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Glucocorticoid receptor expression after the return of spontaneous circulation in patients who experienced cardiac arrest: A prospective observational study

Journal:	BMJ Open
Manuscript ID	bmjopen-2021-060246
Article Type:	Original research
Date Submitted by the Author:	04-Jan-2022
Complete List of Authors:	Yu, Yanan; Capital Medical University, Department of Emergency Medicine Tang, Ziren; Capital Medical University, Department of Emergency Medicine Xie, Miaorong; Capital Medical University, Department of Emergency Medicine Li, Jiabao; Capital Medical University, Department of Critical Care Hang, Chen-Chen; Beijing Chao-Yang Hospital, Emergency Medicine An, Le; Capital Medical University, Department of Emergency Medicine Li, Chunsheng; Beijing Chaoyang Hospital, Department of Emergency Medicine
Keywords:	ACCIDENT & EMERGENCY MEDICINE, INTENSIVE & CRITICAL CARE, Adult intensive & critical care < INTENSIVE & CRITICAL CARE
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1	Glucocorticoid receptor expression after the return of spontaneous circulation in
2	patients who experienced cardiac arrest: A prospective observational study
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15	Word count: 3199
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23 Abstract

Objectives: Rapid changes in glucocorticoid (GC) levels and adrenal insufficiency are related to the development of post-cardiac arrest (CA) syndrome. However, changes in GC receptor (GR) expression have not been studied. Hence, the aim of this study was to investigate the association of early changes in GR expression and prognosis and immune response in patients who experienced CA.

- 29 **Design:** Prospective observational study.
- 30 **Setting:** Emergency department.

Participants: Patients (85) who were in the early period of return of spontaneous
circulation (ROSC) after CA and were admitted between October 2018 and October
2019. Age- and sex-matched healthy individuals (40) were recruited for the control
group after a physical examination.

Primary and secondary outcome measures: GR expression and cell counts of
circulatory T and B lymphocytes, natural killer cells, and regulatory T (Treg) cells were
assessed. Plasma total cortisol and adrenocorticotrophic hormone (ACTH) levels were
also tested.

Results: All cell counts were lower, and plasma total cortisol levels were higher (P<0.001), in patients who experienced CA than in the healthy control group. GR expression in Treg cells and CD3⁺CD4⁺ T lymphocytes was not significantly different, but the mean fluorescence intensity and GR expression in other cells were lower in patients who experienced CA (P<0.05) than in the healthy control group. ACTH levels were not different. There were no significant differences between survivors and non45 survivors.

46 Conclusions: This study revealed that GR expression and cell counts rapidly decreased, 47 whereas plasma total cortisol levels increased, in the early period after ROSC among 48 patients who experienced CA. Our findings provide insights into GC sensitivity and 49 immunosuppressive status in these patients, and a new perspective for GC targeted 50 treatment.

51 Strengths and limitations of this study

52 1. Explore whether controversy over glucocorticoid use is associated with different
53 levels of glucocorticoid receptor expression in cardiac arrest patients for the first
54 time.

Glucocorticoid receptor expression rapidly decreased in the early period following
 restoration of spontaneous circulation among patients who experienced cardiac arrest .
 We only observed changes in glucocorticoid receptor expression of cardiac
 arrest patients at the early period following restoration of spontaneous circulation, and
 long-term dynamic observation would be helpful to understand the significance of
 clinical steroid therapy.

61 Introduction

62 Cardiac arrest (CA) is an important health problem globally; about 356,500 people 63 experience medical emergencies due to CA in the United States, and over 544,000 64 people die from sudden CA in China annually. [1, 2] The systemic ischemia-reperfusion 65 response in patients who have experienced CA can present as post-CA syndrome 66 (PCAS) or systematic inflammatory response syndrome (SIRS), which increases the

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risk of multiple organ failure and infection and affects the inflammatory response and prognosis of patients after the return of spontaneous circulation (ROSC). [3-6] CA is the most intense among acute stress events, which seriously affect the function of the pituitary and adrenal axis. [7] Studies have shown that abnormal cortisol levels and relative adrenocortical insufficiency after ROSC in patients who experienced CA are related to their prognosis. [8-11] However, the clinical application of glucocorticoids (GCs) is controversial. In the 2015 International Cardiopulmonary Resuscitation Guidelines, the routine use of GCs is not recommended for the resuscitation of patients with in-hospital or out-of-hospital CA. [12] Recent clinical studies have shown that early administration of corticosteroids after CA can improve the success rate of ROSC, nervous system functional outcome, and prognosis, which is speculated to be related to its influence on hemodynamics, SIRS response, and other mechanisms. [12-17] Therefore, the role of GCs in the occurrence and development of

