antioxidant enzymes and glutathione in T2DM-rats. ***p < 0.001, **p < 0.01 vs NC; ##p < 0.01, #p < 0.05 vs DC. (n = 5). NC = Nondiabetic control; NE = nondiabetic plus L-egt; DC = diabetic control; DE = diabetic plus L-egt; DM = diabetic plus metformin and DEM = diabetic plus L-egt plus metformin (supplemental data).

Effects on liver inflammatory biomarkers: TNF- α , MCP-1, and TGF- β 1.

The concentration of TNF- α , MCP-1, and TGF- β 1 in the liver homogenates after treatment with L-egt, with or without metformin, for seven weeks, was presented in Fig. 2a-c. DC group had a significant increase (MCP1: p < 0.01; TNF- α : p < 0.001; TGF- β 1: p < 0.01) in the concentration of inflammatory biomarkers vs. NC group. Diabetic groups treated with L-egt alone, or in combination with metformin caused a significant reduction in MCP1, TGF- β 1 (p < 0.05), and TNF- α (DE: p < 0.05; DEM: p < 0.01) vs. DC group.

Figure 2a-b: Effect of L-egt, with or without metformin, on liver inflammatory biomarkers (A) MCP1, (B) TNF- α and TGF- β 1 in T2DM-rats. *** $p < 0.001$, ** $p < 0.01$ vs NC; ## $p < 0.01$, # $p < 0.05$ vs DC. (n = 5). NC = Nondiabetic control; NE = nondiabetic plus L-egt; DC = diabetic control; DE = diabetic plus L-egt; DM = diabetic plus metformin and DEM = diabetic plus L-egt plus metformin. (supplemental data).

Effect on liver mRNA relative expression of SREBP1c, FAS, NF-kB, TGF β 1, Nrf2, and Sirt1

The transcriptional levels of liver SREBP1c, FAS, NF-kB, ($p < 0.001$) and TGF- β 1 ($p < 0.01$) significantly increase in the DC group vs. NC group. Administration of L-egt, with or without metformin, to diabetic rats significantly reduced the expression of SREBP1c (DE: $p < 0.05$; DEM: $p < 0.01$), FAS (DEM: $p < 0.01$), NF-kB (DE: $p < 0.05$; DEM: $p < 0.001$) and TGF- β 1 ($p < 0.05$) vs. DC rats, (Fig. 3A and B). There was a significant decrease in mRNA expression of liver Nrf2 and Sirt1 ($p < 0.01$) in the DC rats vs. NC rats (Fig. 3C). However, administration of L-egt, with or without metformin to diabetic rats significantly increased (DE: $p < 0.05$; DEM: $p < 0.01$) Nrf2 and Sirt1 mRNA expression vs. DC rats. Interestingly, co-administration of L-egt with metformin to diabetic rats (DEM) significantly ($p < 0.05$) increased Nrf2 mRNA expression vs. diabetic rats treated with metformin alone (DM).

Figure 3a-c: Relative mRNA expression level of (A) SREBP1c and FAS (B), NF-kB and TGF- β 1 (C), Nrf2 and Sirt1 relative gene expression in liver tissues of diabetic animals treated with L-egt with or without metformin for 7 weeks. *** $p < 0.001$, ** $p < 0.01$ vs NC groups; # $p < 0.05$, ## $p < 0.01$ vs DC group and § $p < 0.05$ vs DM (n = 5). NC = Nondiabetic control; NE = nondiabetic plus L-egt; DC = diabetic control; DE = diabetic plus L-egt; DM = diabetic plus metformin and DEM = diabetic plus L-egt plus metformin (supplemental data).

Effect on liver histopathological changes.

