

1 **Steroid Metabolomic Signature of Liver Disease in Nonsyndromic Childhood Obesity**

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34 Abstract

35 **Objective:** Analysis of steroids by gas chromatography-mass spectrometry (GC-MS) defines
36 a subject's steroidal fingerprint. Here, we compare the steroidal fingerprints of obese children
37 with or without liver disease to identify the 'steroid metabolomic signature' of childhood
38 non-alcoholic fatty liver disease.

39 **Methods:** Urinary samples of 85 children age 8.5-18.0 with BMI>97% were quantified for
40 31 steroid metabolites by GC-MS. The fingerprints of 21 children with liver disease (L1) as
41 assessed by sonographic steatosis (L1^L), elevated alanine aminotransferases (L1^A) or both
42 (L1^{AL}), were compared to 64 children without markers of liver disease (L0). The steroidal
43 signature of the liver disease was generated as the difference in profiles of L1 against L0
44 groups.

45 **Results:** L1 comparing to L0 presented higher fasting triglycerides (p=0.004), insulin
46 (p=0.002), INS/GLU (p=0.003), HOMA-IR (p=0.002), GGTP (p=0.006), AST/SGOT
47 (p=0.002), postprandial glucose (p=0.001) and insulin (p=0.011). L1^{AL} showed highest level
48 of T-cholesterol and triglycerides (p=0.029; p=0.044). Fasting insulin, postprandial glucose,
49 INS/GLU and HOMA-IR were highest in L1^L and L1^{AL} (p=0.001; p=0.017; p=0.001;
50 p=0.001). The liver disease steroidal signature was marked by lower DHEA and its
51 metabolites, higher glucocorticoids (mostly tetrahydrocortisone) and lower mineralocorticoid
52 metabolites than L0. L1 patients showed higher 5 α -reductase and 21-hydroxylase activity
53 (the highest in L1^A&L1^{AL}) and lower activity of 11 β HSD1 than L0 (p=0.041, p=0.009,
54 p=0.019). **Conclusions:** The 'steroid metabolomic signature' of liver disease in childhood
55 obesity provides a new approach to the diagnosis and further understanding of its metabolic
56 consequences. It reflects the derangements of steroid metabolism in NAFLD that includes
57 enhanced glucocorticoids and deranged androgens and mineralocorticoids.

59 **Introduction**

60 Nonsyndromic childhood obesity is associated with nonalcoholic fatty liver disease
61 (NAFLD), a spectrum of conditions, ranging from steatosis to nonalcoholic steatohepatitis
62 (NASH), and various degrees of fibrosis and cirrhosis (1). NAFLD is regarded as the hepatic
63 manifestation of the metabolic syndrome (2). However, childhood obesity with no NAFLD is
64 also complicated by the metabolic syndrome. Despite the growth of knowledge regarding
65 obesity-related NAFLD in children, we still rely mostly on circulating levels of liver enzymes
66 and ultrasonography imaging and some non-invasive tests (3,4,5,6,7). Liver biopsy in
67 children with suspected NAFLD is recommended only for “those where the diagnosis is
68 unclear, where there is possibility of multiple diagnoses, or before starting therapy with
69 potentially hepatotoxic medications” (8).

70 The consequences of obesity-related NAFLD on liver metabolism are insufficiently
71 understood (8). As steroid hormones are partially catabolized and conjugated by liver
72 enzymes, we have anticipated that NAFLD would have its metabolic impact on steroid
73 metabolism.

74 Here, we utilized our previously reported concept, arguing that an individual’s urinary steroid
75 metabolite profile represents a subject’s unique metabolic fingerprint and offers means of
76 metabolomic phenotyping at the individual level (9,10). Thus, each individual has a unique
77 ‘steroidal fingerprint’. A cluster of similar “steroidal fingerprints” related to a disease would
78 be regarded as a ‘steroid metabolomic disease signature’ (10,11), which represents the impact
79 of a disease in people who differ in their phenotypes or have other health problems. We have
80 previously clustered steroidal fingerprints of children with nonsyndromic obesity into five
81 clusters with distinctive steroidal signatures (11).

82 Here, we analyzed the clinical data of a group of 85 patients with well-phenotyped non-
83 syndromic childhood obesity and defined those affected and those unaffected by NAFLD
84 and/or elevated activities of liver enzymes. We generated steroidal disease signatures of the
85 two groups and suggest that it might shed light on steroid-related metabolic sequelae of liver
86 disease in childhood obesity.

87

88 **Subjects and Methods**

89 Between March 2012 and August 2013, we examined a consecutive series of 117 obese
90 Caucasians children and adolescents (BMI > 97th centile). They were recruited from the
91 patients referred to the Department of Pediatric Endocrinology, Medical University of Silesia,
92 Katowice, Poland. After exclusion of younger participants (<8 years), syndromic obesity,
93 chronic diseases, pharmacotherapy (also metformin) or precocious puberty, we included the
94 remaining 85 patients (43 girls), aged 8.5-18.0 years (mean age 14.4, SD 2.33, median 14.5
95 years).