80 PCAS needs to be studied further.

GCs combine with intracellular GC receptors (GRs) to exert anti-inflammatory and immunosuppressive effects and reduce the production as well as release of inflammatory cytokines. [18, 19] The affinity of GRs to GCs in circulating monocytes is decreased in patients with acquired immunodeficiency syndrome. [20] The expression of GR is decreased in patients with critical illness, [21] pediatric septic shock, and high serum cortisol level. [22] However, hitherto, no study has reported the GR expression after ROSC in patients who experienced CA. Previous studies have found that the counts of circulating B and T lymphocytes, regulatory T (Treg) cells, and

89 monocytes and expression of human leukocyte antigen DR (HLA-DR) on circulatory 90 monocytes and B and T lymphocytes are reduced. [23, 24] Hence, the aim of this study 91 was to investigate the relationship between GR expression and immune alteration in the 92 early period after ROSC in patients who experienced CA by observing GR expression 93 in circulatory T and B lymphocytes, NK cells, and Treg cells, their cell counts, and 94 plasma total cortisol and adrenocorticotrophic hormone (ACTH) levels.

96 MATERIALS AND METHODS

97 Study participants

This was an observational study conducted in the Emergency Department (ED). Following the 2015 International Cardiopulmonary Resuscitation Guidelines, [25] we enrolled patients who were in the early period of ROSC after CA and were admitted to the ED between October 2018 and October 2019. The inclusion criteria were (a) ROSC 6 h after CA and (b) Glasgow Coma Scale score <8 after ROSC. The exclusion criteria were (a) ≤ 18 years of age, (b) terminal stage of disease (such as cancer of any type, acquired immunodeficiency syndrome), (c) corticosteroid treatment within the past 3 months, (d) administration of corticosteroids, and (e) adrenal insufficiency. All patients were treated according to the 2015 International Cardiopulmonary Resuscitation Consensus. [13] Age- and sex-matched healthy individuals were recruited for the control group after a physical examination.

110 Data collection

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We collected data on demographics, resuscitation (initial heart rhythm, ROSC time, and cumulative adrenaline epinephrine dose), and laboratory findings (routine blood cell counts, blood gas analysis, and blood biochemical tests performed 6 h after ROSC). Acute Physiology and Chronic Health Evaluation (APACHE) II and the Sequential Organ Failure Assessment (SOFA) were used to determine disease severity. Residual samples of blood, with heparin anticoagulant, from routine clinical tests or physical health examinations were collected, maintained at 4 °C during transport and storage, and used to determine GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells and their cell counts. The plasma was maintained at -80 °C during storage and used to determine total cortisol and ACTH levels. During follow-up, 28-day survival data were also collected. Supplemental Figure 1 shows the workflow of this LICZ study.

Flow cytometry

GR expression in T and B lymphocytes, NK cells, and Treg cells was measured. Briefly, a 100- μ L peripheral blood sample was stained for 20 min with surface antibodies (CD3, CD4, CD8, CD19, CD16, CD56, CD25, and CD127) in a dark place. Erythrocytes were lysed for 15 min, and the debris was washed away. Before intracellular GR staining, surface-stained cells were fixed and permeabilized using the BD Transcription Factor Buffer Set (BD Pharmingen, San Diego, USA, Catalogue No. 562574). Monoclonal antibodies and their isotype controls were all purchased from BD Biosciences (San Jose, CA, USA). Details of all antibodies are shown in Supplemental Table 1.