Morphological evaluation of the liver sections after seven weeks of treatment showed that L-egt, with or without metformin, alleviates hepatic injury in T2DM animals, as shown in Fig. 4a-f. The photomicrograph of liver sections in NC and NE groups showed normal liver histoarchitecture with normal morphology of the central vein and hepatic sinusoids (Fig. 4a and b). The liver section in the DC animal (Fig. 4c) showed liver injury characterized by disrupted hepatic sinusoids, congested central vein with mild hepatocyte degeneration compared to the NC rats. Administration of L-egt (DE) or metformin (DM), reduced sinusoid disruption and congestion of the central vein (Fig. 4d and 4e resp). In contrast, the liver section in DEM animals shows similar histoarchitecture with NC animals (Fig. 5f).

Discussion

This study aimed to examine the benefits of L-ergothioneine, with or without metformin, on liver injury in a type-2 diabetic rat model. Both hyperglycemia and IR increase the production of reactive oxygen species (ROS) that alter the structure and functions of vital organs (including the liver) with the resultant pathogenesis of diabetic complications. In this regard, the use of natural compounds with significant

bioactive potential have attracted greater interest due to their reduced side effect, increased accessibility and efficacy against the molecular and cellular triggers involved in diabetic complications (Choudhury et al., 2017, Gothai et al., 2016).

A significant reduction in body weight has been reported in poorly managed diabetes (Magalhães et al., 2019). A similar observation was reported in this study where the DC rats show substantial weight reduction compared to NC. This reduction could be correlated with metabolic derangements associated with poor glucose utilization, thereby resulting in excess catabolism of adipose tissue, breakdown of structural proteins and reduced protein synthesis in all tissues, thereby causing muscle wasting. In this study, synergistic administration of L-egt and metformin to diabetic rats improves body weight. This suggests that this treatment regimen could halt some metabolic derangements associated with muscle wasting and loss of adipose tissues, to reduce weight loss during diabetes.

In this study, liver hypertrophy was seen in the DC rats compared to the NC rats, and this was in accordance with other studies where increased liver weight was reported in diabetic rats (Ayepola et al., 2013, Zhang et al., 2019). The increased liver weight may result from IR and hypertriglyceridemia that promote influx and accumulation of fatty acids into the liver cells (Mohamed et al., 2016, Zhang and Lu, 2015). Administration of L-egt, with or without metformin, to diabetic rats significantly reduced liver hypertrophy, suggesting that L-egt may reduce the influx of fatty acids into the liver or promote β -oxidation of fatty acids in the hepatic mitochondrial. Thus, preventing excessive accumulation of fatty acids and triacylglycerol in the liver. Effective regulation of fasting and postprandial blood glucose level plays a significant role in reducing diabetes-induced organ damage. In this study, co-administration of L-egt with metformin to diabetic rats enhanced the antihyperglycemic efficacy of metformin (a conventional drug used in the management of type-2 diabetes). This could be responsible for the reduced blood glucose observed in the diabetic rat. Also, the reduced IR assessed by HOMA-IR in the DEM group indicates improved insulin sensitivity and glycemic control. Metformin reduces blood glucose by increasing insulin sensitivity and reducing hepatic gluconeogenesis (Foretz et al., 2019). The hepatoprotective effect of L-egt may improve the antihyperglycemic efficacy of metformin because the liver is a major site of metformin activity where it reduces hepatic lipogenesis and improves fatty acid oxidation (Zheng et al., 2015). Thus, L-egt supplementation during metformin therapy may enhance glycemic control and prevent hyperglycemia-induced liver damage.

Liver enzymes, including Alanine aminotransferase (ALT), Aspartate aminotransferase (AST), and alkaline phosphatase (ALP), are common biomarkers used to evaluate liver injury. These enzymes help catalyze crucial chemical reactions (e.g., amino transfer) and their serum concentration increases during liver damage (Kwo et al., 2017, Fu et al., 2020). An increase in serum aminotransferase is mostly reported in type-2 diabetes due to membrane damage in the hepatocytes, while increased ALT level may also indicate fatty liver resulting from reduced insulin sensitivity by the hepatocytes (Mandal et al., 2018, Islam et al., 2020). The increased serum level of ALP, AST, and ALT recorded in this study indicate significant liver injury in the DC rats. Interestingly, L-egt with or without metformin attenuated liver damage, as evidenced by reduced serum level of liver enzymes. This result suggests hepatoprotective function, which

may result from the ability of L-egt to enhance membrane integrity in the hepatocytes to reduce leakage of liver enzymes into circulation. Notably, there was no significant change in the serum level of liver enzymes in the NE rats when compared with NC rats suggesting that L-egt does not alter liver functions in the normal rats and provides further credence to its safety evaluation.

