

Lomatogonium Rotatum for Treatment of Acute Liver Injury in Mice: A Metabolomics Study

Renhao Chen ¹, Qi Wang ², Lanjun Zhao ¹, Shinlin Yang ¹, Zhifeng Li ^{1,*}, Yulin Feng ², Jiaqing Chen ³, Choon Nam Ong ⁴ and Hui Zhang ^{5,*}

¹ National Pharmaceutical Engineering Center for Solid Preparation in Chinese Herb Medicine, Jiangxi University of Traditional Chinese Medicine, Nanchang 330002, China

² State Key Laboratory of Innovative Drug and Efficient Energy-Saving Pharmaceutical Equipment, Nanchang 330006, China

³ NUS Graduate School for Integrative Sciences and Engineering, National University of Singapore, 119077, Singapore

⁴ Saw Swee Hock School of Public Health, National University of Singapore, 117549, Singapore

⁵ NUS Environmental Research Institute, National University of Singapore, 117411, Singapore

* Correspondence: Hui Zhang, zhanghui@u.nus.edu; Zhifeng Li, lizhifeng1976@hotmail.com

Figure Captions:

Figure S1. Total ion chromatograms of QC samples and selected top 12 peaks, A: GC-MS, B: LC-MS (ESI+), C: LC-MS (ESI-).

Figure S2. Plots of PCA and OPLS-DA. A: PCA plot of GC-MS data (R^2X : 0.774; Q^2 : 0.603); B: OPLS-DA plot of GC-MS data (R^2X : 0.717; R^2Y : 0.948; Q^2 : 0.631); C: PCA plot of LC-MS data (R^2X : 0.62; Q^2 : 0.53); D: OPLS-DA plot of LC-MS data (R^2X : 0.737; R^2Y : 0.878; Q^2 : 0.815).

Figure S3. MS/MS spectra (ESI+ & ESI-) of identified metabolites and the comparison with major fragments of metabolites in HMDB database.

Figure S4. Result of metabolic pathway analysis through MetPA software.

Table Captions:

Table S1. The peak areas and retention times of top 12 peaks in QC samples (GC-MS).

Table S2. The peak areas and retention times of top 12 peaks in QC samples (LC-MS).

Table S3. Identified metabolites and their relative levels in the control, model and LR groups.

