**Ruscogenin Protects against High-Fat Diet-Induced Nonalcoholic Steatohepatitis in Hamsters**

**Authors**

Hung-Jen Lu1, Shorong-Shii Liou2, Chia Ju Chang2, Sheng Da Lin2, Cheng Yang2, Ming-Chang Wu1, I-Min Liu2

1 Department of Food Science, College of Agriculture, National Pingtung University of Science and Technology, Neipu Township, Pingtung County, Taiwan, ROC
2 Department of Pharmacy & Graduate Institute of Pharmaceutical Technology, Tajen University, Yanpu Township, Pingtung County, Taiwan, ROC

**Key words**

- fatty acid β-oxidation
- inflammatory cytokines
- lipogenic gene
- nonalcoholic steatohepatitis
- ruscogenin

**Abstract**

The protective effects of ruscogenin on nonalcoholic steatohepatitis in hamsters fed a high-fat diet were investigated. Ruscogenin (0.3, 1.0, or 3.0 mg/kg/day) was orally administered by gavage once daily for eight weeks. A high-fat diet induced increases in plasma levels of total cholesterol, triglycerides, and free fatty acids, while the degree of insulin resistance was lowered by ruscogenin. High-fat diet-induced hepatic steatosis and necroinflammation were improved by ruscogenin. Gene expression of inflammatory cytokines and activity of nuclear transcription factor-κB were also increased in the high-fat diet group, which were attenuated by ruscogenin. Ruscogenin decreased hepatic mRNA levels of sterol regulatory element-binding protein-1c and its lipogenic target genes. The hepatic mRNA expression of peroxisome proliferator-activated receptor α, together with its target genes responsible for fatty acid β-oxidation were upregulated by ruscogenin. In conclusion, these findings suggest that ruscogenin may attenuate high-fat diet-induced steatohepatitis through anti-inflammatory mechanisms, reducing hepatic lipogenic gene expression, and upregulating proteins in the fatty acid oxidation process.

