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1 **Cholestatic liver injury model of bile duct ligation and the protection of**
2 **Huang-Lian-Jie-Du decoction by NMR metabolomic profiling**

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25 **Abstract**

26 Cholestatic liver injury has been increasingly recognized as a cause for high morbidity
27 and mortality of some diseases in human. This model could be established by bile duct
28 ligation (BDL), which led to the toxic accumulation of bile acids in animals, resulting
29 in cholestatic liver injury. In this study, rats were intragastrically administrated with an
30 extract of Huang-Lian-Jie-Du decoction once a day for seven consecutive weeks to
31 study its therapeutic effect. Serum and urine samples were collected and subjected to
32 ¹H NMR-based metabolomic analysis. Perturbations on energy metabolism, amino
33 acid metabolism, gut bacteria metabolism and oxidative stress were observed in BDL
34 rats. The metabolomic pattern showed a distinct biphasic feature of BDL model. Most
35 of these metabolic disturbances occurred in acute phase (week 1) were greatly
36 attenuated in the long run. HLJDD ameliorated the disturbed metabolism throughout
37 this model, showing bilateral adjustment of some metabolites varied in opposite
38 direction in the two phases. This study demonstrated that ¹H NMR-based
39 metabolomics approach is a powerful and feasible tool to study the pathological
40 changes of a disease model dynamically and holistically and for the understanding of
41 the therapeutic effects of complex Chinese herbal medicine formula.

42

43 **1. Introduction**

44 Chronic cholestatic liver disease is one of the major risks responsible for the
45 development of liver cirrhosis and end-stage liver disease culminating in liver failure.

46 Cholestatic liver disease occurs when there is a decrease in bile flow,^{1,2} which is a

47 common pathological condition that can be reproduced in rodents by common bile
48 duct ligation (BDL) during surgical laparotomy.³ Bile acids (BAs) are the active
49 constituents of bile and essential for absorption and solubilization of dietary lipids in
50 the digestive tract.⁴ Of them, hydrophobic bile acids could induce damage of
51 mitochondrial membrane structure and increase of oxidative stress, leading to
52 apoptosis in liver cells.^{5, 6} The obstruction of bile flow results in an increased
53 accumulation of potentially harmful hydrophobic bile acids in the liver and blood, and
54 liver dysfunction.⁷

55 Cholestasis is associated with many liver diseases, and a well-established experimental
56 animal model is BDL in rodents, in which hydrophobic bile acid mediated liver injury.
57 This model develops in a biphasic manner, including acute (phase 1) and chronic
58 (phase 2) cholestasis.⁸

59 Huang-Lian-Jie-Du decoction (HLJDD) is a herbal formula of Traditional Chinese
60 Medicine, consisting of *Coptidis rhizoma*, *Radix Scutellariae*, *Cortex Phellodendri*
61 and *Frucuts Gardeniae*. With marked anti-inflammatory activities and the ability to
62 reduce oxidative stress,⁹⁻¹¹ it has been used to treat hepatitis and liver dysfunction.¹²

63 The conventional clinical chemistry and histopathology methods are not
64 region-specific and the sensitivity is relatively low. However, NMR-based
65 metabolomics revealed a global profile of endogenous metabolites, thus performing an
66 overall assessment of the global metabolic state of the entire organism. In this study,
67 an animal model of cholestasis was constructed by BDL and the treatment effect of
68 HLJDD in the BDL model was investigated by a NMR-based metabolomics approach

69 complemented with histological inspection and biochemical evaluation.

70 **2 Materials & Methods**

71 **2.1 Chemicals and kits**

72 Component herbs of HLJDD (*Rhizoma Coptidis*, *Radix Scutellariae*, *Cortex*
73 *Phellodendri* and *Fructus Gardeniae*) were obtained from Jiangsu Medicine Company
74 (Nanjing, China) and authenticated by Professor Mian Zhang, Department of
75 Medicinal Plants, China Pharmaceutical University, Nanjing. The voucher specimens,
76 deposited at the herbarium of the Department of Natural Medicinal Chemistry, China
77 Pharmaceutical University, were 2012066-RC, 2012067-RS, 2012068-CP and
78 2012069-FG for *Rhizoma Coptidis*, *Radix Scutellariae*, *Cortex Phellodendri* and
79 *Fructus Gardeniae*, respectively. 3-Trimethylsilylpropionic acid (TSP) was obtained
80 from Sigma-Aldrich (St. Louis, MO) and deuterium oxide (D₂O, 99.9%) was
81 purchased from QingDao TengLong WeiBo Technology Co. Ltd (QingDao, China).
82 Ultra-pure distilled water was prepared from a Milli-Q purification system. The serum
83 clinical enzymatic chemistry kits of aspartate aminotransferase (AST), alanine
84 aminotransferase (ALT), alkaline phosphatase (ALP), creatinine (CR), total protein
85 (TP), albumin (ALB) and globulin (GLB) were commercially available from Beckman
86 Coulter Inc (Harbor Boulevard, Fullerton, California, 92834 USA), while the serum
87 radioimmunoassay kits of hyaluronic acid (HA), type IV collagen (CIV), type III
88 precollagen (PCIII) and laminin (LN) were bought from Beijing North biology
89 technique institute (Beijing, China).

90 **2.2 Preparation of HLJDD**

91 *Rhizoma Coptidis*, *Radix Scutellariae*, *Cortex Phellodendri* and *Fructus Gardeniae*
92 were mixed in a ratio of 3:2:2:3, reaching a total weight of 500 g. Then they were
93 extracted with 70% ethanol (1:10, 1:10 and 1:5, w/v) under reflux for three times, 1h
94 each, and the extracted solution was filtered through 5 layer gauzes. The decoction
95 was concentrated to dryness to afford 142.5 g HLJDD (yield: 28.5%) using a rotary
96 vacuum evaporator, then stored in refrigerator at 4 °C. The dried extracts were
97 suspended in 0.5% (w/v) sodium carboxy-methylcellulose (CMC-Na) before
98 intragastric administration and the doses were calculated as raw material weights for
99 the animal experiments.

100 **2.3 Bile duct ligation (BDL) operation**

101 BDL was performed using a standard technique.¹³ Briefly, animals were anesthetized
102 by injected intraperitoneally (ip) with chloral hydrate (350 mg/kg) and kept under
103 anesthesia with additional ip injection throughout the experiment. After a midline
104 incision under sterile conditions, a single ligature with silk suture was done with 4-0
105 nylon sutures, followed by careful suturing of the peritoneum and muscle layers as
106 well as the skin wound. The sham-operated rats underwent the same surgical operation
107 except for ligation of bile duct.

108 **2.4 Animals handling procedure**

109 Forty-two adult male Sprague-Dawley rats (220-240 g), of Specified-Pathogens Free
110 (SPF) grade, were obtained from the Experimental Animal Center of Yangzhou
111 University (Yangzhou, China). Rats were group-housed in polysulfone cages (5 rats to
112 one cage) with bedding material, and were housed in a room with controlled humidity

113 (50 ± 10%) and temperature (25 ± 3 °C) under a 12/12-h light/dark cycle. The animals
114 were given free access to standard diet and water and were allowed to acclimate for 7
115 days before treatment. The studies were in accordance with the standard guidelines for
116 the Care and Use of Laboratory Animal from the National Institute of Health (NIH)
117 and were approved by the Animal Ethics Committee of the China Pharmaceutical
118 University.

119 Rats were randomly divided into three groups (n=14): sham-operated (NC), bile duct
120 ligation (BDL) and BDL with HLJDD treatment (BHD). BHD rats were
121 intragastrically administered with HLJDD at doses of 2.7 g/kg body weight, and NC
122 and BDL rats were administered with the same volume of 0.5% CMC-Na for each
123 administration once a day for seven consecutive weeks.

124 **2.5 Collection of serum and urine**

125 On weeks 1, 3, 5 and 7 after the treatment, blood samples were taken from the ocular
126 vein of rats after 12 h fasting. The serum samples were obtained by centrifugation at
127 13282 g for 10 min, and stored at -80 °C before next experiments. Centrifugation was
128 performed on Beckman Coulter Microfuge® 22R refrigerated microcentrifuge using
129 F241.5 rotor, with a radius of 8.25 cm and the maximum RCF at 21591 g.

130 Rats were housed in metabolic cages for a 24 hour interval and the urine samples were
131 collected at weeks 1, 3, 5 and 7. The samples were then centrifuged at 13282 g for 10
132 min to afford the supernatants and stored at -80 °C before NMR spectroscopic
133 analysis.

