

Metabolomics of Ginger Essential Oil against Alcoholic Fatty Liver in Mice

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Supporting Information

ABSTRACT: Fatty liver is significantly associated with hepatic cirrhosis and liver cancer. Excessive alcohol consumption causes alcoholic fatty liver disease (AFLD). Ginger has been reported to exhibit antioxidant potential and hepatoprotective activity. In the present study, a mouse model for AFLD was developed by employing male C57BL/6 mice that were fed an alcohol-containing liquid diet (Lieber–DeCarli diet) ad libitum. In the treatment groups, ginger essential oil (GEO) and citral were orally administered every day for 4 weeks. Serum biochemical analysis, antioxidant enzyme activity analysis, and histopathological evaluation revealed that GEO and citral exhibited hepatoprotective activity against AFLD. Metabolites in serum samples were profiled by high-performance liquid chromatography coupled with quadrupole time-of-flight mass spectrometry (HPLC-QTOF-MS). Metabolomic data indicated the amounts of metabolites such as D-glucurono-6,3-lactone, glycerol-3-phosphate, pyruvic acid, lithocholic acid, 2-pyrocatechuic acid, and prostaglandin E1 were increased after alcohol administration, but the levels were recovered in treatment groups. The analysis indicated that ginger possesses hepatoprotective properties against AFLD. Furthermore, these metabolites can serve as early noninvasive candidate biomarkers in the clinical application of AFLD for health management.

KEYWORDS: AFLD, ginger, Lieber–DeCarli diet, metabolomics, biomarker

INTRODUCTION

The liver is the largest organ in the body, and it plays important roles in metabolism. Liver diseases are mainly caused by viruses, excessive alcohol consumption, and chemicals. Repeated damage-induced inflammation causes hepatitis, hepatic fibrosis, cirrhosis, and even death due to hepatic failure.^{1–3}

Fatty liver, a chronic liver disease, is a common global problem. Fatty liver can result in hepatitis, hepatic fibrosis, hepatic cirrhosis, liver cancer, or cardiovascular diseases. The pathogenesis of fatty liver is generally divided into nonalcoholic and alcoholic types. Excessive ethanol consumption can cause alcoholic fatty liver disease (AFLD).^{4,5} The mechanisms of AFLD are mediated by increased expression of cytochrome P450-2E1 (CYP2E1),⁶ modulation of inflammatory cytokines TNF- α , IL-1, and IL-6,⁷ and decreased hepatocyte retinoid x receptor levels.^{8,9}

To analyze the relationship among alcohol, lipid, and therapeutic agents, the Lieber–DeCarli liquid diet model is frequently used to induce AFLD in animals, especially in the

early stages of AFLD, including liver injury, steatosis, and oxidative stress.^{10,11}

Ginger (*Zingiber officinale* R.) and its extracts have been used widely as Chinese medicine for a long time; it has various physiological functions such as decreasing nausea,¹² enhancing functions of the intestines and stomach,¹³ preventing cardiovascular diseases,¹⁴ promoting liver functions,¹⁵ and inhibiting cancer cell proliferation.¹⁶ Among a variety of applications, ginger essential oil (GEO) is commonly used in the food, cosmetics, and beverage industries. Notably, citral, the main constituent of GEO, has been demonstrated to exhibit antioxidant, anticancer, and anticandida effects.^{17–19}

Metabolomics studies analyze the metabolome, which are the metabolites synthesized by a biological system and end-

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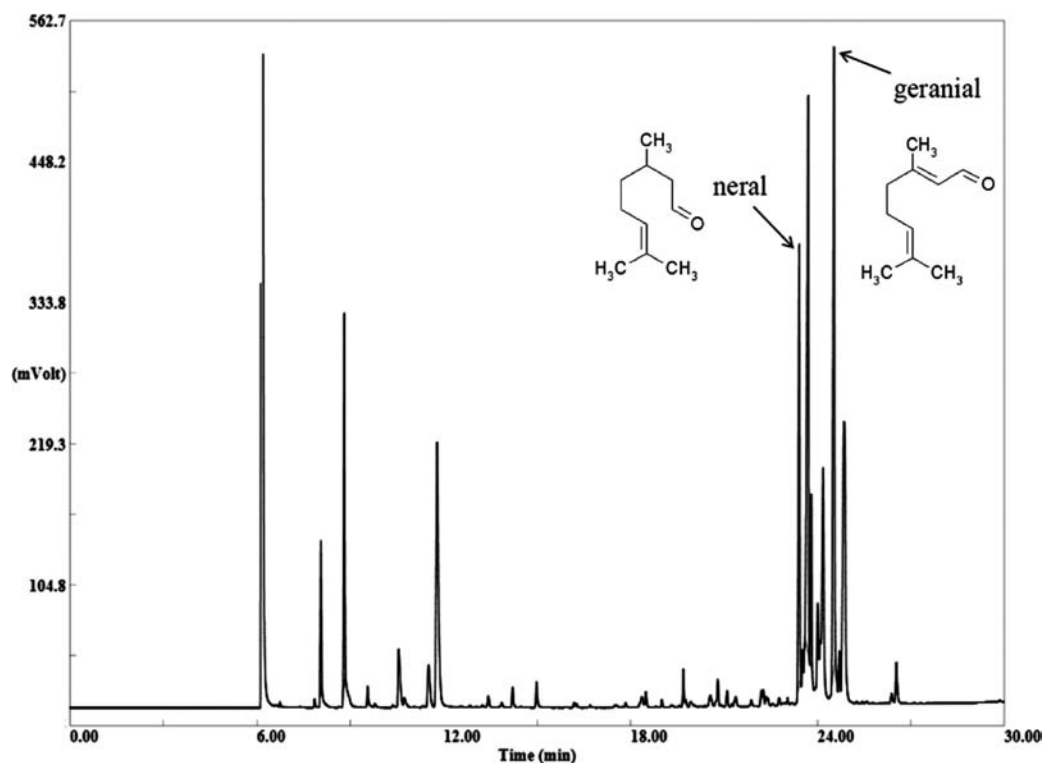


Figure 1. Volatile components of ginger essential oil (GEO).

products of gene expression in a sample.^{20,21} In metabolomics, blood, urine, and tissues are evaluated to generate individual metabolite profiles such as cholesterol and triglyceride profiles. In this study, we investigated the preventive effects of ginger on the formation of alcoholic fatty liver by using a metabolomics approach.