In this study, hypertriglyceridemia observed in DC rats indicates a substantial alteration in fatty acid metabolism, which is another risk factor for diabetic complications. This result is in accordance with similar studies that reported a significant association between increased serum triglycerides (TG) and type-2 diabetes (Thambiah et al., 2016, Rašković et al., 2019). The altered metabolism increased serum TG, which promotes the influx of free fatty acids into the systemic circulation with subsequent infiltration into the hepatocytes with resultant steatosis or NAFLD. Besides, IR can inhibit lipoprotein lipase (LPL), which enhances the mobilization of TG to skeletal muscles and adipose tissue, thereby reducing the TG clearance from circulation (Arguello et al., 2015, Jiang et al., 2013). From this study, the significant reduction in serum and liver TG in diabetic animals treated with L-egt alone, or in combination with metformin may result from the inhibition of *de novo* biosynthesis as well as enhance lipoprotein lipase activity to increase the TG clearance from the circulation or thereby reducing the risk of fatty liver and related injuries (Alves-Bezerra and Cohen, 2017, Kawano and Cohen, 2013). Furthermore, the reduced TG infiltration into the hepatocyte in the L-egt treated groups may result from the downregulation of SREBP1c and FAS genes that promote TG synthesis.

Chronic hyperglycemia has been reported to increase free radical production by the neutrophils and activated kupffer cells. Elevated free radicals (e.g., ROS, H₂O₂, OH⁻) compromises the integrity of the cells via lipid peroxidation in the cell membrane, upregulate inflammatory signals and induce hepatic apoptosis as well as irreversible damage to other biomolecules in the body (e.g., protein, RNA). These biochemical events may result in significant structural and functional damage in the liver (Masarone et al., 2018, Lucchesi et al., 2015). In the present study, the reduced MDA in the L-egt treated groups suggests that L-egt may inhibit oxidative degradation of the lipid bilayers in the cell membrane to enhance cellular integrity and the viability of membrane proteins that helps in cellular communication. The increased antioxidant enzymes (SOD, CAT, and GSH) work synergistically to reduce the deleterious effect of free radicals in the tissues. An increased SOD level in the liver homogenates facilitates the detoxification of free radicals by enhancing the conversion of superoxide into hydrogen peroxide, while CAT helps to neutralize the free radicals by degrading hydrogen peroxide into water and oxygen molecule. Increased GSH reduces hydrogen peroxides to water and other lipid peroxides to alcohol, usually in the mitochondrial and cytosol (Ighodaro and Akinloye, 2018). In this study, administration of L-egt, with or without metformin, increased the production and efficacy of these antioxidant enzymes in diabetic rats, thereby protects against oxidative damage, enhance cell integrity, and complement the efficacy of antioxidant enzymes against ROS. The improved antioxidant defense system observed in the L-egt treated group may result from the upregulation of Sirt1 and Nrf2 genes, which are the major transcription factors in the antioxidant signaling pathways. Nrf2 is a master regulator of cellular antioxidant response that stimulates the production of phase II cytoprotective antioxidant genes (e.g., heme-oxygenase-1 and

NAD(P)H oxidase-1) and ROS-detoxifying enzyme (e.g., GSH, SOD, CAT GPx) to mediate redox balance (Ma, 2013, Tonelli et al., 2017). Also, Sirt1 exerts a wide variety of biological functions, acting as a metabolic sensor that regulates lipid metabolism, inhibits free radical production from the respiratory chain, increases the detoxification of ROS by activating antioxidant enzymes, and downregulates the NF-kB signaling pathway (Li, 2014, Singh et al., 2018).