96 All patients underwent a clinical assessment and diagnostic procedures that included general
97 physical examination, anthropometric measurements of height, weight, waist and hip
98 circumference and puberty assessment, as previously described (11). Morning fasting venous
99 blood samples were collected to measure lipids, glucose (GLU), insulin (INS), TSH, fT4,
100 cortisol and aminotransferases. Plasma total cholesterol (T Chol), high-density lipoprotein
101 cholesterol (HDL-Chol) and triglyceride (TG) levels were analyzed enzymatically (Beckman
102 Coulter, USA). GLU and INS levels were also measured in an oral glucose tolerance test
103 (OGTT, 1.75 g/kg, max 75 g). Enzymatic test (hexokinase method) was used for the
104 quantitative determination of glucose (Beckman Coulter, USA). INS was determined using a
105 chemiluminescence immunoassay on Immulite 2000 analyzer (DPC, USA). Fasting

106 INS/GLU ratio (FIGR) and homeostatic model assessment of INS resistance (R-HOMA,
107 fasting GLU [mmol/L] x fasting INS [mIU/L]/22.5 were calculated as indices of insulin
108 resistance. Cortisol was measured in the morning (8am) and midnight using
109 chemiluminescent immunoassay by Immulite 2000 analyzer (DPC, USA). Serum
110 concentrations of FT4 and TSH were measured with a chemiluminescent immunometric assay
111 (Siemens, Immulite 2000 Free T4 , Immulite 2000 Third Generation TSH, USA). Gamma-
112 glutamyl transpeptidase (GGTP), alanine (ALT/GPT) and aspartate aminotransferases
113 (AST/SGOT) activity in the serum were assessed according to International Federation in
114 Clinical Chemistry (Beckman Coulter, USA).

115

116 *Assessment of Liver Disease / NAFLD*

117 Abdomen ultrasonography to evaluate the liver for hepatic steatosis features was performed
118 with 5 MHz convex transducer (Logiq 5, GE Healthcare GmbH, Germany), according to
119 standards in pediatric population. (12). The evidence of hepatic steatosis by abdominal
120 ultrasound (hepatic echogenicity increased above the echogenicity of the adjacent right renal
121 cortex and increase in fine echoes of liver parenchyma compared with intrahepatic vessel
122 borders (7,13)), and no causes for secondary hepatic fat accumulation, defined here
123 nonalcoholic fatty liver disease (NAFLD) diagnosis (5,7,12). Any markers of liver
124 dysfunction (elevated ALT >45 U/L – L1^A, NAFLD based on ultrasonography – L1^L, or both
125 – L1^{AL}) were defined as liver disease – L1 as compared to L0 – without markers of liver
126 disease.

127 *Gas chromatography-mass spectrometry (GC-MS) of urinary steroids*

128 Steroid metabolites in 24-h-urine samples were analyzed by quantitative targeted GC-MS
129 (9,10,11). Briefly, free and conjugated urinary steroids were extracted by solid phase
130 extraction and conjugates were enzymatically hydrolyzed. After recovery of hydrolyzed
131 steroids by solid phase extraction, known amounts of internal standards (5 α -androstane-
132 3 α ,17 α -diol, stigmasterol) were added to each extract before formation of methyloxime-
133 trimethylsilyl ethers. GC was performed using an Optima-1 fused silica column (Macherey-
134 Nagel, Dueren, Germany) housed in an Agilent Technologies 6890 series GC that was
135 directly interfaced to an Agilent Technologies 5975 inert XL mass selective detector. After
136 calibration, values for the excretion of individual steroids were determined by measuring the
137 selected ion peak areas against the internal standard areas.

138 Steroid metabolites' ratios, as described in our previous paper (11), were used to calculate the
139 activity of the enzymes: 5 α reductase (An/Et; 5 α THF/THF, 5 α THB/THB), 11 β -
140 hydroxysteroid dehydrogenase type 1 -11 β HSD1 ([THF+ α THF]/THE), 3 β -hydroxysteroid
141 dehydrogenase- 3 β HSD ([THE+THF+ α THF]/P5T-17 α) and 21-hydroxylase
142 ([THE+THF+ α THF]/PT, [THE+THF+ α THF]/ PO5 α 3 α)

143

144 The study was conducted according to Helsinki declaration, and approved by the Ethics
145 Committee of the Medical University of Silesia. Informed consent was obtained from each
146 patient over age 16, a parent or a legal guardian, after full explanation of the purpose and
147 nature of all procedures.

