Individuals with isolated congenital GH deficiency due to a GHRH receptor gene mutation appear to cope better with SARS-CoV-2 infection than controls

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Abstract

Purpose Several interactions exist between the GH/IGF axis and the immune system, including effects on innate immunity and humoral and cellular response. Acquired GH deficiency (GHD) has been recently proposed as a risk factor for severity of COVID-19 infections. However, acquired GHD is often associated to other factors, including pituitary tumors, surgery, radiotherapy, and additional pituitary hormones deficits and their replacements, which, together, may hinder an accurate analysis of the relationship between GHD and COVID-19. Therefore, we decided to assess the seroprevalence of SARS-CoV-2 antibodies and the frequency of symptomatic cases of COVID-19 in adults subjects with untreated isolated GHD (IGHD) due to a homozygous null mutation in the GHRH receptor gene.

Methods A cross-sectional study was carried out in 27 adult IGHD subjects and 27 age- and gender-matched local controls. Interview, physical examination, bio-impedance, hematological and SARS-CoV-2 IgM and IgG antibodies were analyzed.

Results There was no difference in the prevalence of positivity of anti-SARS-CoV-2 IgM and IgG antibodies between the two groups. Conversely, no IGHD individual had a previous clinical diagnosis of COVID-19 infection, while 6 control subjects did ($p = 0.023$).

Conclusion The production of anti-SARS-CoV-2 antibodies was similar between IGHD subjects due to a GHRH receptor gene mutation and controls, but the evolution to symptomatic stages of the infection and the frequency of confirmed cases was lower in IGHD subjects than in GH sufficient individuals.

Keywords GH · GHRH receptor · IGF-I · COVID-19 · SARS-CoV-2

Introduction

The function of the GH/Insulin-Like Growth Factors (IGFs) system is extremely complex. It involves two circuits: one critical for body size, the somatotropic axis, including GH releasing hormone (GHRH)/pituitary GH/and circulating IGF-I; and other, the extra-pituitary GH/IGF-I and IGF-II production in several sites, more relevant for specialized body functions including immunity [1, 2]. There is a crosstalk between the GH/IGFs system and the immune system, although its exact implications are not fully understood. These interactions may include effects on innate immunity and humoral and cellular response [3, 4]. The generation of IGF-I requires binding of GH to the GH
receptor, a class I cytokine receptor that triggers phosphorylation of Janus Kinase 2, and activation of the transcriptional-5b gene (STAT5b) [5, 6]. Prolactin, gamma interferon and several interleukins (IL) also bind to cytokine receptors and activate the STAT family [6]. The binding of IL-2 to its receptor and the activation of STAT5b lead to the activation of T lymphocytes and inhibition of its apoptosis (protection against infections), and to the development and proliferation of regulatory T cells (attenuating autoimmunity) [7]. It is possible that local effects of GH, IGF-I, IGF-II, IGF binding proteins are more relevant for the immune system than the activity of pituitary GH and circulating IGF-I [3, 8, 9].

The immune consequences of GH deficiency (GHD) are still debated. Although subclinical changes in laboratory parameters (such as chemotaxis and phagocytosis by granulocytes, macrophages, and natural killer cell activity) have been described in GHD, this condition does not appear to induce relevant alteration of the immune system in humans [10]. The COVID-19 pandemic, with 127 million confirmed cases and a death toll exceeding 2.7 million as of February 2021 [11], constitutes a public health emergency of international concern [12]. Male gender, old age, and obesity are, among others, known risk factors for mortality during this pandemic. As these conditions are associated with decreased GH secretion, it has been hypothesized that GHD may be a risk factor for the severity of COVID-19 [13, 14]. Acquired GHD has been proposed as a model to study the interactions between COVID-19, GH, and immunity [13, 14]. However, acquired GHD is often associated to several confounders such as pituitary tumors, surgery, radiotherapy, and several other pituitary hormone deficits and their replacements, which, together, may hinder an accurate analysis of the relationship between GHD and COVID-19.