The suppression of inflammatory molecules in the liver cells in response to tissue injury has attracted significant interest in managing liver disease. Low-grade chronic inflammation in metabolic diseases, IR, and their deleterious effect on liver function have been reported in previous studies (Ning et al., 2015, Hou et al., 2018). The concentration of TNF- α (a proinflammatory cytokine) varies directly with the severity of the liver disease (Seo et al., 2013, Yang and Seki, 2015), while MCP-1 and TGF- β 1 are responsible for monocyte activation, fatty acid accumulation, and fibrosis in the liver cells (Mandrekar et al., 2011, Nair and Nath, 2020). Taken together, the reduced inflammatory and fibrotic markers in the liver homogenates of diabetic animals treated with L-egt, with or without metformin showed that this treatment regimen may prevent hepatic inflammation and fibrosis, immune cell activation and apoptosis, as well as macrophage infiltration into the hepatocytes to reduce structural and functional damage (Gehrke and Schattenberg, 2020, Del Campo et al., 2018). This result is further supported by the downregulation of NF-kB (a major transcription factor in the inflammatory pathway) and TGF- β 1 gene expression in the L-egt treated groups. It has been established that activation of NF-kB causes the transcription of several genes that mediate cellular inflammation, while L-ergothioneine has been demonstrated to modulate NF-kB signaling to attenuate hepatic injury (Zeng et al., 2016, Salama and Omar, 2021). Thus, L-egt may protect against liver injury associated with type-2 diabetes by reducing hepatic inflammation and fibrosis

The histopathological examination of liver tissue provides essential information on the structural integrity of the liver. The disruption of the hepatic sinusoids and congested central vein with mild degeneration of the hepatocytes observed in the DC rats suggests significant liver damage. This result is consistent with other studies (Rodríguez et al., 2018, Ogar et al., 2019). However, improvement in the structural integrity of the diabetic liver treated with L-egt and metformin can be associated with the hepato-protective function of this treatment regimen. The regular appearance of hepatic sinusoids and central veins in the DEM rats may enhance adequate blood supply from the hepatic artery and portal vein towards the central veins, thus preventing ischemic cell death.

Conclusion

In conclusion, this study showed that L-egt can alleviate oxidative damage by upregulating Sirt1/Nrf2 expression and its downstream antioxidant molecules, downregulate NF-kB and TGF- β 1 expression to reduce hepatic inflammation and fibrosis as well as reduce SREBP1c and FAS expression to lower hypertriglyceridemia and attenuate hepatocyte fatty acid accumulation to protect liver function during diabetes. Thus, supplementation of L-egt could be used as an adjuvant regimen with metformin therapy in the early stage of diabetes to prevent the development or alter the progression of liver complications associated with diabetes. However, further studies to evaluate the status of L-egt transporters and protein

expression of these transcription factors are required to provide the detailed mechanism of action of L-egt.