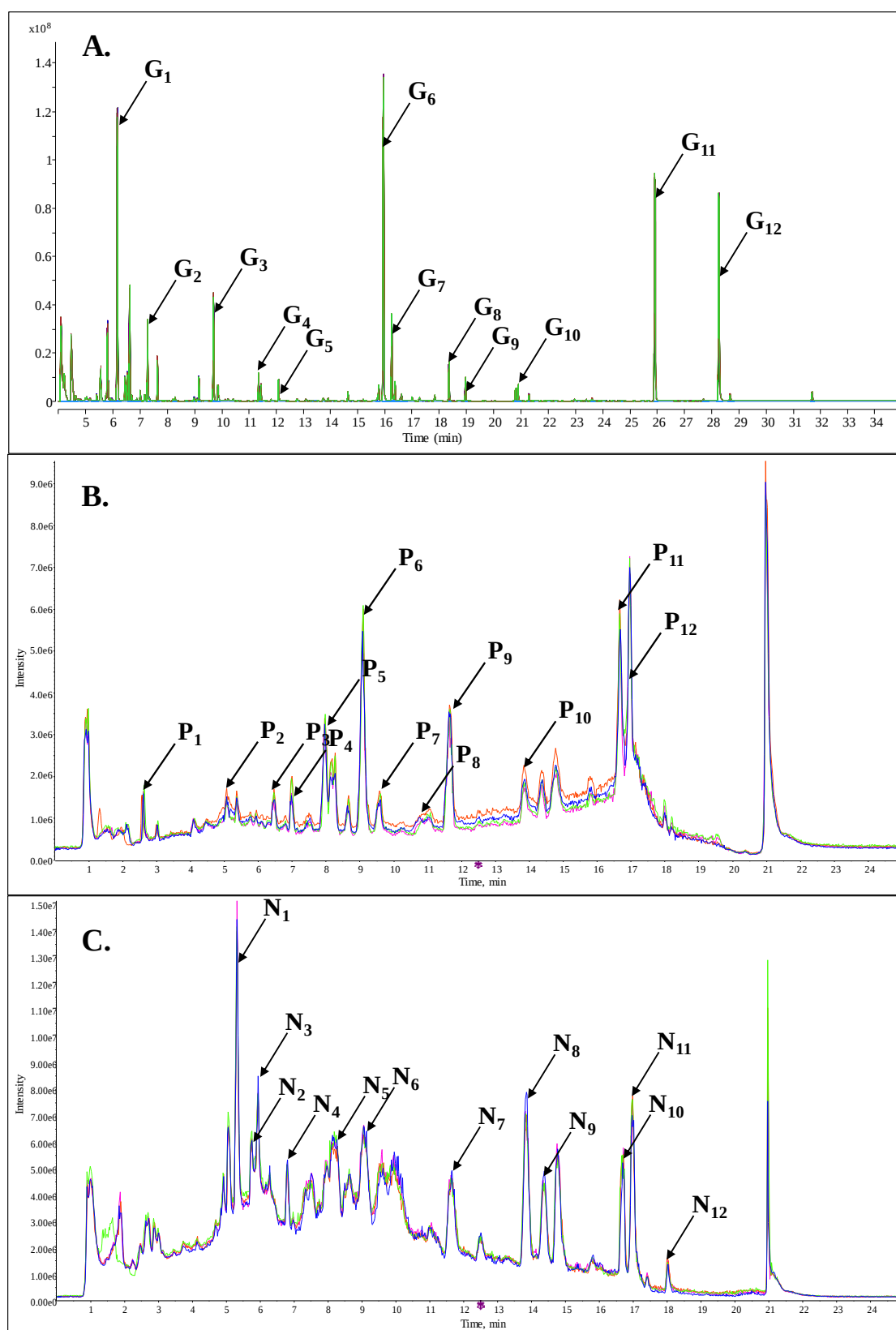


Figure S1. Total ion chromatograms of QC samples and selected top 12 peaks, A: GC-MS, B: LC-MS (ESI+), C: LC-MS (ESI-).

