Increased Glucocorticoid Receptor Alpha Expression and Signaling in Critically III Coronavirus Disease 2019 Patients*

OBJECTIVES: Critical illness is characterized by increased serum cortisol concentrations and bioavailability resulting from the activation of the hypothalamicpituitary-adrenal axis, which constitutes an essential part of the stress response. The actions of glucocorticoids are mediated by a ubiquitous intracellular receptor protein, the glucocorticoid receptor. So far, data on coronavirus disease 2019 and glucocorticoid receptor alpha expression are lacking.

DESIGN: Prospective observational study.

SETTING: One academic multidisciplinary ICU.

SUBJECTS: Twenty-six adult coronavirus disease 2019 patients; 33 adult noncoronavirus disease 2019 patients, matched for age, sex, and disease severity, constituted the control group. All patients were steroid-free.

INTERVENTIONS: None.

MEASUREMENTS AND MAIN RESULTS: Glucocorticoid receptor alpha, glucocorticoid-inducible leucine zipper expression, and serum cortisol were measured on ICU admission. In coronavirus disease 2019 patients, glucocorticoid receptor alpha and glucocorticoid-inducible leucine zipper messenger RNA expression were upregulated (4.7-fold, p < 0.01 and 14-fold, p < 0.0001, respectively), and cortisol was higher (20.3 vs 14.3 µg/dL, p < 0.01) compared with the control group.

CONCLUSIONS: ICU coronavirus disease 2019 patients showed upregulated glucocorticoid receptor alpha and glucocorticoid-inducible leucine zipper expression, along with cortisol levels, compared with ICU noncoronavirus disease 2019 patients. Thus, on ICU admission, critical coronavirus disease 2019 appears to be associated with hypercortisolemia, and increased synthesis of glucocorticoid receptor alpha and induced proteins.

KEY WORDS: coronavirus disease 2019; corticosteroid response; critical illness; glucocorticoid receptor alpha; glucocorticoid-inducible leucine zipper

The clinical spectrum of coronavirus disease 2019 (COVID-19) is heterogeneous, ranging from a flu-like syndrome to severe pneumonia, requiring monitoring and support in an ICU (1). Critical illness is an acute stress condition, characterized by increased concentration of cortisol due to activation of the hypothalamic-pituitary-adrenal (HPA) axis (2). Most of the immunologic, metabolic, and hemodynamic actions of glucocorticoids (GCs) are mediated by a ubiquitous intracellular receptor protein, the glucocorticoid receptor (GCR). In humans, glucocorticoid receptor alpha (GCR- α) is the predominant and functionally active receptor (3). The cortisol-GCR complex translocates to the nucleus and exerts transcriptional activation or repression of GC-target genes, such as the glucocorticoid-inducible leucine zipper (*GILZ*). The end-result is inhibition of the proinflammatory response. Alice G. Vassiliou, PhD¹ Nikolaos Athanasiou, MD¹ Chrysi Keskinidou, MSc¹ Edison Jahaj, MD¹ Stamatios Tsipilis, MD¹ Alexandros Zacharis, MD¹ Efthimia Botoula, BSc² Aristidis Diamantopoulos, MD² Ioannis Ilias, MD³ Dimitra A. Vassiliadi, MD² Stylianos Tsagarakis, MD² Anastasia Kotanidou, MD¹ Ioanna Dimopoulou, MD¹

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A previous study in a non-ICU COVID-19 cohort showed activation of the HPA axis; patients exhibited an increase in cortisol, which was significantly higher than in those without COVID-19 infection (4). However, so far, data on COVID-19 and *GCR-\alpha* expression are lacking.

In view of the above, in the present study, we measured whole blood expression of the key receptor, $GCR-\alpha$, the expression of a GC-target gene, *GILZ*, and the levels of the signaling molecule, cortisol, in ICU COVID-19 patients.

MATERIALS AND METHODS

This prospective, observational study was approved by the "Evangelismos" Hospital Research Ethics Committee (170/24-4-2020), and all procedures carried out were in compliance with the Helsinki Declaration. Informed written consent was obtained from all patients' next-ofkin. Enrolment took place between May and October 2020. A Consolidated Standards of Reporting Trials flow diagram shows the enrolment process (**Supplementary Fig. S1**, http://links.lww.com/CCM/G486). A control group of 33 adult ICU non-COVID-19 patients, matched for age, sex, and disease severity, was also included. None of the patients had received corticosteroids, including dexamethasone, prior to blood sampling.

