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[Ricky Jensch](#)*, Ronny Baber, Antje Körner, [Wieland Kiess](#), [Uta Ceglarek](#), [Antje Garten](#), [Mandy Vogel](#)

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Association of Whole Blood Amino Acid and Acylcarnitine Metabolome with Anthropometry and IGF1 Serum Levels in Healthy Children and Adolescents

Ricky Jensch ^{1,2,*}, Ronny Baber ^{1,3}, Antje Koerner ^{1,2}, Wieland Kiess ^{1,2}, Uta Ceglarek ^{1,3}, Antje Garten ^{2,#} and Mandy Vogel ^{1,2#}

¹ LIFE Leipzig Research Center for Civilization Diseases, University of Leipzig, Philipp-Rosenthal-Strasse 27, 04103 Leipzig, Germany

² Hospital for Children and Adolescents and Center for Pediatric Research (CPL), University of Leipzig, Liebigstrasse 20a, 04103 Leipzig, Germany

³ Institute of Laboratory Medicine, Clinical Chemistry and Molecular Diagnostics (ILM), University Hospital Leipzig, Paul-List Str.13/15, 04103 Leipzig, Germany

* Correspondence: ricky.jensch@medizin.uni-leipzig.de

both authors contributed equally

Abstract: Background: Physiological changes of blood amino acids and acylcarnitines during healthy child development are not well studied. The LIFE Child study offers a platform with a large cohort of healthy children to investigate these dynamics. We aimed to assess the intra-person variability of these blood metabolites and associations with anthropometric parameters related to excess body fat and IGF1 serum levels. Methods: Concentrations of 23 amino acids, 5 acylcarnitines and free carnitine of 2213 children aged between 3 months and 19 years were analyzed using liquid chromatography/tandem mass spectrometry. Standard deviation score values were used to account for physiological sex- and age-related differences. The stability of metabolites was assessed through the coefficient of determination. Associations with parameters for body composition and IGF1 serum levels were determined by Pearson correlation and linear regression. Results: Our research reveals substantial fluctuations in metabolite concentrations in the same person during childhood and adolescence. Most metabolites had a positive correlation with parameters of body composition with notable influence of sex, pubertal status and weight group specific differences. Glycine exhibited negative associations with parameters of body fat distribution especially in normal weight girls, overweight/obese boys and during puberty. Conclusions: The observed alterations in metabolite concentrations contribute to our knowledge on the complex interplay between blood amino acid and acylcarnitine levels and metabolic health in children and adolescents.

Keywords: pediatrics; metabolomics; amino acids; acylcarnitines; anthropometry; body composition

1. Introduction

The assessment of blood amino acid (AA) and acylcarnitine (AC) concentrations has long been established as part of newborn screening to identify congenital disorders [1]. Furthermore, blood AA and AC concentrations are also informative on physiological and disease-associated changes in human metabolic pathways, such as fatty acid oxidation, the carnitine shuttle system, the urea cycle, and the metabolism of branched-chain amino acids. Previously, AAs and ACs were examined as potential biomarkers or predictors of complex metabolic diseases in adults (e.g., type 2 diabetes, cardiovascular disease, obesity) [2–4] as well as in children and adolescents [2,5–8]. However, there is still a lack of studies examining physiological AA and AC blood levels and the influence of factors such as weight and excess body fat in healthy pediatric populations. The LIFE Child study, a large-scale longitudinal pediatric cohort study, aims to identify factors related to lifestyle-related diseases. A previous study based on the LIFE Child cohort examined AA and AC concentrations and the

influence of age, sex, weight, and pubertal status as well as laboratory parameters of carbohydrate, fat, liver, kidney, and thyroid metabolism. They found associations between both weight and pubertal status and AA and AC levels under physiological conditions and suggested that AA and AC levels are potential biomarkers for metabolic alterations in overweight and obese pediatric individuals [9]. In this follow-up study, we asked how stable blood AA and AC concentrations are during repeated measurements and whether they differ in their stability. Furthermore, we wanted to examine the association between weight status and blood AA/AC values in more detail and asked if associations exist with different parameters for excess body fat, body fat distribution and IGF1 serum levels and whether sex and pubertal status would have an impact on these associations. With this study, we want to further contribute to understanding the complex changes of AA and AC metabolism during childhood and adolescence and aid in the identification of potential biomarkers and early predictors for pathological processes that lead to metabolic disorders like type 2 diabetes and hypertriglyceridemia.