134 **2.6 Histopathology and serum biochemical analysis**

135 At the end of week 7, rats were fasted overnight and sacrificed after deep
136 anesthetization with chloral hydrate (350 mg/kg, i.p.). The livers and kidneys were
137 removed at the time of death, flushed with cold phosphate buffer solution to remove
138 residual blood. The liver and kidney tissues obtained were immediately immersed in
139 10% neutral-buffered formaldehyde and embedded in paraffin to be stained with
140 hematoxylin eosin (HE).

141 Serum samples harvested at different time-points were used for clinical chemistry
142 evaluation. To assess liver and renal function, the concentrations of AST, ALT, ALP,
143 TP, GLB, ALB and Cr were measured using commercially available kits from Nanjing
144 Jiancheng Biotech Inc. On week 7, HA, PCIII, LN and CIV were detected by
145 radioimmunoassay (RIA).

146 **2.7 ^1H NMR spectroscopic measurement of serum and urine**

147 The serum and urine samples were thawed at room temperature and 300 μL of each
148 was added with 300 μL D_2O ($0.2 \text{ mol L}^{-1} \text{Na}_2\text{HPO}_4$ and $0.2 \text{ mol L}^{-1} \text{NaH}_2\text{PO}_4$, pH 7.4,
149 containing 0.05 % TSP). TSP acted as a chemical shift reference (δ 0.0) and D_2O
150 provided a lock signal. The samples were vortexed and centrifuged at 13282 g for 10
151 min at 4 $^\circ\text{C}$ to remove insoluble material. The supernatants were then pipetted out into
152 5 mm NMR tubes for NMR recording.

153 ^1H NMR spectra of the samples were recorded on a Bruker AV 500 MHz spectrometer
154 at 300 K. For each serum sample, the transverse relaxation-edited Carr–Purcell–
155 Meiboom–Gill (CPMG) spin-echo pulse sequence ($\text{RD-}90^\circ\text{-(}\tau\text{-}180^\circ\text{-}\tau\text{) n-ACQ}$) with a
156 total spin-echo delay ($2n\tau$) of 40 ms was used to suppress broad signals from

157 macromolecules, therefore the signals of micromolecules were clearly observed. ^1H
158 NMR spectra were measured with 128 scans into 32 K data points over a spectral
159 width of 10000 Hz. Prior to Fourier transformation, an exponential window function
160 with a line broadening of 0.5 Hz was used to the free induction decays (FIDs). For
161 urine, a nuclear overhauser effect spectroscopy (NOESY) pulse sequence (relaxation
162 delay- 90° - μs - 90° -tm- 90° -acquire-FID) was used to attenuate the residual water signal.
163 FIDs were collected into 32 k data points over a spectral width of 10000 Hz with an
164 acquisition time of 2.04 s. The FIDs were weighted by an exponential function with a
165 0.3 Hz line-broadening factor prior to Fourier transformation.

166 **2.8 Spectral pre-processing and data analysis**

167 The spectra for all samples were manually phased and baseline corrected, and
168 referenced to TSP at 0.0 ppm, using Bruker Topspin 3.0 software (Bruker GmbH,
169 Karlsruhe, Germany). The ^1H NMR spectra were automatically exported to ASCII
170 files using MestReNova (Version 8.0.1, Mestrelab Research SL), which were then
171 imported into “R” (<http://cran.r-project.org/>), and aligned with an in-house developed
172 R-script to further reduce phase and baseline distortions. The one-dimensional (1D)
173 spectra were converted to an appropriate format for statistical analysis by
174 automatically segmenting each spectrum into 0.005 ppm integrated spectral regions
175 (buckets) between 0.2 and 10 ppm. The region of the residual water and affected
176 signals (4.2–5.7 ppm) was removed. To account for different dilutions of samples, all
177 binned spectra were probability quotient normalized and then mean-centered before
178 further multivariate analysis.

179 The mean-centered and Pareto-scaled NMR data were analyzed by principal
180 component analysis (PCA) and orthogonal partial least-squares discriminant analysis
181 (OPLS-DA). PCA is an exploratory unsupervised method to maximize the separation
182 by providing model-free approaches for determining the latent or intrinsic information
183 in the dataset.¹⁴ However, no clustering was observed when variables were not
184 selected. OPLS-DA determines PLS components that are orthogonal to the grouping
185 and was used to concentrate group discrimination into the first component with
186 remaining unrelated variations contained in subsequent components.¹⁵ All OPLS-DA
187 models were validated by a repeated two-fold cross-validation method and
188 permutation test. The parameters of R^2 and Q^2 reflected the goodness of fitness and the
189 predictive ability of the models, respectively. The p value of the permutation test
190 denoted the number of times that the permuted data yielded a better result than the
191 one using the original labels. The fold change values of metabolites among different
192 groups were calculated. The Benjamini & Hochberg method¹⁶ was used to adjust the
193 related p-values for controlling the false discovery rate in multiple comparisons
194 applying scripts written in R language, which is available freely, open-source software
195 package.

196 **3 Results**

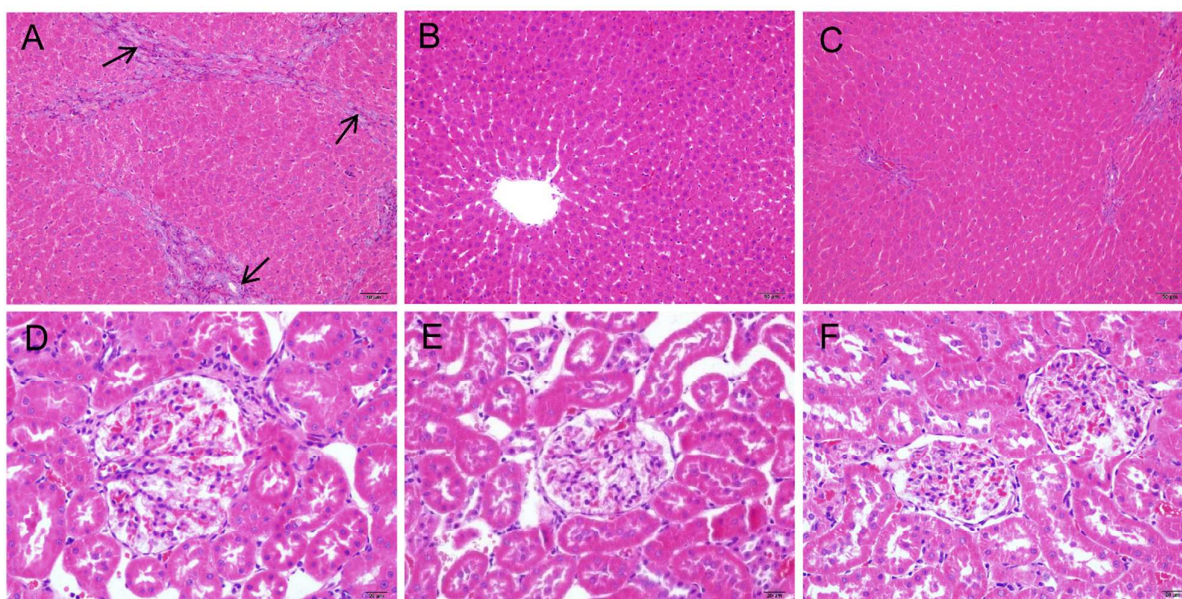
197 **3.1 Animal monitoring**

198 There was no significant difference in the weight of rats in NC, BDL and BHD group
199 on week 7. The serum and urine collected from BDL rats were visually yellow, and
200 after dissection rats undergoing BDL had yellow obstructive jaundice skin, thicken

201 liver especially in the ventral lobe and enlarged bile duct.

202 **3.2 Liver and kidney histopathology**

203 H&E staining of representative liver sections were observed and shown at 200 ×
204 magnifications. The most obvious character of liver impairment after BDL was the
205 severe proliferation of bile duct around the pre-existing interlobular ducts and
206 surrounded by a connective-tissue sheath. Additional diffuse collagen fiber was also
207 noted in interstitial and the portal zones with formation of pseudo lobe. Inflammatory
208 cell infiltration localized specifically to the periportal zones of the livers of BDL rats,
209 was observed in the vicinity of proliferating bile ducts (Figure 1A), as compared with
210 normal livers in the NC group (Figure 1B). Consecutive administration of HLJDD for
211 7 weeks resulted in remarkable amelioration in these pathological changes (Figure 1C).
212 Histopathological inspection revealed a slight increase of the volume of glomerular in
213 BDL group (Figure 1D) as compared with the NC group (Figure 1E); BHD group
214 showed almost no difference from the normal group (Figure 1F).