We have previously described extended kindred with severe, congenital isolated GHD (IGHD) due to the c.57 +1G→A mutation in the GHRH receptor gene (GHRHR, OMIM n.618157) residing in Itabaianinha county, in the northeastern Brazilian state of Sergipe [15]. These subjects exhibit throughout life extremely low GH levels and very low to undetectable serum IGF-I levels [16]. They have severe short stature, visceral obesity [17], but increased insulin sensitivity [18], and normal longevity [19]. We have shown that these IGHD subjects do not exhibit increased frequency of infections or a significant alteration in humoral immune response [20]. However, they exhibit reduction of total IgG levels, smaller papule diameter after streptokinase test, and a non-significant tendency to reduced cellular response to the three combined tests: streptokinase, protein purified derived and Candida albicans antigen. These differences seem without clinical relevance in normal conditions, but may contribute to an unfavorable outcome in situations of more severe infections by extracellular pathogens [20]. Conversely, In vitro and in vivo studies in experimental models of an intracellular pathogen, *Leishmania amazonensis* have shown a positive effect of low serum IGF-I on this infection [21, 22]. Indeed, macrophages from the IGH subjects of Itabaianinha are less prone to Leishmania infection compared to GH sufficient controls [23].

This population represent an ideal model to study whether GHD influences the prevalence or severity of SARS-CoV-2 infections. The seroprevalence of SARS-CoV-2 antibodies in an asymptomatic population in Itabaianinha county was quite high in May 2020 (13.2% for IgM and 10.2 % for IgG) [24]. Therefore, in October 2020 we measured the seroprevalence of SARS-CoV-2, as well as the frequency of confirmed cases of COVID-19 in this period in a group of asymptomatic individuals with IGHD and compared with local controls. In addition, we analyzed some demographic and clinical variables that may influence this infection.

### Subjects and methods

#### Subjects

In a cross-sectional study, IGHD and age- and sex-matched control subjects were recruited by advertising placed in the local Dwarfs Association building, and by word of mouth among the inhabitants of Itabaianinha, a homogeneous miscegenated population in rural northeastern Brazil. Inclusion criteria for IGHD were homozygosity for the c.57 +1G→A *GHRHR* mutation, while homozygosity for the wild-type *GHRHR* allele was required for the control group. Exclusion criteria were age less than 12 years, GH treatment in the past ten years, and current signs and symptoms of COVID 19, namely cough, sore throat, fever, muscle or joint pain, fatigue, or headache [25]. Presently, we are aware of 54 living IGHD individuals in this cohort [2]. With the obvious difficulties of the time of the pandemic, especially fear of contamination, we decided to study half of this number, 27 IGHD subjects. From our database of genotyped homozygous normal subjects, we randomly enrolled the first 27 controls with similar age and sex distribution to the IGHD group, living in the same household or neighborhoods. Both groups are followed in the same primary care unit by one of the authors (HTS-J). They also have direct and rapid access to tertiary medical care at the University hospital. None of the 54 IGHD living individuals was diagnosed or hospitalized for Covid 19.

Collection was accomplished in three successive days in October 2020, with all the standard safety procedures. The Federal University of Sergipe Institutional Review Board approved these studies, and all subjects gave informed consent.
Interview, physical examination, bio-impedance, and laboratory assessment