Here, we evaluated the hepatoprotective efficacy for GEO and citral by using the Lieber–DeCarli diet-induced AFLD model. Moreover, we adopted metabolomics approaches to identify differentially regulated metabolites in healthy mice and mice with AFLD. These metabolites may serve as candidate biomarkers for noninvasive detection during the early stages of AFLD, and this approach could be useful for humans.

MATERIALS AND METHODS

Chemicals. Solvents used in high-performance liquid chromatography coupled with quadrupole time-of-flight mass spectrometry (HPLC-QTOF-MS) were of MS grade. Water and acetonitrile were obtained from Scharlau (Sentmenat, Spain) and J. T. Baker (Phillipsburg, NJ, USA), respectively. Reagents were of analytical grade and were obtained from Sigma-Aldrich (Steinheim, Germany). Hexane, chloroform, and dichloromethane were procured from Merck (Darmstadt, Germany).

Preparation of Ginger Essential Oil Extracts and Citral. GEO was prepared by the steam distillation method.²² Briefly, 1 kg of ginger (*Z. officinale* R.) was added to 3 L of distilled water and distilled to extract the essential oil. The constituents of GEO were analyzed by a Thermo Scientific Focus GC equipped with an AI 3000 II autosampler, a flame ionization detector, and a Stabilwax (crossbond Carbowax-PEG, Restek) column (60 m × 0.32 mm; 1.0 μm). Standard citral (purity > 95%) was used. The composition of GEO analyzed by gas chromatography (GC) and the representative profile (Figure 1) indicates citral (neral and geranial) as the major component in GEO (approximately 30%). The composition of GEO was similar to previous research,²³ which employed the same species of ginger used

in the present study. Citral that was used for the animal experiments was obtained from Sigma-Aldrich.

In Vivo Animal Model of Alcoholic Fatty Liver Disease. Six-week-old male C57BL/6 (B6) mice (BioLASCO Taiwan Co.) were maintained at the animal housing facility of the Institute of Food Science and Technology, National Taiwan University, at a temperature of 23 ± 2 °C and relative humidity of 50–70% with a 12 h light/dark cycle. Mice were divided into a normal control group, an AFLD group, and four treatment groups. Mice in the normal control group were fed a normal liquid diet, whereas those in the AFLD and treatment groups were fed ethanol-containing Lieber–DeCarli diet. Each treatment group received either GEO [2.5 or 12.5 mg/kg body weight (BW)] or citral (0.375 or 1.875 mg/kg BW), both of which were mixed with olive oil (vehicle) and administered orally every day for 4 weeks. Normal and AFLD groups were given the same volume of only olive oil (vehicle). The BW of each mouse was recorded every week during the entire period of the experiment. Mice were sacrificed after 4 weeks; serum samples were then collected, and livers were sampled. Mice were handled according to the guidelines issued by the National Taiwan University Institutional Animal Care and Use Committee (Approval No. NTU-IACUC-99-53).

Serum Biochemical and Enzyme Analysis. Serum was extracted by centrifuging coagulated blood at 1000g for 5 min at 4 °C. The biochemical parameters of liver function indicators such as aspartate aminotransferase (AST), alanine aminotransferase (ALT), triglyceride (TG), and total cholesterol (TC) were estimated with commercial test strips (Commercial ALT and AST Spotchem II reagent strips, Arkray Inc., Kyoto, Japan) in an automatic blood analyzer (Spotchem EZ).

Liver Enzyme Activities. Liver homogenate was prepared by homogenizing liver (0.5 g) in 10 volumes of ice-cold homogenization buffer (8 mM KH₂PO₄, 12 mM K₂HPO₄, and 1.5% KCl, pH 7.4) at 4 °C. The homogenate was then centrifuged at 10000g for 30 min at 4 °C. The supernatant was stored at –80 °C and used to detect liver TG and hepatic antioxidant enzyme activity. Protein content in the homogenate was estimated spectrophotometrically by measuring the absorbance at 595 nm using the Bio-Rad protein Assay Kit (Hercules, CA, USA). Commercial kits purchased from the Cayman Chemical Co. (Ann Arbor, MI, USA) were used to examine the hepatic

Table 1. Effect of Samples on Serum Biochemical and Liver Biochemical Parameters in Mice with AFLD^a

measurement	control diet	alcohol diet	alcohol diet + (2.5 mg/kg BW) GEO	alcohol diet + (12.5 mg/kg BW) GEO	alcohol diet + (0.375 mg/kg BW) citra	alcohol diet + (1.875 mg/kg BW) citra
Serum						
AST (IU/L)	138 ± 34b	185 ± 33a	128 ± 22b	115 ± 42b	147 ± 52b	120 ± 37b
ALT (IU/L)	23 ± 4c	65 ± 8a	39 ± 9b	27 ± 5c	46 ± 18b	42 ± 16b
cholesterol (mg/dL)	141 ± 15b	175 ± 19a	122 ± 14c	114 ± 12cd	111 ± 31cd	97 ± 14d
triglyceride (mg/dL)	99 ± 19b	134 ± 16a	79 ± 25bc	57 ± 22c	92 ± 58b	76 ± 20bc
Liver						
triglyceride (mg/g liver)	39 ± 19c	91 ± 17a	63 ± 20b	39 ± 21c	69 ± 17b	65 ± 28b
GSH (nmol/mg protein)	38 ± 3b	29 ± 2c	40 ± 3b	49 ± 11a	38 ± 11b	39 ± 4b
GPx (nmol/min/mg protein)	737 ± 63ab	582 ± 51c	710 ± 90b	812 ± 96a	670 ± 86b	682 ± 42b
GRd (nmol/min/mg protein)	34 ± 5ab	23 ± 6d	29 ± 6bc	35 ± 4a	24 ± 3cd	29 ± 3b
CAT (nmol/min/mg protein)	326 ± 47b	253 ± 18c	329 ± 35b	395 ± 33a	295 ± 26b	333 ± 55b
SOD (U/mg protein)	52 ± 4b	42 ± 5c	52 ± 4b	60 ± 6a	50 ± 6b	51 ± 6b

^aResults are expressed as the mean ± SD ($n = 8$). Statistical analysis was performed among treatments within each group. Different letters (a–d) indicate a significant difference among groups according to the one-way ANOVA coupled with the Duncan's multiple-comparison test ($p < 0.05$). AFLD, alcoholic fatty liver disease; GEO, ginger essential oil; AST, aspartate aminotransferase; ALT, alanine aminotransferase; GSH, glutathione; GPx, glutathione peroxidase; GRd, glutathione reductase; CAT, catalase; SOD, superoxide dismutase.

antioxidant systems of glutathione (GSH; item 703002), glutathione peroxidase (GPx; item 703102), glutathione reductase (GRd; item 703202), catalase (CAT; item 707002), and superoxide dismutase (SOD; item 706002).

