Daily physical activity, fasting glucose, uric acid, and body mass index are independent factors associated with serum fibroblast growth factor 21 levels

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Abstract

Objective: Fibroblast growth factor 21 (FGF21) levels have been linked with beneficial effects on glucose and lipid metabolism in animals. It is elevated in humans with the metabolic syndrome. This study investigates independent factors associated with serum FGF21 levels.

Design: Cross-sectional study done in healthy blue-collar workers.

Methods: A medical history was taken, and FGF21 (measured using an ELISA commercial kit), glucose, uric acid, plasma lipids, total/high-molecular weight (HMW) adiponectin, and retinal-binding protein 4 (RBP4) were measured in 210 individuals with (n = 81) and without (n = 129) metabolic syndrome.

Results: The median of serum FGF21 levels were higher in subjects with metabolic syndrome (339.5 vs 276.4 ng/l, P = 0.01). Serum FGF21 levels correlated positively with body mass index (BMI; r = 0.23, P = 0.001) and age (r = 0.17, P = 0.01). After adjusting for age and BMI, a significant positive correlation persisted for fasting glucose, uric acid, and physical activity in both males (r = 0.21, r = 0.11, and r = 0.19, all P < 0.05) and females (r = 0.20, r = 0.19, and r = 0.14, all P < 0.05). In addition, FGF21 also correlates negatively with RBP4 (r = −0.27, P = 0.02), total (r = −0.26, P = 0.03), and HMW adiponectin (r = −0.30, P = 0.01) in women. A multiple linear regression model analysis identified that BMI (standardized β (SB) = 0.247; P = 0.008), glucose (SB = 0.226; P = 0.003), uric acid (SB = 0.191; P = 0.04), and physical activity (SB = 0.223; P = 0.004) are independent factors influencing serum FGF21 levels (R² = 0.105, P < 0.001). In addition, fasting hyperglycemia ≥ 100 mg/dl, excess body weight with BMI ≥ 25 kg/m², and uric acid ≥ 5.5 mg/dl predicted higher serum FGF21 levels.

Conclusion: Serum FGF21 levels are influenced by BMI, fasting glycemia, uric acid, and physical activity.

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Introduction

Fibroblast growth factor 21 (FGF21), a member of the FGF family, has recently emerged as a novel regulator of metabolism (1). FGF21 is mainly produced by the liver (2) and functions as a potent activator of glucose uptake in adipocytes (1). FGF21 causes paracrine effects, such as induction of hepatic ketogenesis, and endocrine actions, such as the promotion of lipolysis in white adipose tissue (3, 4), although this last effect is controversial (5). Moreover, when FGF21 is over expressed in transgenic mice, it protects animals from diet-induced obesity, and its administration in diabetic rodents (1) and monkeys (6) lowers blood glucose and triglyceride levels. FGF21 has been shown to produce a significant increment in total adiponectin and a reduction in leptin levels as well (6). The expression and release of FGF21 from liver are regulated by the peroxisome proliferator-activated receptor-α (PPAR-α or PPARα as listed in the HUGO Database) in liver (3, 4). Accordingly, free fatty acids appear to be major regulators of FGF21 expression, acting via PPAR-α-dependent activation of the FGF21 gene (7). This effect of fatty acids may explain the induction of FGF21 after fasting, and the paradoxically high levels of FGF21 reported recently in patients with type 2 diabetes and obesity (8, 9). However, other clinical studies in humans have shown inconsistent results. Positive correlations between FGF21 levels and some obesity traits such as body mass index (BMI), waist circumference, waist-to-hip ratio, and fat percentage have been reported (9, 10). On the other hand, similar correlations were not found in patients with a recently diagnosed type 2 diabetes mellitus (11). While some...
studies suggest a role for FGF21 in insulin sensitivity (1, 5), other studies using more reliable methodology (i.e. hyperglycemic clamp) do not confirm this association (10). Although current data support the role of FGF21 as a metabolic regulator of glucose uptake (1), adiposity, and lipid metabolism (3, 4), more investigation is needed to clarify the action, regulation, and clinical relevance of FGF21. The relationships between FGF21 and physical activity, serum adipokines (such as retinol-binding protein 4 (RBP4) and high-molecular weight (HMW) adiponectin) and other traits of the metabolic syndrome besides glycemia and lipid parameters have not been investigated in humans. An increased serum RBP4 concentration has been reported in subjects with the metabolic syndrome (12). However, not all studies are concordant, and some did not observe a relationship between RBP4 and insulin resistance (13). Adiponectin modulates a number of metabolic processes, including glucose regulation and fatty acid catabolism (14). Adiponectin self-associates into larger structures, and studies on multimer formation in human blood have demonstrated that HMW adiponectin is the active form of the protein (15). The levels of both total and HMW adiponectin are lower when metabolic syndrome is present and usually higher in women than in men (16).