The subjects were first submitted to an interview including demographic data; previous signs and symptoms of COVID-19 from May to October 2010 registered in medical files, and with COVID-19 diagnosis confirmed by real-time reverse transcription polymerase chain reaction (qRT-PCR) [21]. Risk factors for COVID-19 complications, such as cardiovascular disease, diabetes, arterial hypertension, and dyslipidemia were recorded. Body weight and height were measured, and body mass index was calculated. Blood pressure was calculated by the average of three measurements obtained in the left arm after 10 minutes of rest in the sitting position using a mercury sphygmomanometer with a cuff appropriate for the size of the arm. Bioimpedance was performed using the HBF-514C Full Body Digital Bioimpedance Scale, Omron Health Care Co., Ltd, Kyoto, 617–0002, Japan. Blood was collected for blood count (red, white blood cells and platelets) and for IgM and IgG measurements performed by immunofluorescence assays at the Laboratory of Biochemistry and Clinical Immunology (LaBiC-Imm) of the Federal University of Sergipe (UFS). Anti-SARS-CoV-2 IgM and IgG antibodies were detected in sera using an in vitro diagnostic test system based on lateral flow sandwich detection immunofluorescence technology (Ichroma™ COVID-19 Ab in conjunction with an Ichroma™ II Reader, Boditech Med Inc., South Korea). The immunofluorescence method applied showed a sensitivity of 95.8% and a specificity of 97%. A validation study was performed by qRT-PCR, which confirmed 60 cases of COVID-19, and 60 negative subjects [24].

Statistical analysis

The continuous variables were expressed as mean (standard deviation) and were compared by Student’s t test. Categorical variables were compared by the Fisher’s exact test. We also provide the 95% confidence interval (95% CI) for the difference of means in each comparison of the continuous variables [26]. The 95% CI that does not contain zero implies a significant difference between the groups. The effect size was estimated by the Cohen’s test in small (0.20 to 0.30), medium (0.40 to 0.7) and large (≥0.80). The software IBM SPSS® Statistics version 20 was used, and the statistical significance was set at $p < 0.05$.

Results

There was no significant difference in gender (12 females in IGHD and 17 in controls, $p = 0.137$), number of diabetics (three IGHD, four in control, $p = 0.698$), dyslipidemia (three in IGHD and two in control, $p = 1$), arterial hypertension (five in IGHD and seven in controls $p = 0.417$), cardiac disease (three in IGHD and one in controls, $p = 0.611$), and respiratory disease (one in IGHD and three in controls, $p = 0.341$). Table 1 shows the comparison of the anthropometric, hematological, and immunological data between the two groups. As expected, height, weight, body surface, and lean mass were lower in IGHD ($p < 0.0001$ in all cases). Body mass % had a trend to being higher ($p = 0.07$), and visceral fat mass was higher in IGHD group ($p = 0.006$). There were also no differences in BMI, blood pressure, and in any of the hematology data. Similarly, there was no difference in the prevalence of Anti-SARS-CoV-2 IgM and IgG antibodies between the two groups. However, no IGHD individual had a history of symptoms or diagnosis of COVID-19 infection, while 6 control subjects had it ($p = 0.023$), with a mean difference of 22.8%, 95% CI (4.7–38.1%), and an effect size of 0.535. Of the six subjects who had symptoms of Covid disease, four were positive for both IgM and IgG, one was positive for IgG and the other for IgM.

Discussion

This study suggests an unexpected advantage of an endocrine genetic defect causing IGHD [23]. IGHD individuals with severe short stature and extremely low GH and undetectable IGF-I levels [16], in an environment with a high transmission rate of SARS-CoV-2, appear to cope better with this infection than GH sufficient controls. It is noteworthy that the positivity of anti-SARS-CoV-2 IgM and IgG antibodies was similar between IGHD and controls, and about three times greater than that recorded, with the same method, in Itabaianinha five months earlier, indicating an acceleration of the pandemic [24]. Despite this acceleration, in this period no IGHD subject had a clinical diagnosis of COVID-19, or reported symptoms that would be consistent with it, while six people had it in the control group. No differences in age, sex and associated comorbidities were identified between the groups that are known to influence the clinical outcome of SARS-CoV-2 infection. Similarly, no hematological data was different between the two groups. Mask use, social distancing, and personal hygiene are likely similar in the two groups, because most IGHD and controls live in the same house or neighborhood.