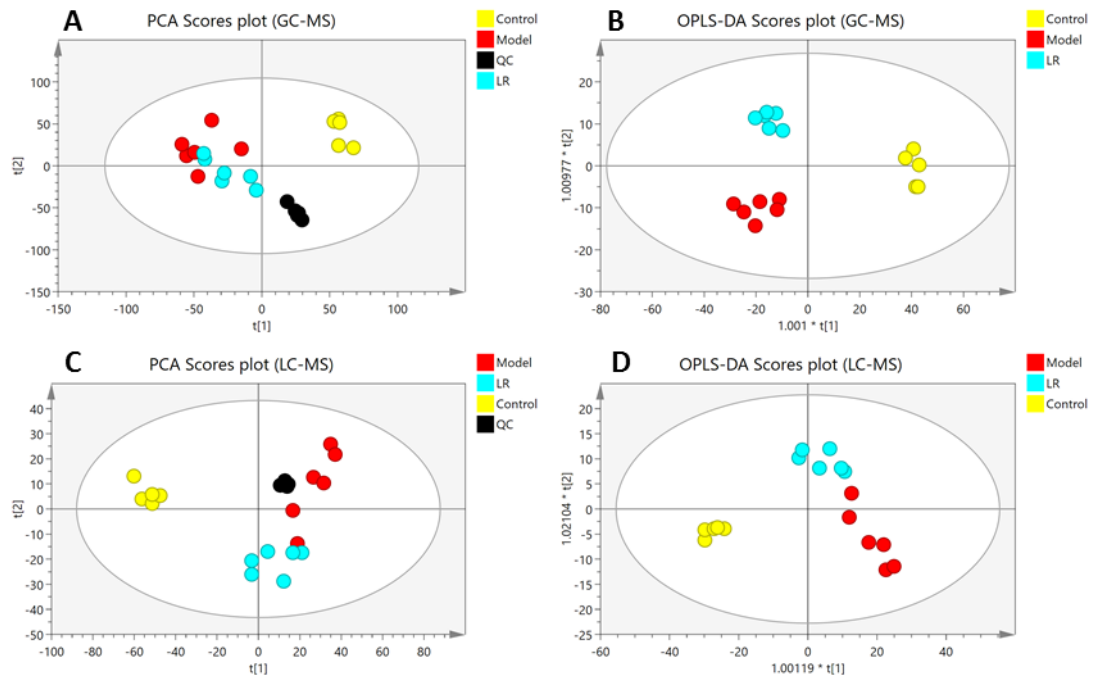


Figure S2. Plots of PCA and OPLS-DA. A: PCA plot of GC-MS data (R^2X : 0.774; Q^2 : 0.603); B: OPLS-DA plot of GC-MS data (R^2X : 0.717; R^2Y : 0.948; Q^2 : 0.631); C: PCA plot of LC-MS data (R^2X : 0.62; Q^2 : 0.53); D: OPLS-DA plot of LC-MS data (R^2X : 0.737; R^2Y : 0.878; Q^2 : 0.815).

Supplementary Materials

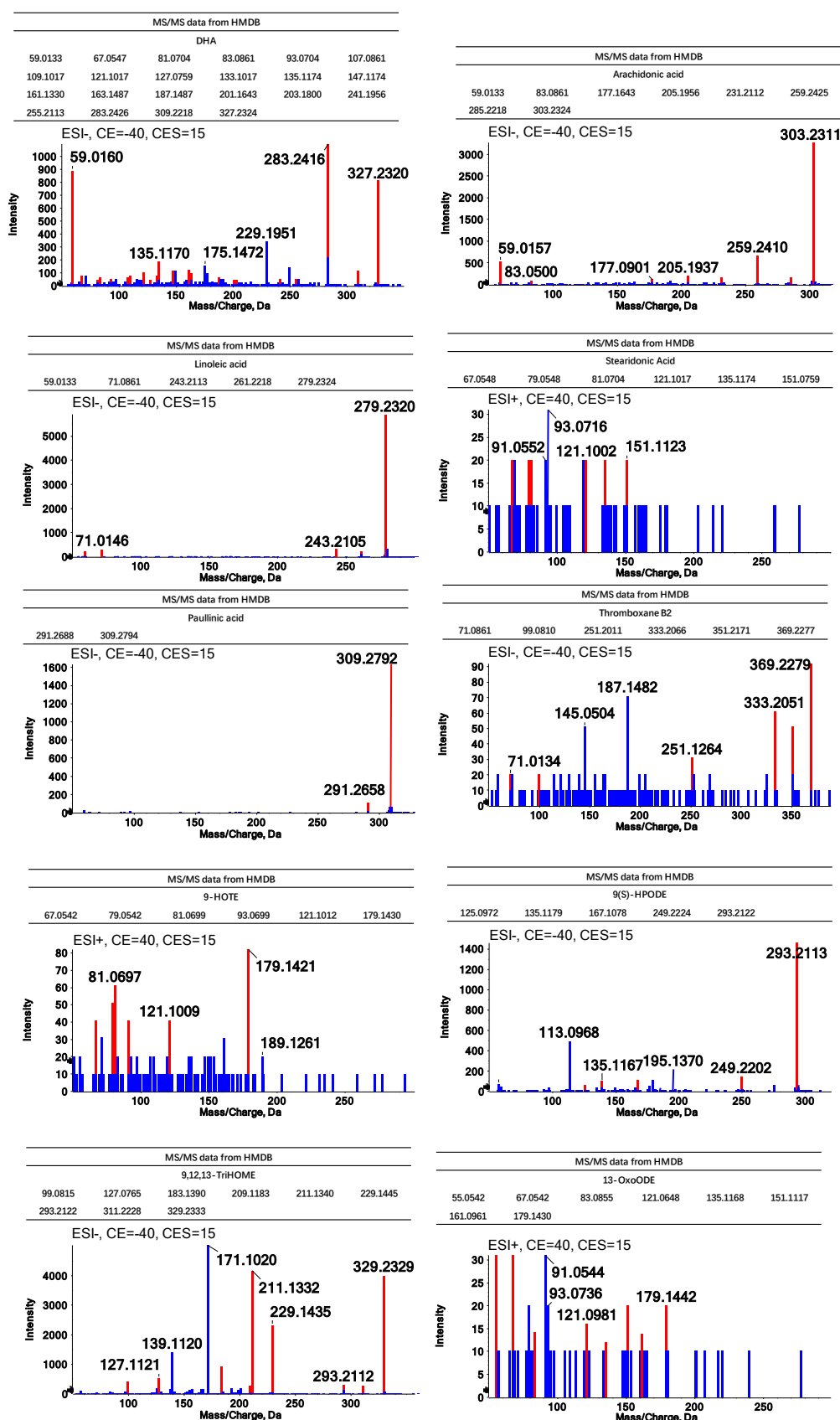


Figure S3. MS/MS spectra (ESI+ & ESI-) of identified metabolites and the comparison with major fragments of metabolites in HMDB database.