148

149 *Statistical analysis and visualization of metabolomic data.*

150 Steroid metabolites quantities were z-transformed based on sex and age-adjusted normal
151 reference groups, as described elsewhere (11). Per each of 31 z-transformed steroid
152 metabolites and per each one of the groups L0, L1 and subgroups L1^A, L1^L, L1^{AL} the median
153 was computed. The standard R-function 'matplot' (<https://www.R-project.org/>) (14) was used
154 to depict the steroidal signatures of each subgroup as the difference between the above
155 medians versus the median of L0 group (11).

156 Clinical and chemical data as well as steroid metabolites' concentrations ratios of patients in
157 each group were analyzed, and t-student test, t-test with separate variance estimation,
158 ANOVA or Kruskal-Wallis ANOVA where appropriate were utilized to assess the difference
159 between groups. p-value <0.05 was considered statistically significant.

160

161 **Results**

162 *Clinical phenotype*

163 Out of 85 obese children, a liver disease was diagnosed in 21 (21/85, 24.7%; L1); in 5
164 patients by elevated ALT activity (L1^A), in 10 by sonographic liver steatosis (L1^L) and in 6
165 by both markers (L1^{AL}). The clinical phenotype is presented in Tables 1a and 2a.

166 The mean age, BMI, BMI z-score, hSDS, waist and hip circumference as well as blood
167 pressure values were not significantly different between patients of groups L0 and L1 and
168 among L0, L1^A, L1^L, L1^{AL} subgroups. There were relatively more males in L1 group than in
169 L0 (14/21, 67% vs. 28/64, 44%, Table 1a).

170 *Chemical phenotype*

171 At the biochemical level, patients of L1 comparing to L0 group presented higher
172 concentration of fasting triglycerides and insulin, postprandial glucose and insulin. Both
173 indices of insulin resistance, insulin/glucose ratio (INS/GLU) and HOMA-IR, GGTP,
174 AST/SGOT activity were significantly higher in L1 group. Comparison of three liver affected
175 subgroups L1^A, L1^L, L1^{AL} and L0 (ANOVA) confirmed significantly the highest level of T-
176 cholesterol and triglycerides in L1^{AL} patients. Fasting INS and postprandial GLU levels were
177 higher in L1^L and L1^{AL} patients than in L1^A and L0, postprandial INS was the highest in L1^L
178 group. INS/GLU ratio and HOMA-IR values were also the highest in both groups with liver
179 steatosis features in the ultrasonography - L1^L and L1^{AL}. The highest GGTP and AST/SGOT
180 activities were observed in L1^A group (Table 2b).

181 *Steroid signature of liver disease*

182 Obese patients of the L0 group presented higher midnight plasma cortisol concentration
183 ($p < 0.001$) than the L1 group (Table 1b). Comparing the z-transformed values of steroid
184 metabolites, significantly higher tetrahydrocortisone (THE) concentration were found in L1
185 group ($p = 0.046$).

186 ‘Steroidal signature’ of liver disease is presented as the difference between z-transformed
187 concentrations of steroid metabolites in L0 and L1 patients (Figure 1). Liver-affected patients
188 have shown significantly enhanced 5 α -reductase and 21-hydroxylase activity and lower
189 activity of 11 β HSD1 than L0 subjects (Table 3).

190 ‘Steroid metabolomic disease signature’ of L1^A, L1^L, L1^{AL} are presented in Figure 2 a, b
191 &c). Liver affected patients L1^L presented significantly enhanced activity of 21-hydroxylase,
192 and those with elevated ALT (L1^A & L1^{AL}) showed enhanced 5 α -reductase activity (Table 4).

193 **Discussion**

194 Based on our previous definition of ‘steroid metabolomic disease signature’ by quantitative
195 urinary steroidal GC-MS data (10,11), here we define the steroidal signature of liver disease
196 in non-syndromic childhood obesity.

197 The results emphasize the fact that the clinical picture of obese children with liver disease is
198 not different from that of obese children with no liver disease; they have comparable age,
199 height, weight, BMI, waist and hip circumference and blood pressure. They have comparable
200 serum TSH and 8 am cortisol, while their midnight cortisol is lower. Those with liver disease
201 have higher circulating triglycerides, though their lipoproteins are comparable, as previously
202 reported (15). We confirm that obesity and insulin resistance play important roles in the
203 development of NAFLD (16). The insulin sensitivity indices of obese children with liver
204 disease are marked by higher postprandial glucose and insulin, higher insulin/glucose ratio
205 and higher HOMA-IR (17) than those with no liver disease.