**Introduction**

Nonalcoholic fatty liver disease (NAFLD) is defined as the accumulation of liver fat exceeding 5% of hepatocytes in the absence of significant alcohol intake (significant alcohol intake being defined as 20 g/10 g per day for men/women, respectively) or other specific etiology [1]. Nonalcoholic steatohepatitis (NASH), the extreme form of NAFLD, is characterized by an increase in fat deposition, fibrosis, oxidative stress, and chronic inflammation in the liver [2]. It is also a common cause of cirrhosis and could later progress to hepatocellular carcinoma [3]. A “multi-hit” theory has been proposed to depict the key pathogenic processes of NASH development. In this theory, obesity and insulin resistance is thought to be the “first hit” to the liver because it induces a net accumulation of lipid in hepatocytes which may cause the occurrence of following “hits”, such as oxidative stress, inflammation, fibrosis, and apoptosis [4]. Therapies that limit hepatic injury and the related occurrence of inflammation and fibrosis are particularly appealing for this condition. Ruscogenin [(1β,3β,25R)-spirost-5-en-1,3-diol; ○ Fig. 1], first isolated from Ruscus aculeatus, is also a major steroidal sapogenin of the traditional Chinese herb Radix Ophiopogon japonicus, which has been clinically used for a long time to treat acute and chronic inflammatory and cardiovascular diseases [5]. In Chinese medicine, Radix Ophiopogon japonicus is the cardinal herb for yin deficiency. According to the Chinese Herbal Medicine Materia Medica, the herb is sweet, slightly bitter and slightly cold, enters the heart, lungs and stomach channels, and nourishes the yin of the stomach, spleen, heart and lungs, clears heat, and quiets irritability [6]. Ruscogenin acts as an elastase inhibitor, decreases capillary permeability, and is widely used to treat chronic venous insufficiency and vasculitis [7,8]. It has been previously found to exert significant antithrombotic activity, and the possible mechanism of ruscogenin could be attributed to its inhibition of intercellular adhesion molecule-1 upregulation via nuclear factor (NF)-κB p65 translocation induced by tumor necrosis factor (TNF)-α [9,10]. It was also found that ruscogenin significantly attenuated lipopolysaccharide-induced acute lung injury in a model with mice, which was possibly linked to the inhibition of NF-κB activation [11]. Such activities of ruscogenin indicate its potential protec-
tissue, and perirenal adipose tissue were significantly lower in weight gain. The relative weights of the liver, epididymal adipose plate (AST) activities in HFD-fed hamsters were higher than those in the normal diet (ND)-fed group (Table 1). High doses of ruscogenin (3.0 mg/kg/day) suppressed body weight gain. The relative weights of the liver, epididymal adipose tissue, and perirenal adipose tissue were significantly lower in ruscogenin (3.0 mg/kg/day)-supplemented groups than those in the HFD-fed group (Table 1). Similar results were seen in HFD-fed hamsters treated with fenofibrate (Table 1). No significant differences in daily food or water intake were observed between the groups over the experimental period (Table 1). Plasma alanine aminotransferase (ALT) and aspartate aminotransferase (AST) activities in HFD-fed hamsters were higher than those in the ND-fed group (Table 1). The ALT and AST activities were markedly reduced in HFD-fed hamsters treated for eight weeks with a high dose (3.0 mg/kg/day) of ruscogenin (Table 1). The HFD caused elevated concentrations of plasma total cholesterol (TC), triglycerides (TG), and low-density lipoprotein cholesterol (LDL-C). Oral administration of ruscogenin at a dose of 0.3, 1.0, or 3.0 mg/kg/day significantly reduced plasma TC levels (22.8%, 31.4%, and 39.7% reduction, respectively; Fig. 2A). All doses of ruscogenin decreased plasma TG levels in HFD-fed hamsters; the reduction of plasma TG induced by ruscogenin at 3.0 mg/kg/day was nearly 32.1% (Fig. 2A). Ruscogenin at an oral dose of 0.3, 1.0, or 3.0 mg/kg/day significantly reduced total plasma LDL-C levels (32.1%, 42.2%, and 52.8% reduction, respectively) compared to that of vehicle-treated HFD-fed hamsters (Fig. 2A). Plasma TC, TG, and LDL-C concentrations were reduced significantly by 46.7%, 37.8%, and 60.4%, respectively, in fenofibrate-treated HFD-fed hamsters compared with vehicle-treated HFD-fed hamsters (Fig. 2A). The plasma concentration of high-density lipoprotein cholesterol (HDL-C) in HFD-fed hamsters was reduced to 60.4% of the level observed in the ND-fed group (Fig. 2A). After eight weeks of treatment with ruscogenin (3.0 mg/kg/day) or fenofibrate, the plasma HDL-C concentration in HFD-fed hamsters was elevated to nearly that of the ND-fed group (Fig. 2A). Plasma free fatty acid (FFA) levels in vehicle-treated HFD-fed hamsters were about 2.0-fold of that observed in the ND-fed group (Fig. 2A). The plasma FFA levels decreased by 38.1% in HFD-fed hamsters treated with ruscogenin (3.0 mg/kg/day) compared to their vehicle-treated counterparts (Fig. 2A). Fenofibrate treatment reduced FFA concentrations in HFD-fed hamsters by 38.3% relative to the level in vehicle-treated HFD-fed hamsters (Fig. 2A). The hepatic TC level was significantly higher in HFD-fed hamsters than in hamsters from the ND-fed group, which was reduced by 37.2% in HFD-fed hamsters treated with ruscogenin (3.0 mg/kg/day; Fig. 2B). Ruscogenin treatment (3.0 mg/kg/day) also produced a significant reduction in the hepatic TG concentration by 27.2% of that in the vehicle-treated HFD-fed hamsters (Fig. 2B). Hepatic TC and TG levels were significantly reduced (by 44.5% Table 1 The effects of treatments on body weight (BW), liver and visceral fat weight, feeding behaviors, and plasma transaminase activity of hamsters.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>ND-fed Vehicle</th>
<th>HFD-fed Vehicle</th>
<th>Ruscogenin (mg/kg/day)</th>
<th>0.3</th>
<th>1.0</th>
<th>3.0</th>
<th>Fenofibrate (100 mg/kg/day)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Initial BW (g)</td>
<td>93.86 ± 3.52</td>
<td>94.12 ± 2.98</td>
<td>93.71 ± 3.07</td>
<td>92.82 ± 2.78</td>
<td>93.64 ± 4.11</td>
<td>93.42 ± 3.62</td>
<td></td>
</tr>
<tr>
<td>Final BW (g)</td>
<td>227.33 ± 8.26</td>
<td>301.02 ± 7.42</td>
<td>285.21 ± 6.33</td>
<td>279.32 ± 7.09</td>
<td>260.35 ± 6.12</td>
<td>246.13 ± 7.83</td>
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<tr>
<td>Epididymal WAT (mg/100 g BW)</td>
<td>161.92 ± 18.26</td>
<td>242.39 ± 20.31</td>
<td>233.47 ± 17.82</td>
<td>207.85 ± 16.29</td>
<td>173.69 ± 17.37</td>
<td>167.58 ± 15.33</td>
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</tr>
<tr>
<td>Perirenal WAT (mg/100 g BW)</td>
<td>80.72 ± 6.41</td>
<td>111.43 ± 7.32</td>
<td>101.12 ± 7.23</td>
<td>88.84 ± 8.16</td>
<td>86.36 ± 7.32</td>
<td>84.17 ± 8.08</td>
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<tr>
<td>Mesenteric WAT (mg/100 g BW)</td>
<td>75.27 ± 5.26</td>
<td>104.25 ± 6.17</td>
<td>94.82 ± 4.38</td>
<td>87.32 ± 5.17</td>
<td>81.24 ± 6.21</td>
<td>77.39 ± 5.86</td>
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<tr>
<td>Hepatic weight coefficient (g/100 g BW)</td>
<td>4.81 ± 0.83</td>
<td>5.83 ± 0.76</td>
<td>5.62 ± 0.61</td>
<td>5.23 ± 0.69</td>
<td>5.17 ± 0.53</td>
<td>4.91 ± 0.66</td>
<td></td>
</tr>
<tr>
<td>Food intake (g/day)</td>
<td>10.16 ± 1.76</td>
<td>11.05 ± 1.83</td>
<td>10.76 ± 1.97</td>
<td>10.96 ± 1.58</td>
<td>9.92 ± 1.64</td>
<td>10.08 ± 2.01</td>
<td></td>
</tr>
<tr>
<td>Water intake (mL/day)</td>
<td>12.78 ± 4.83</td>
<td>13.16 ± 5.68</td>
<td>12.87 ± 6.11</td>
<td>13.11 ± 5.87</td>
<td>13.06 ± 4.36</td>
<td>12.97 ± 5.26</td>
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<tr>
<td>Plasma ALT (U/L)</td>
<td>52.61 ± 7.13</td>
<td>340.05 ± 15.17</td>
<td>276.31 ± 12.33</td>
<td>200.49 ± 11.86</td>
<td>138.28 ± 9.12</td>
<td>95.24 ± 6.13</td>
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<tr>
<td>Plasma AST (U/L)</td>
<td>121.27 ± 9.38</td>
<td>489.32 ± 18.73</td>
<td>406.35 ± 17.42</td>
<td>294.85 ± 18.26</td>
<td>218.21 ± 19.37</td>
<td>175.42 ± 10.86</td>
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The vehicle (distilled water) used to prepare the tested medication solution was given at the same volume. Values (mean ± S.D.) were obtained from each group of eight animals after eight weeks of the experimental period. *P < 0.05 and **P < 0.01 compared to vehicle-treated HFD-fed hamsters in each group, respectively. †P < 0.05 and ‡P < 0.01 compared to vehicle-treated HFD-fed hamsters in each group, respectively.