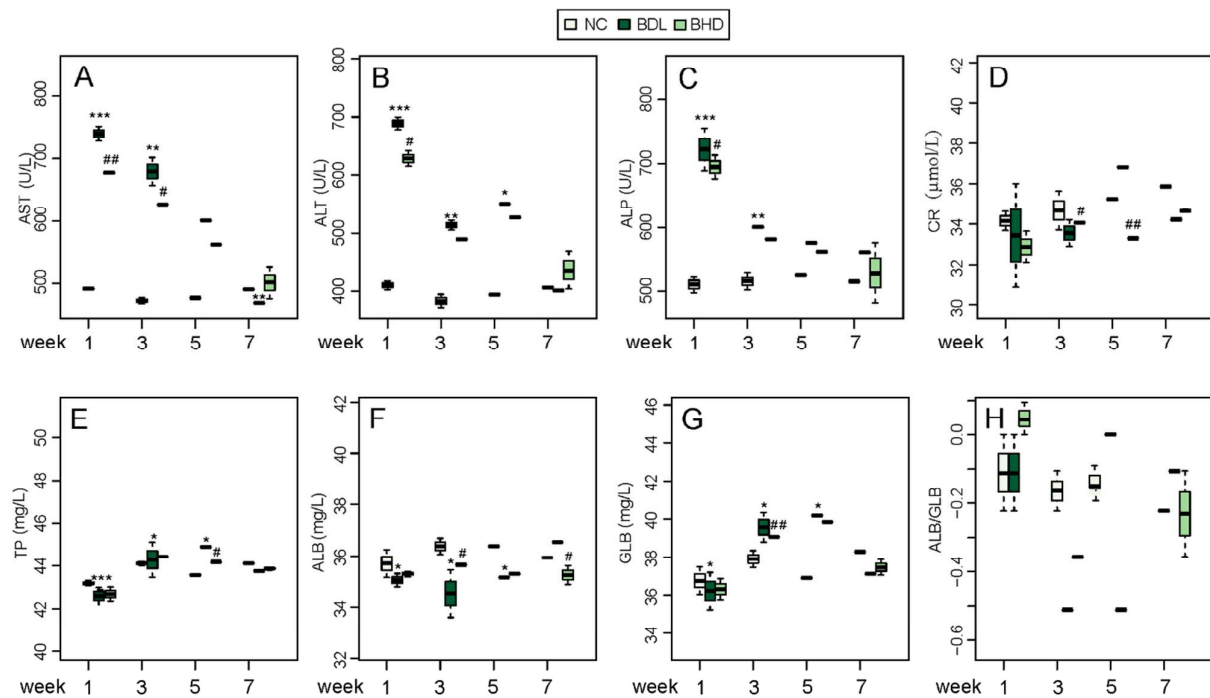
215

216 **Figure 1.** Histopathological study of liver and kidney of NC, BDL and BHD rats at 200 ×
217 magnifications. (A) Liver of BDL rats 49 days after operation: exhibiting severe hyperplasia of
218 bile-ducts, inflammatory cell infiltration, and additional diffuse collagen fiber. (B) Livers in sham
219 operation rats: showing no pathological changes. (C) Livers in BHD rats 49 days after intragastric
220 administration: showing no obvious change without any sign of cell degeneration or necrosis. (D)
221 Kidneys of BDL rats after 7 weeks, with slight increase in the volume of glomerular. (E) Kidneys in
222 sham operation group rats, with no pathological changes. (F) Kidneys in BHD group rats after 7
223 weeks, with no pathological changes.

224 **3.3 Serum biochemical parameters**

225 Serum levels AST, ALT, ALP, CR, TP, ALB and GLB were determined on week 1, 3, 5
226 and 7, and the results of the clinical chemistry are presented in Fig. 2. Levels of AST,
227 ALT and ALP in BDL group increased significantly on week 1 as compared with the
228 NC group, but the increase attenuated from week 1 onwards, reaching to a minimum
229 on week 7. The elevated AST, ALT and ALP levels indicated that cholestasis and liver
230 cell necrosis were already present one week after BDL¹⁷. The ALB concentration of
231 BDL rats significant decrease on week 1, 3 and 5, but without significant disturbance
232 on week 7 as compared with NC group. The TP and GLB in BDL group showed some
233 fluctuations throughout the experiments: markedly decreasing on week 1, but
234 markedly increasing on week 3 and 5 and finally kept at normal levels. ALB and GLB,
235 synthesized by liver cells, are indicators of hepatocellular function.^{18, 19} The increase
236 of AST, ALT, ALP and decrease of ALB in BDL rats on week 1 indicated a severe
237 cholestatic liver injury in the model.²⁰ However, variations of these parameters were

238 attenuated and reversed towards the normal status. On week 7, showing a partial
 239 recovery of liver function in BDL rats. The significantly decreased levels of TP, ALB
 240 and GLB on week 1 indicated the impaired hepatocellular function in synthesis, and
 241 also featuring an acute liver injury. In contrast, on week 3 and 5, both the decrease of
 242 ALB and increase of GLB were observed, characterizing chronic liver damage.
 243 Similarly, both of them reversed to normal on week 7, indicating the alleviation of
 244 liver injury in the long run. The levels of those index indicated liver dysfunction, but
 245 HLJDD could restore most of the fluctuations of AST, ALT, ALP, TP, ALB and GLB
 246 in the serum. CR in BDL group did not show any significant difference at all time
 247 periods.

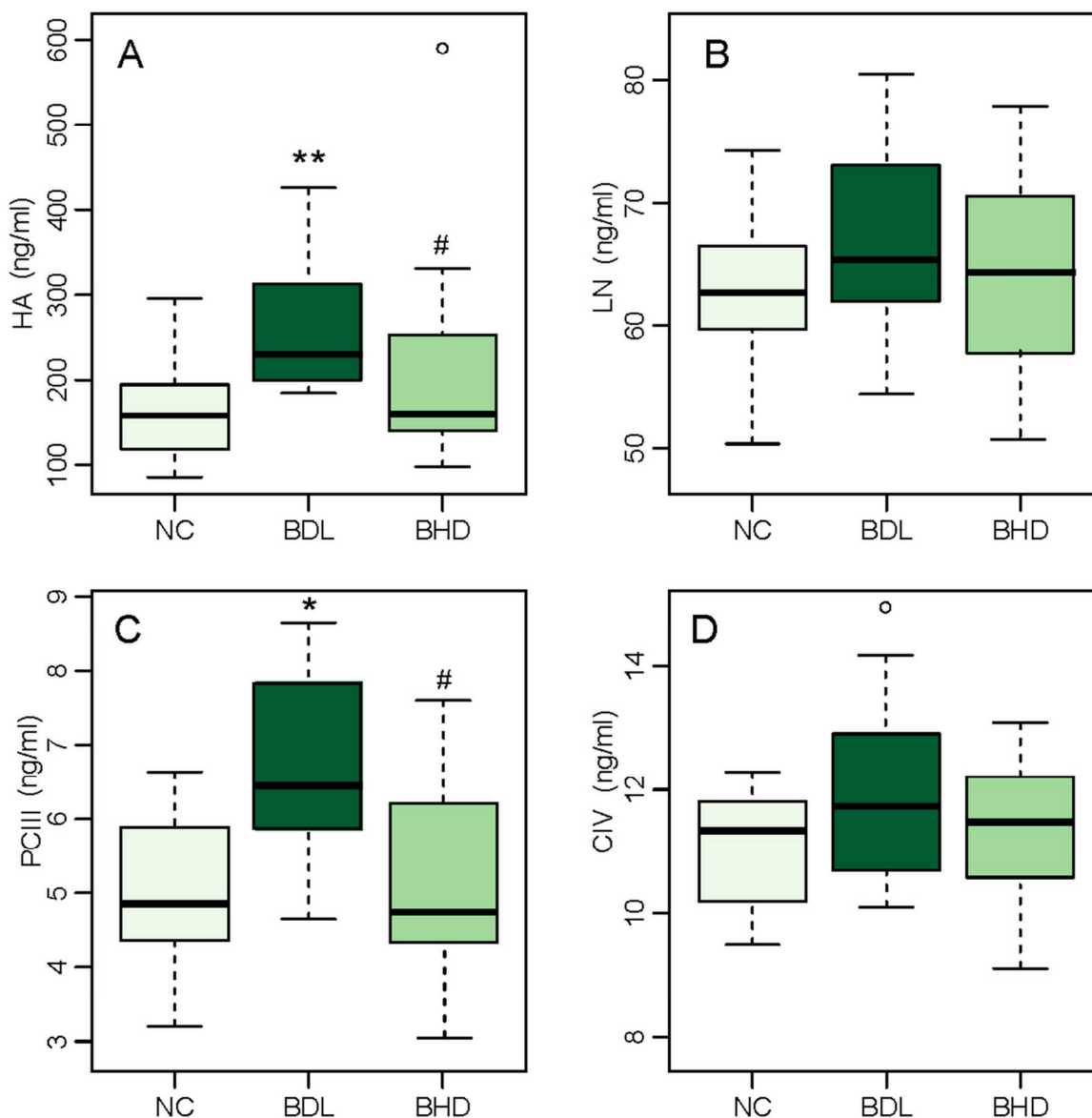


248
 249 **Figure 2.** Boxplots of serum levels of creatinine (Cr), total protein (TP), albumin (ALB), alanine
 250 aminotransferase (ALT), aspartate aminotransferase (AST), alkaline phosphatase (ALP), globulin
 251 (GLB) and the levels of ratio of albumin and globulin (ALB/GLB). The boxes cover 25% quartile