206 This complex ‘steroidal signature’ of liver disease reflects previously published single
207 observations. The steroidal disease signature is marked by low urinary DHEA (18,19) and its
208 metabolites, higher glucocorticoid metabolites, due to increased glucocorticoid production
209 rate (20), and lower mineralocorticoid metabolites. It is characterized by derangement of the
210 cortisol/cortisone shuttle generated by 11β hydroxysteroid dehydrogenase (HSD) type 1 (20),
211 as is evident from the lower $(\text{THF}+\alpha\text{THF})/\text{THE}$ ratio, enhanced $3\beta\text{HSD}$ activity
212 $([\text{THE}+\text{THF}+\alpha\text{THF}]/5\text{PT}-17\alpha \text{ ratio})$ and enhanced 21-hydroxylase activity
213 $([\text{THE}+\text{THF}+\alpha\text{THF}]/\text{PT})$. These findings may suggest lesser hepatic recycling (reduction) of
214 cortisone to cortisol in liver steatosis, which is compensated for by increased adrenal cortisol
215 generation and further metabolic consequences resulting from higher glucocorticoids
216 concentrations - this mechanism resulting in a model of a vicious circle.

217 Therefore, it is not surprising that higher tetrahydrocortisone concentration in L1 patients
218 corresponds with unfavorable biochemical profile: higher triglycerides and insulin resistance.
219 The clinical profile, however, defined by BMI z-score or waist circumference, is not useful in
220 the prediction of liver disease as well as other obesity complications.

221 A previous study focused on the measurement of circulating DHEAS, and found low DHEAS
222 in NASH patient. The authors assumed that this might have resulted from reduced
223 sulphonation of DHEA (19). Low sulphonation of steroids has been also found in a study in
224 obese children (21). A further important feature of the obesity-associated liver disease
225 signature is the low urinary DHEA excretion rate and its metabolites. It was previously
226 suggested that DHEA treatment reduced hepatic injury in experimental animals by inhibiting
227 several inflammatory mediators such as tumor necrosis factor- α and macrophage mitogen
228 inhibitory factor, and preventing the increase in serum ALT levels (22). Thus, we speculate
229 that DHEA might have a protective effect against hepatotoxicity. It has been shown that
230 DHEA inhibits 11 β -hydroxysteroid dehydrogenase -1 expression in liver and adipose tissues
231 (23) - another component of the steroidal signature. The liver is also the site of greatest
232 activity of 11 β HSD (24), and as such responds to liver disease with decreasing activity.
233 Obesity *per se* tends to enhance 11 β HSD-1 activity (25), but insulin resistance, a prominent
234 manifestation of the metabolic syndrome in obesity and the group of children presented here,
235 inhibits 11 β HSD-1 activity (26). Moreover, insulin resistance and the metabolic syndrome are
236 involved in the development and progression of NAFLD (15).

237 In conclusion, we present the ‘disease signature’ of liver disease in childhood obesity. We are
238 aware of the limitations of our study as our results may be biased by observational cross-
239 sectional character of the study and the relatively small number of participants in subgroups
240 with liver dysfunction. Moreover, we did not quantify the ultrasonographic steatosis, other
241 than visually. However, our findings suggest a new approach to the diagnosis and further

242 understanding of the metabolic consequences of liver disease as part of the metabolic
243 syndrome of obesity. They reflect the derangements of steroid metabolism in NAFLD that
244 includes enhanced glucocorticoid production and deranged androgens and
245 mineralocorticoids, and suggests a protective effect of DHEA on the liver in childhood
246 obesity. Knowledge of these sequels may provide ways for personalized medicine in obese
247 children with liver disease. Future prospective intervention study is also needed to verify if
248 obtained findings are only reversible consequences of obesity or whether they reflect non-
249 modifiable individual genetic predisposition.

250

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257 **Author contribution statement**

258 Conceptualization, AG, ZH, MS, MFH and SAW.; Methodology, ZH, AG, MS, MFH, SAW,
259 ZO and KG.; Software, MS and AG.; Validation, AG and MS.; Formal Analysis, MS;
260 Investigation, AG.; Resources AG and SAW; Data Curation, AG.; Writing – Original Draft
261 Preparation, ZH, AG, MS, MFH and SAW.; Writing – Review & Editing, ZH, AG, MS,
262 MFH, SAW, ZO, KG.; Visualization, ZO and KG.; Supervision, ZH and AG.; Project
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268

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362

363

365 **Figure legends**

366 Figure 1. Steroidal signature of liver disease in childhood obesity: differences between z-
367 transformed concentrations of steroid metabolites (androgens, glucocorticoids and
368 mineralocorticoids) in liver diseases (L1) and with non-liver disease features (L0) patients.