Fig. 1 Structure of ruscogenin.
and 35.5%, respectively) in fenofibrate-treated HFD-fed hamsters compared with their vehicle-treated counterparts (Fig. 2B). The hepatic FFA levels were significantly increased in hamsters fed an HFD, which were decreased by 37.3% and 41.2% in HFD-fed hamsters treated with ruscogenin (3.0 mg/kg/day) and fenofibrate, respectively (Fig. 2B).

Representative histological photomicrographs of liver specimens are shown in Fig. 3. Hamsters fed a ND had normal liver histological findings; however, numerous macrovascular fat droplets and mild necroinflammatory foci were present in the livers of those fed an HFD. The treatment of HFD-fed hamsters with ruscogenin (3.0 mg/kg/day) and fenofibrate, respectively (Fig. 2B).

Plasma glucose and insulin concentrations as well as the homeostasis model assessment of insulin resistance (HOMA-IR) were higher in vehicle-treated HFD-fed hamsters than those in the vehicle-treated ND-fed group (Fig. 4). After eight weeks of treatment with ruscogenin (3.0 mg/kg/day), plasma glucose concentrations were approximately 9.3% lower in HFD-fed hamsters compared with their vehicle-treated counterparts; corresponding plasma insulin and HOMA-IR values were 27.7% and 33.2% lower, respectively (Fig. 4). After eight weeks of treatment with fenofibrate, plasma glucose, plasma insulin, and HOMA-IR values were 7.6%, 32.5%, and 38.4% lower, respectively, in HFD-fed hamsters compared with their vehicle-treated counterparts (Fig. 4).

The hepatic levels of monocyte chemoattractant protein (MCP)-1, TNF-α, and interleukin (IL)-6 in HFD-fed hamsters were significantly higher (2.4-, 2.3-, and 2.9-fold, respectively) than those of the ND-fed group (Fig. 5). Administration of ruscogenin (3.0 mg/kg/day) to HFD-fed hamsters for eight weeks significantly downregulated the hepatic levels of MCP-1, TNF-α, and IL-6 to 51.1%, 63.3%, and 53.3%, respectively, compared with those in the vehicle-treated counterparts (Fig. 5). Fenofibrate treatment reversed the HFD-induced increase of hepatic levels of MCP-1, TNF-α, and IL-6 to 36.1%, 57.0% and 44.9%, respectively, compared with their vehicle-treated counterparts (Fig. 5).
An HFD significantly increased NF-κB activity (by 3.2-fold relative to that of the vehicle-treated ND-fed group) in the livers of hamsters (Fig. 6A). The HFD-induced upregulation of NF-κB activity was reduced 39.1% and 48.1% relative to that of the HFD-fed hamsters after eight weeks of treatment with ruscogenin (3.0 mg/kg/day) or fenofibrate, respectively (Fig. 6A). The phosphorylation degree of hepatic NF-κB in HFD-fed hamsters was also higher by 2.5-fold relative to their vehicle-treated

Table 2  Summary of steatosis and necroinflammation scores of ND- or HFD-fed hamsters receiving eight weeks of treatment.