252 and 75% quartile of the data. The line in the box represents the median value. The extended whiskers
253 show the extent of the rest of the data. (A) AST. (B) ALT. (C) AST_ALT. (D) ALP. (E) CR. (F) TP. (G)
254 ALB. (H) GLB. (I) ALB/GLB. Outliers are shown as open circle. Values were expressed as mean \pm
255 SD (n = 10-14). * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$ vs. NC rats; # $p < 0.05$, ## $p < 0.01$ and ###
256 $p < 0.001$ vs. BDL rats.

257

258 To assess liver fibrosis, we measured the levels of hyaluronic acid (HA), laminin (LN),
259 type III procollagen (PCIII) and type IV collagen (CIV) in serum (Fig. 3). HA, LN,
260 PCIII and CIV were markers for liver fibrosis.²¹ The results revealed that levels of HA
261 and PCIII in BDL rats significantly increased on week 7 ($p < 0.05$), compared with
262 NC rats. The levels of LN and CIV in the serum of BDL group augmented slightly.
263 The results suggested the initial formation of liver fibrosis on week 7. HLJDD could
264 restore increased levels of HA, LN and PCIII in serum.



265

266 **Figure 3.** Boxplots of serum levels of hyaluronic acid (HA), type IV collagen (CIV), type III
 267 precollagen (PCIII) and laminin (LN) . The boxes and whiskers represent the same as in Figure 2.

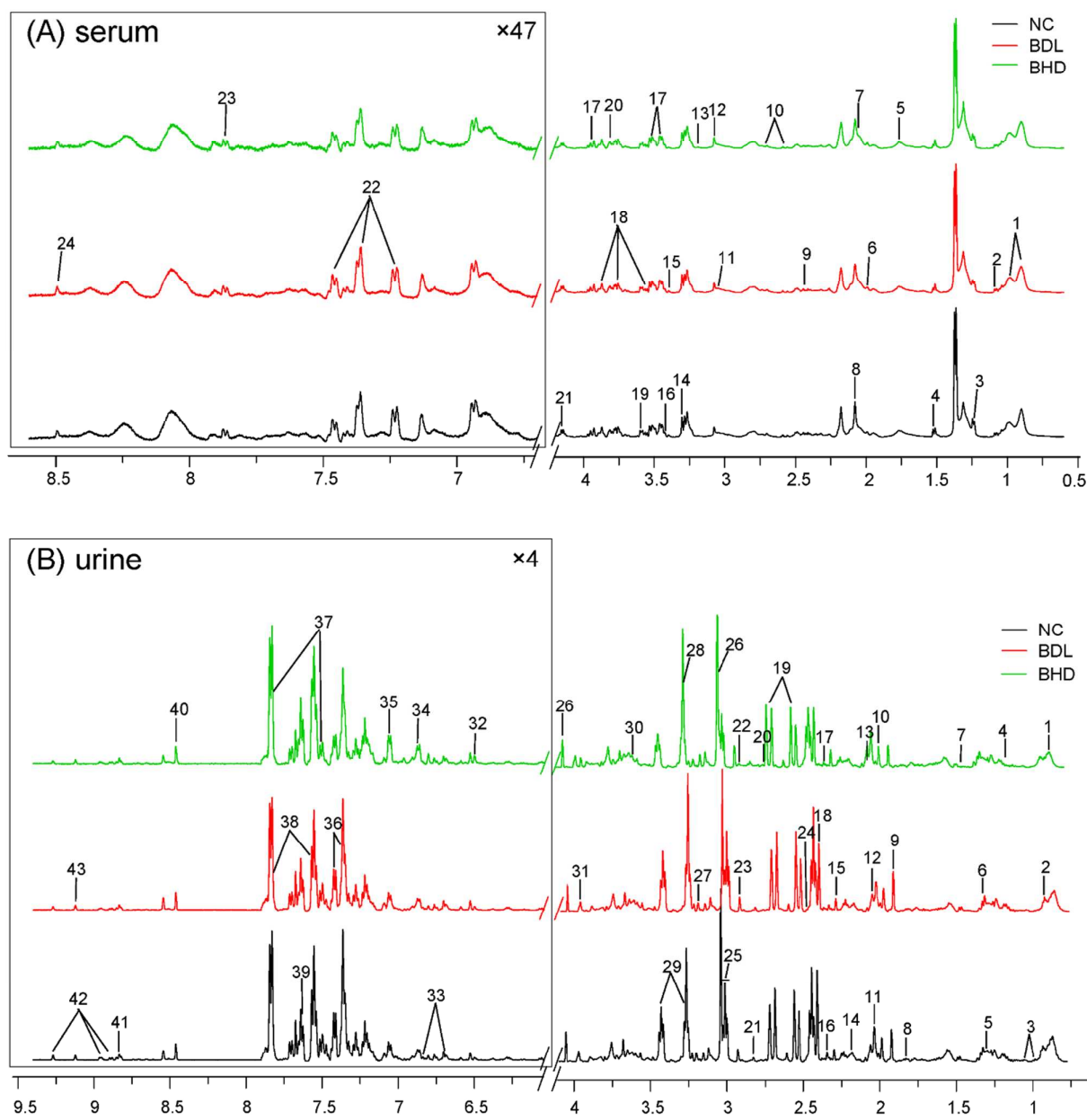
268 Boxplots of HA (A), LN (B), PCIII (C), and CIV (D), respectively. Outliers are shown as open circle.

269 Values were expressed as mean \pm SD. * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$ vs. NC rats; # $p <$
 270 0.05 , ## $p < 0.01$ and ### $p < 0.001$ vs. BDL rats.

271

272 **3.4 Metabolites identification in serum and urine**

273 Representative 500 MHz ^1H NMR spectra of serum and urine samples from NC, BDL
274 and BHD rats were shown in Figures 4 A and B with the assignment of metabolites.
275 Aided by Chenomx NMR suit (Version 8.1, Chenomx, Inc.) and a statistical total
276 correlation spectroscopy (STOSCY) technique,²² their assignments were made by
277 referencing reported data^{16, 23} and querying publicly accessible metabolomic databases,
278 such as HMDB (<http://www.hmdb.ca>), KEGG (<http://www.kegg.jp>), METLIN
279 (<http://metlin.scripps.edu>). The detailed information of the metabolites was listed in
280 Tables S1 and S2.



281

282 **Figure 4.** (A) Typical 500 MHz CPMG ^1H NMR spectra of the serum samples from NC, BDL and
 283 BHD rats on week 1. 1. very low density lipoprotein (VLDL) / low density lipoprotein (LDL); 2.
 284 valine; 3. β -Hydroxybutyrate (HB); 4. alanine; 5. lysine; 6. acetate; 7. adipate; 8. N-Acetyl
 285 glycoproteins (NAGP); 9. O-Acetyl glycoproteins (OAGP); 10. glutamate; 11. citrate; 12. cysteine;
 286 13. creatine; 14. tyrosine; 15. trimethylamine-N-oxide (TMAO); 16. glycerophosphorylcholine
 287 (GPC); 17. taurine; 18. β -glucose; 19. α -glucose; 20. glycine; 21. glutamine; 22. lactate; 23.