2. Materials and Methods

2.1. Study Population and Design

LIFE Child is a prospective, longitudinal, population-based childhood cohort study carried out at the Leipzig Research Center for Civilization Diseases (LIFE) in the city of Leipzig (Saxony, Germany). Participants join the study at any age from 3 months to 16 years and attend annual follow-up visits until the age of 20. For participants recruited during the first year of life, data is collected at 3, 6, and 12 months of age. Most participants are German residents living in Leipzig and its vicinity. The LIFE Child study is designed to investigate the impact of metabolic, genetic, and environmental factors on health and development in children and adolescents [10,11]. The sub-study presented includes all visits where metabolite assessments were performed (4029 measurements from 2213 children aged between 3 months and 19 years). Measurements of skinfold thickness (biceps, triceps, iliac crest, subscapular) started from the age of 2. Children aged 4 years and older were instructed to fast overnight for at least 8 hours. From age 8 to age 15, for each year more than 100 measurements for the follow-up visits on year later were collected, whereas for lower and older ages, the number of available measurements was lower (Supplement Table S1).

2.2. Preanalytics and Analytics

22 AAs, 6 ACs and free carnitine were determined by liquid chromatography/tandem mass spectrometry of dried whole blood samples collected by venous blood sampling. IGF1 was measured by immunoassay from serum blood samples. The detailed procedures have been described elsewhere [9,12–14]. The data collection for this study was carried out between May 2011 and December 2014.

2.3. Other Measures

Qualified and certified staff conducted all measurements during anthropometric assessments of the participants. Measurements of height, weight, skinfold thicknesses, waist circumference and hip circumference were used in this study. The categorization into pre-pubertal and pubertal was done based on Tanner Stages (Stage 1=prepubertal, Stage 2-5=pubertal).

2.3. Statistical Analysis

To reduce the confounding of age and sex regarding the investigated correlations, AAs, ACs, free carnitine and anthropometric measurements were transformed to age- and sex-adjusted standard deviation scores (SDS) [9,12,15,16]. Subjects were categorized into normal weight: BMI-SDS -1.28 to 1.28 and overweight/obese: BMI-SDS > 1.28 [17]. Extreme outliers with SDS values of more than ± 9 standard deviations were excluded. For statistical analysis, hierarchical linear regression analysis and the Pearson correlation coefficient were applied. To estimate the stability of metabolites, the age of the participants was rounded (e.g. age 3,5-4,49 = age 4) and then the coefficient

of determination (r^2) was calculated through linear regression for every pair of consecutive visits. Data analyses were carried out using R software (version 4.2.2; R Foundation for Statistical Computing, Vienna, Austria) [18]. ggplot2 was used for visualization [19]. The statistical significance level was set to $\alpha = 0.05$.

3. Results

The distribution of sex, weight, and pubertal status of the study population at the time of their first visit is presented in Table 1.

Table 1. Characteristics of the children during their first visit.

| | [ALL] N=2213 | female N=1088 | male N=1125 |
|--|---------------------|----------------------|--------------------|
| Age (years) \pm SD | 8.11 (5.13) | 8.46 (5.21) | 7.77 (5.04) |
| Weight group, n% | | | |
| Underweight | 200 (9.17%) | 91 (8.47%) | 109 (9.85%) |
| Normal weight | 1764 (80.8%) | 865 (80.5%) | 899 (81.2%) |
| Overweight | 139 (6.37%) | 80 (7.44%) | 59 (5.33%) |
| Obese | 79 (3.62%) | 39 (3.63%) | 40 (3.61%) |
| Puberty status | | | |
| Pre-pubertal | 1215 (68.6%) | 577 (60.0%) | 638 (78.8%) |
| Pubertal | 556 (31.4%) | 384 (40.0%) | 172 (21.2%) |

¹ Weight groups defined by BMI-SDS = body mass index standard deviation score: underweight: BMI-SDS < -1.28; normal weight: BMI-SDS -1.28 to <1.28; overweight: BMI-SDS > 1.28 to 1.88; obese: BMI-SDS > 1.88. Pre-pubertal=Tanner 1, Pubertal=Tanner 2-5.

