## 1 Supplementary materials

- 2 Untargeted metabolomic analysis using LC–TOF/MS and LC–MS/MS for revealing
- 3 metabolic alterations linked to alcohol-induced hepatic steatosis in rat serum and
- 4 plasma
- 5 Huan Wu<sup>ab</sup> and Fang Feng<sup>\*abc</sup>
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## 8 Affiliation

9 a Department of Pharmaceutical Analysis, China Pharmaceutical University, Nanjing

10 210009, China

- 11 <sup>b</sup> Key Laboratory of Drug Quality Control and Pharmacovigilance (Ministry of
- 12 Education), China Pharmaceutical University, Nanjing 210009, China
- 13 ° State Key Laboratory of Natural Medicine, China Pharmaceutical University, Nanjing

14 210009, China

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## 17 Correspondence

- 18 Prof. Fang Feng. Postal address: Department of Pharmaceutical Analysis, China
- 19 Pharmaceutical University, TongJiaXiang No. 24, Nanjing 210009, China.
- 20 Tel.: +86 025 83271301. E-mail addresses: fengfang1@hotmail.com (F. Feng)

- 21 Supporting information Text S1
- 22 Parameter settings for XCMS processing depend on the instrument platform in which
- 23 the data were acquired. The parameters as follows: centWave for feature detection
- 24 ( $\Delta m/z$ = 5 ppm, minimum peak width = 10 s, and maximum peak width = 60 s);
- 25 obiwarp settings for retention-time correction (profStep = 0.5); and parameters for
- 26 chromatogram alignment, including mzwid = 0.025, minfrac = 0.5, and bw = 5.



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28 Fig.S1. The %RSD distribution of all features detected from QCP sample in positive

29 mode and negative mode, respectively. The 80%-rule was applied on the data and

30  $\,$  features having %RSD values higher than 60% were excluded because these are likely  $\,$ 

31 to have large RSDs and may not be a true reflection of the variation between

32 subjects.



34 Fig. S2. Score plots of OPLS-DA in positive-ion mode (A) with the statistical

35 parameters ( $R^2X = 0.597$ ,  $R^2Y = 0.989$ ,  $Q^2 = 0.902$ ,  $p[CV-ANOVA] = 4.3 \times 10^{-4}$ ) and in

36 negative-ion mode (C) with the statistical parameters ( $R^2X = 0.760$ ,  $R^2Y = 0.984$ ,  $Q^2 =$ 

37 0.957, p[CV-ANOVA] = 1.2 × 10<sup>-5</sup>) between the normal rat serum (blue squares) and

38 AHS rat serum (yellow inverted triangle). The corresponding S-plot from OPLS-DA

39 model of the two groups in positive-ion mode (B) and in negative-ion mode (D).



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41 Fig. S3. Score plots of OPLS-DA in positive-ion mode (A) with the statistical

42 parameters ( $R^2X = 0.539$ ,  $R^2Y = 0.996$ ,  $Q^2 = 0.978$ , p[CV-ANOVA] = 4.6 × 10<sup>-9</sup>) and in

43 negative-ion mode (B) with the statistical parameters ( $R^2X = 0.741$ ,  $R^2Y = 0.994$ ,  $Q^2 =$ 

44 0.967, p[CV-ANOVA] = 3.5 × 10<sup>-6</sup>) between the normal rat plasma (green dot) and

45 AHS rat plasma (brown triangle). The corresponding S-plot from OPLS-DA model of

46 the two groups in positive-ion mode (B) and in negative-ion mode (D).



49 Fig. S4. HPLC-ESI-TOF/MS chromatograms of normal rat plasma and AHS rat plasma 50 in positive-ion mode (A, B) and in negative-ion mode (C, D), respectively.