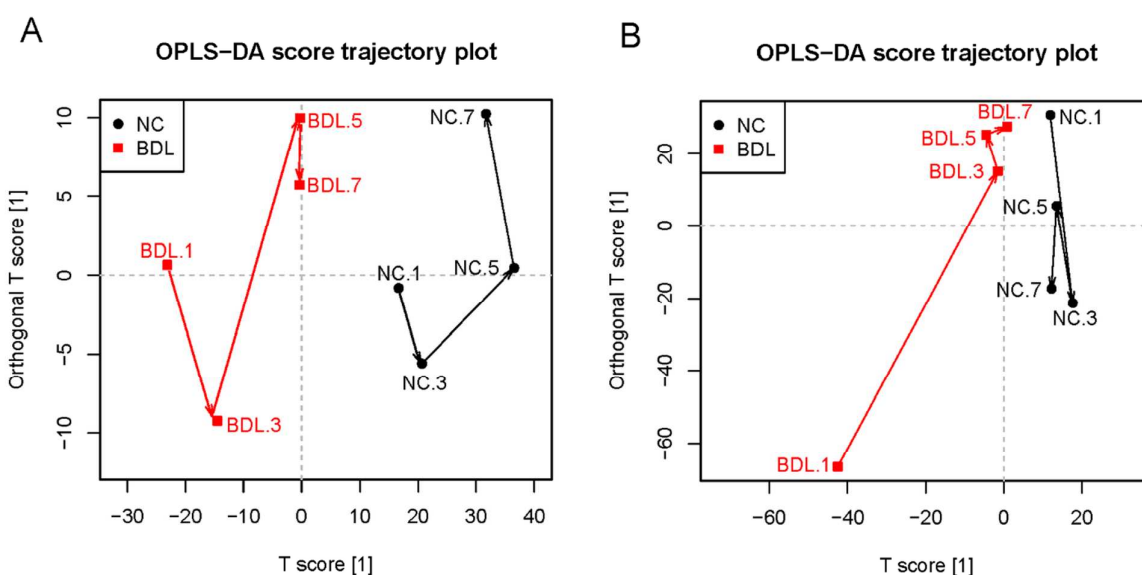
288 phenylalanine; 24. histidine; 25. formate. (B) Typical 500 MHz NOESY ^1H NMR spectra of the
289 urine samples from NC rats, BDL rats and BHD rats on week 1: 1. very low density lipoprotein
290 (VLDL) / low density lipoprotein (LDL); 2. isoleucine/leucine; 3. valine; 4. β -Hydroxybutyrate; 5.
291 α -hydroxyisobutyrate; 6. lactate; 7. alanine; 8. ornithine; 9. acetate; 10. proline; 11. N-Acetyl
292 glycoproteins (NAGP); 12. O-Acetyl glycoproteins (OAGP); 13. methionine; 14. adipate; 15.
293 acetoacetate; 16. glutamate; 17. oxalacetate; 18. succinate; 19. citrate; 20. dimethyl amine (DMA);
294 21. methylguanidine; 22. trimethyl amine (TMA); 23. dimethyl glycine (DMG); 24. pyruvate; 25.
295 2-oxoglutarate ; 26. creatinine; 27. choline; 28. phosphocholine; 29. trimethylamine-N-oxide
296 (TMAO); 30. taurine; 31. glycine; 32. creatine/phosphocreatine; 33. fumarate; 34.
297 3-hydroxymandelate; 35. 4-aminohippurate; 36. gallate; 37. phenylalanine; 38. tryptophan; 39.
298 benzoate; 40. hippurate; 41. formate; 42. nicotinate; 43. N-methylnicotinamide. 44. NAD^+ .

299

300 **3.5. OPLS-DA score trajectory plot at all time points**

301 The metabolic profiles were first subjected to PCA analysis. The three groups showed
302 partial separation from each other in PCA score plots (data not shown) due to the
303 unsupervised nature of PCA. The variations of ^1H NMR signals among groups were
304 not only arisen from interested grouping but also from other factors that were group
305 unrelated. To concentrate group discrimination into the first component and filter those
306 variations unrelated to class discrimination, supervised OPLS-DA was further
307 performed.¹⁶ The dynamic metabolic events in the rats were visualized by the
308 OPLS-DA score trajectory plots (Fig.2), where each spot represented the mean
309 position of each group at one time. The direction denoted by arrows represented the

310 trend of the changing metabolite pattern. The shift of the metabolic patterns in both
 311 serum and urine were similar in that BDL group showed the furthest deviation to NC
 312 group on week 1, and then moved towards the normal (Fig. 5 A and B). The score
 313 trajectory showed a distinct biphasic course of the BDL model. Data on week 1 (acute
 314 stage) and week 3 to week 7 (recovery stage) were analyzed independently.



315
 316 **Fig. 5** OPLS-DA score trajectory plots of rats serum (A) and urine (B) of different groups on week 1,
 317 3, 5 and 7 after BDL operation. Symbols of ● (black filled circles), ■ (red filled squares) represented
 318 NC and BDL group respectively.

319

320 3.6 ¹H NMR metabolomics profiles on week 1

321 3.6.1 Metabolic changes in BDL rats

322 On week 1, the BDL and NC group showed a complete separation in score plots of
 323 serum and urine (Fig. 6 A and C), suggesting a severe metabolic disturbance in BDL
 324 groups by a supervised OPLS-DA with a well goodness of fit ($R^2 = 0.93$, $Q^2 = 0.79$; R^2

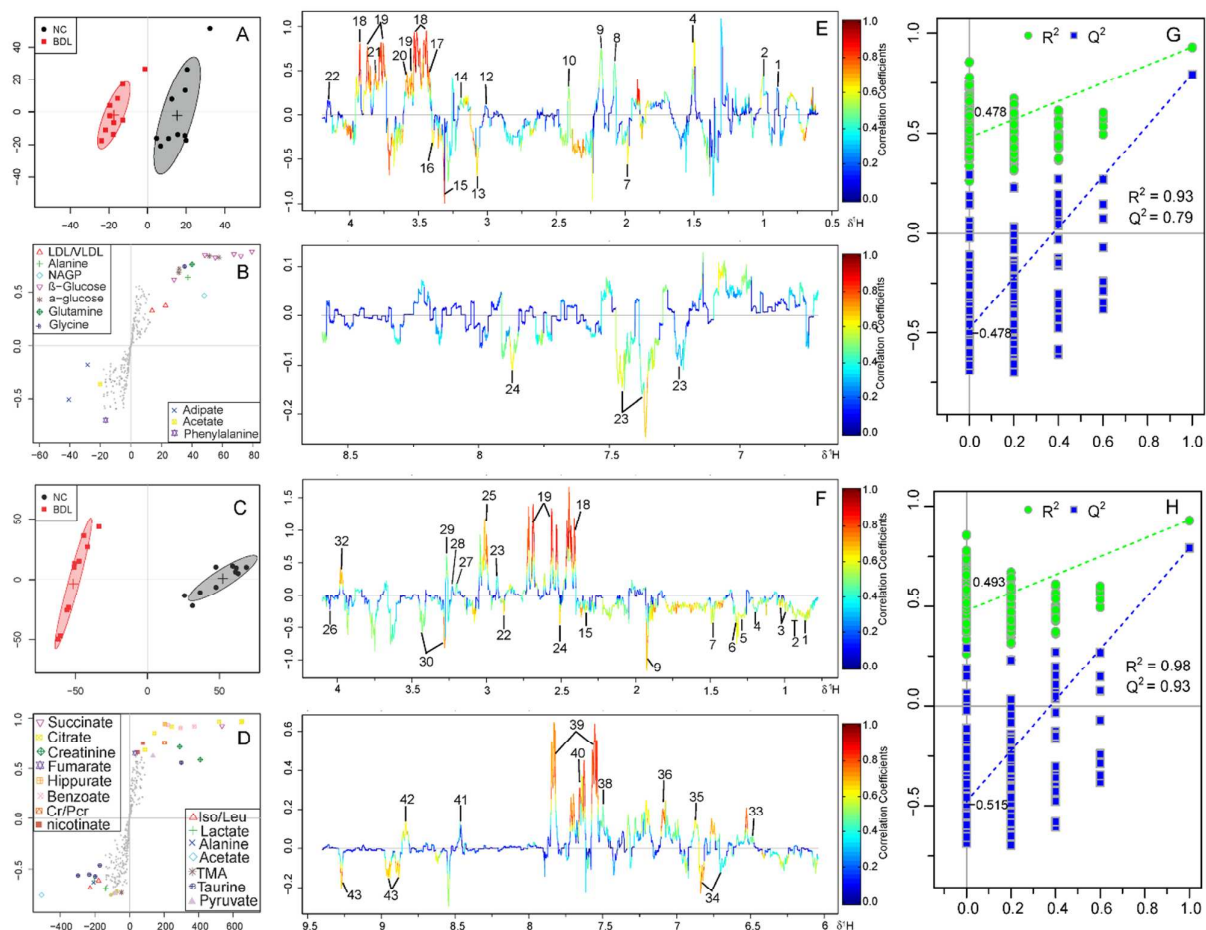
325 = 0.98, $Q^2 = 0.93$ for the serum and urine, respectively) displayed in Fig. 6 G and H.
326 The corresponding S-plots and loading plots revealed the differential metabolites in
327 serum and urine. In the S-plots, where points in different color and shape represented
328 variables (metabolites), the more further away from the center of a variable, the more
329 contribution of the variable to the class separation of the groups. The loading plots
330 were color-coded with the absolute value of correlation coefficients where red (high
331 coefficients) indicates more marked contribution to the separation than blue (low
332 coefficients) one. The S-plots (Fig.6 B and D) and loading plots (Fig.6 E and F)
333 revealed elevated levels of acetate, creatine, trimethylamine-N-oxide (TMAO),
334 glycerophosphorylcholine (GPC), histidine, phenylalanine in serum; elevated levels of
335 3-methylglutarate, α -hydroxyisovalerate, isoleucine, leucine, valine, isobutyrate,
336 α -hydroxyisobutyrate, alanine, acetate, trimethylamine (TMA), taurine,
337 N-methylnicotinamide, 3-hydroxymandelate, NAD^+ , pyruvate in urine, and lower
338 levels of High density lipoprotein (HDL), alanine, β -glucose, α -glucose, glycine in
339 serum; lower levels of succinate, citrate, dimethylglycine (DMG),
340 creatine/phosphocreatine, creatinine, 2-oxoglutarate, 4-aminohippurate, nicotinate,
341 gallate, hippurate, benzoate, acetylsicylate in urine of BDL rats. These important
342 differential metabolites were further tested for their between-group difference using
343 univariate analysis, and found to be mostly significant as visualized in the heat map
344 (Fig. 8) and fold change plots (Fig. S1).