Liver Biopsy Examination. Liver histological sections for pathological staining and semiquantitative analysis were performed from the right lobe of the liver to avoid observational bias. For histopathological observations, formalin-fixed paraffin-embedded (FFPE) liver sections were observed for liver fatty accumulation, necrosis, fibrosis, and other changes in liver tissues, after staining with hematoxylin and eosin (H&E). Scoring and interpretation of biopsy results were carried out by an experienced pathologist, College of Veterinary Medicine Animal Disease Diagnostic Center, National Chung Hsing University, Taichung, Taiwan.

Metabolic Profiling. Sample Preparation. Aliquots of 100 μ L of serum were added to extraction solvents (1:4) and extracted by the Geno Grinder 2010 (SPEX, Pittsburg, CA, USA) at 1000 rpm for 2 min. Extracts were centrifuged at 15000g for 5 min at 4 °C. The supernatant was collected and evaporated by a centrifugal vaporizer (EYELA, Tokyo, Japan) for 2 h. The residue was reconstituted in 200 μ L of 50% methanol and centrifuged at 15000g for 5 min. The supernatant was filtered with a 0.2 μ m Minisart RC 4 filter (Sartorius, Goettingen, Germany) and subjected to HPLC-QTOF-MS analysis. The same amounts of sample aliquots were pooled for the QC sample.

Chromatographic and Mass Spectrometric Analysis. All serum samples were analyzed by an Agilent 1290 U-HPLC system coupled with 6540-QTOF (Agilent Technologies, Santa Clara, CA, USA). The sample (2 μ L) was injected into an ACQUITY UPLC HSS T3 column (2.1 \times 100 mm; 1.8 μ m) (Waters, Milford, MA, USA). The mobile phase consisted of 0.1% formic acid (solvent A) and acetonitrile (solvent B). The gradient profile used was as follows: 0–1.5 min, 2% B; 1.5–9 min, linear gradient from 2 to 50% B; 9–14 min, linear gradient from 50% to 95% B; 14–15 min, 95% B; the column was then re-equilibrated. The flow rate was maintained at 0.3 mL/min. The column oven was set at 40 °C, and the autoinjection system was set at 10 °C. A Jet Stream electrospray ion source with capillary voltage of 4 kV in positive mode and 3.5 kV in negative mode was used for sample ionization. MS parameters were set as follows: dry gas temperature, 325 °C; dry gas flow, 5 L/min; nebulizer, 40 psi; sheath gas temperature, 325 °C; sheath gas flow, 10 L/min; and fragmentor, 120 V. The scan range was m/z 50–1700.

Data Analysis. The Kyoto Encyclopedia of Genes and Genomes (KEGG) database was used to identify metabolite biomarkers by comparing molecular weights. The molecular structures of candidate compounds were retrieved by comparison and then confirmed by MS scans for characteristic ions and fragmentation patterns of metabolites.

Three-dimensional principal component analysis (3D-PCA) was performed, and the values of PC1, PC2, and PC3 were plotted with designation of cases as control, AFLD, or treatment groups.

Statistical Analysis. Statistical differences between groups were compared by one-way analysis of ANOVA software with Duncan's multiple-comparison test by SPSS software (Statistical Package for Social Sciences, SPSS Corporation, Chicago, IL, USA). All values are expressed as the mean ± SD.

RESULTS AND DISCUSSION

In Vivo Model for Alcohol-Induced Fatty Liver.

Establishing an ideal rodent model for the study of alcoholic fatty liver is a prerequisite for this experiment. Because metabolic rates considerably differ among species, optimization and validation of the model are necessary. In our analysis, C57BL/6 mice were employed to develop a model for alcoholic fatty liver using the Leiber–DeCarli liquid diet. The ability to induce visible phenotypic symptoms of alcohol-induced liver disease such as steatosis, inflammation, and necrosis is a benchmark for the model's success. Many studies have failed to induce hepatitis coupled with fatty liver.^{24,25} However, consumption of alcohol-containing Leiber–DeCarli liquid diet for 28 days has been shown to result in alcoholic fatty liver, inflammation, and necrosis in C57BL/6 mice.²⁶

Hepatoprotective Activity of Ginger on AFLD. Here, we induced similar effects in C57BL/6 mice after feeding for 28 days. The AFLD group recorded a significant ($p < 0.05$) increase in the serum AST and ALT activity levels after 28 days of feeding, whereas mice fed control Leiber–DeCarli liquid diet (normal control group) exhibited no liver damage. The AFLD group exhibited a significant ($p < 0.05$) increase in TC and TG levels when compared to the normal control group. Moreover, AST, ALT, TC, and TG levels were recovered significantly in the treatment groups compared to the AFLD group ($p < 0.05$), especially in mice gavaged with high doses of GEO (Table 1). GSH is an important indicator of the liver antioxidant system. GSH levels were significantly decreased in the AFLD group compared to the normal control group. Similarly, liver antioxidant enzyme activities, namely, GPx, GRd, catalase, and SOD, were lower in the AFLD group than in the control group. However, a significant increase was observed in the antioxidant activity of these enzymes in the treatment groups (p

Table 2. Histopathological Analysis for the Fatty Liver^a

score	control diet	alcohol diet	alcohol diet + (2.5 mg/kg BW) GEO	alcohol diet + (12.5 mg/kg BW) GEO	alcohol diet + (0.375 mg/kg BW) citral	alcohol diet + (1.875 mg/kg BW) citral
fatty liver	1.0 ± 0.5c	3.5 ± 0.5a	2.0 ± 0.8b	1.3 ± 0.7c	2.8 ± 1.0b	2.3 ± 0.7b

^aResults are expressed as the mean ± SD ($n = 8$). Statistical analysis was performed among treatments within each group. Different letters (a–c) indicate a significant difference among groups according to the one-way ANOVA coupled with the Duncan's multiple-comparison test ($p < 0.05$). GEO, ginger essential oil. Degree of lesions was graded from 1 to 5 depending on severity: 1 = minimal (<1%); 2 = slight (1–25%); 3 = moderate (26–50%); 4 = moderate/severe (51–75%); 5 = severe/high (76–100%).