One variable that may have an impact in the apparent protection against the COVID-19 is stature. Although most people believe that being taller is a sign of higher social status and privilege, shorter, smaller bodies have numerous advantages in terms of health and longevity. With adequate nutrition and lifestyle, and good medical care, short people are less likely to suffer from age-related chronic diseases
and more likely to reach advanced age [27]. Several biological factors can explain these benefits, including reduced cell replication, lower DNA damage and reduced cancer incidence, higher sex hormone binding globulin, lower IGF-I, IGF binding protein, and insulin [27]. Several findings are present in the Itabaianinha IGHD subjects [1, 2]. The results of a survey of 2000 people in the UK and the USA indicated that height above 1.8 m is a significant predictor of risk SARS-CoV-2 infection for men suggesting that downward droplet transmission may be less important than aerosol transmission https://www.medrxiv.org/content/10.1101/2020.07.13.20152819v1. Because the positivity of Anti-SARS-CoV-2 antibodies was similar between the IGHD and controls, we can infer that stature, in the studied range, did not influence the rate of infection with SARS-CoV-2. However, short stature may be a protective factor through a reduced viral load, which could influence the lower rates of COVID-19 symptoms.

The second aspect that can have an impact in our findings is body composition. IGHD subjects have reduced lean mass and higher visceral fat mass than controls [17, 28]. Obesity is a strong risk factor for the severity of COVID19 disease. The reasons of this increased risk may involve mechanical alterations of the airways and lung parenchyma, systemic and airway inflammatory, and metabolic dysfunction [29, 30]. It is known that visceral obesity characterizes a chronic low-grade inflammation, with abnormal secretion of adipokines and cytokines like interleukin 6 (IL-6), TNF-α and interferon, as it has been described in H1N1 influenza and other infections, as well as in asthma [31, 32]. This inflammatory pattern could cause an impaired immune response to the infection by SARS-CoV-2. Our data suggest that visceral obesity in the context of IGHD, does not confer a greater infectious risk, similarly to the previously described lack of risk of metabolic [17, 18] and cardiovascular disease [33, 34]. This surprising association of metabolic and immune benefits with increased visceral adiposity is also shown in GH resistant (GHRKO) mice, that exhibit favorable changes in visceral fat secretory activity, like an increase in adiponectin [35, 36], also present in our IGHD subjects [37]. Accordingly, the visceral fat of GHRKO mice has reduced infiltration of inflammatory cells and therefore less intense inflammatory potential [35]. Interestingly, hyperactivation of the mechanistic target of rapamycin (mTOR) pathway seems a possible contributor to the higher rate and severity of COVID-19 in obese patients [38].

Table 1 Comparison of the anthropometric, hematological, and immunological data in 27 IGHD and 27 controls