This study is aimed to investigate independent factors associated with serum FGF21 levels and the possible association of FGF21 with other parameters of insulin resistance, physical activity, and adiposity.

Materials and methods

Study subjects

A total of 210 Mexican Mestizos were recruited for this study. The population consisted of medium-income healthy workers without a personal history of metabolic abnormalities. Evaluations were scheduled in advance. The individuals were studied at their place of work in a properly conditioned area. We included all individuals who agreed to be evaluated, of both genders, between 18 and 65 years of age. We excluded subjects with current treatment for diabetes, dyslipidemia, hypertension, thyroid diseases, or obesity in order to eliminate the possible effect of medications on clinical and biochemical parameters. In addition, we also excluded subjects with other chronic disease (such as HIV infection or viral hepatitis, lupus or rheumatoid arthritis, seizures, major depression with treatment, hospitalization in last 6 months, active cancer or under treatment, and pregnant women). Individuals with severe hypertriglyceridemia and a lipemic serum were also excluded since it could interfere with biochemical results.

The metabolic syndrome was defined according to the US 2004 National Cholesterol Education Program (NCEP) Adult Treatment Panel III guidelines (17) and modified as recommended in the latest American Heart Association/National Heart, Lung, and Blood Institute Scientific Statement for fasting glucose (18). The metabolic syndrome was defined as the presence of three or more of the following risk factors: i) central obesity (waist circumference ≥88 cm in women and ≥102 cm in men); ii) hypertriglyceridemia (fasting triglycerides ≥150 mg/dl (≥1.69 mmol/l); iii) low high-density lipoprotein (HDL) cholesterol <50 mg/dl (<1.29 mmol/l) in women and <40 mg/dl (1.04 mmol/l) in men; iv) hyperglycemia (fasting glucose ≥100 mg/dl (≥5.6 mmol/l); and v) hypertension (sitting blood pressure of 130/85 mmHg or more, taken as a mean of two readings obtained after a resting for at least 10 min). The human research ethics committee of our institution approved the study. All procedures were done in accordance with the Declaration of Helsinki. Informed consent was obtained from all subjects.