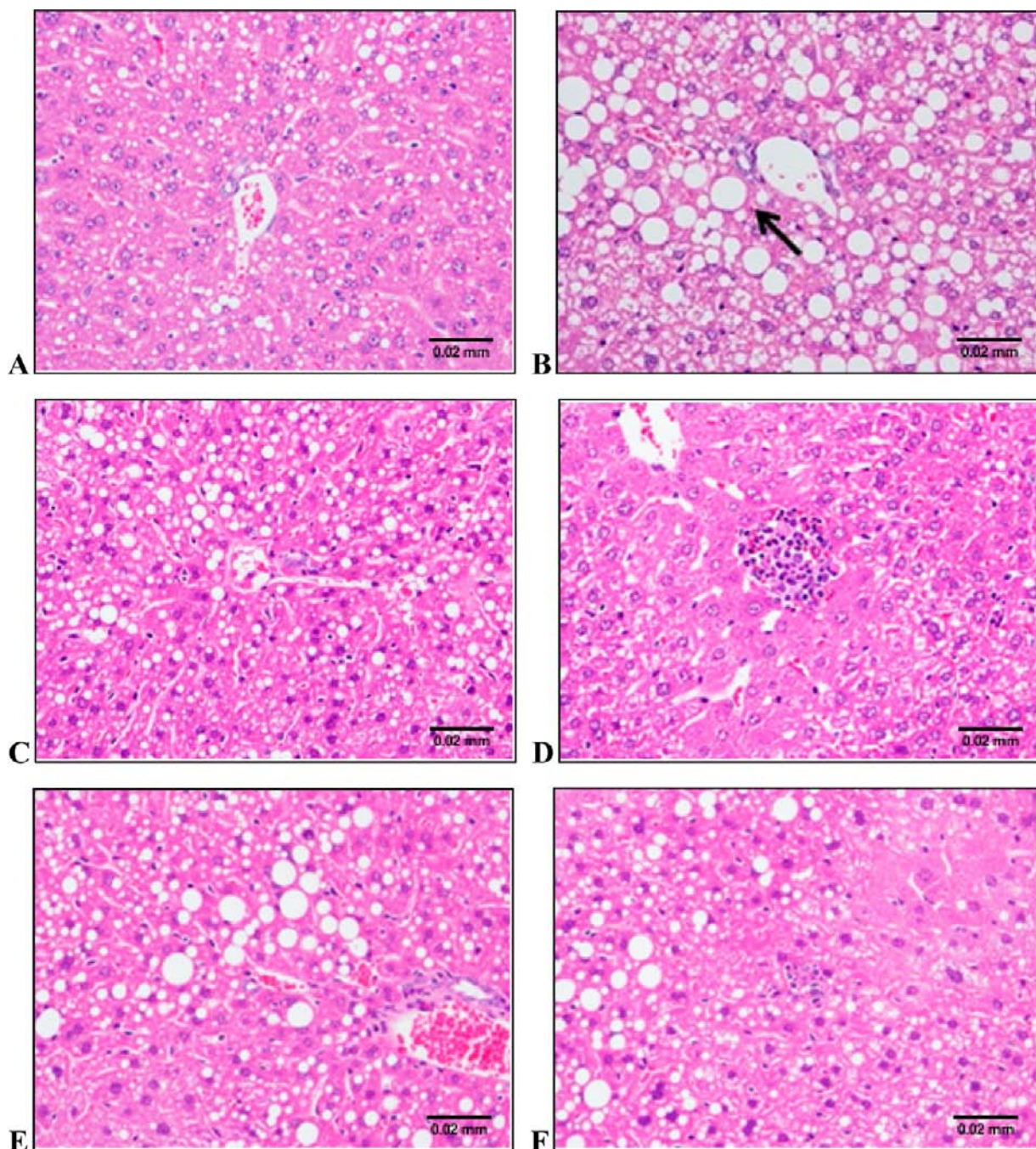


Figure 2. Representative histological features of liver section stained with H&E (400×) depicting the effect of samples on liver histopathology of mice with AFLD. Control liquid diet induced minimal (1) fatty infiltration with macrovesicles of hepatocytes in the normal control group (A). Alcohol liquid diet-induced fatty liver with macrovesicles (indicated by arrowhead) of hepatocytes was graded as moderate in AFLD group (B); minimal (1) in high dose of GEO (12.5 mg/kg BW) group (D), and slight (2) in low dose of GEO (2.5 mg/kg BW) (C), low dose of citral (0.375 mg/kg BW) (E), and high dose of citral (1.875 mg/kg BW) groups (F).