Supplementary Materials

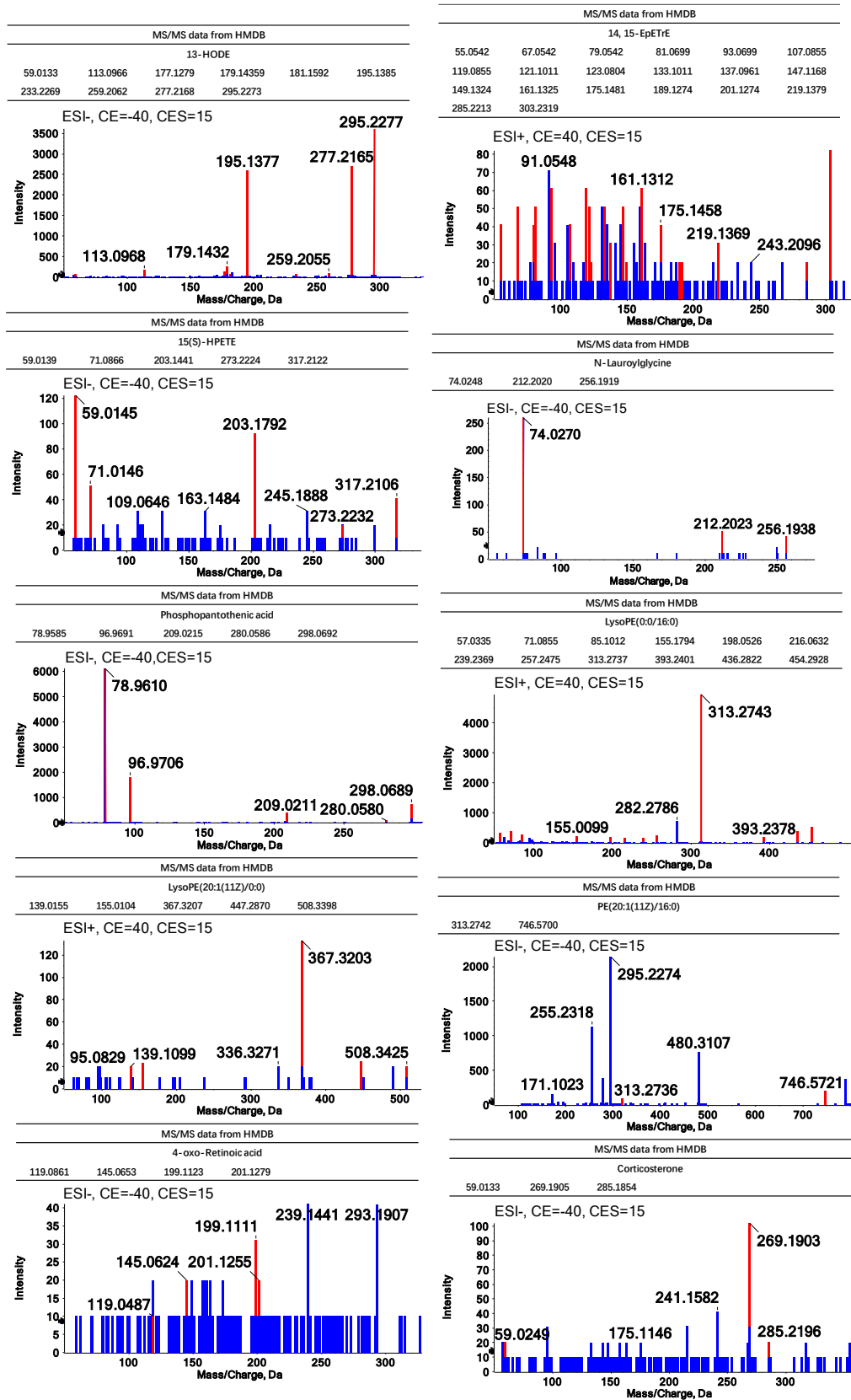


Figure S3. Continued.