3.1. Variance of metabolites

AAs and ACs showed low year-to-year correlations, indicating low longitudinal stability. As an example, we present the trend of yearly correlations across age for aromatic AAs and ACs in Figure 1. The mean r^2 -values of the aromatic AAs histidine ($r^2=0.020$), phenylalanine ($r^2=0.072$), tryptophan: ($r^2=0.086$) and tyrosine: ($r^2=0.048$) showed no significant change with age ($p>0.05$ for all), as did the mean values of free carnitine ($r^2=0.129$), and the ACs acetylcarnitine ($r^2=0.032$) propionylcarnitine ($r^2=0.075$) palmitoylcarnitine ($r^2=0.079$) stearoylcarnitine ($r^2=0.087$) and methylmalonyl carnitine: ($r^2=0.067$) ($p>0.05$ for all ACs). Results for the other AAs are provided in Supplement Figure S1.

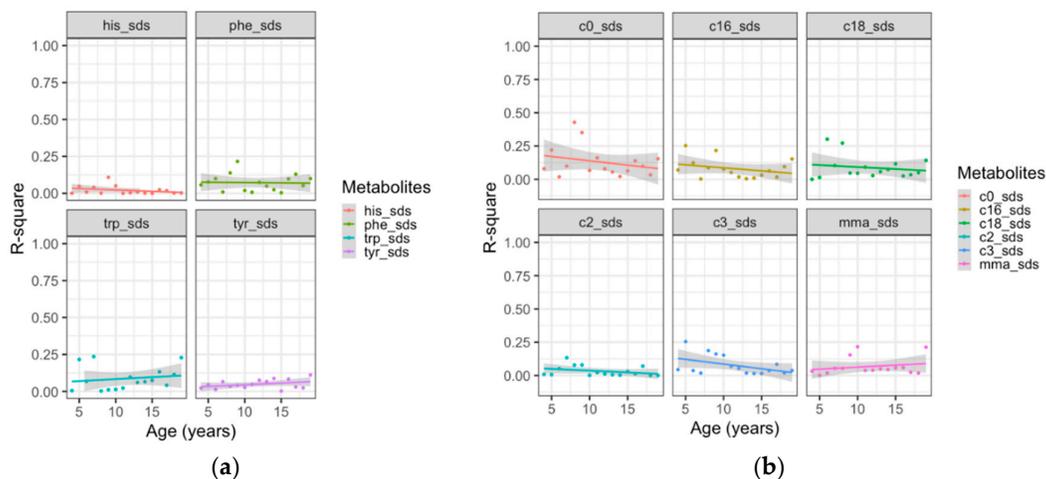


Figure 1. Linear trends of year-to-year correlations of consecutive metabolite concentration SDS across age for aromatic amino acids (a) his=histidine, phe=phenylalanine, trp=tryptophan, tyr=tyrosine and ACs (b) c0= free carnitine, c2= acetylcarnitine, c3= propionylcarnitine, c16= palmitoylcarnitine, c18= stearoylcarnitine, mma= methylmalonyl carnitine.

3.2. Metabolites and anthropometric parameters

Most AAs and ACs were positively correlated with BMI, WHR, waist circumference, skinfold thicknesses and IGF1. In contrast, glycine showed negative correlations in female individuals with skinfold thicknesses of biceps ($r = -0.12$, $p < 0.001$), triceps ($r = -0.11$, $p < 0.001$), iliac crest ($r = -0.09$, $p < 0.001$), and subscapular ($r = -0.11$, $p < 0.001$), as well as waist circumference ($r = -0.07$, $p = 0.007$) and BMI ($r = -0.09$, $p < 0.001$). For males, the only significant correlation of glycine with an anthropometric parameter existed for subscapular skinfold thickness ($r = -0.05$, $p = 0.045$). The association between metabolite concentrations and parameters of body fat distribution showed a weaker correlation for WHR in comparison to waist circumference. These findings are presented in Figure 2.