Table 3. Metabolomics Profiles in Serum Samples in Mice^a

metabolite name	m/z	mean intensity										Duncan's test					significance level
		A	B	C	D	E	F	A	B	C	D	E	F				
1-methylhistamine 3-methylhistamine	126.102	10659	3752	3469	3795	4529	4099	a	b	b	b	b	b	b	b	b	****
D-glucurono-6,3-lactone ascorbic acid	175.024	38727	93648	63269	37745	55137	39898	c	a	b	c	b	c	b	c	c	****
glycerol-3-phosphate	171.006	2332	2911	2363	1829	2172	1657	b	a	b	cd	bc	d	bc	d	d	****
pyruvic acid	87.0088	14133	30848	24365	13062	19996	14509	c	a	ab	c	bc	c	bc	c	c	****
ascorbic acid	175.024	102	341	177	178	1201	212	b	b	b	b	a	a	b	a	b	****
galacturonic acid glucuronate	193.035	1798	5399	5262	4256	4276	4760	b	a	a	a	a	a	a	a	a	****
FAD	784.149	727	611	409	599	699	671	a	a	b	a	a	a	a	a	a	****
2-hydroxy-2-methylbutyric acid 3-hydroxyisovaleric acid 2-hydroxy-3-methylbutyric acid	117.055	11387	8479	7396	8008	8577	9406	a	bc	c	bc	bc	bc	bc	b	b	****
trans-aconitic acid cis-aconitate dehydroascorbic acid	173.009	12583	9869	8788	7926	10347	9151	a	bc	bc	c	b	b	bc	bc	bc	****
2-methylmaleate ribonolactone	147.029	14290	13723	11837	10310	12902	11011	a	ab	bcd	d	abc	cd	abc	cd	cd	****
phenylacetyl-glycine	194.081	5219	3262	2727	2264	4050	2693	a	bc	bc	c	ab	bc	bc	bc	bc	****
methylhippuric acid	192.066	54283	32177	29028	24111	43614	28490	a	bc	bc	c	ab	bc	bc	bc	bc	****
citric acid D-threo-isocitric acid	191.019	883175	668160	607577	559617	648578	658819	a	b	b	b	b	b	b	b	b	****
traumatic acid	227.128	18536	9155	4570	4088	7934	6697	a	b	b	b	b	b	b	b	b	****
sebacic acid	201.113	86573	35405	17468	15409	25673	24122	a	b	b	b	b	b	b	b	b	****
lithocholic acid	375.290	1492	2736	1773	1227	1237	1402	b	a	b	b	b	b	b	b	b	****
uridine	243.062	68254	73599	70307	74810	85627	88374	c	c	c	bc	ab	a	a	a	a	****
1-methyladenosine	282.119	5665	2136	1983	3476	1143	1763	a	b	b	ab	b	b	b	b	b	****
N-formyl-L-methionine	176.038	3447	2280	2141	2517	2881	2953	a	bc	c	bc	abc	ab	ab	ab	ab	****
2-pyrocatechuic acid	153.019	2065	54810	20888	29058	55998	29297	b	a	b	ab	a	a	ab	ab	ab	****
o-hydroxyphenylacetic acid mandelic acid	151.040	2719	321	697	606	513	1096	a	b	b	b	b	b	b	b	b	****
dihydrobiopterin	240.109	905	427	424	694	581	715	a	b	b	ab	b	b	b	b	b	****
N-acetyl-leucine	172.097	179066	112453	100609	104616	107266	121865	a	b	b	b	b	b	b	b	b	****
3-hydroxymethylglutaric acid	161.045	25685	16836	15197	13152	20819	18660	a	bc	bc	c	ab	bc	bc	bc	bc	****
prostaglandin E1	353.233	5199	8116	7485	5888	5832	6591	c	a	ab	bc	bc	bc	bc	abc	abc	****
uracil	113.034	3387	3609	3055	3616	4030	4267	bc	abc	c	abc	ab	a	a	a	a	****
methionine	150.058	229039	287861	211902	197358	230186	226654	b	a	b	b	b	b	b	b	b	****
methylmalonate succinate	117.019	398174	450430	399451	449559	488715	517680	b	ab	b	ab	a	a	a	a	a	****
1-methylhistidine 3-methylhistidine	170.092	1236	297	607	574	164	1424	ab	c	bc	bc	c	a	a	a	a	****
cortisolone corticosterone	347.221	692	1123	2741	1738	3356	2863	c	bc	ab	abc	a	a	a	a	a	****
L-citrulline	176.103	142	1544	1559	604	1683	1308	c	ab	ab	bc	a	ab	ab	ab	ab	****
phenylglyoxylic acid	149.024	1415	652	440	466	517	506	a	b	b	b	b	b	b	b	b	**
spermine	203.223	702	406	672	875	508	827	ab	b	ab	a	b	a	b	a	b	**
glycine	76.0393	6948	5479	5699	4682	4254	5755	a	ab	ab	b	b	b	ab	ab	ab	**
S-methylthioadenosine	298.096	3966	3312	2829	2940	3489	3658	a	abc	c	bc	abc	ab	ab	ab	ab	**
deoxycholic acid	391.285	6537	14721	26156	40787	39183	29743	c	bc	abc	a	ab	abc	ab	abc	abc	**
L-acetylcarbitine	204.123	1128499	1167555	1118265	781735	977344	1043685	a	a	a	b	ab	ab	ab	ab	ab	**
N6-acetyl-L-lysine	189.123	2484	2394	1353	1174	1938	1932	a	a	b	b	ab	ab	ab	ab	ab	**
L-alloisoleucine L-norleucine L-isoleucine	132.101	1077930	1392815	1211970	1166625	1267987	1302868	c	a	abc	bc	abc	ab	ab	ab	ab	**
pipecolic acid	130.086	27139	24107	19822	17008	22216	21653	a	ab	bc	c	abc	abc	abc	abc	abc	**

Table 3. continued

metabolite name	m/z	mean intensity										significance level				
		A	B	C	D	E	F	A	B	C	D		E	F		
l-gulonol-1,4-lactone	177.040	584	2413	1555	721	1506	1113	b	a	ab	b	ab	b	ab	b	**
deoxycholic acid hyodeoxycholic acid P6-24	391.285	321	1082	2082	3721	3330	3492	b	ab	ab	a	a	a	a	a	**
hydroxyoctanoic acid	159.102	21629	20514	18400	13306	11885	16101	a	ab	abc	bc	c	c	abc	abc	**
D-glutamine	147.076	5962	8954	6705	13895	11544	13970	b	ab	b	a	ab	a	a	a	**
acetylglucine	118.049	5032	4144	4446	3435	3024	4160	a	ab	ab	b	b	b	ab	ab	**
N-methylhydantoin dihydrouracil	115.050	1522	959	138	1091	896	937	a	ab	b	a	ab	ab	ab	ab	**
2-hydroxycaproic acid 2-ethyl-2-hydroxybutyric acid	131.071	229125	199443	191813	161109	156370	195000	a	ab	ab	b	b	b	ab	ab	*
pyrrolidonecarboxylic acid	128.035	110399	154555	146918	156875	151455	142229	b	a	a	a	a	a	ab	ab	*
hydroxysocaproic acid 2-hydroxycaproic acid 2-ethyl-2-hydroxybutyric acid	131.071	11210	7748	6688	4714	7428	8249	a	ab	ab	b	ab	ab	ab	ab	*
lactate	89.0244	114149	134677	129118	116490	115226	119702	b	a	ab	b	b	b	ab	ab	*
pyroglutamic acid	130.049	5740	7782	7506	10878	11913	9759	b	ab	ab	a	a	a	ab	ab	*
glutaric acid ethylmalonic acid methylsuccinic acid dimethylmalonic acid	131.035	723	678	312	279	3042	1111	b	b	b	b	a	a	b	b	*
levulinic acid	115.040	18741	16205	14669	17107	13575	18391	a	ab	ab	ab	b	b	a	a	*
N-methylhydantoin dihydrouracil	115.050	908	718	276	646	708	567	a	ab	b	ab	ab	ab	ab	ab	*
deoxyribose	133.050	955	1430	856	447	1367	1125	ab	a	ab	b	a	a	ab	ab	*
glutaconic acid	129.019	13983	13064	13013	9791	13438	11325	a	ab	ab	b	a	a	ab	ab	*