Declarations

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Figures

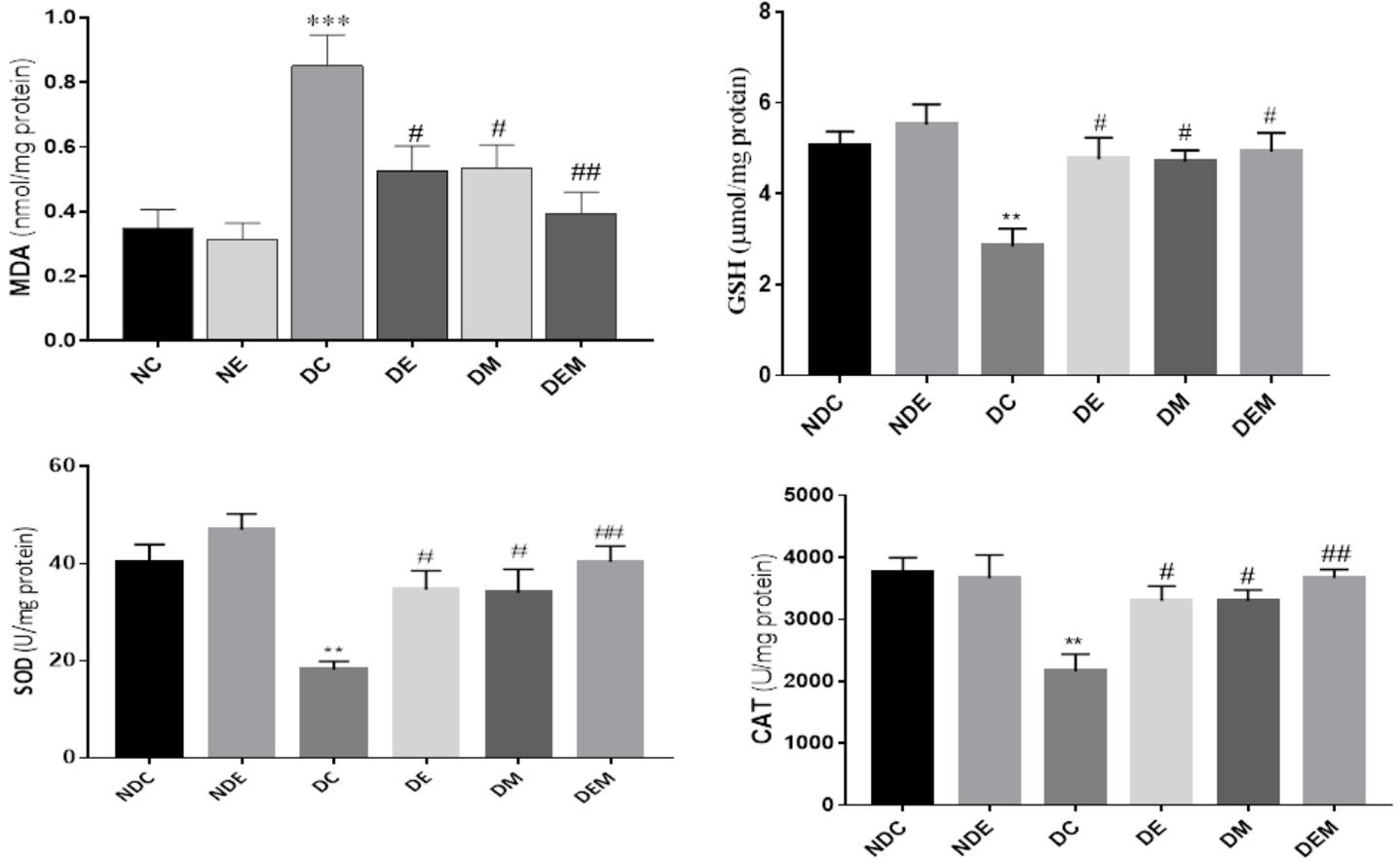


Figure 1

a: MDA levels in L-egt, with or without metformin, treated T2DM rats
 b: SOD levels in L-egt, with or without metformin, treated T2DM rats
 c: GSH levels in L-egt, with or without metformin, treated T2DM rats
 d: CAT levels in L-egt, with or without metformin, treated T2DM rats