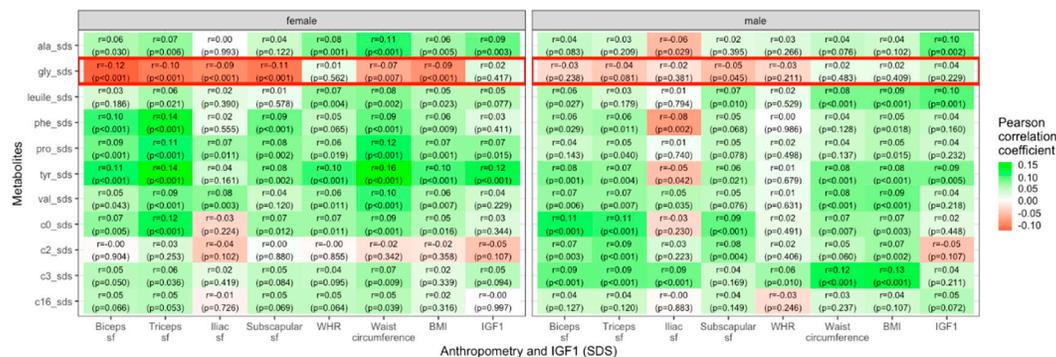


Figure 2. Heatmap showing Pearson correlation values (green=positive, red=negative) between selected metabolite concentrations and anthropometric parameters in female and male children and adolescents. (sf=skinfold).

Alanine and proline showed stronger associations with almost all anthropometric measurements in females. The differences were highest for waist circumference (alanine: $\beta = 0.107$, $p < 0.01$ in females vs. $\beta = 0.045$, $p = 0.079$ in males; proline $\beta = 0.115$, $p < 0.001$ in females vs. $\beta = 0.035$, $p = 0.196$ in males – Figure 3).

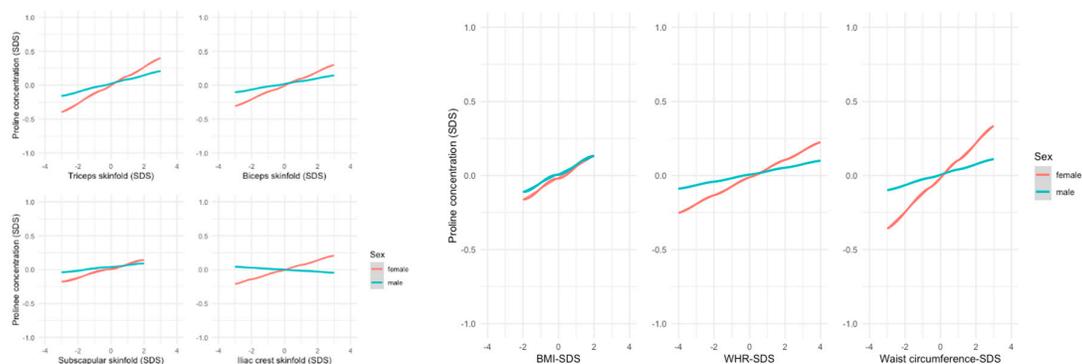


Figure 3. Linear regression model - Association between anthropometric parameters and mean SDS levels of proline; Proline has a stronger positive association with anthropometric parameters in females.

In contrast, the associations between the short-chain ACs acetylcarnitine (C2) resp. propionylcarnitine (C3) and various anthropometric parameters were stronger in males, e.g., biceps skinfold thickness (C2: $\beta=0.084$, $p=0.005$ in males vs. $\beta=-0.004$, $p=0.884$ in females; C3 $\beta=0.095$, $p=0.002$ in males vs $\beta=0.045$, $p=0.904$ in females). The associations for C3 measurements are presented in Figure 4.