*Statistical analysis was carried out among the treatment within each groups. Different letters (a-d) indicate a significant difference among groups according to the one-way ANOVA coupled with Duncan's multiple-comparison test (*****, $p < 0.001$; ***, $p < 0.005$; **, $p < 0.01$; *, $p < 0.05$; *, $p < 0.1$). A, normal control group; B, AFLD group; C, 2.5 mg/kg BW GEO group; D, 12.5 mg/kg BW GEO group; E, 0.375 mg/kg BW citral group; F, 1.875 mg/kg BW citral group.

< 0.05; Table 1) compared to the AFLD group. It has been reported that ginger ameliorates fatty liver and hypertriglyceridemia in rats.²⁷ In addition, results of other studies have confirmed that ginger exerts hepatoprotective properties against liver damage by bromobenzene, acetaminophen, carbon tetrachloride, and ethanol.^{28,29} In the present study, the hepatoprotective activity of ginger on AFLD was found to be similar to that reported previously. Histopathological observation of liver sections from the control group exhibited a normal liver architecture, whereas fatty infiltration with macrovesicles was observed in the livers of the AFLD group. However, compared to the AFLD group, there was a significant improvement in treatment groups gavaged with a high dosage of GEO or citral (Table 2; Figure 2). Recently, more research has focused on herbal remedies for alcoholic liver diseases.³⁰ Garlic and ginger are familiar, traditional Chinese herbs; it has been reported that garlic exhibits liver protection properties in alcohol-induced fatty liver.³¹ Our study results show that ginger provided hepatoprotection to alcohol-induced fatty liver. Hence, ginger may be a potential hepatoprotective candidate against AFLD.

Metabolic Profiling of Serum Samples of AFLD.

Metabolomes of serum samples from control, AFLD, and treatment groups were profiled by U-HPLC/Q-TOF MS. The metabolites with a significant difference among groups are listed in Table 3. We investigated the overall metabolic differences by using the 3D-PCA method. 3D-PCA was used as an unsupervised statistical method to study the metabolome differences between AFLD, treatment mice, and healthy controls. The score plots of 3D-PCA are displayed in Figure 3. Results revealed that PC1 accounted for 54% of variations in the 3D-PCA model, PC2 for 22%, and PC3 for 15%; these extracts clearly enabled differentiation. These separation results indicate the potential diagnostic value of metabolic profiles that reflect biochemical differences among the control, AFLD, and treatment groups. A one-way ANOVA followed by Duncan's

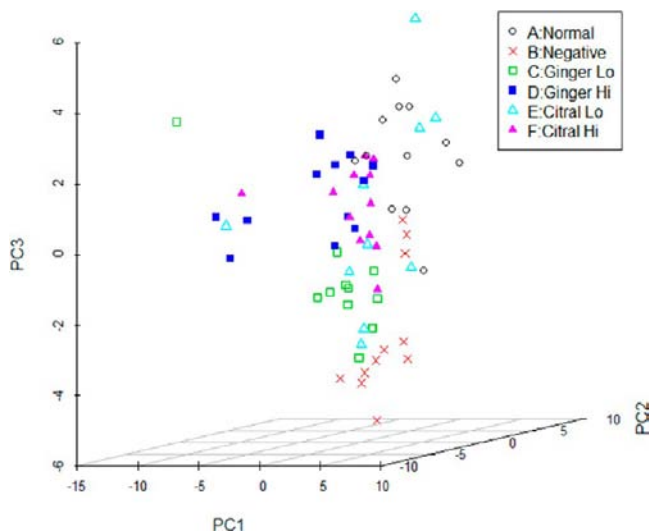


Figure 3. Three-dimensional principal component analysis (3D-PCA) of serum samples obtained from the control, AFLD, and treatment mice. PC1 is 54%, PC2 is 22%, PC3 is 15%. (Open black circles): A, normal control group; (red times signs) B, AFLD group; (open green squares) C, 2.5 mg/kg BW GEO group; (solid blue squares) D, 12.5 mg/kg BW GEO group; (open teal triangles) E, 0.375 mg/kg BW citral group; (solid pink triangles) F, 1.875 mg/kg BW citral group.

multiple-comparison test was used to analyze the statistically significant differences of metabolites among the control, AFLD, and treatment groups. Levels of D-glucurono-6,3-lactone, glycerol-3-phosphate, and pyruvic acid, which are important metabolites in many metabolic pathways, were significantly increased ($p < 0.001$) in the AFLD group compared to those in the normal control group (Figure 4). However, levels of these metabolites were significantly decreased ($p < 0.001$) in the treatment groups compared to the levels in the AFLD group. Similar changes in the levels of other metabolites involved in key metabolic pathways, including lithocholic acid ($p < 0.005$), 2-pyrocatechuic acid ($p < 0.005$), prostaglandin E1 ($p < 0.005$), methionine ($p < 0.01$), L-gulonolactone ($p < 0.05$), and lactate ($p < 0.1$; Figures 5 and 6), were observed upon treatment (Figure 4–6). The levels of these metabolites in the AFLD group were the highest among the six groups of mice, but these metabolites were down-regulated in the treatment groups. Our results suggest that these metabolites are related to AFLD. These metabolites can be used as potential candidate biomarkers for healthy management and clinical application after they are validated in humans.