369

370 Figure 2 Steroidal signatures of liver disease in childhood obesity: differences between z-
371 transformed concentrations of steroid metabolites (androgens, glucocorticoids and
372 mineralocorticoids) in non-liver disease features (L0) patients and (a) patients with ALT+
373 (L1^A), (b) patients with sonographic liver steatosis (L1^L) and (c) patients with ALT+ and
374 sonographic liver steatosis (L1^{AL})

375

Table 1 Comparison of clinical (a) and chemical (b) phenotype of non-liver disease patients (L0) and patients with liver disease features (L1)

Legend:; L0 – non-liver disease patients. L1- liver disease patients (ALT+ or sonographic liver steatosis or both). f-female; m – male; hSDS- height standard deviation score; BMI – body mass index; WHR- waist to hip ratio; BP – blood pressure; TSH - thyroid-stimulating hormone; TChol – total cholesterol; HDL-Chol – HDL-cholesterol; TG- triglycerides; GLU- glucose; INS- insulin; ALT/GPT -alanine aminotransferases; AST/SGOT - aspartate aminotransferases; GGTP - Gamma-glutamyl transpeptidase; NS- not significant. Values are means and SD; Significance by t-student test.

| | Mean L0 | SD L0 | Mean L1 | SD L1 | <i>p-value</i> |
|-------------------------------|----------------|--------------|----------------|--------------|-----------------------|
| | (n=64) | | (n=21) | | |
| <i>a. Clinical phenotype</i> | | | | | |
| sex [f/m] | 36/28 | | 7/14 | | <i>0.001</i> |
| age [years] | 14.4 | 2.3 | 14.0 | 2.4 | NS |
| weight [kg] | 87.0 | 20.0 | 91.2 | 21.8 | NS |
| height [cm] | 164.0 | 11.4 | 166.1 | 11.5 | NS |
| hSDS | 0.4 | 1.4 | 0.8 | 1.6 | NS |
| BMI [kg/m²] | 32.0 | 5.2 | 32.8 | 5.4 | NS |

| | | | | | |
|-----------------------------|-------|------|-------|------|-------|
| BMI- z score IOTF | 2.7 | 0.5 | 2.8 | 0.5 | NS |
| waist [cm] | 100.6 | 11.5 | 102.1 | 11.6 | NS |
| WHR | 0.9 | 0.1 | 1.0 | 0.0 | 0.022 |
| BP systolic [mm Hg] | 125 | 11 | 130 | 12 | NS |
| BP diastolic [mm Hg] | 78 | 8 | 76 | 10 | NS |

| | Mean L0 | SD L0 | Mean L1 | SD L1 | <i>p-value</i> |
|-------------------------------|----------------|--------------|----------------|--------------|-----------------------|
| | (n=64) | | (n=21) | | |
| <i>b. Chemical phenotype</i> | | | | | |
| TSH [μIU/ml] | 2.7 | 1.2 | 3.0 | 1.3 | NS |
| cortisol 8.00 [μg/dl] | 17.4 | 6.2 | 19.7 | 6.1 | NS |
| cortisol 24.00 [μg/dl] | 3.4 | 3.9 | 1.6 | 1.0 | <0.001 |
| T Chol [mg/dl] | 171 | 35 | 178 | 31 | NS |
| HDL-Chol [mg/dl] | 50 | 10 | 46 | 12 | NS |
| TG [mg/dl] | 134 | 60 | 182 | 76 | 0.004 |
| GLU 0' [mg/dl] | 90 | 9 | 92 | 10 | NS |

| | | | | | |
|---|--------------|------|--------------|-------|------------------|
| GLU 120' [mg/dl] | 112 | 20 | 129 | 21 | <i>0.001</i> |
| INS 0' [μIU/ml] | 16.7 | 9.2 | 35.1 | 24.1 | <i>0.002</i> |
| INS 120' [μIU/ml] | 91.6 | 67.9 | 163.6 | 114.2 | <i>0.011</i> |
| INS/GLU 0' | 0.2 | 0.1 | 0.4 | 0.3 | <i>0.003</i> |
| INS/GLU 0' [%>0.3] | 10/64, 15.6% | | 10/21, 47.6% | | <i>0.004</i> |
| HOMA-IR | 3.7 | 2.2 | 8.1 | 5.6 | <i>0.002</i> |
| ALT/GPT [U/l] | 24 | 9 | 54 | 29 | <i><0.001</i> |
| AST/SGOT [U/l] | 25 | 8 | 36 | 13 | <i>0.002</i> |
| GGTP [U/l] | 22 | 10 | 33 | 12 | <i>0.006</i> |

Table 2 Comparison of clinical (a) and chemical (b) phenotype of non-liver disease patients (L0) and patients with liver disease features: L1^A (ALT+), L1^L (sonographic liver steatosis) and L1^{LA} (both ALT+ and sonographic liver steatosis).