345 **3.6.2 Effect of HLJDD on the metabolic profiles of BDL rats**

346 To explore the influence of HLJDD on BDL rats, 1H NMR data of serum and urine of

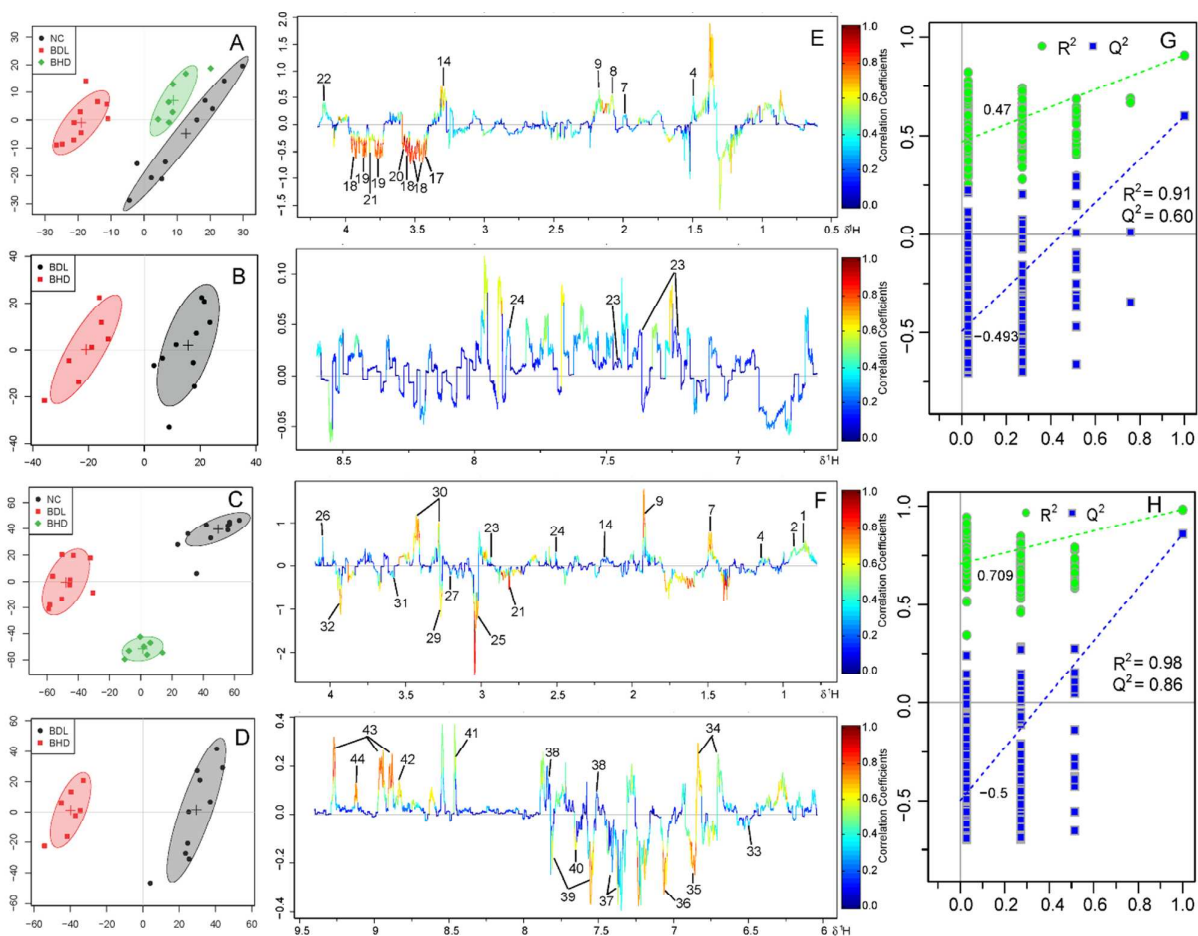
347 NC, BDL and BHD group were analyzed together. They were clearly separated in the
348 OPLS-DA score plots (Fig. 7 A and C), and BHD groups were in the middle and close
349 to the NC groups, declaring HLJDD can reduce the changes of metabolites caused by
350 the administration of BDL. To find out metabolites that were directly associated with
351 the treatment effect of HLJDD on BDL rats, the metabolic profiles of BDL and BHD
352 groups were analyzed by OPLS-DA (Fig. 7 B and D). The BHD rats were clearly
353 separated from BDL rats with a well goodness of fit (Fig. 7 G and H).

354 The loading plots (Fig. 7 E and F) revealed great increase of β -glucose, α -glucose,
355 glycine, caprate, glutamine in serum; higher levels of isobutyrate, N-Acetyl
356 glycoproteins (NAGP), O-Acetyl glycoproteins (OAGP), methylguanidine,
357 creatine/phosphocreatine, trimethylamine-N-oxide (TMAO), creatinine,
358 2-oxoglutarate, threonine, glycine, creatinine, lactate, acetoacetate, β -hydroxybutyrate
359 (3-HB), fumarate, phenylalanine, 4-aminohippurate, 2-oxoglutarate, gallate, NAD^+ ,
360 benzoate, threonine in urine, and lower levels of LDL, VLDL, adipate,
361 trimethylamine-N-oxide (TMAO) in serum; lower levels of α -hydroxyisovalerate,
362 isoleucine, leucine, valine, alanine, acetate, proline, adipate, dimethyl glycine (DMG),
363 taurine, tryptophan, hippurate, nicotinate, formate, N-methylnicotinamide,
364 3-hydroxymandelate, NAD^+ in urine of BHD groups. The important differential
365 metabolites selected based on loading plots of OPLS-DA, and found to be mostly
366 significant as visualized in the heat map (Fig. 8) and fold change plots (Fig. S1).