133	According to the manufacturer's recommendations, all antibodies and their isotype
134	controls were used at a concentration of 1 μL per 100 μL of whole blood. Samples were
135	measured using the Gallios flow cytometer (Beckman Coulter, Brea, CA, USA) and
136	analyzed using Gallios Software version 1.0 (Beckman Coulter). The flow cytometer
137	was periodically calibrated by an engineer. Cells were stained for 20 min; thresholds
138	were defined using the manufacturer's recommended isotype controls. T cells were
139	gated by CD3 ⁺ CD4 ⁺ or CD3 ⁺ CD8 ⁺ , B cells were gated by CD3 ⁻ CD19 ⁺ , NK cells were
140	gated by CD16 ⁺ CD56 ⁺ , and Tregs were gated by CD4 ⁺ CD25 ^{high} CD127 ^{low} . At least
141	10,000 events were collected in the lymphocyte cell gate for each sample. Results are
142	expressed as percentages and mean fluorescence intensity (MFI) values.
143	Absolute CD3 ⁺ and CD4 ⁺ lymphocyte, NK cell, and Treg cell counts were obtained
144	using Flow-Count fluorospheres (Beckman Coulter, Catalogue No. 7547053),
145	according to the manufacturer's instructions. B, CD3 ⁺ CD4 ⁺ T, CD3 ⁺ CD8 ⁺ T, and Treg
146	cell counts were calculated by their percentages in CD3 ⁺ or CD4 ⁺ lymphocytes
147	multiplied by CD3 ⁺ or CD4 ⁺ lymphocyte counts.
148	
149	Determination of plasma total cortisol and ACTH levels after ROSC

Venous blood samples were collected in ethylenediaminetetraacetic acid tubes,
centrifuged 10 min at 3000 rpm, and then stored at -80 °C. Plasma total cortisol
(IMMULITE 2000 Cortisol, L2KCO2, UK) and ACTH (IMMULITE 2000 ACTH,
L2KAC2, UK) levels were assayed using a chemiluminescent immunoassay on a
Siemens automated analyzer (IMMULITE 2000 XPi; Siemens Healthcare Diagnostics,

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Erlangen, Germany). The equipment and reagents were calibrated by engineers before
use. The lower detection limit of total cortisol was 2.00 ng/mL and that of ACTH was
5.00 pg/mL.

159 Statistical analyses

All data were analyzed using SPSS version 22.0 (IBM Corp., Armonk, NY, USA). For normally distributed data, continuous variables are expressed as means with standard deviations. Since the data for total cortisol and ACTH levels had a skewed distribution, we compared our results with the natural logarithmic conversion values after adding 1 (ln [total cortisol+ 1], ln [ACTH+ 1]). Measurement data with a skewed distribution are expressed as medians (25th and 75th percentiles). The Mann–Whitney U test was used to compare variables between groups. The qualitative parameters in the 2×2 contingency table were used for analysis. All statistical tests were two-tailed, and a P-value of <0.05 was considered statistically significant.

170 Follow-up

Patients who experienced CA were classified into survivor and non-survivor groups according to the 28-day survival endpoint. Those with all-cause mortality within the follow-up period were considered non-survivors. If data were lost, the corresponding candidate was excluded.

Patient and public involvement

This study was approved by the Medical Ethics Committee (2013-KE-1). Patient consent to participate was obtained prior to enrolment in this study. Results **Patient characteristics** In total, 40 healthy individuals and 85 patients who experienced CA were analyzed. The demographics and clinical characteristics of both groups are shown in Table 1. In this study, acute cardiac and brain events were the main causes of CA. Other causes of CA included poisoning (including carbon monoxide poisoning) and hypokalemia. Sex and age were not significantly different between the CA and healthy control groups. The comparisons of clinical characteristics of the survivor and non-survivor groups based on 28-day survival are shown in Supplemental Table 2. The APACHE II and SOFA scores were significantly different between the CA and healthy control groups (P<0.001 for all) and survivor and non-survivor groups (P<0.001 and P=0.011, respectively).

Table 1. Patient Characteristics at Admission

	Healthy Control	Successful	
Characteristics	Group (n=40)	Resuscitation Group	<i>P</i> -value
		(n=85)	
Age (years), median [IQR]	64.0 (54.3, 69.8)	65.0 (55.0, 74.0)	0.209
Male/Female (n)	23/17	58/27	0.241
Previous medical history, n (%)			