<table>
<thead>
<tr>
<th></th>
<th>ND-fed Vehicle</th>
<th>HFD-fed Vehicle</th>
<th>Ruscogenin (mg/kg/day)</th>
<th>Fenofibrate (100 mg/kg/day)</th>
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<tr>
<td></td>
<td></td>
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<td>0.3</td>
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<td>Steatosis</td>
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<td>0</td>
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<td>0</td>
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<td>1</td>
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<td>5</td>
<td>5</td>
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<td>0</td>
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<tr>
<td>Necroinflammation</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>1</td>
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<td></td>
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<td>6</td>
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<td>2</td>
<td>3</td>
<td>2</td>
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<td>3</td>
<td>1</td>
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There were eight animals in each group. Data are expressed as the number of hamsters exhibiting the grade of steatosis or necroinflammation indicated.

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counterparts (Fig. 6B). Treatment for eight weeks with ruscogenin (3.0 mg/kg/day) or fenofibrate decreased phosphorylated NF-κB in the livers of HFD-fed hamsters to 59.3% and 49.1%, respectively, relative to the vehicle-treated controls (Fig. 6B).

An HFD markedly increased the hepatic mRNA levels of sterol regulatory element-binding protein (SREBP)-1c in hamsters 3.2-fold relative to those in the ND-fed group (Fig. 7A). Ruscogenin (3.0 mg/kg/day) suppressed the HFD-induced stimulation in hepatic mRNA levels of SREBP-1c to 63.5% relative to those in the vehicle-treated group (Fig. 7A). Hepatic mRNA levels of SREBP-1c were reduced (by 46.9%) in fenofibrate-treated HFD-fed hamsters compared with their vehicle-treated counterparts (Fig. 7A).

An HFD caused a 2.8-fold induction of hepatic acetyl-CoA carboxylase (ACC) mRNA and a 2.5-fold induction of hepatic fatty acid synthase (FAS) mRNA relative to those in the ND-fed group. The hepatic mRNA levels of ACC and FAS were downregulated by ruscogenin (3.0 mg/kg/day) treatment, with a decrease of 23.5% and 36.7%, respectively, when compared with those observed in the vehicle-treated group (Fig. 7A). The HFD-induced mRNA levels of ACC and FAS in the liver were reversed after fenofibrate treatment (a 48.6% and 47.1% decrease, respectively) compared to those in the vehicle-treated group (Fig. 7A).

The mRNA levels of peroxisome proliferator-activated receptor (PPAR)α in the livers of HFD-fed hamsters were lowered to 55.1% compared with those from the ND-fed group (Fig. 7B). HFD-fed hamsters administered ruscogenin (3.0 mg/kg/day) or fenofibrate for eight weeks upregulated the hepatic PPARα mRNA levels to 1.6- and 2.0-fold relative to those in their vehicle-treated counterparts, respectively (Fig. 7B). Compared with the ND-fed group, the hepatic mRNA levels of carnitine palmitoyl transferase (CPT)-1 and acyl-CoA oxidase (ACO) in HFD-fed hamsters decreased obviously, and were upregulated by fenofibrate treatment (a 1.9- and 2.3-fold increase, respectively). The hepatic mRNA levels of CPT-1 and ACO in HFD-fed hamsters receiving ruscogenin (3.0 mg/kg/day) supplementation were increased 1.6- and 2.0-fold relative to the expression levels in their vehicle-treated counterparts, respectively (Fig. 7B).
In our model, HFD-fed hamsters developed hepatic steatosis, visceral obesity, hyperlipidemia, and increased FFA and HOMA-IR values, mimicking almost all of the clinical features of human NAFLD [12]. Fenofibrate, a fibric acid derivative, is a commercially available drug in the management of hyperlipidemia and NAFLD [13]. Fenofibrate was used to compare the efficacy of ruscogenin. Similar to the effects of fenofibrate, we found that ruscogenin supplementation significantly decreased body and liver weights, and plasma and liver contents of TG, TC, and FFA. In addition, ruscogenin attenuated the development of HFD-induced hepatic steatosis and injury as assessed by microscopic analysis. These data demonstrated that ruscogenin shows biological activity towards the regulation of lipid metabolism and steatosis development in the liver.