Biochemical and anthropometric measurements

The Endocrinology and Metabolism Department of the Instituto Nacional de Ciencias Médicas y Nutrición Salvador Zubirán (INCMNSZ) performed all biochemical laboratory measurements using standardized procedures. The measurements were performed with commercially available standardized methods. This laboratory is certified for standardization of the tests by the External Comparative Evaluation of Laboratories Program of the College of American Pathologists. Glucose, total cholesterol, HDL cholesterol, triglycerides, and uric acid were measured using the Synchon CX analyzer (Beckman Systems, Fullerton, CA, USA). The coefficients of variation for cholesterol and HDL cholesterol were 3.3 and 2.5% respectively. Low-density lipoprotein cholesterol concentration was estimated by the Friedewald formula. Plasma insulin concentrations were estimated using a RIA method (MEIA, Abbott Laboratories). High-sensitivity C-reactive protein (CRP) was measured in plasma in duplicate using a particle-enhanced immuno-turbidimetric assay (Roche Diagnostics). A human FGF21 ELISA kit was used (BioVendor Laboratory Medicine, Modrice, Czech Republic) that do not showed cross-reactivity with other members of the human FGF family, basic FGF and adipokines such as adiponectin, leptin, and RBP4. For the measurement of FGF21, serum samples were diluted 1:3 before the assay, and then 100 μl diluted sera, calibrators, and quality controls were added to 96-well microtiter plates coated with an affinity-purified polyclonal anti-human FGF21 antibody. The intra- and inter-assay variations were 5.1 and 6.6% respectively. Serum total and HMW adiponectin (Millipore, Billarica, MA, USA), leptin (Millipore), and RBP4 (AdipoGen, Inc., Incheon, Korea) were determined by a sandwich ELISA kit respectively. Insulin resistance was estimated using homeostasis model assessment index-insulin resistance (HOMA-IR) (19). Anthropometric measurements were done after participants removed their shoes and upper garments.
Body weight and fat was quantified with the UM-026 Tanita Body Composition Analyzer (Tanita, Tokyo, Japan). All subjects were instructed to stand in the center of the scale during weight assessment. Height was obtained using the floor scale’s stadiometer. Height was measured to the nearest 0.5 cm. BMI was calculated as weight (kg) divided by height (m²). Abdominal circumference was measured to the nearest 0.1 cm at the level of the greatest frontal extension of the abdomen between the bottom of the rib cage and the top of the iliac crest. All subjects underwent physical examinations and routine biochemical analyses after overnight fasting between 8 and 12 h.

**Physical activity**

Subjects were instructed to answer a physical activity questionnaire developed by Tremblay and colleagues at Laval University (Canada). This questionnaire had been validated previously in a Mexican population (20). The questionnaire quantifies the physical activity level (kcal/day or in kJ if kcal is multiplied by 4.1855) throughout a 24-h period. Every subject answered three questionnaires distributed over two labor days and one day of the weekend. The individual chose from 1 (lowest) to 9 (highest) depending on the intensity of physical activity done in periods of 15 min, and wrote down the information in a specified table. Each number is multiplied by a constant that represents the calculated energetic expenditure in those 15 min. These results were added together, and the kcal/day estimate was obtained. The questionnaire has a sensitivity to detect changes of 4.7 kcal/kg per day (20 kJ/kg per day). The intra-individual reproducibility ($r=0.88$, $P<0.001$) and interclass correlation coefficient (0.86; $P<0.001$) were higher when these were compared with similar instruments (21, 22). Physical activity was measured in 167 (79.5%) of the 210 patients.

**Statistical analysis**

Normally distributed data, determined using Kolmogorov–Smirnov test, were expressed as means and s.d. (±S.D.), whereas variables with a skewed distribution were reported as median (interquartile range) and