Legend: L0 – non-liver disease patients. L1- liver disease patients. L1^A – patients L1 with ALT+. L1^L – patients L1 with sonographic liver steatosis. L1^{AL} – patients with ALT+ and sonographic liver steatosis); f- female; m-male; hSDS- height standard deviation score; BMI – body mass index; WHR- waist to hip ratio; BP – blood pressure; TSH - thyroid-stimulating hormone; TChol – total cholesterol; HDL-Chol – HDL-cholesterol; TG- triglicerydes; GLU- glucose; INS- insulin; ALT/GPT -alanine aminotransferases; AST/SGOT - aspartate aminotransferases; GGTP - Gamma-glutamyl transpeptidase; NS- not significant. Values are means and SD, Significance by ANOVA.

| | L0 Mean ±SD (n=64) | L1^A Mean ±SD (n=5) | L1^L Mean ±SD (n=10) | L1^{LA} Mean ±SD (n=6) | p-value | All |
|-------------------------------|---------------------------------|--|---|---|----------------|------------|
| a. Clinical phenotype | | | | | | |
| sex [f/m] | 36/28 | 2/3 | 4/6 | 1/5 | | 43/42 |
| age [years] | 14.4±2.3 | 15.4±1.5 | 13.0±2.3 | 14.6±2.6 | NS | 14.3±2.3 |
| weight [kg] | 87.0±20.0 | 110.3±21.6 | 81.0±11.9 | 92.3±26.5 | NS | 88.1±20.4 |
| height [cm] | 164.0±11.4 | 171.2±5.9 | 162.6±7.7 | 167.6±18.4 | NS | 164.5±11.4 |
| hSDS | 0.4±1.4 | 0.4±0.5 | 1.2±1.2 | 0.5±2.6 | NS | 0.5±1.5 |
| BMI [kg/m²] | 32.0±5.2 | 37.5±5.9 | 30.6±3.8 | 32.4±5.5 | NS | 32.2±5.2 |
| BMI- z score IOTF | 2.7±0.5 | 3.1±0.5 | 2.7±0.5 | 2.8±0.4 | NS | 2.7±0.5 |
| waist [cm] | 100.6±11.5 | 115.8±16.3 | 98.6±7.1 | 98.6±10.1 | NS | 101.0±11.5 |
| WHR | 0.93±0.1 | 0.98±0.00 | 0.97±0.0 | 1.00±0.03 | NS | 0.95±0.09 |

| | | | | | | |
|-------------------------------------|-----------|------------|-------------|------------|--|------------|
| BP systolic [mmHg] | 125±11 | 136±15 | 129±11 | 124±5 | NS | 126±11 |
| BP diastolic [mmHg] | 78±8 | 80±10 | 76±11 | 74±5 | NS | 77±8 |
| <i>b. Chemical phenotype</i> | | | | | | |
| TSH [μIU/ml] | 2.7±1.2 | 3.3±1.0 | 3.0±1.6 | 2.8±1.1 | NS | 2.8±1.2 |
| cortisol 8.00 [μg/dl] | 17.4±6.1 | 22.3±3.2 | 19.3±4.0 | 18.2±10.0 | NS | 18.0±6.2 |
| cortisol 24.00 [μg/dl] | 3.4±3.9 | 1.7±1.2 | 1.5±1.2 | 1.4±0.2 | NS | 2.9±3.5 |
| T-Chol [mg/dl] | 171±35 | 182±32 | 158±18 | 208±23 | <i>0.029</i> | 172±34 |
| | | | | | L1 ^{LA} >L1 ^L | |
| HDL-chol [mg/dl] | 50±10 | 49±19 | 43±9 | 49±10 | NS | 49±10 |
| TG [mg/dl] | 134±60 | 176±86 | 184±67 | 184±94 | <i>0.044</i> | 146±67 |
| | | | | | L1 ^{LA} & L1 ^L >L0 | |
| GLU 0' [mg/dl] | 90±9 | 89±9 | 90±7 | 97±13 | NS | 90±9 |
| GLU 120' [mg/dl] | 112±20 | 126±5 | 130±22 | 129±30 | <i>0.017</i> | 116±21 |
| | | | | | L1 ^L >L0 | |
| INS 0' [μIU/ml] | 16.7±9.2 | 23.1±8.7 | 39.1±29.9 | 38.5±21.6 | <i>0.001</i> | 21.3±16.3 |
| | | | | | L1 ^L , L1 ^{LA} >L0 | |
| INS 120' [μIU/ml] | 91.6±67.9 | 139.6±85.4 | 186.1±140.5 | 146.0±93.9 | <i>0.033</i> | 109.6±87.0 |
| | | | | | L1 ^L >L0 | |
| INS/GLU | 0.2±0.1 | 0.3±0.1 | 0.4±0.3 | 0.4±0.2 | <i>0.001</i> | 0.2±0.2 |