As inflammation plays a pivotal role in NAFLD, its modulation represents an important pharmacological target of inflammatory activation [14]. We therefore investigated whether ruscogenin could improve fatty liver changes by decreasing inflammatory cytokines in the HFD-induced NAFLD hamsters. Administration of ruscogenin during NAFLD development significantly attenuated the expression of MCP-1, TNF-α, and IL-6 in the liver of HFD-fed hamsters. These data indicate that ruscogenin might be effective for preventing and reversing the inflammatory response, which may accelerate the lipid metabolism disorder and/or more severe liver injuries. Actually, these inflammatory cytokines are shown to be involved in the development of insulin resistance [15].
observed that not only hyperinsulinemia but also higher HO-
MA-IR values were alleviated by ruscogenin treatment in
HFD-fed hamsters. These results further suggest that ruscogenin
could ameliorate fatty liver in the insulin-resistant state.
NF-κB is a ubiquitous and well-known transcription factor, which
undergoes phosphorylation on serine 276 in its p65 subunit and
associates with surrounding chromatin components. It sub-
sequently binds with DNA and promotes the transcription of proin-
flammatory cytokines, chemokines, and adhesion molecules
[16]. Thus, detection of the phosphorylated p65 subunit of NF-
κB was effective for evaluating NF-κB activation [17]. Ruscogenin
has been reported to have anti-inflammatory activity through in-
hibiting NF-κB activity [10, 11]. The inhibition of NF-κB by rusco-
genin might be a critical step for prevention of the cascading in-
flammatory response during NAFLD development. In the present
study, ruscogenin significantly inhibited hepatic NF-κB activation
in HFD-fed hamsters, as evidenced by a decrease in NF-κB activity
and downregulation of phosphorylated NF-κB. These results sug-
gest that ruscogenin inhibits the activation of NF-κB, leading to
the downregulation of inflammatory mediators and, therefore,
shows a promising effect in preventing NAFLD. Whether rusco-
genin can directly target NF-κB or act as the upstream signaling
molecules for its activation requires further investigation.

To explore the possible mechanisms whereby ruscogenin de-
creases hepatic lipid accumulation, we then investigated the ex-
pression levels of several genes related to lipid metabolism, in-
cluding lipogenesis and β-oxidation. SREBP-1c has been shown
to regulate the transcription of genes in the lipogenic pathway in-
cluding ACC and FAS [17]. Liver-specific ACC knockout mice
showed a decreased hepatic triglyceride accumulation, suggest-
ing that ACC plays a crucial role in the regulation of lipogenesis
[18], FAS catalyzes the last step in fatty acid biosynthesis and is
thus believed to be a major determinant of the maximal hepatic
capacity to generate fatty acids by de novo lipogenesis [19]. Low
levels of SREBP-1c mRNA in the livers of HFD-fed hamsters re-
ceiving ruscogenin were accompanied by a concomitant reduc-
tion in the expression of ACC and FAS mRNA. Ruscogenin is thus
likely to have an inhibitory effect on SREBP-1c expression, which
in turn decreases transcription of its target lipogenic genes,
thereby reducing enzyme activity and resulting in a low rate of
lipid synthesis. Ruscogenin appeared to mediate its beneficial ef-
ects on HFD-induced hepatic steatosis by downregulating ex-
pression of hepatic genes related to lipid synthesis.
PPARα mediates β-oxidation of fatty acids, and agonists of PPARα
possess the property of reducing lipid dystrophy and obesity
[20]. PPARα activates some of the key enzymes of β-oxidation
and β-oxidation of fatty acids in the liver [21]. HFD-fed hamsters
displayed a reduced expression of PPARα in the liver. A multitude
of studies have postulated that during diet-induced obesity, fatty
acid oxidation is left incomplete accompanied by mitochondrial
lipid overload and dysfunction. To explore whether the effect of
ruscogenin on the attenuation of HFD-induced hepatic steatosis
was related to the PPARα-mediated pathway, mRNA expression
of PPARα and its target genes responsible for β-oxidation of fatty
acids was measured. Ruscogenin markedly increased the HFD-in-
duced low expression of hepatic PPARα mRNA. Furthermore,
ruscogenin treatment upregulated the PPARα-mediated transcrip-
tion of CPT-1 and ACO mRNA in the liver of HFD-fed hamsters.
CPT-1 and ACO are major enzymes that catalyze fatty acid β-oxi-
dation and are responsive to PPARα activation [22, 23]. The simi-
lar trend in expression among these genes suggests that rusco-
genin enhanced β-oxidation of fatty acids via the pathway involving
PPARα-mediated gene transcription.
ALT and AST are considered to be sensitive indicators of hepato-
cellular damage and, within limits, can provide a quantitative
evaluation of the degree of damage to the liver [24]. We found
that the ALT and AST activities were markedly reduced in
HFD-fed hamsters treated for eight weeks with ruscogenin. It
seems that animals treated with ruscogenin at the chosen doses
exhibited no evidence of hepatotoxicity. For a more reliable
safety evaluation performed on the basis of the acceptable daily
intake concept, data on the chronic toxicity, reproductive toxic-
ity, genotoxicity, and carcinogenicity of ruscogenin would also
be required.