<table>
<thead>
<tr>
<th>Variable</th>
<th>Controls <em>(n=129)</em></th>
<th>Metabolic syndrome <em>(n=81)</em></th>
<th>$P$ value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (years)</td>
<td>36.2±10.3</td>
<td>41.5±9.1</td>
<td>0.001</td>
</tr>
<tr>
<td>Sex (M/F)</td>
<td>44/85</td>
<td>37/44</td>
<td>0.116</td>
</tr>
<tr>
<td>BMI (kg/m²)</td>
<td>22.8±1.8</td>
<td>29.0±3.8</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Serum FGF21* (ng/l)</td>
<td>276.4 (169.0–404.4)</td>
<td>339.5 (248.0–492.7)</td>
<td>0.01</td>
</tr>
<tr>
<td>Serum RBP4* (ng/ml)</td>
<td>M: 70.7 (50.1–83.5)</td>
<td>M: 53.3 (31.9–74.0)</td>
<td>0.04</td>
</tr>
<tr>
<td>Serum adiponectin* (ng/ml)</td>
<td>F: 47.8 (33.4–70.4)</td>
<td>M: 46.1 (38.5–67.3)</td>
<td>0.77</td>
</tr>
<tr>
<td>Serum HMWA* (ng/ml)</td>
<td>M: 7.8 (5.8–10.2)</td>
<td>M: 6.8 (5.4–8.4)</td>
<td>0.02</td>
</tr>
<tr>
<td>Leptin* (ng/ml)</td>
<td>M: 8.4 (6.9–10.0)</td>
<td>M: 12.7 (7.2–21.0)</td>
<td>0.01</td>
</tr>
<tr>
<td>Serum hsCRP* (mg/l)</td>
<td>2.7 (1.9–3.2)</td>
<td>F: 25.7 (16.6–35.6)</td>
<td>0.18</td>
</tr>
<tr>
<td>Waist circumference (cm)</td>
<td>89.7±4.7</td>
<td>M: 97.7±9.0</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Waist-to-hip ratio</td>
<td>97.8±5.9</td>
<td>F: 93.6±9.7</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Fat percentage (%)b</td>
<td>20.7±4.5</td>
<td>F: 27.5±4.5</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Systolic blood pressure (mmHg)</td>
<td>103.6±12.4</td>
<td>111.5±14.5</td>
<td>0.001</td>
</tr>
<tr>
<td>Diastolic blood pressure (mmHg)</td>
<td>69.47±9.9</td>
<td>74.9±11.2</td>
<td>0.002</td>
</tr>
<tr>
<td>Cholesterol (mg/dl)</td>
<td>202.0±40.2</td>
<td>209.0±41.8</td>
<td>0.29</td>
</tr>
<tr>
<td>Triglycerides* (mg/dl)</td>
<td>113.0 (92–148.0)</td>
<td>161 (116–247)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>LDL cholesterol (mg/dl)</td>
<td>122.8±32.3</td>
<td>130.8±30.6</td>
<td>0.13</td>
</tr>
<tr>
<td>HDL cholesterol (mg/dl)</td>
<td>43±5.8</td>
<td>37.9±9.2</td>
<td>0.009</td>
</tr>
<tr>
<td>Fasting insulin* (mIU/l)</td>
<td>4.5 (4.0–7.8)</td>
<td>10.8 (8.0–15.5)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>HOMA-IR*</td>
<td>1.26 (0.91–1.6)</td>
<td>2.3 (1.7–3.5)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Fasting glucose (mg/dl)</td>
<td>84.6±8.2</td>
<td>89.0±9.4</td>
<td>0.003</td>
</tr>
<tr>
<td>Physical activity (kcal/kg per day)</td>
<td>55.2 (47.7–69.6)</td>
<td>50.9 (46.8–55.7)</td>
<td>&lt;0.01</td>
</tr>
<tr>
<td>Physical activity (kcal/day)</td>
<td>2894.5 (2349–3378)</td>
<td>3905.2 (3312.8–4502)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Uric acid (mg/dl)</td>
<td>M: 6.2±1.3</td>
<td>M: 6.6±1.19</td>
<td>0.22</td>
</tr>
<tr>
<td>Creatinine (mg/dl)</td>
<td>0.89±0.15</td>
<td>0.90±0.15</td>
<td>0.85</td>
</tr>
</tbody>
</table>

HMWA, high-molecular weight adiponectin; RBP4, retinal-binding protein 4; hsCRP, high-sensitivity C-reactive protein; M, male; F, female.

*Log-transformed before analysis.

bIncluded 160 subjects with available data (63 men and 97 women).