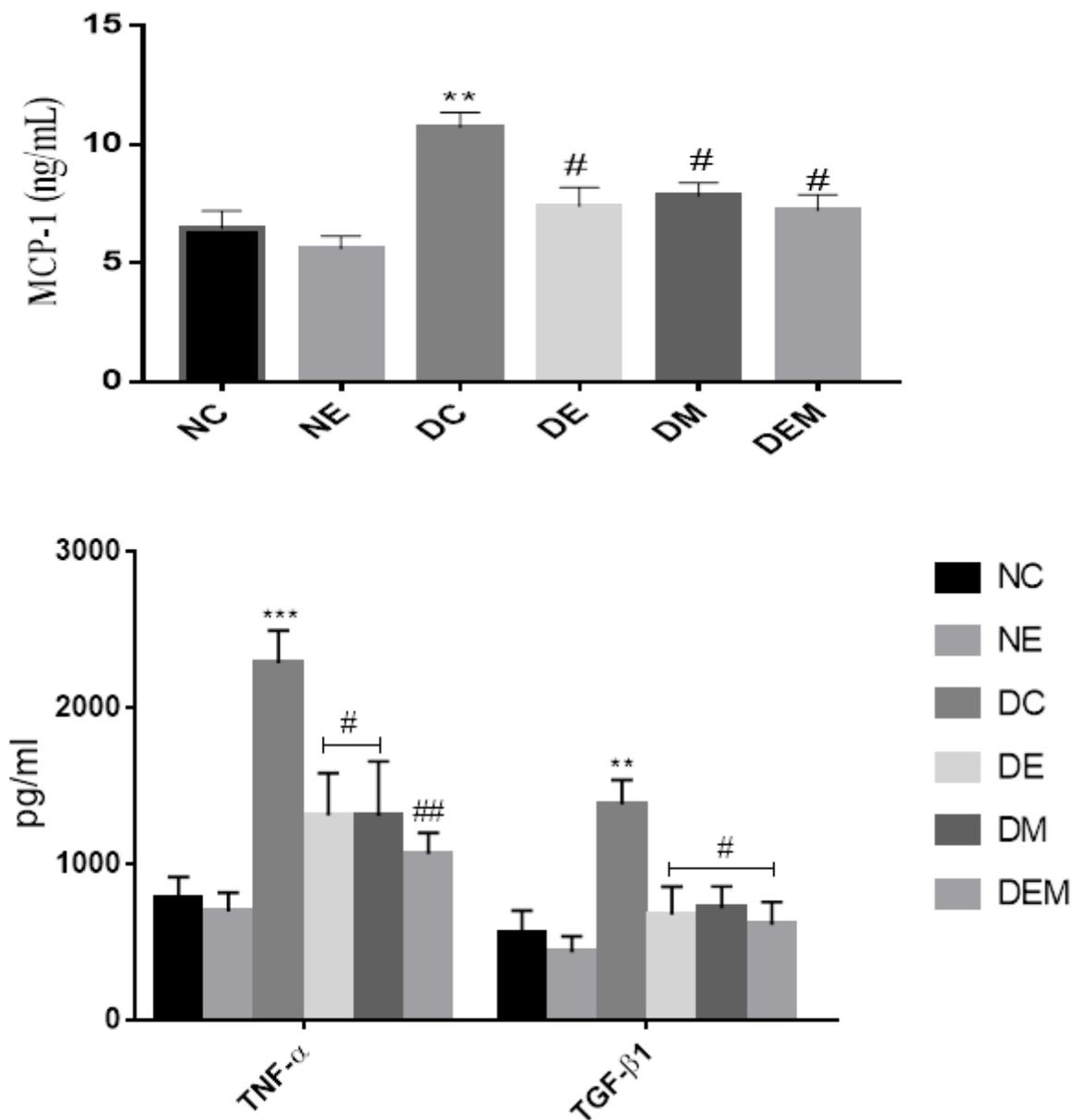


Figure 2

a-b: Effect of L-egt, with or without metformin, on liver inflammatory biomarkers (A) MCP1, (B) TNF- α and TGF- β 1 in T2DM-rats. *** $p < 0.001$, ** $p < 0.01$ vs NC; ## $p < 0.01$, # $p < 0.05$ vs DC. (n=5). NC= Nondiabetic control; NE=nondiabetic plus L-egt; DC= diabetic control; DE=diabetic plus L-egt; DM=diabetic plus metformin and DEM= diabetic plus L-egt plus metformin. (supplemental data).