In conclusion, the results of this study demonstrated that rusco-
genin exerted a protective effect on HFD-induced hepatic steato-
sis in hamsters by modulating lipid metabolism-associated
genes. It is also possible that the inhibition of NF-κB activation
thereby led to the downregulation of inflammatory mediators
and this is integral to the protective functions of ruscogenin in
HFD-induced hepatocellular injury and its subsequent progres-
sion to steatohepatitis.

Materials and Methods

Animal models
Male Golden Syrian hamsters, eight weeks old and weighing
90 ± 10 g, were obtained from the National Laboratory Animal
Center. They were maintained in a temperature-controlled room
(25 ± 1 °C) on a 12-h:12-h light–dark cycle (lights on at 06:00 h)
in our animal center. Food and water were provided ad libitum.
An ND (#D12450B, Research Diets) with 20 kcal% protein,
70 kcal% carbohydrate, and 10 kcal% fat from lard was used as
the maintenance and control diet. A purified ingredient HFD
20 kcal% protein, 35 kcal% carbohydrate, and 45 kcal% fat primar-
ily from lard (#D12451, Research Diets) was used to induce
weight gain [25]. All animal procedures were performed accord-
ing to the Guide for the Care and Use of Laboratory Animals of
the National Institutes of Health, as well as the guidelines of the Ani-
mal Welfare Act. These studies were conducted with the approval
of the Institutional Animal Care and Use Committee (IACUC) at
Tajen University (approval number, IACUC 101–16; approval
date, December 24, 2012).

Treatment protocols
After being fed an HFD for two weeks, hamsters were dosed by
oral gavage once per day for eight weeks with ruscogenin (doses
of 0.3, 1.0 or 3.0 mg/kg) in a volume of 1.5 ml/kg distilled water.
The dosage regime was selected based on a previous report dem-
onstrating that ruscogenin (≥98%; Chengdu Biopurify Phyto-
chemicals Ltd.; Cat. No. 472–11–7) at the indicated dosage regi-
men was potentially effective in inhibiting inflammation in mice
[11]. Another group of HFD-fed hamsters was treated orally
for eight weeks with 100 mg/kg/day fenofibrate (isopropyl 2-[4-
(4-chlorobenzoyl)phenoxyl]-2-methylpropionate; Laboratories
Fournier SA) with a purity greater than 99.5%. Two g of powdered
fenofibrate was dissolved in 20 ml of distilled water, so that 1 ml
will contain 100 mg of fenofibrate. The dose of fenofibrate was
based on a study reporting that long-term fenofibrate treatment
could ameliorate hepatic insulin resistance and steatosis in high-
fructose-fed mice [26]. The control groups received water by ga-
vage instead of ruscogenin or fenofibrate. Water consumption,
food intake, and body weight were measured individually at the same time (08:00 h) on each day throughout the experiment. Food cups containing fresh food were weighed at the beginning and end of each 24-h period. Food intake was calculated by determining the difference in food cup weights, adjusting for any spillage that occurred. Water intake was calculated by measuring the difference in water bottle weights at the beginning and end of the daily change of water.

After eight weeks of treatment, animals were weighed and sacrificed by being deeply anesthetized with ketamine (50–75 mg/kg) after fasting for 12 hours. Blood samples were taken from the inferior vena cava for analysis. After blood collection, the left lobe of the liver, and visceral and subcutaneous white adipose tissue (WAT) were removed, rinsed with a physiological saline solution, and immediately stored at −80°C in liquid nitrogen until assayed. The hepatic weight coefficient was also calculated as liver weight (g) divided by body weight (100 g). The right hepatic lobe was fixed in 10% neutralized formalin for histology.

Determination of plasma biochemical parameters and insulin sensitivity
Blood samples were centrifuged at 2000 × g for 10 min at 4°C. The plasma was then removed and placed into aliquots for the respective analyses. Kits for determining plasma glucose (Cat. No. 10009582) concentration was purchased from Cayman Chemical Company. Commercial enzyme-linked immunosorbent assay (ELISA) kits were used to quantify plasma insulin concentration (LINCO Research, Inc.; Cat. #EZRMI-13K). Whole-body insulin sensitivity was estimated using HOMA-IR with the following formula: [fasting plasma glucose (mmol) × fasting plasma insulin (mU/mL)]/22.5 [27]. Diagnostic kits for determination of plasma levels of TC (Cat. # 10007640) and TG (Cat. # 10010303) were purchased from Cayman Chemical Company. The diagnostic kit for determination of plasma levels of HDL-C was purchased from Bio-Quan Diagnostics (Cat. # BQ 019CR). LDL-C concentrations in plasma were determined by a commercial ELISA kit (anti-bodies-online, Inc.; Cat. #ABIN416222). The plasma FFA level was determined using an FFA quantification kit obtained from Abcam plc (Cat. # ab65341). Kits for determination of plasma ALT (EC 2.6.1.2) (Cat. No. A524-780TM) and AST (EC 2.6.1.1) (Cat. No. A559-780TM) concentrations were purchased from Tec-Diagnostics. All analyses were performed in triplicate in accordance with the manuals provided by the manufacturers.