 $GCR-\alpha$ and GILZ expression, and serum cortisol were measured on ICU admission. Detailed methods are given in the **Online Supplementary Material** (http://links.lww.com/CCM/G486).

RESULTS

Characteristics of the Study Population

In Greece, the first pandemic wave did not hit hard the country, possibly due to the early lockdown on March 10, 2020. Between March and April, there were between 30 and 130 daily new cases, which declined in May. In August, cases started rising again, reaching up to 1,000 in October.

Twenty-six adult ICU COVID-19 patients were enrolled (Supplementary Fig. S1, http://links.lww. com/CCM/G486). The ICU non-COVID-19 patients consisted of trauma (n = 17), medical (n = 7; CNSrelated pathologies), and surgical (n = 9) cases. Eleven of the patients (33%) had sepsis or septic shock on ICU admission. Characteristics of the two patient groups are presented in **Table 1**. In ICU COVID-19 patients, comorbidities were more frequent, oxygenation was worse, whereas fibrinogen and lactate dehydrogenase were higher. ICU mortality in COVID-19 was similar to our overall ICU mortality (~30%). This mortality compares favorably with the 40–50% mortality rate reported worldwide for ICU admissions during this period and could be possibly explained by the fact that not all patients were mechanically ventilated.

GCR- α Expression, Signaling, and Cortisol Levels in ICU COVID-19 Patients

At admission and compared with the non-COVID-19 controls, COVID-19 patients exhibited a 4.7-fold higher *GCR-* α messenger RNA expression (interquartile range [IQR], 2.1–7.8; *p* < 0.01) (**Fig. 1***A*). Furthermore, *GILZ* expression was upregulated by 14.1-fold (IQR, 6.2–27.8; *p* < 0.0001) (**Fig. 1***B*). Cortisol was also higher (20.3 vs 14.3 µg/dL; *p* < 0.01) (**Fig. 1***C*), and none of the patients had cortisol less than 10 µg/ dL (Fig. 1*C*). Interleukin (IL)-6 and IL-10 levels were comparable among the groups (**Fig. 1**, *D* and *E*).

GCR- α and GILZ messenger RNA (mRNA) expression did not correlate with cortisol ($r_s = -0.028$, p = 0.9 and $r_s = 0.159$, p = 0.4, respectively). Furthermore, GILZ expression did not correlate with IL-6 or IL-10 ($r_s = 0.172$, p = 0.4 and $r_s = 0.362$, p = 0.1, respectively). However, GILZ expression strongly correlated with GCR- α expression ($r_s = 0.551$, p = 0.004).

DISCUSSION

In this prospective study, we measured $GCR-\alpha$ expression, signaling, and serum cortisol levels, in ICU COVID-19 patients relatively to a general ICU population of similar disease severity. The novel finding is that receptor expression and signaling, and cortisol were higher in ICU COVID-19 patients. Furthermore, according to baseline cortisol, none of the patients with COVID-19 exhibited critical-illness–related corticosteroid insufficiency (CIRCI).

Nearly every tissue in the body expresses GCR, highlighting its critical role in homeostasis and survival. Data on *GCR-* α expression and critical illness in humans point toward GC resistance in sepsis; specifically, results have shown lower *GCR-* α expression in septic patients, thereby limiting the anti-inflammatory effects of GCs (5). In our ICU COVID-19 patients, *GCR-* α expression was elevated, indicating that GC

TABLE 1. Clinical Characteristics and Laboratory Data of Critically III Patients on ICU Admission