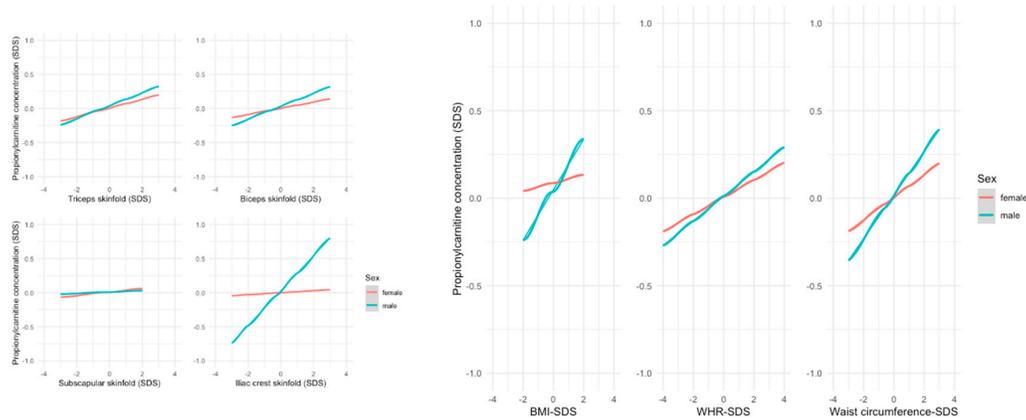
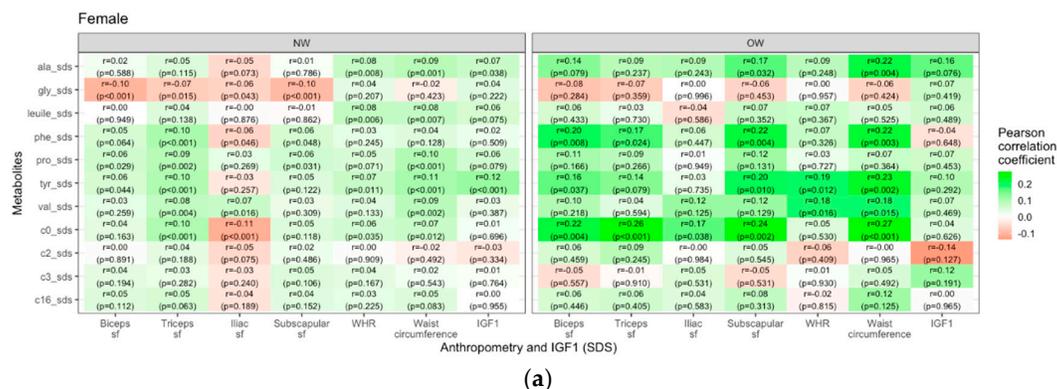


Figure 4. Linear regression model showing the relationship between anthropometric parameters and mean SDS levels of C3; Male individuals showed a stronger association particularly in biceps and iliac crest skinfold as well as BMI and waist circumference.

Lastly, we examined the interaction of weight group and puberty status on the relationship between metabolites and anthropometric parameters in boys and girls. Generally, in girls the overweight/obese group showed stronger positive correlations between metabolites and weight- and body-fat-related measures than the normal weight group, most prominently for C0. In contrast, for boys, there were no similar consistent patterns. Most prominently, glycine displayed negative correlations with skinfold thicknesses ($r=-0.24$, $p=0.006$ for biceps sf, $r=-0.23$, $p=0.009$ for subscapular sf and $r=-0.19$, $p=0.03$ for triceps sf). WHR only showed a few strong correlations with different metabolites, primarily in the female overweight/obese group (e.g. valine: $r=0.18$, $p=0.016$; tyrosine: $r=0.19$, $p=0.12$) and males (e.g. C3: $r=0.27$, $p=0.01$). In overweight/obese boys we found negative correlations of IGF1 with tyrosine ($r=-0.22$, $p=0.041$) and palmitoylcarnitine ($r=-0.24$, $p=0.026$). The correlations are visualized as heatmaps showing the Pearson correlation coefficients (Figure 5).