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	Hypertension	5 (12.5%)	38 (44.7%)	< 0.001
	Diabetes	3 (7.5%)	27 (31.8%)	0.003
	Coronary heart disease	2 (5.0%)	29 (34.1%)	< 0.001
	Chronic lung disease	1 (2.5%)	9 (10.6%)	0.230
	Chronic kidney disease	0	9 (10.6%)	0.077
	Cardiac arrest cause (n, %)			
	Cardiac		34 (40.0%)	
	Respiratory		20 (23.5%)	
	Cerebral		23 (27.1%)	
	Others		7 (8.2%)	
	Unknow		1 (1.2%)	
	Initial resuscitation			
	Time to ROSC (min), median		20.0 (10.0, 30.0)	
	[IQR]			
	Adrenaline (mg), median [IQR]		2.0 (0.0, 5.0)	
	Initial rhythm VF/VT, n (%)		30 (35.3%)	
	MAP (mmHg), median [IQR]	95.7 (86.0,	74.3 (56.2, 97.2)	< 0.001
		103.2)		
	White cell count ($\times 10^{9}/L$), median	5.81 (4.85, 6.53)	13.56 (10.84, 18.29)	< 0.001
	[IQR]			
	APACHE II score, mean±SD	0	32.9±6.5	< 0.001
	SOFA score, median [IQR]	0	11.5 (8.5, 14.0)	< 0.001
	28-day mortality, n (%)		65 (76.5%)	
	28-day CPC 1–2, n (%)		14 (16.5%)	
4	Abbreviations: IQR: interquartile	e range; ROSC: re	eturn of spontaneous c	irculatior
	VF: ventricular fibrillation; VT:	ventricular tachyo	cardia; MAP: mean art	erial pres
	APACHE II: acute physiology a	nd chronic health	evaluation; SOFA: se	quential
7	organ failure assessment; SD: sta	undard deviation;	CPC: cerebral perform	nance
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198 category.

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200	Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts after
201	ROSC
202	The T and B lymphocyte, NK cell, and Treg cell counts were significantly lower after
03	ROSC in patients who experienced CA than in healthy controls (P<0.001 for all).
04	Additionally, the CD3+CD4+/T lymphocyte, CD3+CD8+/T lymphocyte, and Treg
)5	cell/CD4 ⁺ T lymphocyte ratios were significantly lower after ROSC in patients who
)6	experienced CA than in healthy controls (P<0.001 for all) (Fig. 1; Supplemental Table
07	3). However, there were no significant differences in these cell counts and ratios
08	between survivors (n=20) and non-survivors (n=65) (P>0.05 for all) (Supplemental
)9	Table 4).
10	
1	GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells after
2	ROSC
13	The MFI and percentages of GR expression in B and T lymphocytes, NK cells, and
4	CD3 ⁺ CD8 ⁺ T lymphocytes were significantly lower after ROSC in patients who
15	experienced CA than in healthy individuals (P<0.01 for all) (Fig. 2A–D, G, H, K, L).
16	There were also significant reductions in the MFI in Treg cells and $\text{CD3}^+\text{CD4}^+$ T
17	lymphocytes (P<0.05 for all) (Figs. 2E, I) but not in the percentages of GR expression
18	(P>0.05 for all) (Figs. 2F, J; Supplemental Table 5). However, there were no significant
19	differences in the MFI and percentages of GR expression in these cells between
20	survivors and non-survivors (P>0.05 for all) (Supplemental Table 6).

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222	Changes in plasma total cortisol and ACTH levels after ROSC
223	We measured the plasma total cortisol and ACTH levels of the 40 healthy individuals
224	and 85 patients who experienced CA (two samples were excluded because their total
225	cortisol levels were not measured). Plasma total cortisol levels were significantly higher
226	in patients who experienced CA than in healthy controls (P<0.001) but ACTH levels
227	were not (Figs. 3A, C). No significant differences in ln (total cortisol+1) and ln
228	(ACTH+1) values were observed between survivors and non-survivors (P>0.05 for all)
229	(Fig. 3B, D).
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230 231	Discussion
	Discussion In this study, the relationship between GR expression and immune alteration in the early
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231 232	In this study, the relationship between GR expression and immune alteration in the early
231 232 233	In this study, the relationship between GR expression and immune alteration in the early period after ROSC in patients who experienced CA was explored by observing GR
231232233234	In this study, the relationship between GR expression and immune alteration in the early period after ROSC in patients who experienced CA was explored by observing GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells and changes
 231 232 233 234 235 	In this study, the relationship between GR expression and immune alteration in the early period after ROSC in patients who experienced CA was explored by observing GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells and changes in cell counts and plasma total cortisol and ACTH levels. We found that GR expression,
 231 232 233 234 235 236 	In this study, the relationship between GR expression and immune alteration in the early period after ROSC in patients who experienced CA was explored by observing GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells and changes in cell counts and plasma total cortisol and ACTH levels. We found that GR expression, cell counts, and ratios rapidly decreased, and plasma total cortisol levels increased, in