Characteristics	ICU COVID-19	ICU Non-COVID-19	p	Reference Values
Number of patients, <i>n</i>	26	33		
Age (yr), median (IQR)	62 (55–75)	57 (43–74)	0.2	
Sex, n (%)				
Male	19 (73.1)	20 (60.6)	0.3	
Female	7 (26.9)	13 (39.4)		
Comorbidities, n (%)	18 (69.2)	12 (36.4)	0.01	
Hypertension	13	7		
Hyperlipidemia	6	0		
Diabetes	3	1		
Coronary artery disease	2	2		
Chronic obstructive pulmonary disease	0	1		
Chronic kidney disease	0	1		
Acute Physiology and Chronic Health Evaluation II score, mean ± sp	15 ± 5	17 ± 6	0.2	
Pao ₂ /Fio ₂ (mm Hg), median (IQR)	178 (127–261)	245 (189–353)	0.003	
Vitals signs				
Heart rate (beats/min), median (IQR)	92 (82–106)	82 (64–104)	0.05	
Mean arterial pressure (mm Hg), median (IQR)	75 (71–86)	75 (69–97)	0.9	
Temperature (°C), median (IQR)	37.9 (37.4–38.2)	37.0 (35.5–37.9)	0.005	
Laboratory data				
Hemoglobin, mean ± s⊳	13.1 ± 1.8	10.0 ± 2.3	<0.0001	12-17.5
WBC count (per μ L), mean ± sp	$10,480 \pm 5,000$	$14,170 \pm 5,850$	0.02	$4-10.5 \times 10^{3}$
Neutrophils (%), mean ± sp	81.6 ± 6.7	81.2 ± 6.7	0.8	40-70
Lymphocytes (%), mean ± sp	13.0 ± 6.2	10.1 ± 4.7	0.08	25-45
Platelets (per μ L), median (IQR)	220,000 (143,000–270,000)	163,000 (126,000–230,000)	0.04	140-450 × 10 ³
Urea (mg/dL), median (IQR)	31 (25–53)	31 (21–37)	0.5	10-50
Creatinine (mg/dL), median (IQR)	0.9 (0.7-1.2)	0.7 (0.6–0.9)	0.05	0.6-1.4
Glucose (mg/dL), median (IQR)	137 (119–187)	122 (100–183)	0.1	70-110
Total bilirubin (mg/dL), median (IQR)	0.7 (0.4–1.1)	0.6 (0.4–1.0)	0.5	<1
C-reactive protein (mg/dL), median (IQR)	9.7 (4.4–20.8)	9.9 (5.8–17.5)	0.9	<0.5
Fibrinogen (mg/dL), mean ± s⊳	631 ± 161	421 ± 167	<0.0001	200-400
Lactate dehydrogenase (U/L), median (IQR)	465 (345–632)	290 (206–460)	0.001	<225
Inotropes (mean dose, μg/kg/min), median (IQR)	0.2 (0.0-0.3)	0.3 (0.1–0.5)	0.2	
Outcomes				
Length of stay in the ICU (d), median (IQR)	23 (15–40)	21 (10–31)	0.2	
Mechanical ventilation, n (%)	21 (80.8)	33 (100)	0.01	
Mechanical ventilation duration (d), median (IQR)	15 (5–37)	10 (5–23)	0.2	
ICU mortality, n (%)	8 (30.8)	11 (33.3)	0.8	

COVID-19 = coronavirus disease 2019, IQR = interquartile range.

Data are expressed as percentages of total related variable (%) and mean \pm sp for normally distributed variables and median (IQR) for skewed data.

Patients' outcome and demographics are also shown.

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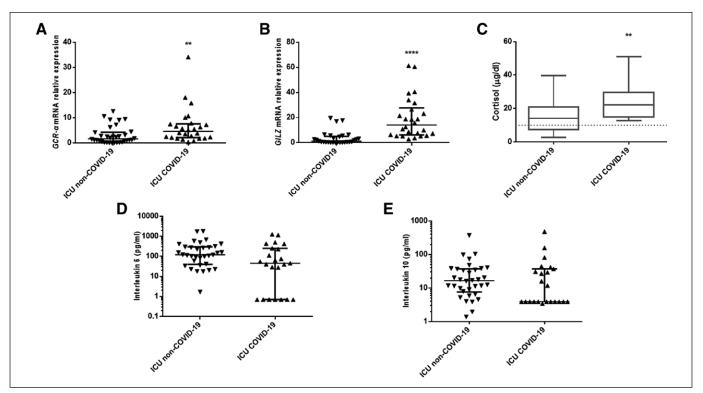