The table shows the clinical and biochemical characteristics of the study sample stratified by the presence of the metabolic syndrome (n=210). Data are expressed as mean ± s.d. or as median (interquartile range).
log-transformed to approximate normality before analysis. The \( \chi^2 \), Student unpaired \( t \)-test, or Mann–Whitney \( U \) test was used as appropriate for comparison between the two groups. Homogeneity of variance was evaluated with Levene’s test. Correlation coefficients between FGF21 and metabolic features were evaluated in all participants, and were calculated using Spearman’s, Pearson’s \( r \) tests or using partial correlation analysis when adjusted for age and BMI. One-way ANOVA was used for comparison between serum levels of FGF21 and number of components of metabolic syndrome and physical activity level. Serum FGF21 levels were adjusted for age, BMI, and physical activity. Stepwise linear regression model was used to examine the impact of variables in serum FGF21 level. Homogeneity and linearity were confirmed through the pattern of studentized residuals plotted against adjusted predicted values. No correlation was seen between them \( (r = 0.062, P = 0.42) \). The variables selected to enter into regression analyses were those that correlated significantly with serum FGF21 \( (\text{Table 2}) \). The mean of uric acid values in both groups was used as cutoff \( (5.5 \text{ mg/dl}) \). All reported \( P \) values are based on two-sided tests considering \( %0.05 \) as significant. All analyses were performed with SPSS 15.0 (Chicago, IL, USA).

Results

The characteristics of the study population are shown in Table 1. The metabolic syndrome group included 81 individuals (44 females and 37 males) with a mean age of 41.5 ± 9.1 years and BMI of 29.0 ± 3.8 kg/m\(^2\). The control group included 129 individuals (85 females and 44 males) with a mean age of 36.2 ± 10.3 years and BMI of 22.8 ± 1.8 kg/m\(^2\). Serum FGF21 levels ranged from 47 to 1839 ng/l. Serum FGF21 levels were significantly higher \( (P < 0.01) \) in subjects with the metabolic syndrome \( (339.5 \text{ ng/l} (248.0–492.7)) \) in comparison with patients without it \( (276.4 \text{ ng/l} (169.0–404.4)) \). There was no sex difference in serum FGF21 levels. In the whole population, serum FGF21 levels had significant positive association with age \( (r = 0.17, P = 0.01) \), glucose \( (r = 0.29, P < 0.001) \), BMI \( (r = 0.23, P = 0.001) \), fat percentage \( (r = 0.19, P = 0.01) \), insulin \( (r = 0.13, P = 0.05) \), HOMA-IR \( (r = 0.16, P = 0.01) \), triglycerides \( (r = 0.15, P = 0.02) \), HDL cholesterol \( (r = 0.15, P = 0.02) \), systolic blood pressure \( (r = 0.19, P = 0.006) \), uric acid \( (r = 0.17, P = 0.008) \), and physical activity (expressed as kcal/day, \( r = 0.29, P = 0.04) \). No association was identified between FGF21 and high-sensitivity CRP (hsCRP). After adjusting for age and BMI, a significant positive correlation persisted only for fasting glucose, uric acid, and physical activity \( (\text{Fig. 1}) \). This was true also in males or females \( (\text{Table 2}) \). The association between FGF21 and glucose was stronger in the subgroup of patients with metabolic syndrome \( (n = 81, r = 0.30, P = 0.006) \), physical activity, glucose, uric acid, and BMI influence serum FGF21 levels

Physical activity, glucose, uric acid, and BMI influence serum FGF21 levels

Stepwise linear regression model identified that BMI, glucose, uric acid, and physical activity (kcal/day) were independent factors associated with serum FGF21 levels \( (F = 10.05; r^2 = 0.199, P < 0.001; \text{Table 3}) \).
There was a progressive increment in serum FGF21 levels (adjusted for age and BMI) with increasing intensity of physical activity ($F = 4.8$, $P = 0.009$; Fig. 2). We also examined the combined effects of BMI, glucose, and uric acid on serum FGF21 levels according to the level of physical activity. As summarized in Table 4, fasting hyperglycemia $>100$ mg/dl, excess body weight with BMI $>25$ kg/m$^2$, and uric acid $>5.5$ mg/dl predicted higher serum FGF21 levels in comparison with subjects without these abnormalities. Moreover, a further increment in serum FGF21 levels was observed when...
the clinical or biochemical abnormality coexisted with higher intensity of daily physical activity.