After ingestion, alcohol is metabolized to acetaldehyde by ADH, the microsomal ethanol-oxidizing enzyme CYP2E1, and bacterial enzymes of intestinal microbiota. Acetaldehyde is then oxidized to acetate by ALDH and introduced into the citric acid cycle (Krebs cycle) as acetyl-CoA. Alcohol metabolism is known to significantly slow the TCA cycle when fatty acids are available in mice²⁶ and rats.³² We observed changes in the metabolite lactate, which is consistent with the data presented by Bradford et al. Higher levels of lactate correlate with hypoxia.³³ Chronic alcohol administration increases the rate of alcohol metabolism,³⁴ induces oxidative stress, the production of reactive oxygen species (ROS), cytokine release, mitochondrial dysfunction, and endoplasmic reticulum stress. Alcohol also stimulates liver metabolism and increases oxygen uptake, leading to hypoxia in the pericentral regions in liver lobules.³⁵ ROS initiate lipid peroxidation that directly damages plasma and intracellular membranes and leads to the production of reactive aldehydes with potent pro-inflammatory and profibrotic properties. Acute alcohol exposure results in increased ketogenesis and increased lipid peroxidation.³⁶ Chronic treatment of liver with alcohol induces liver inflammation through key arachidonic acid pathway metabolites including prostaglandins E1, prostacyclins, thromboxanes, and leukotrienes.

Liver injury also influences phenylalanine metabolism by affecting the following metabolites: succinate, pyruvate, and 2-hydroxyphenylacetate. An alcohol diet also affects arginine and proline metabolism. In addition, major amino acid metabolism pathways, including phenylalanine metabolism, glycine, serine, and threonine metabolism, arginine and proline metabolism, tryptophan metabolism, and valine, leucine, and isoleucine biosynthesis, as well as alanine, aspartate, and glutamate metabolism, are influenced by alcohol administration. All of these are inter-related with fatty acid, nucleotide, and carbohydrate metabolism. Phenylalanine, tyrosine, leucine, and tryptophan have been considered serum biomarkers for liver injury induced by alcohol.³⁷ In this study, we found that all pathways involving metabolites were influenced by alcohol administration. However, plasma lipid metabolic profiling can give a reasonable understanding of the pathogenesis of ethanol-mediated liver injury.³⁸ Furthermore, L-acetylcarnitine concentration was increased in the sera of humans with liver cirrhosis compared with that of healthy subjects.³⁹ Our results show that

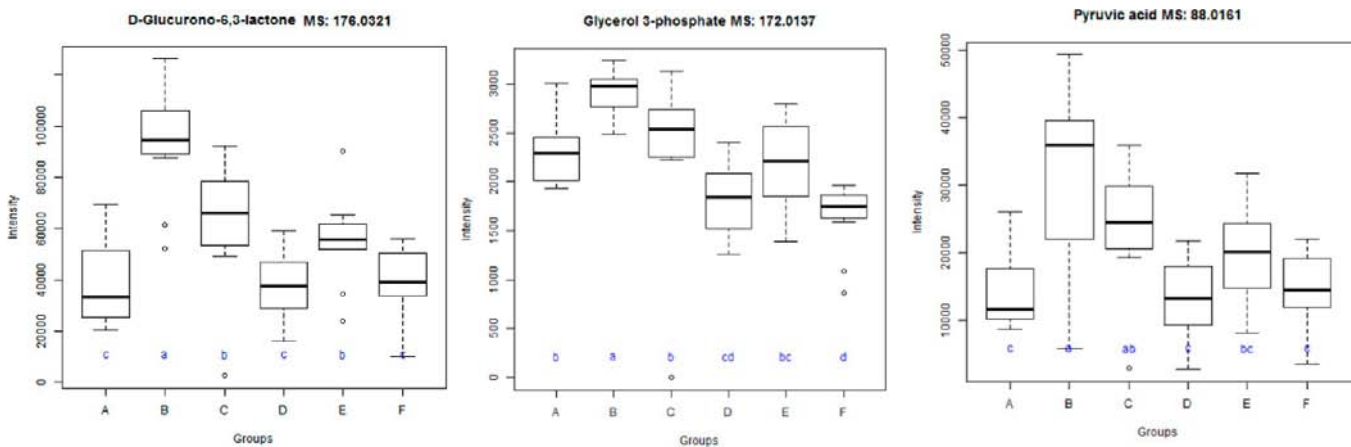


Figure 4. Metabolites of mice with AFLD that changed significantly among the six groups of mice (D-glucurono-6,3-lactone, glycerol-3-phosphate, and pyruvic acid; $p < 0.001$): A, normal control group; B, AFLD group; C, 2.5 mg/kg BW GEO group; D, 12.5 mg/kg BW GEO group; E, 0.375 mg/kg BW citral group; F, 1.875 mg/kg BW citral group. Statistical analysis was performed among treatments within each group. Different letters (a–d) indicate a significant difference among groups according to one-way ANOVA coupled with the Duncan's multiple-comparison test ($p < 0.001$).

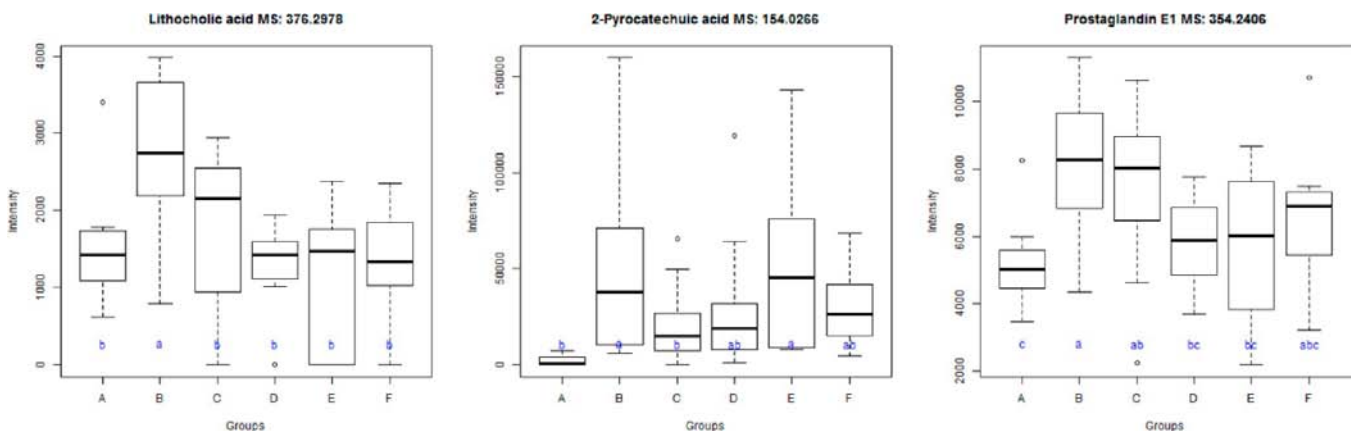


Figure 5. Metabolites of mice with AFLD that changed significantly among the six groups of mice (lithocholic acid, 2-pyrocatechic acid, and prostaglandin E1; $p < 0.005$): A, normal control group; B, AFLD group; C, 2.5 mg/kg BW GEO group; D, 12.5 mg/kg BW GEO group; E, 0.375 mg/kg BW citral group; F, 1.875 mg/kg BW citral group. Statistical analysis was performed among the treatment within each group. Different letters (a–c) indicate a significant difference among groups according to one-way ANOVA coupled with the Duncan's multiple-comparison test ($p < 0.005$).