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Table S1: OPLS-DA markers ( $t_{R}$ - $m/z$ ) of serum and plasma samples in positive-ion mode and negative-ion mode										
Features (t <sub>R</sub> - <i>m/z</i> )	Adduct	Formula	M <sub>w</sub> (Da)	Diff. (ppm)	Metabolite	Chemical classe	VIP	Intensity response		
Positive-ion mode										
9.77-120.0811	[M+H] <sup>+</sup>	$C_8H_{10}N$	120.0813	-1.6	Frag of Phe-Phe	Peptide	3.65	Δ		
31.21-453.1721	-	-	-	-	DCP	Peptide	3.32	Δ		
30.10-437.1983	-	-	-	-	DCP	Peptide	3.31	Δ		
9.79-313.1531	[M+H] <sup>+</sup>	$C_{18}H_{21}N_2O_3$	313.1552	-6.7	Phe-Phe	Peptide	2.72	Δ		
41.48-542.3242	[M+Na] <sup>+</sup>	$C_{26}H_{50}NO_7PNa$	542.3223	3.5	LysoPC (18:2)	Phospholipid	2.68	$\uparrow^*$		
45.18-522.3556	[M+H] <sup>+</sup>	$C_{26}H_{53}NO_7P$	522.3560	-0.8	LysoPC (18:1)	Phospholipid	2.36	$\uparrow^*$		
31.76-513.1842	-	-	-	-	DCP	Peptide	2.32	Δ		
31.26-504.3089	-	-	-	-	DCP	peptide	2.09	Δ		
38.10-539.3192	-	-	-	-	DCP	peptide	1.83	Δ		
59.02-782.5683	[M+H] <sup>+</sup>	$C_{44}H_{81}NO_8P$	782.5700	-2.2	PC (36:4)	Phospholipid	1.55	$\uparrow^*$		
43.71-518.3235	[M+Na] <sup>+</sup>	$C_{24}H_{50}NO_7PNa$	518.3223	2.3	LysoPC (16:0)	Phospholipid	1.51	$\uparrow^*$		
50.63-524.3713	[M+H] <sup>+</sup>	$C_{26}H_{55}NO_7P$	524.3716	-0.6	LysoPC (18:0)	Phospholipid	1.33	$\uparrow^*$		
50.81-303.2313	[M+H] <sup>+</sup>	$C_{20}H_{31}O_2$	303.2324	-3.6	Eicosapentaenoic acid	Fatty acid	1.27	$\uparrow^*$		
3.21-175.0244	[M+Na]+	$C_5H_4N_4O_2Na$	175.0232	6.9	Xanthine	Heterocyclic compound	1.17	个*		
5.94-159.0274	[M+Na] <sup>+</sup>	$C_5H_4N_4ONa$	159.0283	-5.7	Hypoxanthine	Heterocyclic compound	1.14	<b>^</b> *		
59.01-758.5691	[M+H] <sup>+</sup>	$C_{42}H_{81}NO_8P$	758.5700	-1.2	PC (34:2)	Phospholipid	1.06	$\uparrow^*$		
Negative-ion mod	de									

ty acid 2.43	$\uparrow^*$	
ospholipid 2.33	$\uparrow^*$	
ty acid 2.07	/ 个*	
ty acid 2.02	个*	
ty acid 1.87	/ 个*	
ty acid 1.82	↓*	
ty acid 1.72	$\downarrow^*$	
ospholipid 1.62	$\uparrow^*$	
ospholipid 1.57	/ 个*	
ty acid 1.55	$\uparrow^*$	
ospholipid 1.36	个*	
ospholipid 1.29	$\uparrow^*$	
ospholipid 1.12	个*	
ospholipid 1.02	$\uparrow^*$	
	ospholipid 2.33 ity acid 2.07 ity acid 2.01 ity acid 1.87 ity acid 1.87 ity acid 1.81 ity acid 1.72 ospholipid 1.62 ospholipid 1.57 ity acid 1.55 ospholipid 1.36 ospholipid 1.29 ospholipid 1.11 ospholipid 1.02	ospholipid2.33 $\uparrow^*$ :ty acid2.07 $\uparrow^*$ :ty acid2.01 $\uparrow^*$ :ty acid1.87 $\uparrow^*$ :ty acid1.87 $\downarrow^*$ :ty acid1.72 $\downarrow^*$ ospholipid1.62 $\uparrow^*$ ospholipid1.57 $\uparrow^*$ ity acid1.36 $\uparrow^*$ ospholipid1.29 $\uparrow^*$ ospholipid1.11 $\uparrow^*$ ospholipid1.11 $\uparrow^*$

Abbreviations: *M<sub>w</sub>*, monoisotopic weight; VIP, variable importance on projection; Phe, phenylalanine; DCP, doubly charged peptide; LysoPC, lysophosphatidylcholine; PC, phosphatidylcholine; HHT, hydroxyheptadecatrienoic acid; HEPE, hydroxyeicosapentaenoic acid; HETE, hydroxyeicosatetraenoic acid; diHETE, dihydroxyeicosatetraenoic acid; diHETrE, dihydroxyeicosatetraenoic acid; diHETrE, dihydroxyeicosatetraenoic acid; diHETE, dihydroxyeicosatetraenoic acid; diHETrE, dihydroxyeicosatet

\*: Response compared to heparinized plasma samples.

 $\Delta$ : Detected only in serum.