Supplementary Materials

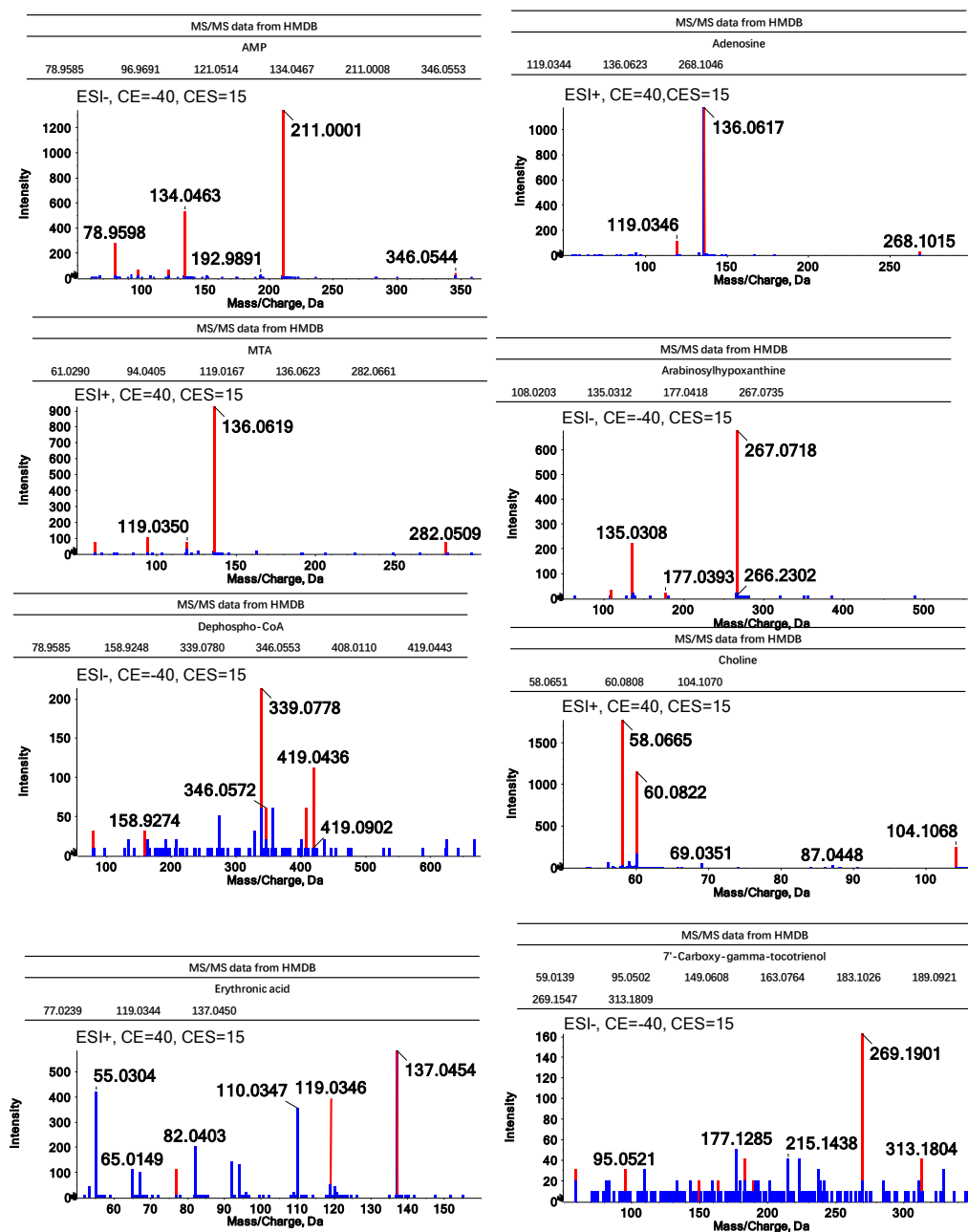


Figure S3. Continued.