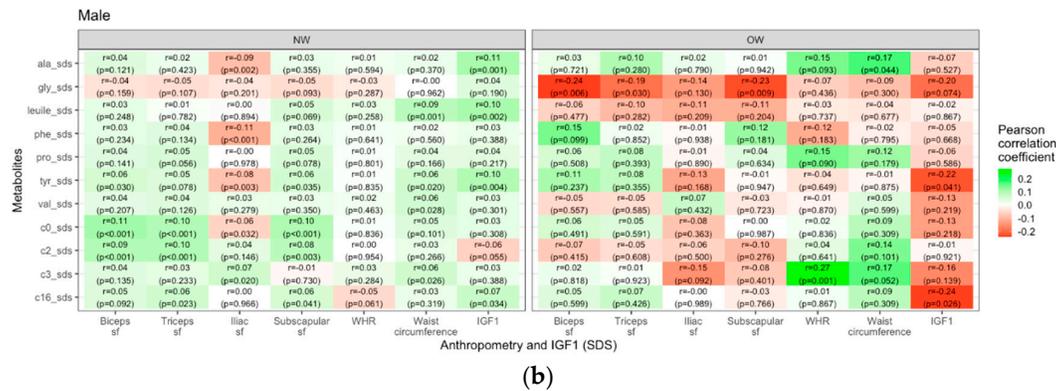


Figure 5. Heatmaps showing the Pearson correlations (green=positive, red=negative) between selected metabolite concentrations and anthropometric parameters in female (a) and male (b) children and adolescents. (sf=skinfold).

The differences in the associations between pre-pubertal and pubertal status were most prominent for glycine and ACs. The negative association between glycine and the anthropometric measures was stronger during puberty than before puberty in boys, whereas for girls the negative association existed before and during puberty. For ACs, there was a stronger association between metabolite concentrations and anthropometric parameters (except for BMI) during puberty in girls. Here, we found a positive correlation for prepubertal and pubertal girls (Supplementary Figure S3).

4. Discussion

The investigation of AA and AC concentrations in children and adolescents presents a unique set of challenges due to the inherent complexity of influences during growth and development. Our study shows that metabolite concentrations underlie substantial within-person variations. Despite reducing the impact of confounding factors like age and sex by using SDS values, these changes were still present. This could be explained by the role of metabolites as being both regulated by and acting as a regulation tool for physiological processes, the varying demands of the body during adolescence, as well as changes in hormones and dietary intake, which all impact blood metabolite concentrations. An investigation in adults of the importance of pre-collection factors, such as time of day of blood collection, season, hours of fasting, and physical activity, showed only negligible differences regarding AA and AC concentrations [20]. Another adult study showed WHR, sex, application of sex hormones, age, and hematocrit to be factors influencing metabolite concentrations [21].

The influence of overweight and excess body fat on metabolite concentrations was a major topic in our study. A previous study found significant correlations between BMI and branched-chain (leucine/isoleucine, valine) and aromatic (phenylalanine, tyrosine) amino acids [9] which is in line with our findings. We further assessed the influence of excess body fat and body fat distribution on metabolites by considering body-fat and body-fat-distribution related parameters. The measuring of subcutaneous fat through skinfold thickness measurements provides an estimation of total body fat percentage, whereas waist circumference is an indicator of core body fat and WHR serves a tool for evaluating body fat distribution, higher values suggesting a higher amount of abdominal/visceral fat that is associated with greater health risks [22,23]. The increase of many metabolite concentrations (BCAAs, aromatic amino acids, acylcarnitines) with increasing body-fat could be linked to diseases like hypertriglyceridemia and insulin resistance in adults and children [7,24,25]. Our study confirmed these findings, showing that, in general, there were positive correlations between metabolite concentrations and the different skinfold thicknesses, waist circumference and BMI. We detected significant correlations between WHR and metabolite concentrations only for a few metabolites, supporting the hypothesis that in childhood, waist circumference might be a better predictor for high trunk fat mass than WHR [26].

Glycine was an exception from the general trend of positive associations between metabolite concentrations and parameters of excess body fat in our study. We found glycine negatively associated with the markers of increased body fat. This was especially true for skinfold thicknesses in overweight/obese boys, whereas in girls, the association was found only in the normal weight group. The result is in line with other studies which found decreased glycine concentrations associated with metabolic disorders such as obesity, type 2 diabetes, and non-alcoholic fatty liver disease. Further, the improvement of insulin resistance also led to normalization of glycine concentration.