| | | | | | | |
|------------------------------|--------------|----------|-----------|----------|--------------------------|--------------|
| | | | | | $L1^L, L1^{LA}>L0$ | |
| INS/GLU [n, %>0.3] | 10/64, 15.6% | 1/5, 20% | 6/10, 60% | 3/6, 50% | <i>0.014</i> | 20/85, 23.5% |
| | | | | | $L1^L, L1^{LA}>L0$ | |
| HOMA-IR | 3.7±2.2 | 5.1±2.2 | 8.8±6.7 | 9.4±5.5 | <i>0.001</i> | 4.8±3.9 |
| | | | | | $L1^{LA}, L1^L>L0$ | |
| ALT/GPT [U/l] | 24±9 | 77±33 | 34±9 | 69±28 | <i><0.0001</i> | 31±21 |
| | | | | | $L1^A, L1^{LA}>L1^L>L0$ | |
| AST/SGOT [U/l] | 25±8 | 47±14 | 27±4 | 43±15 | <i>0.0001</i> | 28±11 |
| | | | | | $L1^A, L1^{LA}>L1^L, L0$ | |
| GGTP [U/l] | 22±10 | 39±10 | 26±12 | 35±12 | <i>0.003</i> | 25±12 |
| | | | | | $L1^A, L1^{LA}>L0, L1^L$ | |

Table 3 Ratio of steroid metabolites (enzyme activity): differences between patients with non-liver diseases (L0) and with liver disease features (L1) patients.

Legend: Values are means and SD; ; L0 – non-liver disease patients. L1- liver disease patients (ALT+ or sonographic liver steatosis or both). NS- not significant. Significance by t-student test.

**Ratio of steroid metabolites was calculated based on steroid metabolites concentrations.*

| | Mean L0 | SD L0 | Mean L1 | SD L1 | <i>p-value</i> |
|---|----------------|--------------|----------------|--------------|-----------------------|
| | (n=64) | | (n=21) | | |
| An/Et (5α reductase) | 2.1 | 0.9 | 2.5 | 0.9 | <i>0.041</i> |
| 5αTHF/THF (5α reductase) | 1.3 | 0.6 | 1.3 | 0.5 | NS |
| 5αTHB/THB (5α reductase) | 3.4 | 1.8 | 3.3 | 1.3 | NS |
| (THF+αTHF)/THE (11βHSD1) | 0.9 | 0.3 | 0.7 | 0.2 | <i>0.019</i> |
| (THE+THF+αTHF)/P5T-17α (3βHSD) | 19.1 | 15.4 | 24.2 | 21.3 | NS |
| (THE+THF+αTHF)/PT (21-OHase) | 10.1 | 3.9 | 13.6 | 5.1 | <i>0.009</i> |
| (THE+THF+αTHF)/PO5α3α (21-OHase) | 300.9 | 160.4 | 320.7 | 193.3 | NS |

Table 4 Ratio of steroid metabolites (enzyme activity): differences between non-liver disease patients (L0) and patients with liver disease features: L1^A (ALT+), L1^L (sonographic liver steatosis) and L1^{LA} (both ALT+ and sonographic liver steatosis).

Legend: Values are means and SD; L0 – non-liver disease patients. L1- liver disease patients. L1^A – patients L1 with ALT+. L1^L – patients L1 with sonographic liver steatosis. L1^{AL} – patients with ALT+ and sonographic liver steatosis. NS- not significant. Significance by ANOVA.

**Ratio of steroid metabolites was calculated based on steroid metabolites concentrations.*

| Ratio | L0 Mean ±SD (n=64) | L1^A Mean ±SD (n=5) | L1^L Mean ±SD (n=10) | L1^{LA} Mean ±SD (n=6) | <i>p-value</i> | All |
|--|-----------------------------------|--|---|---|--|-------------|
| An/Et (5α-reductase) | 2.1±0.9 | 2.9±1.1 | 2.0±0.7 | 3.1±0.8 | 0.011 | 2.2±0.9 |
| | | | | | <i>L1^{LA}, L1^A>L0, L1^L</i> | |
| 5αTHF/THF (5α-reductase) | 1.3±0.6 | 1.4±0.4 | 1.1±0.4 | 1.7±0.5 | NS | 1.3±0.5 |
| 5αTHB/THB (5α-reductase) | 3.4±1.8 | 3.6±0.8 | 2.7±1.1 | 4.0±1.6 | NS | 3.4±1.7 |
| (THF+αTHF)/THE (11βHSD1) | 0.9±0.3 | 0.8±0.1 | 0.7±0.2 | 0.8±0.2 | NS | 0.8±0.3 |
| (THE+THF+αTHF)/P5T-17α (3βHSD) | 19.1±15.4 | 15.1±8.1 | 27.5±17.1 | 26.4±33.5 | NS | 20.4±17.0 |
| (THE+THF+αTHF)/PT (21-OH) | 10.1±3.9 | 12.8±6.0 | 14.4±4.6 | 12.7±5.9 | 0.016 | 11.0±4.5 |
| | | | | | <i>L1^L>L0</i> | |
| (THE+THF+αTHF)/PO5α3α (21-OH) | 300.9±160.4 | 228.5±99.0 | 416.1±230.8 | 238.4±104.0 | NS | 305.8±168.1 |