Measurement of hepatic lipids
Liver tissue (0.5 g) was homogenized for 5 min in 1 mL of ice-cold phosphate-buffered saline (PBS) using a tissue homogenizer (T-10 basic Ultra-Turrax; IKAWerke GmbH & Co. KG). A 2-ml aliquot of chloroform/methanol (2:1, v/v) was added to the homogenate was centrifuged at 2300 × g for 15 min. The bottom of organic phase was dried under nitrogen gas and the residue was suspended in 1 mL of 3% Triton X-100, and an aliquot was used for the measurement of TG, TC, and FFA using the same commercial kits used for plasma analysis. Samples were assayed in triplicate according to the manufacturer’s instructions.

Hepatic cytokines determination
Liver tissue (0.5 g) was homogenized with a teflon Potter-Elvejhem tissue homogenizer in 10 mmol/L Tris-HCl buffered solution (pH 7.4) containing 2 mol/L NaCl, 1 mmol/L EDTA, 0.01% Tween 80, and 1 mmol/L PMSF, and centrifuged at 9000 × g for 30 min at 4°C. The resultant supernatant was used for cytokine determination. ELISA kits for the determination of MCP-1 (Cat. No. ab100777), TNFα (Cat. No. ab46070), and IL-6 (Cat. No. ab100772) were obtained from Abcam, Inc. Samples were assayed in triplicate according to the manufacturer’s instructions. The protein concentrations of liver filtrate were determined using a Bio-Rad protein assay kit (Bio-Rad Laboratories) and bovine serum albumin (BSA) as a standard.

Hepatic histology
After the liver samples were fixed in 10% formalin solution at room temperature, they were processed by the standard method. Briefly, tissues were embedded in paraffin, sectioned at 5 μm, and stained with hematoxylin and eosin (H&E), and then picked up on glass slides for light microscopy. All samples were evaluated by an experienced pathologist who was blinded to the experiment. A semiquantitative scoring system was used to assess the severity of hepatic steatosis and necroinflammation in 10 microscopic fields examined at 200× magnification as described previously [28]. Steatosis was scored as the percentage of parenchymal cells containing fat: 0 = no parenchymal cells containing fat, 1 = <20% of parenchymal cells containing fat, 2 = 20–39% of parenchymal cells containing fat, 3 = 40–50% of parenchymal cells containing fat, and 4 = >51% of parenchymal cells containing fat. Inflammation and necrosis were scored by the number of foci of inflammation and necrosis identified under a low-power field of a light microscope: 0 = no inflammation and necrosis, 1 = 1 focus of inflammation and necrosis per low-power field, 2 = 2 foci of inflammation and necrosis per low-power field, and 3 = 3 foci of inflammation and necrosis per low-power field.

Nuclear transcription factor-κB activity measurement
Nuclear extracts of livers from the above-mentioned groups were prepared using the nuclear extraction kit (Active Motif; Cat. No. 40010). In brief, the liver (0.5 g) was pulverized under liquid nitrogen. Powdered tissue was suspended in hypotonic buffer containing a protease inhibitor cocktail, phosphatase inhibitors, DTT, and detergent (Active Motif). Samples were dounce homogenized, incubated on ice for 15 min, and then centrifuged (850 g, 10 min, 4°C). The pellet was suspended in hypotonic buffer containing detergent and then centrifuged (14000 × g, 30 s, 4°C). The resulting pellet was suspended in lysis buffer and incubated on ice for 30 min, and then nuclear extracts were obtained by centrifugation (14000 × g, 10 min, 4°C). Twenty micrograms of nuclear extract were used for the determination of NF-κB activity with the TransAM® NF-κB p65 transcription factor assay kit (Active Motif; Cat. No. 40096), according to the manufacturer’s instruction.