367

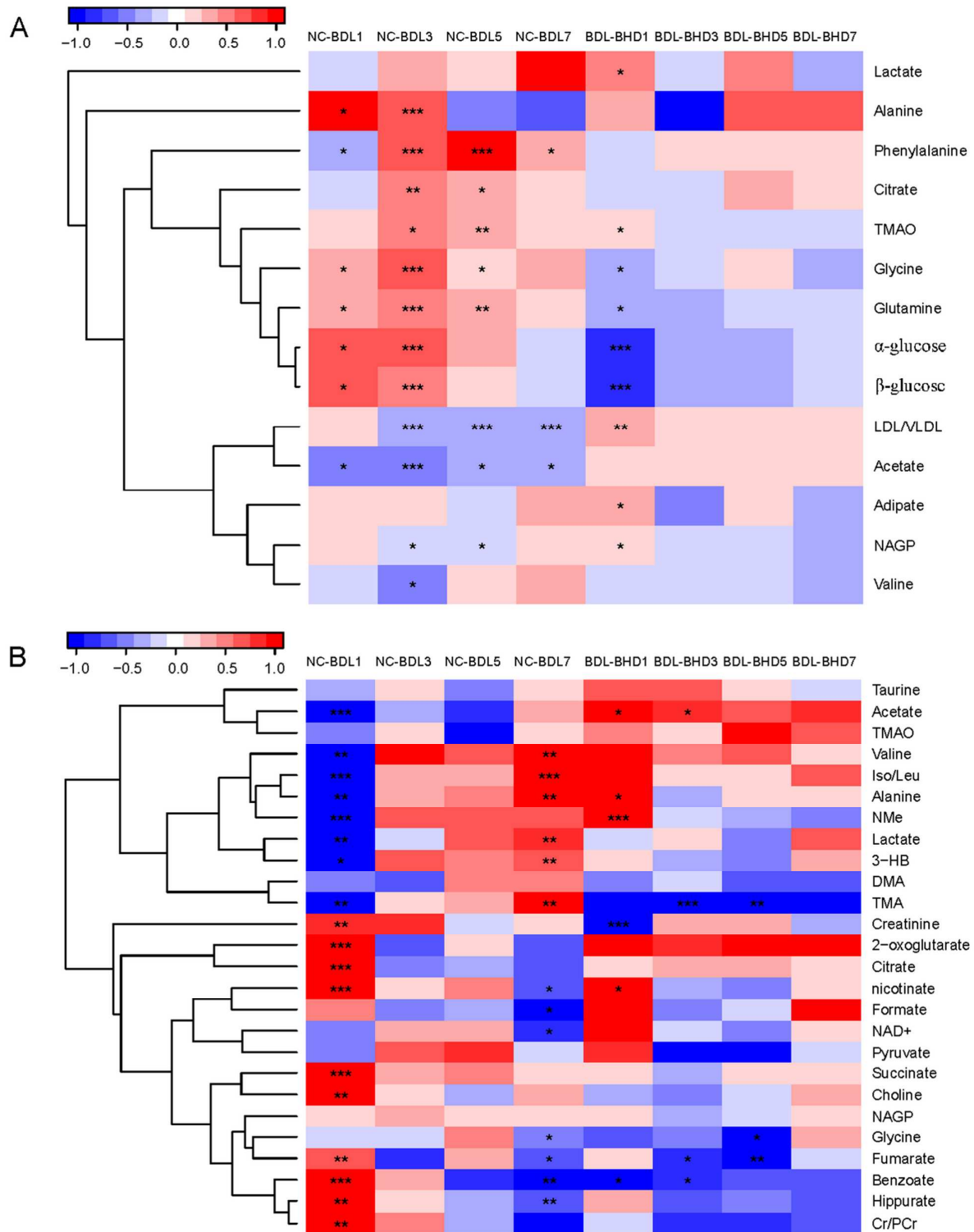
368 **Fig. 6** On week 1, cross-validated OPLS-DA scores plots (A for serum and C for urine), the
 369 corresponding S-plots (B for serum, D for urine) and loadings plots (E for serum, F for urine)
 370 derived from ^1H NMR spectra for NC, BDL and BHD rats. OPLS-DA scatter plot from serum (G)
 371 and urine (H) of the statistical validations obtained by 200 times permutation tests, with R^2 and Q^2
 372 values in the vertical axis, the correlation coefficients (between the permuted and true class) in the
 373 horizontal axis, and OLS line representing the regression of R^2 and Q^2 on the correlation coefficients.



374

375 **Fig. 7** On week 1, OPLS-DA analysis of ^1H NMR data in serum and urine for NC, BDL and BHD
 376 group. Scores plots on NC, BDL and BHD rats (A and C), and on BDL and BHD rats (B and D),
 377 loadings plots (E and F) for OPLS-DA. OPLS-DA scatter plot from serum (G) and urine (H) of the
 378 statistical validations obtained by 200 times permutation tests, with R^2 and Q^2 values in the vertical
 379 axis, the correlation coefficients (between the permuted and true class) in the horizontal axis, and
 380 OLS line representing the regression of R^2 and Q^2 on the correlation coefficients.

381



382

383 **Fig. 8** Heatmap visualization of the z-scored levels of metabolites in serum (A) and urine (B) with

384 stars denoting the differential significance. Row represent metabolites and column represent groups.

385 “BDL” and “BHD” mean “BDL group compare with NC group” and “BHD group compare with
386 BDL group”, the number “1”, “3”, “5” and “7” mean week one, three, five and seven. Color key
387 indicates metabolite quantities value, white: no significant change, deep blue: highest, deep red:
388 lowest, $P < 0.05$ represented statistically significant threshold. * $P < 0.05$, ** $P < 0.01$ and *** $P <$
389 0.001.

390

391 **3.7 ^1H NMR metabolomics profiles on week 3, 5, 7**

392 On week 3, 5 and 7, OPLS-DA method was carried out in the BDL, NC, BHD groups.
393 From the score plots of BDL and NC groups (Fig. S2 C and D), revealed a good
394 separation. Compared with NC group, these findings according to the corresponding
395 loading plots (Fig. S3 A, B and C) were observed in BDL group: elevated levels of
396 very low density lipoprotein (VLDL)/low density lipoprotein (LDL), caprate,
397 3-methyladipate/succinyl acetone in serum; elevated levels of citrate, 2-OG in urine,
398 and reduced levels of alanine, lysine, adipate, acetate, glycine in serum; reduced levels
399 of 3-methylglutarate, isoleucine, leucine, valine, α -hydroxyisobutyrate, glutamate in
400 urine of BDL groups. What's more, the OPLS-DA score plots of the serum and urine
401 (Fig. S2 A and B) showed a partial separation of BDL group from NC and BHD group,
402 suggesting damage caused by BDL is gradually restored on week 3, 5 and 7, and NC
403 and BHD group were much overlap in the score plots indicated HLJDD can reduce the
404 changes of metabolites caused by the administration of BDL. Most of the disturbances
405 of metabolites in BDL rats were reversed after treatment of HLJDD (Fig. S2 E,F and
406 Fig. S3 D, E, F).

407

408 **4 Discussion**

409 In this study, ¹H NMR-based metabolomics approach combined with clinical
410 chemistry and histopathology inspection was used to investigate cholestatic liver
411 damage caused by bile duct ligation (BDL) and the treatment effects of HLJDD on
412 BDL rats. Histopathology inspection indicated initial formation of fibrosis in the liver
413 which was testified by the increase of ALT, AST, ALP and decrease of ALB in serum
414 of BDL rats, and the increase of hyaluronic acid (HA), laminin (LN), type III
415 precollagen (PCIII) and type IV collagen (CIV). Slight edema of glomerular was
416 found in the kidney of BDL rats. OPLS-DA analyses of the serum and urinary NMR
417 data of the three groups on week 1, 3, 5 and 7 were performed. The metabolic profiles
418 in BDL rats were severely disturbed, the furthest away from the control on week 1
419 (acute phase) and then gradually recovered on week 3, 5 and 7 (chronic phase).

420 One week after BDL operation, rats showed a series of metabolic perturbations,
421 including energy and amino acid metabolism, oxidative stress and intestinal flora
422 disruption.

423 **4.1 Acute phase**

424 **4.1.1 Energy metabolism**

425 Significantly decreased levels of urinary citrate, 2-OG, succinate and serum β -glucose,
426 α -glucose together with slightly increased pyruvate level in urine were observed in
427 BDL group as compared with the NC group. Pyruvate is a key intermediate that takes
428 part in both glycolysis and the tricarboxylic acid (TCA) cycle. Generated by

429 decomposition of glucose, pyruvate can be converted into acetyl-CoA by
430 decarboxylation and enter the TCA cycle under aerobic conditions.²⁴ The increased
431 level of pyruvate in urine suggested a hampered conversion of pyruvate to acetyl-CoA,
432 which together with notably decreased other intermediates of TCA cycle, citrate, 2-OG
433 and succinate, suggested an inhibition of the TCA cycle.^{25, 26} TCA cycle is the most
434 efficient and major source of energy supply. Its inhibition caused energy deficiency, so
435 other means, such as glycolysis, come to rescue. By glycolysis, glucose is converted to
436 lactate by lactate dehydrogenase (LDH) or to alanine by alanine aminotransferase
437 (ALT), resulting in increased lactate and alanine levels.²⁷ The marked decrease of
438 serum glucose, and increase of lactate and alanine in serum of BDL rats demonstrated
439 an enhanced glycolysis in BDL group.