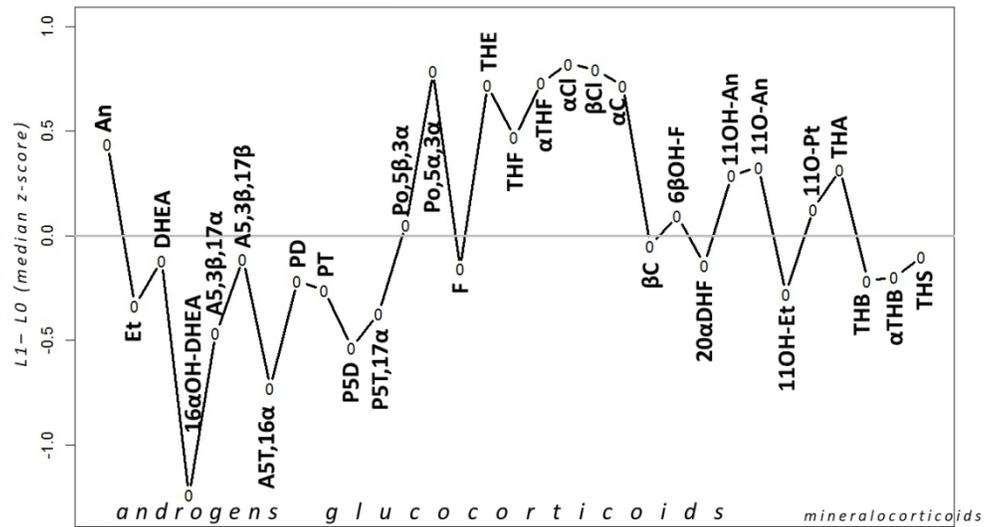


Figure 1. Steroidal signature of liver disease in childhood obesity: differences between z-transformed concentrations of steroid metabolites (androgens, glucocorticoids and mineralocorticoids) in liver diseases (L1) and with non-liver disease features (L0) patients.

338x190mm (96 x 96 DPI)

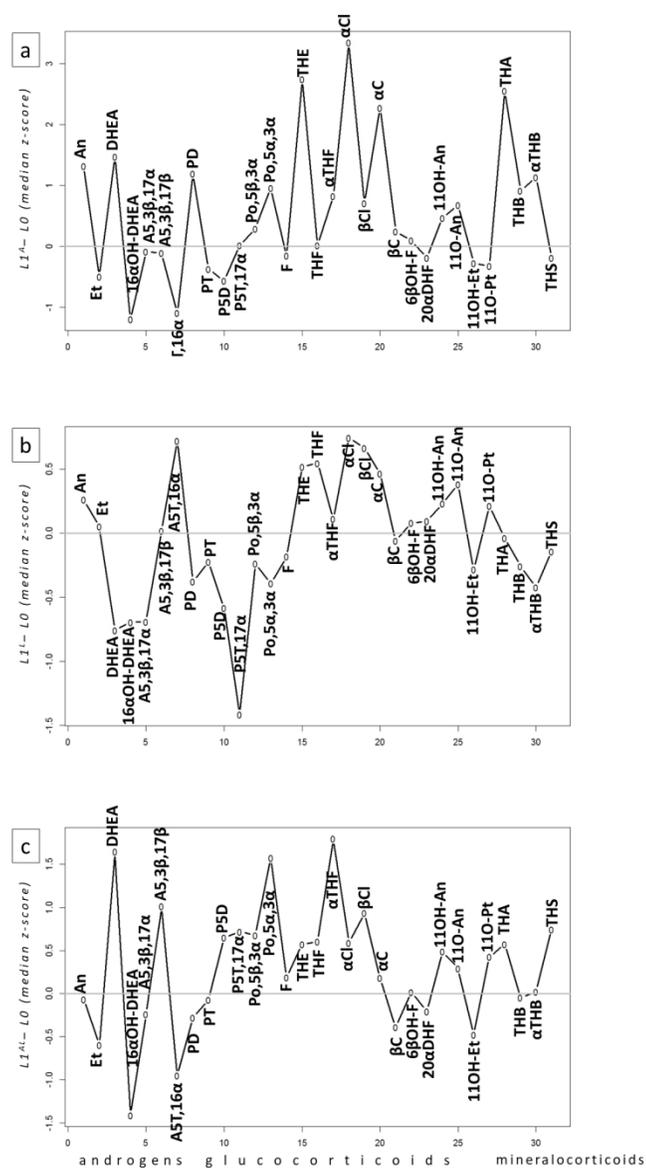


Figure 2 Steroidal signatures of liver disease in childhood obesity: differences between z-transformed concentrations of steroid metabolites (androgens, glucocorticoids and mineralocorticoids) in non-liver disease features (L0) patients and (a) patients with ALT+ (L1A), (b) patients with sonographic liver steatosis (L1L) and (c) patients with ALT+ and sonographic liver steatosis (L1AL)

225x381mm (144 x 144 DPI)