We found differences in the associations of metabolite concentrations and anthropometric measures between normal weight vs. overweight/obese individuals as well as between boys and girls. The association between higher metabolite concentrations and obesity was stronger in females [29]. The increase of metabolite concentrations such as ACs and BCAAs and aromatic AAs was also observed in overweight/obese adults [3,30]. However, a defective fatty acid or amino acid metabolism, which can lead to incomplete fatty acid oxidation and further metabolic defects associated with obesity and diabetes, as seen in adults, was not seen in children. [31–35] How a child's fatty acid and glucose metabolism differs from those in adults is still not exactly understood. An explanation for the observed discrepancies might be that in children, metabolite concentrations can still be regulated more easily by the body, whereas in adults, these mechanisms are compromised by aging and prolonged co-morbidities, resulting in stronger alterations of metabolite concentrations in overweight adults. For example, higher mitochondrial adaptability and plasticity in young people are hypothesized to play a critical protective role. [3,31]

In adults, a negative association between IGF1 and obesity has been described in several studies [36,37]. However, there was also a study in which no significant correlation was observed [38]. Furthermore, differences between men and women in the interplay of IGF1 and metabolite concentrations are described [39]. In children, we could show that in overweight/obese boys, IGF1 correlated negatively with metabolite concentrations of tyrosine and the long-chain AC palmitoylcarnitine, whereas in overweight/obese girls, we found no significant correlation between metabolite concentrations and IGF1.

One strength of our study is the large sample size and homogeneity of our cohort of healthy children. Data collection and measurement of metabolites followed a standardized protocol, thus achieving a high internal validity and minimizing measurement distortions [14]. Because most study participants live in Leipzig and are of German origin, there is only minimal influence of different ethnicities and living conditions. On the other hand, this homogeneity might limit the generalizability of our results for other ethnic or social groups. Additionally, although the children are deeply phenotyped and examined for health, there could still be undiagnosed diseases that might impact metabolite concentrations. Another advantage of our study is the availability of different parameters for body fat measurements like WHR, waist circumference and skinfold thicknesses. We can, therefore, provide a detailed and comprehensive analysis of the interplay between metabolite concentrations and excess body fat.

5. Conclusion

Integrating metabolite profiling with anthropometric parameters like skinfold thicknesses and waist circumference helps to support current research regarding body fat distribution, overweight and metabolic diseases. We showed that metabolite concentrations in children undergo significant changes during childhood and adolescence. Additionally, we identified sex-related differences in the associations between metabolite concentrations and anthropometric parameters associated with excess body fat. Future studies may identify pathways regulating metabolite concentrations in the human body. A deeper understanding of the underlying mechanisms can enhance disease prevention, early diagnosis and personalized treatment plans.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org. Table S1: Number of one year follow-up measurements by age; Figure S1: Linear trends of year-to-year correlations of consecutive metabolite concentration SDS across age for uncategorized

AAs (a), non-proteinogenic AAs (b) and acidic AAs and BCAAs (c); Figure S3: Heatmap showing the Pearson correlations (green=positive, red=negative) between selected metabolite concentrations and anthropometric parameters in pre-pubertal and pubertal female (a) and male (b) children and adolescents. (sf=skinfold).

Author Contributions: Individual contributions to this manuscript are as follows: conceptualization, W.K., M.V.; methodology, W.K., M.V. (creation of models: R.J., M.V.); formal analysis, R.J., M.V. and A.G.; data curation, W.K., U.C.; writing—original draft preparation, R.J.; writing—review and editing, W.K., U.C., M.V., A.G., R.J.; data analysis and visualization, R.J., M.V.; supervision, W.K. and M.V.; project administration, W.K.; funding acquisition, W.K.. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The study was conducted in accordance with the Declaration of Helsinki and approved by the Ethics Committee of the University of Leipzig (reference number: Reg. No. 264-10-19042010). The study is registered at ClinicalTrials.gov (NCT02550236).

Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

Data Availability Statement: The dataset presented in this article cannot be shared publicly because of ethical and legal restrictions. The LIFE Child study is a study collecting potentially sensitive information. Publishing data is not covered by the informed consent provided by the study participants. Furthermore, the data protection concept of LIFE requires all (external as well as internal) researchers interested in accessing data to sign a project agreement. Researchers interested in accessing data from the LIFE Child study may contact the study by writing to forschungsdaten@medizin.uni-leipzig.de.

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Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript; or in the decision to publish the results.

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