|                          | IGHD            | Controls        | 95% CI          | p    |
|--------------------------|-----------------|-----------------|-----------------|------|
| Age (years)              | 47.0 (16.5)     | 46.9 (17.1)     | −9.0 to 9.3     | 0.974|
| Height (m)               | 1.30 (0.1)      | 1.62 (0.1)      | −0.37 to −0.26  | <0.0001|
| Weight (kg)              | 43.8 (10.2)     | 67.4 (10.6)     | −29.5 to −17.7  | <0.0001|
| Body surface (m²)        | 1.21 (0.13)     | 1.71 (0.18)     | −0.59 to −0.42  | <0.0001|
| BMI (kg/m²)              | 24.5 (7.3)      | 25.6 (3.3)      | −4.45 to 2.19   | 0.480|
| Diastolic BP (mmHg)      | 65.2 (11.7)     | 69.5 (10.2)     | −10.69 to 2.14  | 0.186|
| Systolic BP (mmHg)       | 119.2 (23.1)    | 120.3 (21.8)    | −14.16 to 11.9  | 0.862|
| Lean mass (kg)           | 26.7 (9.7)      | 34.0 (7.48)     | −14.11 to 11.8  | 0.028|
| Body fat (%)             | 34.8 (13.1)     | 26.6 (11.4)     | −0.64 to 16.9   | 0.068|
| Visceral fat (kg)        | 10.7 (4.4)      | 7.1 (2.5)       | 1.1 to 6.1      | 0.006|
| Red blood cells (10⁶/mm³)| 4.7 (0.4)       | 4.6 (0.5)       | −0.22 to 0.37   | 0.597|
| Platelets (10⁹/mm³)      | 275 (55.3)      | 259.8 (66.5)    | −23.0 to 53.4   | 0.426|
| White blood cells (10⁹/mm³)| 8.1 (2.0)   | 7.1 (2.6)       | −0.45 to 2.47   | 0.171|
| Hemoglobin (g/dL)        | 13.8 (1.2)      | 13.4 (1.1)      | −0.4 to 1.03    | 0.378|
| Hematocrit (%)           | 41.2 (3.5)      | 40 (3.6)        | −1.05 to 3.4    | 0.295|
| Neutrophils (number/mm³)| 4866 (1755)     | 4039 (1651)     | −236 to 1889    | 0.124|
| Lymphocytes (number/mm³)| 2302 (652)      | 2179 (1001)     | −403 to 651     | 0.638|
| Monocytes (number/mm³)   | 560 (253)       | 570 (304)       | −184 to 165     | 0.913|
| Eosinophils (number/mm³)| 393 (449)       | 261 (256)       | −96.5 to 360.3  | 0.250|
| Basophils (number/mm³)   | 4.8 (6.7)       | 3.3 (4.8)       | −2.2 to 5.1     | 0.437|
| Positive IgM only n (%)  | 5 (18.5)        | 4 (14.8)        | −15 to 22.2     | 0.720|
| Positive IgG only n (%)  | 1 (3.7)         | 5 (18.5)        | −5.2 to 34.8    | 0.190|
| Positive IgM and IgG n (%)| 6 (22.2)   | 5 (18.5)        | −21.5 to 28.9   | 0.735|
| COVID-19 diagnosis n (%) | 0 (0)           | 6 (22.2)        | 4.7−38.1        | 0.023|

The continuous variables were analyzed by Student t test and the categorical data by the Fisher exact test. 95% CI: 95% confidence interval for the difference of means.
enhancing the virus entry into host cells, through its human angiotensin-converting enzyme 2 receptor [39]. Some evidence suggests that the mTOR pathway is down-regulated in GH deficient mice and humans [1], which can exert a protective effect against a deleterious progression of the SARS-CoV-2 infection in the IGHD subjects.

In interpreting these data and in considering a wide application on the effects of GHD on the clinical manifestations of SARS-CoV-2 infection, one should consider many differences between our subjects and patients with acquired GHD. In addition to the previously mentioned differences related to acquired GHD (often seen in panhypopituitarism with history of surgery and/or radiation), one important aspect is the vitamin D status. Acquired GHD patients often exhibit vitamin D deficiency, a condition that has been linked to an increased risk of systemic infections and to immune response impairment [40]. Vitamin D insufficiency in acquired GHD may reflect reduced sun light exposure, sedentary lifestyle, co-morbidities, and drug use (such as anti-convulsant). Conversely, vitamin D deficiency it is not found in quite active IGHD subjects from Itabaianinha, who often work outdoor under direct sun exposure [41].

Our work has some limitations. First, the apparently small number of 27 individuals per group. However, with the effect size of 0.535 in the diagnosis of COVID-19, estimating a power of 0.8 with α of 0.05, we calculated that 20–30 individuals in each group would be needed, making the number appropriate. Second, in the very dynamic process of an expanding the pandemic, we carried out our study in a specific period, without longitudinal data. However, the two groups were studied in the same week, allowing for appropriate comparison. Thirdly, these results were obtained in a group with congenital IGHD, due to a particular mutation in the GHRH receptor gene, which is different from most individuals with GHD in clinical practice.

In conclusion, subjects with congenital untreated IGHD due to a GHRH receptor gene mutation appear to cope better with SARS-2-coV-2 infection than GH sufficient controls, with reduced risk of progression to symptomatic stages, suggesting a protective effect on the progression of the infection.

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Compliance with ethical standards

Conflict of interest R.S. serves on Novordisk Advisory Board. The other authors declare no competing interests.

Ethical approval All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

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