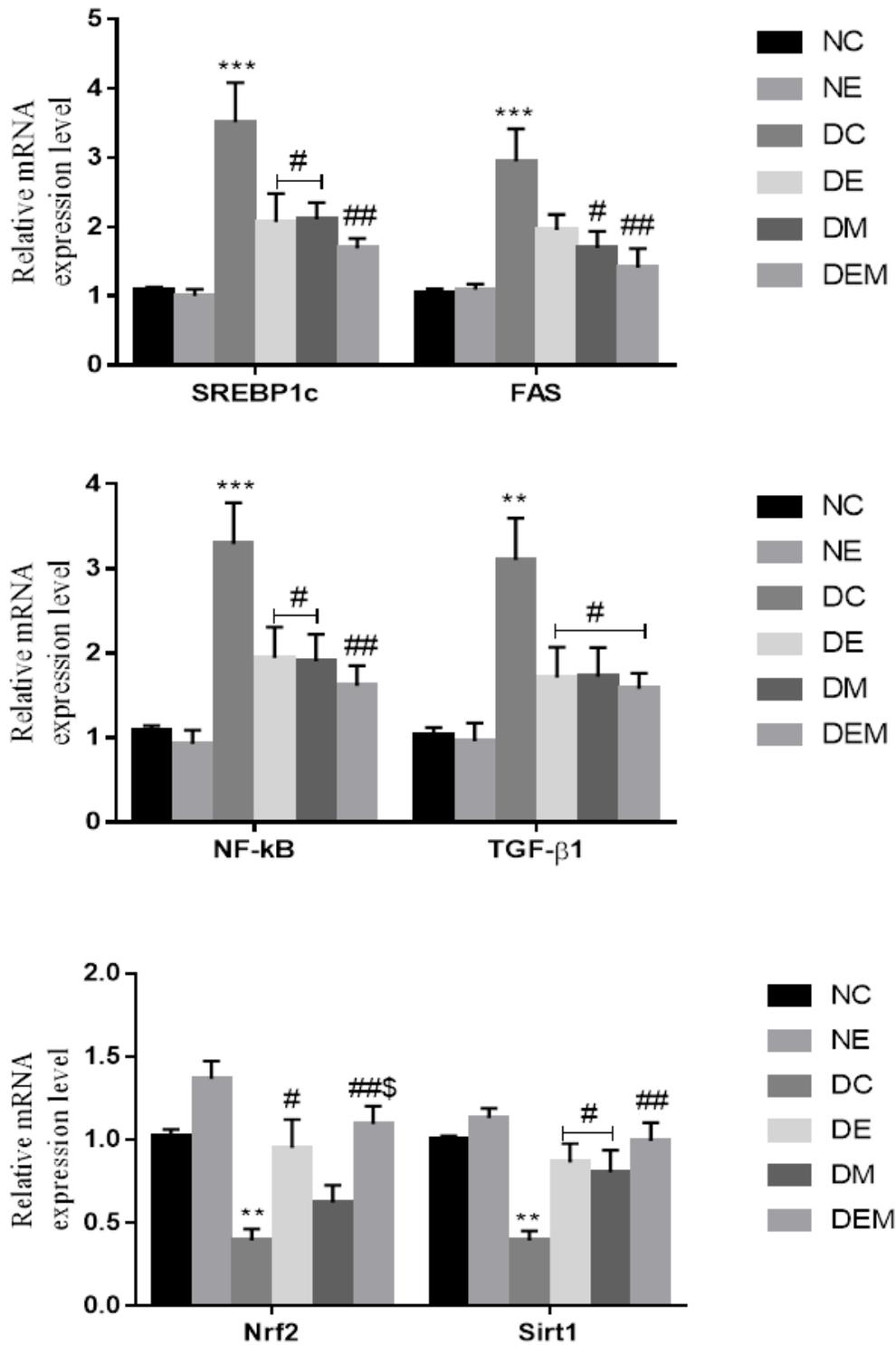


Figure 3

a-c: Relative mRNA expression level of (A) SREBP1c and FAS (B), NF-kB and TGF-β1 (C), Nrf2 and Sirt1 relative gene expression in liver tissues of diabetic animals treated with L-egt with or without metformin for 7 weeks. ***p<0.001, **p<0.01 vs NC groups; #p<0.05, ##p<0.01 vs DC group and \$ p<0.05 vs DM (n=5). NC= Nondiabetic control; NE=nondiabetic plus L-egt; DC= diabetic control; DE=diabetic plus L-egt; DM=diabetic plus metformin and DEM= diabetic plus L-egt plus metformin (supplemental data).

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