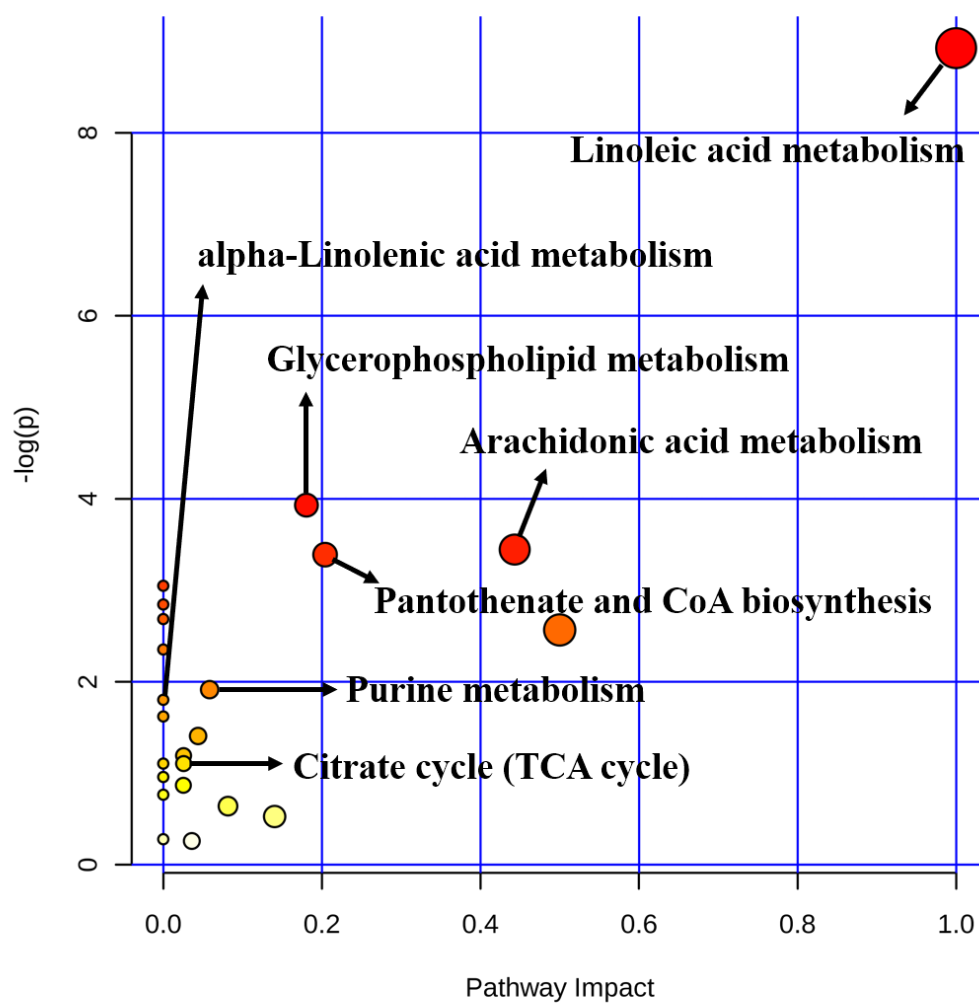


Figure S4. Result of metabolic pathway analysis through MetPA software.

Table S1. The peak areas and retention times of top 12 peaks in QC samples (GC-MS).

NO.	Peak area		Retention time	
	Mean $\pm$ SD ($\times 10^7$)	RSD%	Mean $\pm$ SD	RSD%
G1	26.75 $\pm$ 0.69	2.60	6.17 $\pm$ 0.00	0.06
G2	5.04 $\pm$ 0.17	3.29	7.29 $\pm$ 0.00	0.04
G3	10.08 $\pm$ 0.27	2.68	9.70 $\pm$ 0.00	0.03
G4	1.86 $\pm$ 0.11	5.94	11.35 $\pm$ 0.00	0.02
G5	1.49 $\pm$ 0.10	6.55	12.10 $\pm$ 0.00	0.02
G6	37.69 $\pm$ 0.31	0.83	15.95 $\pm$ 0.00	0.03
G7	7.59 $\pm$ 0.15	1.97	16.25 $\pm$ 0.00	0.02
G8	3.01 $\pm$ 0.07	2.47	18.35 $\pm$ 0.00	0.02
G9	2.08 $\pm$ 0.07	3.29	18.97 $\pm$ 0.00	0.02
G10	1.34 $\pm$ 0.07	4.97	20.89 $\pm$ 0.00	0.01
G11	23.73 $\pm$ 0.49	2.08	25.93 $\pm$ 0.00	0.01
G12	22.88 $\pm$ 0.46	2.00	28.28 $\pm$ 0.00	0.01

Table S2. The peak areas and retention times of top 12 peaks in QC samples (LC-MS).