440 However, glycolysis is inefficient in energy production. Therefore, ketone bodies
441 metabolism, another means of energy production, have to be enhanced to ameliorate
442 the shortage of energy. Ketone bodies, including acetoacetate and 3-HB, are
443 well-known metabolites of fatty acids in liver mitochondria. Ketones could be
444 transported from serum to muscles, tissues and organs, where ketone bodies could be
445 oxidized to produce energy. The obvious increase of ketone bodies in urine as well as
446 slight decrease of LDL or VLDL in serum of BDL group, suggested an enhanced fatty
447 acid oxidation to produce ketone bodies. As a result, the concentration of acetate, the
448 end product of fatty acid oxidation, was significant increase in the BDL rats. Serum
449 creatine and urinary creatinine were increased in BDL rats. Creatine-phosphate can
450 transfer high-energy phosphate to ADP and produce ATP for energy demand.²⁸

451 Therefore, the significant increase of creatine and its degradation product creatinine
452 suggested a facilitated utilization of creatine-phosphate to replenish energy demand.
453 The increase of creatine and creatinine was also deemed as a sign of hepatic injury,²⁴
454 in consistent with pathological result in this study.

455 BDL rats after HLJDD treatment exhibited markedly enhanced levels of 2-OG, and
456 slightly decreased amount of pyruvate, alanine, 3-HB and acetate, which showcased
457 the ability of HLJDD to ameliorate the disturbed energy supply induced by BDL,
458 possibly by restoration of TCA cycle.

459 **4.1.2 Oxidative stress**

460 Bile acids (BAs) were secreted by hepatocytes, transported into the extracellular
461 matrix in liver and then excreted to duodenum for the absorption and solubilization of
462 dietary lipids. BDL prevented bile flow from liver to duodenum, leading to an
463 accumulation of BAs in liver that generate reactive oxygen species (ROS).²⁹ The
464 imbalance between the generation of ROS and antioxidant defenses induce oxidative
465 stress.³⁰ We observed significantly decreased levels of glutamine and glycine and
466 slightly decreased levels of glutamate and cysteine in serum, the precursors of GSH, in
467 BDL rats. GSH as a major natural antioxidant, can react with free radicals directly,³¹
468 thus resisting the damage caused by ROS. The lowered levels of these GSH precursors
469 demonstrated an accelerated GSH synthesis as a consequence of excessive depletion
470 of GSH to counteract ROS.³²

471 Urinary choline and phosphocholine were decreased obviously in BDL group
472 compared with the NC group. Phospholipid, consisting of choline and phosphocholine,

473 is the major component of cell membrane and essential for the maintenance of its
474 integrity.³³ ROS could attack membrane phospholipids, leading to the damage on the
475 construction and function of membranes, and ultimate the rupture of cell and
476 organelles, such as mitochondria.³⁴ As an evidence, serum levels of NAGP and OAGP
477 were decreased slightly in BDL group since that they were synthesized in membranes
478 of endoplasmic reticulum and golgi apparatus. The lowered levels of choline and
479 phosphocholine indicated an accelerated use of them to renovate the membranes
480 damaged by ROS, thus representing a self-repair mechanism.

481 HLJDD significantly decreased the elevated urinary level of taurine, and markedly
482 increased the lower serum levels of glutamine, glycine and taurine and urinary choline
483 and phosphocholine in BDL rats, showcasing its protection on BDL induced oxidative
484 injury.

485 **4.1.3 Amino acid metabolism**

486 Taurine and glycine were decreased in serum but increased in urine of BDL rats. BAs
487 could be conjugated to either taurine or glycine in order to reduce the toxicity caused
488 by accumulated BAs. The conjugated BAs were hydrolyzed into free BAs, liberating
489 taurine or glycine under the activation of intestinal bacteria. Normally, most BAs were
490 reabsorbed and, taurine and glycine were excreted into urine.³⁵ Therefore, the opposite
491 change of taurine and glycine in serum and urine of BDL rats demonstrated
492 accelerated their conjugation with BAs to attenuate the toxicity of accumulated BAs.

493 Leucine, isoleucine and valine (branched-chain amino acids, BCAAs) were significant
494 increased in urine of BDL rats. ROS induced the decomposition of proteins, resulting

495 in the damage of cell membrane.³⁶ BCAAs are important precursors for protein
496 synthesis,³⁷ and thus essential for the repairment of the damaged cell membranes. The
497 increased excretion of BCAAs in urine of BDL rats showed either an inhibited protein
498 synthesis or an enhanced protein degradation.³⁸ In addition, these BCAAs can be
499 reabsorbed by glomerular in normal status and their increase may also suggested the
500 dysfunction of glomeruli reabsorption,^{39,40} which also supported by histopathological
501 examination.

502 Phenylalanine was markedly increased and tyrosine was slightly decreased in serum of
503 BDL rats. phenylalanine is an essential amino acid that has to be obtained from food
504 directly. Tyrosine is a semi-essential amino acid as it can only be synthesized by the
505 hydroxylation of phenylalanine under the catalysis of phenylalanine hydroxylase
506 (PAH). The increased conversion of phenylalanine to tyrosine ratio in BDL group has
507 been observed in patients incurred with hepatitis C virus suffering hepatic damage,⁴¹
508 and thus may also indicated liver injury induced by BDL operation.

509 The increased urinary BCAAs, and enhanced conversion of Phe to tyrosine ratio due
510 to BDL operation exhibited a downward trend towards a normal status by the
511 intervention of HLJDD, showing its ability to ameliorate amino acid metabolism and
512 protect rats from BDL induced liver injury.²⁶

513 **4.1.4 Intestinal flora metabolism**

514 The levels of hippurate and benzoate were significantly decreased in urine of BDL rats.
515 Benzoate was synthesized from plant phenolics and aromatic amino acids, by
516 intestinal microflora.⁴² Benzoate could be absorbed by intestinal tract, eventually

517 entering into the liver through the portal vein by systemic circulation. BDL prevented
518 the entrance of BAs to intestinal tract, leading to deficiency of intestinal bile salts, and
519 inevitably alteration of gut bacteria,⁴³ as supported by the obvious decrease of
520 benzoate in urine of BDL rats. As a sequence, the synthesis of hippurate was decreased
521 in BDL group since that hippurate was synthesized by conjugation of glycine with
522 benzoate in the mitochondrial matrix of liver^{44, 45} A series of studies have also
523 concluded that the decrease of urinary levels of benzoate and hippurate could be
524 ascribed to the disruption of intestinal flora.⁴⁶⁻⁴⁸

525 The disturbance of gut microbes could also be evidenced by the observed significant
526 increase of trimethylamine (TMA) and partial decrease of trimethylamine (TMAO),
527 since that TMAO is the oxidation product of TMA through the action of gut
528 microbes.¹⁷

529 HLJDD significantly increased the levels of urinary TMAO and hippurate,
530 demonstrating a great amelioration of gut microbiota metabolism by HLJDD.

531 **4.2 Chronic cholestasis phase**

532 The severe disturbance in energy metabolism, amino acid metabolism, oxidative stress
533 and gut bacteria in acute phase was greatly attenuated in the long run, characteristic of
534 BDL model. The BDL rats employed a self-repairing process with compensation to
535 address the hampered bile flow by construction of the bypass of bile ducts, which was
536 fully established in BDL rats on week 7.⁴⁹

537 Interestingly, with no good reasons, levels of urinary citrate, 2-OG and adipate in BDL
538 group were significantly higher than those in NC group, suggesting the complex of the

539 body in response to pathological changes. Surprisingly, these metabolites could also be
540 reversed towards normal levels by HLJDD, in opposite direction to its performance in
541 acute phase, exhibiting a bilateral adjustment of HLJDD.

542 **5 Conclusion**

543 ¹H NMR-based metabolomics approach was applied to explore global metabolic
544 features in serum and urine of BDL-induced cholestasis in rats and the treatment
545 effects of HLJDD. The metabolomic pattern showed a distinct biphasic feature of BDL
546 model: acute phase (week 1) and chronic phase (week 3-7). BDL brought severe
547 disturbance in energy and amino acid metabolism, alteration of intestinal flora and
548 oxidative stress in acute phase. The metabolomic results combined with clinical
549 chemistry indicated conspicuous liver dysfunction and the damage of renal glomerular
550 function at acute phase, which were greatly attenuated at chronic phase due to the
551 self-protection mechanism of the body. HLJDD showed bilateral adjustment of some
552 metabolism disturbance. These results demonstrated sensitivity and superiority of
553 metabonomics in disease subtypes and diagnosis, and in the understanding of complex
554 mechanism of a Chinese herbal medicine formula. This integrated metabolomics
555 approach might help to develop a systematic view of BDL-induced injury process and
556 assess its therapy.

557

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