We confirmed a dose–effect relationship between serum FGF21 levels (adjusted for age, BMI, and physical activity) and the number of the metabolic syndrome traits (Fig. 2). The association remained significant even after adjusting for individual components of the syndrome, except for waist circumference (Supplementary Tables 1 and 2, see section on supplementary data given at the end of this article).

Serum FGF21 levels and adipocytokines

In the whole population, no significant correlation was identified between FGF21 and leptin, RBP4, HMW, and total adiponectin. However, after adjusting for age, gender, and BMI, a significant association was identified for RBP4 ($r = -0.35, P = 0.02$), total ($r = -0.23, P = 0.01$), and HMW adiponectin ($r = -0.34, P = 0.03$). After gender stratification, the association between FGF21 and RBP4 ($r = -0.27, P = 0.02$), total ($r = -0.26, P = 0.03$), and HMW ($r = -0.30, P = 0.01$) adiponectin persisted only in women (Table 2). In addition, FGF21 correlated more significantly with RBP4 ($r = -0.48, P = 0.003$), total ($r = -0.39, P = 0.01$), and HMW adiponectin ($r = -0.49, P = 0.002$) in the subgroup of women with central obesity ($n = 106$). The association between FGF21 and these adipokines was not identified in men ($r = 0.09, r = 0.08$, and $r = 0.03$ respectively, all $P > 0.05$).

Discussion

FGF21 is a recently identified player in carbohydrate and lipid metabolism (2, 10, 11). Its physiological relevance in humans is still under study. Here, we report an independent association between serum FGF21 levels and fasting plasma glucose, BMI, uric acid, and physical activity. Additionally, in women, we found that as FGF21 increased, levels of RBP4, total and HMW adiponectin decreased.

In animal models, Fgf21 mRNA concentrations are significantly induced by fasting and in the presence of increased concentrations of free fatty acids. This process appears to be mediated through the activation of the PPAR-α (10, 11, 23). FGF21 in turn has diverse effects including the stimulation of gluconeogenesis (but not glycogenolysis), fatty acid oxidation, lipolysis, and ketogenesis (24); inhibition of lipogenesis (25); the induction of a hibernation-like state of torpor (11); and blunting of the GH-signaling pathway (26). FGF21 also increases glucose utilization (through the AKT pathway) in fat and muscle (27). In summary, the secretion of this hormone improves the utilization of energy substrates (fatty acids, ketones, and glucose), and interferes with energy consuming processes (i.e. lipogenesis and growth). For example, Badman et al. (10) have reported a deficient long-term adaptation to ketosis in FGF21-deficient mice. A similar physiological process may occur in humans, since FGF21 concentrations are increased after a prolonged fast (28).

In line with previous reports, our results confirm that FGF21 concentrations are increased in the metabolic syndrome (6, 7); a condition characterized by a chronic exposition to calorie surplus. Our report extends the available information by showing that BMI, fasting plasma glucose, uric acid, and physical activity are all independently associated with serum FGF21 levels. Increased free fatty acid concentration is also a feature of obesity (29). This abnormality may be the mechanism explaining the observed association between BMI and FGF21 levels. In the metabolic syndrome, the excess lipids enter alternative nonoxidative pathways resulting in the production of toxic reactive lipid species. Therefore, FGF21 by inhibiting lipogenesis and increasing fatty acid utilization may play a compensatory role against the metabolic defects associated with obesity. Some of the FGF21 actions that participate in the adaptation to fasting may limit the ectopic deposition of lipids in obese individuals. However, to confirm this theory, more research is needed. The association between glycemia and FGF21

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**Table 3** Multiple stepwise regression analysis of variables independently associated with serum fibroblast growth factor 21 (FGF21) levels.