NO.	LC-MS ^a	M/Z	Intensity		Retention time	
			Mean $\pm$ SD ($\times 10^4$)	RSD%	Mean $\pm$ SD	RSD%
P1	ESI+	166.09	43.10 $\pm$ 3.50	8.12	2.60 $\pm$ 0.03	0.96
P2	ESI+	287.06	26.11 $\pm$ 2.03	7.76	5.07 $\pm$ 0.00	0.10
P3	ESI+	274.27	46.72 $\pm$ 4.18	8.94	6.44 $\pm$ 0.00	0.08
P4	ESI+	432.24	28.85 $\pm$ 3.35	11.63	6.98 $\pm$ 0.01	0.14
P5	ESI+	520.34	62.24 $\pm$ 9.44	15.17	7.98 $\pm$ 0.01	0.10
P6	ESI+	496.34	229.53 $\pm$ 18.02	7.85	9.09 $\pm$ 0.01	0.06
P7	ESI+	522.35	29.46 $\pm$ 3.94	13.36	9.60 $\pm$ 0.01	0.12
P8	ESI+	601.33	7.14 $\pm$ 0.62	8.72	10.83 $\pm$ 0.05	0.47
P9	ESI+	524.37	117.52 $\pm$ 6.16	5.24	11.67 $\pm$ 0.01	0.08
P10	ESI+	329.25	17.12 $\pm$ 2.80	16.35	13.85 $\pm$ 0.01	0.04
P11	ESI+	760.58	56.53 $\pm$ 5.12	9.06	16.68 $\pm$ 0.01	0.03
P12	ESI+	758.57	111.25 $\pm$ 11.75	10.56	16.96 $\pm$ 0.01	0.03
N1	ESI-	285.04	113.50 $\pm$ 6.33	5.58	5.07 $\pm$ 0.00	0
N2	ESI-	514.28	561.74 $\pm$ 39.28	6.99	5.32 $\pm$ 0.01	0.09
N3	ESI-	453.29	102.40 $\pm$ 3.97	3.87	5.93 $\pm$ 0.01	0.08
N4	ESI-	471.06	61.13 $\pm$ 3.88	6.36	6.79 $\pm$ 0.00	0
N5	ESI-	564.33	93.70 $\pm$ 5.80	6.19	8.28 $\pm$ 0.01	0.06
N6	ESI-	540.33	155.90 $\pm$ 11.07	7.10	9.11 $\pm$ 0.01	0.06
N7	ESI-	480.31	143.84 $\pm$ 4.98	3.46	11.59 $\pm$ 0.01	0.05
N8	ESI-	327.23	282.50 $\pm$ 20.99	7.43	13.84 $\pm$ 0.01	0.04
N9	ESI-	303.23	159.25 $\pm$ 11.68	7.33	14.37 $\pm$ 0.01	0.04
N10	ESI-	255.23	69.30 $\pm$ 8.03	11.58	16.68 $\pm$ 0.01	0.03
N11	ESI-	281.25	133.64 $\pm$ 4.79	3.58	16.99 $\pm$ 0.00	0.03
N12	ESI-	283.26	62.78 $\pm$ 4.22	6.72	18.01 $\pm$ 0.00	0.03

^a ESI+: ESI positive mode; ESI-: ESI negative mode.

Table S3. Identified metabolites and their relative levels in the control, model and LR groups.