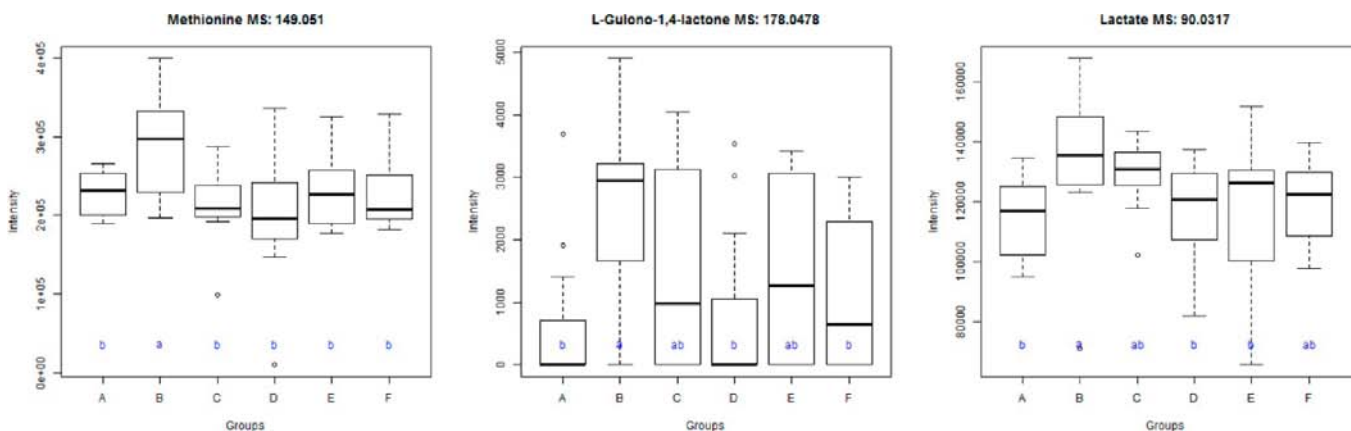


Figure 6. Metabolites of mice with AFLD that changed significantly among the six groups of mice (methionine, L-gulono-1,4-lactone, and lactate; $p < 0.01$, $p < 0.05$, $p < 0.1$, respectively): A, normal control group; B, AFLD group; C, 2.5 mg/kg BW GEO group; D, 12.5 mg/kg BW GEO group; E, 0.375 mg/kg BW citral group; F, 1.875 mg/kg BW citral group. Statistical analysis was performed among treatments within each group. Different letters (a, b) indicate a significant difference among groups according to one-way ANOVA coupled with the Duncan's multiple-comparison test.

the levels of fatty liver induced by consumption of an alcoholic liquid diet were not significant, which may have led to similar concentrations of L-acetylcarnitine in the control and AFLD groups.

Metabolomics results showed changes in the levels of D-glucurono-6,3-lactone and L-gulono-1,4-lactone, which are key metabolites in ascorbate and aldarate metabolism. Moreover, glucuronic acid exhibits hepatoprotective properties in liver detoxification systems, and it metabolizes to glucuronolactone in the liver. Lithocholic acid also plays an important role in bile secretion, which is involved in alcohol-induced liver damage.⁴⁰ In our research, alcohol consumption induced an increase in the NADH/NAD⁺ ratio. It also increased the activity of glycerophosphate dehydrogenase and promoted the conversion of dihydroxyacetone phosphate to glycerol-3-phosphate, which might enhance lipid synthesis in the liver. In our study, 2-pyrocatechuic acid was detected in the metabolites, which could serve as a biological marker for the detection and quantification of OH radicals.⁴¹ The levels of essential amino acids such as leucine, isoleucine, and methionine were elevated in the control, AFLD, and treatment groups; this finding was consistent with the result of a previous study,⁴² which suggests a higher rate of whole-body protein turnover. In addition, methionine is involved in many important pathways such as cysteine and methionine metabolism, protein digestion and absorption, aminoacyl-tRNA biosynthesis, and mineral absorption. Although hepatic steatosis could be observed with the noninvasive magnetic resonance imaging (MRI) technique, liver inflammation could not be detected using this technique.⁴³ Variations of these metabolites, which play very important roles in physiological metabolism in the body, could be used as noninvasive biomarkers of AFLD.

Conclusion. Ginger exhibited hepatoprotective properties to prevent AFLD. In particular, GEO played an important role in hepatoprotection. The metabolomic data obtained in this study provide new, significant mechanistic clues concerning the effects of an alcoholic liquid diet on liver metabolism, which may be crucial for understanding the pathogenesis of alcohol-induced liver disease. We propose that several metabolites could be used as novel noninvasive biomarkers of AFLD induced by alcohol consumption and oxidative stress. Furthermore, these metabolites observed in mice with AFLD need to be compared and validated in human samples to reveal their potential as biomarkers.

■ ASSOCIATED CONTENT

🔍 Supporting Information

Additional table. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Notes

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■ ABBREVIATIONS USED

AFLD, alcoholic fatty liver disease; GEO, ginger essential oil; HPLC-QTOF-MS, high-performance liquid chromatography coupled with quadrupole time-of-flight mass spectrometry; AST, aspartate aminotransferase; ALT, alanine aminotransferase; TG, triglyceride; TC, total cholesterol; GSH, glutathione; GPx, glutathione peroxidase; GRd, glutathione reductase; CAT, catalase; SOD, superoxide dismutase; ADH, alcohol dehydrogenase; ALDH, aldehyde dehydrogenase; ROS, reactive oxygen species; 3D-PCA, three-dimensional principal component analysis; MRI, magnetic resonance imaging

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