<table>
<thead>
<tr>
<th>Independent variables</th>
<th>Standardized β</th>
<th>t</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>BMI</td>
<td>0.247</td>
<td>2.696</td>
<td>0.008</td>
</tr>
<tr>
<td>Glucose</td>
<td>0.226</td>
<td>2.971</td>
<td>0.003</td>
</tr>
<tr>
<td>Uric acid</td>
<td>0.191</td>
<td>2.180</td>
<td>0.04</td>
</tr>
<tr>
<td>Physical activity</td>
<td>0.223</td>
<td>2.943</td>
<td>0.004</td>
</tr>
</tbody>
</table>

The analysis also included age, waist circumference, fat percentage, insulin, leptin, RBP4, and adiponectin, which were all excluded in the final model. Physical activity expressed as kcal/day. Parameters of the model: $F = 10.05; r^2 = 0.199, P < 0.001$.

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**Figure 2** Log serum FGF21 levels according to the number of metabolic syndrome components (adjusted for age, BMI, and physical activity) and tertiles of physical activity (kcal/day) level (adjusted for age and BMI). Data are shown as means ± s.d. Results of one-way ANOVA for metabolic syndrome: $F = 16.5, P < 0.001$; for physical activity level: $F = 4.8, P = 0.009$. 

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levels is in line with this interpretation. The FGF21 overproduction may counterbalance the deleterious effects of free fatty acids on insulin action and glucose disposal. An alternative explanation may be that obesity and hyperglycemia cause an impaired tissue response to FGF21 leading to increased concentrations of this hormone. This alternative explanation implies that FGF21 has a regulatory axis, in which its own plasma level may be a determinant. Hence, we propose that FGF21 may form part of a defense response against the damaging effects of a chronic exposition to increased plasma fatty acid concentrations. Clearly, additional studies in humans and in animal models are required to describe FGF21 physiology.

Our data also show that daily physical activity has a positive association with serum FGF21 levels. Physical activity stimulates lipolysis (30, 31), increases serum free fatty acids, and activates PPAR-α-regulated pathways such as that of FGF21 secretion. This could explain the direct association identified between physical activity and FGF21.

Additionally, we report a novel association between uric acid and FGF21. Studies in healthy volunteers have shown that PPAR-α activation increases urinary uric acid excretion (32). In accordance, fenofibrate (a PPAR-α activator) lowers serum uric acid levels (33). However, it is not clear how uric acid plasma levels may increase FGF21 secretion. Hyperuricemia is frequently associated with the metabolic syndrome (34). This abnormality has been linked to endothelial dysfunction, decreased nitric oxide synthesis, and arterial hypertension (35). Additional studies are needed to confirm the participation of uric acid in FGF21 physiology.

In our study, BMI, physical activity, and plasma levels of glucose and uric acid explained only 20% of the variation in FGF21 concentrations. Other factors must be involved in the regulation of FGF21. For example, we found a negative association between FGF21 and RBP4, total, and HMW adiponectin. However, these associations lost their statistical significance in multivariate analysis. Adiponectin has been associated with insulin-sensitizing actions (36). Patients with insulin resistance and the metabolic syndrome have lower HMW and total adiponectin levels compared with their control peers (15). Based on this, the negative association between total and HMW adiponectin and FGF21 is predictable. Additional studies are required to confirm the potential interrelationships existing between these two hormones. Finally, RBP4 is another adipokine that may be associated with FGF21 physiology; a negative association between RBP4 and FGF21 was found in women. This relationship could be gender specific but needs confirmation. However, an alternative explanation is a sample size effect. The number of women included in our study sample was double that of the number of men.

Some limitations of this study must be recognized. Indirect estimates of insulin sensitivity (instead of the euglycemic clamp technique) were used in our report. This may limit our ability to identify associations between insulin sensitivity and FGF21. In addition, we did not measure circulating FFA levels, or the urinary excretion of uric acid. The prevalence of the metabolic syndrome in our cohort was relatively high, and the results of this study are applicable only to similar populations.

**Supplementary data**

This is linked to the online version of the paper at [http://dx.doi.org/10.1530/EJE-10-0454](http://dx.doi.org/10.1530/EJE-10-0454).

**Declaration of interest**

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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