Class	Metabolites	HMDB	Model/Control ^a	LR/Control ^b	LR/Model ^c	Analytical Instrument ^d
Fatty acids	Docosahexaenoic acid (DHA)	HMDB0002183	↓***	↓	↑***	LC-MS (ESI-)
Fatty acids	Arachidonic acid	HMDB0001043	↓***	↓***	↑**	LC-MS (ESI-)
Fatty acids	Linoleic acid	HMDB0000673	↑***	↑***	↓***	LC-MS (ESI-)
Fatty acids	Stearidonic Acid	HMDB0006547	↑***	↑**	↓	LC-MS (ESI+)
Fatty acids	Paullinic acid	HMDB0035159	↑***	↑***	↓	LC-MS (ESI-)
Fatty acids	Thromboxane B2	HMDB0003252	↑***	↑***	↓	LC-MS (ESI-)
Fatty acids	9-HOTE	HMDB0010224	↑***	↑**	↓	LC-MS (ESI+)
Fatty acids	9(S)-HPODE	HMDB0006940	↑***	↑**	↓	LC-MS (ESI-)
Fatty acids	9,12,13-TriHOME	HMDB0004708	↑**	↑*	↓	LC-MS (ESI-)
Fatty acids	13-OxoODE	HMDB0004668	↑**	↑**	↓	LC-MS (ESI+)
Fatty acids	13-HODE	HMDB0004667	↑***	↑***	↓	LC-MS (ESI-)
Fatty acids	14,15-EpETrE	HMDB0004264	↑**	↑***	↓	LC-MS (ESI+)
Fatty acids	15(S)-HPETE	HMDB0004244	↑***	↑**	↓	LC-MS (ESI-)
Amino acids	L-Tyrosine	HMDB0000158	↓*	↓	↑	GC-MS
Amino acids	N-Lauroylglycine	HMDB0013272	↓***	↓**	↑	LC-MS (ESI-)

Supplementary Materials

Amino acids	Phosphopantothenic acid	HMDB0001016	↑***	↑***	↓***	LC-MS (ESI-)
Lipids	Glycerol 3-phosphate	HMDB0000126	↓**	↓	↑*	GC-MS
Lipids	LysoPE(0:0/16:0)	HMDB0011473	↑***	↑***	↓	LC-MS (ESI+)
Lipids	LysoPE(20:1(11Z)/0:0)	HMDB0011512	↑**	↑***	↓	LC-MS (ESI+)
Lipids	PE(20:1(11Z)/16:0)	HMDB0009253	↑***	↑***	↓	LC-MS (ESI-)
Lipids	4-oxo-Retinoic acid	HMDB0006285	↑***	↑***	↓	LC-MS (ESI-)
Lipids	Corticosterone	HMDB0001547	↑***	↑**	↓	LC-MS (ESI-)
Nucleosides	Adenosine monophosphate (AMP)	HMDB0000045	↓**	↓**	↑*	LC-MS (ESI-)
Nucleosides	Adenosine	HMDB0000050	↓*	↓*	↑	LC-MS (ESI+)
Nucleosides	Inosine	HMDB0000195	↓**	↓**	↑	LC-MS (ESI+)
Nucleosides	5'-Methylthioadenosine (MTA)	HMDB0001173	↓***	↓***	↑	LC-MS (ESI+)
Nucleosides	Arabinosylhypoxanthine	HMDB0003040	↓***	↓***	↑	LC-MS (ESI-)
Nucleosides	Dephospho-CoA	HMDB0001373	↓***	↓***	↑	LC-MS (ESI-)
Others	Succinic acid	HMDB0000254	↑	↑	↓*	GC-MS
Others	Choline	HMDB0000097	↑***	↑*	↓**	LC-MS (ESI+)
Others	Erythronic acid	HMDB0000613	↓*	↓***	↑	LC-MS (ESI+)
Others	3-Hydroxybutyric acid	HMDB0000357	↓***	↓**	↓	GC-MS
Others	N-Acetylmannosamine	HMDB0001129	↑***	↑**	↓	GC-MS
Others	7'-Carboxy-gamma-tocotrienol	HMDB0012851	↑***	↑**	↓	LC-MS (ESI-)

^a Comparison of relative levels of metabolites in the model group and the control group.

^b Comparison of relative levels of metabolites in the LR group and the control group.

^c Comparison of relative levels of metabolites in the LR group and the model group.

^d ESI+: ESI positive mode; ESI-: ESI negative mode.

The hypothesis test for the difference between two means was conducted through t-test. *: $p < 0.05$; **: $p < 0.01$; ***: $p < 0.001$.