Western blotting
Liver tissues were homogenized in 1 ml of ice-cold hypotonic buffer A [10 mmol/L HEPES (pH 7.8), 10 mmol/L KCl, 2 mmol/L MgCl2, 1 mmol/L DTT, 0.1 mmol/L EDTA, 0.1 mmol/L phenylmethylsulfonylfluoride]. The cells were then lysed with 12.5 μL 10% Nonidet P-40. The homogenate was centrifuged and the supernatant containing the cytoplasmic extracts was stored frozen at −80°C. The nuclear pellet was resuspended in 25 μL ice-cold nuclear extraction buffer. After 30 min of intermittent mixing, the extract was centrifuged, and the supernatants containing nuclear extracts were secured. Before immunoblotting, the protein concentration of each tissue was determined using a Bio-Rad protein assay kit (Bio-Rad Labo-
and 5 min, were blocked with 5% nonfat dry milk in Tris-buffered saline Tween (20 mmol/L Tris, pH 7.6, 137 mmol/L NaCl, and 0.1% Tween 20) for 3 h at room temperature, and incubated overnight at 4°C with the following primary antibodies: p-NF-κB p65 (Ser276) (Santa Cruz Biotechnology, Inc.; Cat. No. sc-101749) and NF-κB p65 (Cell Signaling Technology; Cat. No. 4764). The level of lamin A (Santa Cruz Biotechnology, Inc.; Cat. No. sc-20680) was estimated for equal loading of the nuclear sample. After washing three times with Tris-buffered saline Tween 20 (TBST), incubation with appropriate horseradish peroxidase-conjugated secondary antibodies was performed for 1 h at room temperature. After three additional TBST washes, the immunoreactive bands were visualized by enhanced chemiluminescence (Amersham Biosciences), according to the manufacturer’s instructions. Band densities were determined using ATTO Densitograph Software (ATTO Corporation).

Statistical analysis

Data are expressed as the mean ± standard deviation (S.D.). Statistical analysis was performed with one-way analysis of variance (ANOVA). Dunnett range post hoc comparisons were used to determine the source of significant differences, where appropriate. For the histological study, a nonparametric Kruskal-Wallis test was performed and Mann-Whitney’s U test was used to compare data within the groups. The SigmaPlot (Version 11.0) program was used for statistical analysis. A p value < 0.05 was considered statistically significant.

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Conflict of Interest

No competing financial interests exist.

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Actin mRNA levels and calculated according to the delta-delta Ct method [29].

Analysis of mRNA expression of hepatic genes

Total RNA was extracted from 100 mg frozen liver samples using Trizol reagent (Invitrogen). RNA was quantitated by A260 and its integrity verified by agarose gel electrophoresis using ethidium bromide for visualization. For the reverse transcriptase reaction, 1 µg of total RNA per sample and 8.5 µg/µL random hexamer primers were heated at 65°C for 5 min and then quenched on ice. This mixture was combined with 500 µmol/L each of dATP, dTTP, dCTP, and dGTP, 10 mmol/L DTT, 20 mmol/L Tris-HCl (pH 8.4), 50 mmol/L KCl, 5 mmol/L MgCl2, 40 units of RNaseOUT™ recombinant ribonuclease inhibitor (Invitrogen), and 100 units SuperScript III reverse transcriptase (Invitrogen). Samples were subjected to DNase (Promega) treatment at 37°C for 20 min in a GeneAmp 9700 Thermal Cycler (Applied Biosystems) and then held at 4°C. After the aliquots were taken for immediate use in PCR, the remainder of the cDNA was stored at −20°C. mRNA expression was measured by calculating real-time reverse transcription polymerase chain reaction (RT-PCR) in a fluorescent temperature Lightcycler 480 (Roche Diagnostics). The following primer sequences were used: 5′-TCAACAACAAAGACAGTACT-GTCCCTGGCC-3′ (forward) and 5′-GTTCTCCTGGTGGACTTGCT-GTGGTGTTG-3′ (reverse) for SREBP-1c; 5′-CTCTTGGAATCAGTACAT-GTGGT-3′ (forward) and 5′-GCTCTCTGAAAAAGAGAAAG-3′ (reverse) for FAS; 5′-CTCTGAAAACCCCGACAGTAC-3′ (forward) and 5′-GTTGACAGTACATCTCCATGG-3′ (reverse) for ACC; 5′-CGCTCTGGCCTTCTAATAAGTAG-3′ (forward) and 5′-CCTGTA-GATCTCTCTGAGTAGCC-3′ (reverse) for PPAR γ; 5′-ATGGGAGAGCTCCTACCAAGCCG-3′ (forward) and 5′-CTCTCGTGGATCTCCAAAGCC-3′ (reverse) for ACO; 5′-TGTGATGGTGGAAATGGTCA-3′ (forward), and 5′-TGTGATGGTGGAAATGGTCA-3′ (reverse) for β-actin. Primers were designed using Primer Express Software version 2.0 System (Applied Biosystems). The PCR reaction was performed using the following cycling protocol: 95°C for 5 min, followed by 45 cycles of 95°C for 5 s, 58°C for 15 s, and 72°C for 20 s. Dissociation curves were run after amplification to identify the specific PCR products. The mRNA expression levels were normalized to β-
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