Figure 1. Corticosteroid response in coronavirus disease 2019 (COVID-19). Distribution of glucocorticoid receptor alpha (*GCR-a*) (**A**) and glucocorticoid-inducible leucine zipper (*GILZ*) (**B**) expression in 26 COVID-19 critically ill patients and 33 non-COVID-19 critically ill patients on ICU admission. Data are presented as scatter plots. *Line in the middle*: median value; *Iower and upper lines*: 25th–75th percentiles. **p < 0.01, ****p < 0.0001 by Mann-Whitney *U* test. **C**, Morning cortisol levels were measured in 26 COVID-19 critically ill patients and in 33 non-COVID-19 critically ill patients on ICU admission. Data are presented as the provided set of values; *Iower and upper lines*: 25th–75th percentiles: whiskers: range of values; *Iorizontal line*: cortisol levels 10 µg/dL. **p < 0.01 by Mann-Whitney *U* test. **D** in 26 COVID-19 critically ill patients and 33 non-COVID-19 critically ill patients on ICU admission. Data are presented as scatter plots. *Line in the box*: median value; *Iower adges*: 25th to 75th percentiles; *whiskers*: range of values; *Iorizontal line*: cortisol levels 10 µg/dL. **p < 0.01 by Mann-Whitney *U* test. Distribution of interleukin-6 (**D**) and interleukin-10 (**E**) in 26 COVID-19 critically ill patients and 33 non-COVID-19 critically ill patients on ICU admission. Data are presented as scatter plots. *Line in the middle*: median value; *Iower and upper lines*: 25th–75th percentiles. mRNA = messenger RNA.

resistance may not constitute a problem. Furthermore, downstream GCR- α signaling occurred, as confirmed by transactivation of the GC-inducible gene *GILZ*. These might be regarded as beneficial responses to increase the effectiveness of corticosteroid therapy on target cells (6). According to recent data, dexamethasone's beneficial effect on survival is mainly attributed to its potent immunosuppressive effects (7). Dexamethasone at the recommended dose (6 mg) would probably not have a substantial effect if GC resistance was present; our data showing a functional GCR-a support the results of studies, showing increased survival rates when administered while the disease is dominated by immunopathological elements.

Cortisol secretion is essential for surviving acute illness. One immunoevasive strategy used by other viruses, such as the severe acute respiratory syndrome (SARS) coronavirus or the influenza virus, is to inhibit its host's corticosteroid stress response (8). These viruses express amino acid sequences homologous to human adrenocorticotropic hormone (ACTH). The host produces antibodies against viral antigens, which, however, may bind to the host's ACTH, resulting in an inability of ACTH to stimulate cortisol secretion. The overall result is adrenal cortex dysfunction (8). Furthermore, viruses, in particular SARS coronavirus 2, may cause hemorrhage, necrosis, or thrombosis at the adrenal level causing acute hypoadrenalism (9). These assumptions provide compelling evidence that the HPA axis is a potential target of COVID-19. Indeed, in critically ill patients, cortisol was lower in COVID-19 patients compared with non-COVID-19 patients (10), whereas hospitalized patients with more severe disease had lower cortisol and ACTH levels, suggesting a direct link between COVID-19 infection and impaired GC response (11). On the other hand, noncritically ill COVID-19 patients exhibited extremely high cortisol, which was associated with higher mortality rates (4). Similarly, in our ICU COVID-19 patients, cortisol was higher than that in their counterparts without COVID-19, despite similar disease severity. CIRCI is characterized by the inability of the organism to produce sufficient cortisol or resistance of the tissues to its actions, or both (12). We did not assess adrenal function following stimulation with synthetic ACTH; instead, we used the alternative for CIRCI definition (morning baseline cortisol levels <10 μ g/dL) (13). We found that none of our patients fulfilled the CIRCI criteria, contrasting the well-established knowledge that a subset of critically ill patients might exhibit adrenal dysfunction (12). Whether cortisol may be useful as a biomarker to monitor COVID-19 severity and ICU outcomes remains to be investigated.

Critical illness is often accompanied by hypercortisolemia, mainly attributed to stress-induced activation of the HPA axis. However, higher cortisol in critically ill patients may also be a consequence of normal production levels but reduced cortisol metabolism (14, 15). Data on the disturbance of the pulsatile and circadian nature of the HPA axis and its potential clinically important impact on gene expression rhythms of core clock genes are emerging (16).

A major limitation of this article is that *GILZ* measurement in whole blood reflects both neutrophil and monocyte production, perhaps even more the former. Since monocytes are the cells of interest in the immunomodulatory effects of GCs, it would be more appropriate to look at isolated monocytes. Another limitation of our study was the relatively small number of patients but similar to other investigations on the role of *GCR* mRNA expression levels in critical illness and sepsis.

CONCLUSIONS

Our data provide novel evidence on the upregulated expression and signaling of the functionally active GCR- α isoform in the peripheral blood of ICU COVID-19 patients. Circulating baseline cortisol was high, and consequently, our patients did not meet CIRCI criteria. Overall, it seems that critical infection with COVID-19 appears to be associated with hypercortisolemia, and increased synthesis of GCR- α and induced proteins.

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