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lymphocyte, NK cell, and Treg cell counts as well as CD3⁺CD4⁺/T, CD3⁺CD8⁺/T, and

Treg cell/CD4⁺ T lymphocyte ratios were significantly reduced after ROSC. NK cells,

which are special innate immune cells that have cytotoxic functions similar to

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243	CD3 ⁺ CD8 ⁺ T lymphocytes, mainly distinguish infected and stressed cells from healthy
244	cells and eliminate intracellular infection as well as dysfunctional cells. [27, 28] T
245	lymphocytes are also important because of their function as adaptive immune cells for
246	the control and elimination of infection. [27] Moreover, B and T lymphocytes mediate
247	humoral and cellular immunity, respectively. This study was performed at an earlier
248	period and involved a more comprehensive assessment of the immune system of
249	patients who experienced CA, and our findings more substantially supported the rapid
250	emergence of immune dysfunction in these patients after ROSC than previous reports.
251	The effectiveness of GC use in these patients during and after resuscitation has been
252	controversial due to insufficient evidence. However, the use of GCs during resuscitation
253	improves the survival rate of patients who experience CA due to its direct anti-
254	inflammatory, immunosuppressive effects, hemodynamics, and positive inotropic
255	effects. All of this ultimately leads to an increased stress capacity of the body. [18, 19]
256	GCs can activate GRs in cells when the body is under stress, thereby increasing both
257	the effectiveness of resuscitation and discharge survival rate. This study is the first to
258	explore GR expression in circulating immune cells in patients who experienced CA
259	after ROSC. We observed that GR expression in B and T lymphocytes, NK cells, and
260	CD3 ⁺ CD8 ⁺ T lymphocytes decreased significantly in patients who experienced CA,
261	whereas the percentage of GR ⁺ Treg cells and CD3 ⁺ CD4 ⁺ T lymphocytes showed a
262	slight decrease. Moreover, we observed a more significant decrease in the MFI of GR

expression in Treg cells and CD3⁺CD4⁺ T lymphocytes but not in the percentage of GR

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expression. Previous studies have found decreased expression of GRs in peripheral
polymorphonuclear cells in critically ill patients, [21] and antagonism to GRs
aggravates viral and bacterial infections. [29] The results of this study suggest that the
decrease in intracellular GR expression in patients who experienced CA is one of the
causes of GC resistance, due to insufficient binding of GRs and GCs, GC insensitivity
and the inability of GCs to effectively exert anti-inflammatory and immunosuppressive
effects. These findings may also explain why different results regarding the clinical
application of GCs have been reported previously and support the possibility of using
GCs in the clinical treatment of patients who experienced CA.
We also found that the total plasma cortisol levels were significantly higher ir
patients who experienced CA, but ACTH levels were not. High levels of inflammatory
cytokines inhibit ACTH release. [18] During critical illness, the body does not
sufficiently metabolize cortisol. [30] In addition, the continuous increase in plasma
cortisol levels may trigger the negative feedback pathway of the hypothalamic-
pituitary-adrenal axis, inhibiting the release of ACTH and cortisol and eventually
leading to adrenal insufficiency. These factors may explain the opposite trends of
plasma ACTH and cortisol levels in the patients who were included in this study and
experienced CA. Notably, this result suggests that low GR expression levels are not
matched with high plasma total cortisol levels. Previous studies have found that GC use
during resuscitation may benefit patients who experience CA. [13-16] The benefits
such as direct anti-inflammatory and anti-shock effects, improvement of vascular
endothelial permeability, and other mechanisms may be related to the effects of using
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a high dose of GCs, or GCs may work through other non-GR pathways. It is also
possible that the immune function of patients who experience CA is suppressed due to
ischemia-reperfusion injury, which requires a large dose of GCs to stimulate GRs to
function. This study did not provide data on plasma GC levels and GR expression in a
group of patients who were administered GCs and successfully resuscitated; therefore,
further studies are required to explore the exact mechanisms of GCs.

292 Limitations

Our study has several limitations. First, to assess changes, we only enrolled patients who experienced CA and had signs of systemic ischemic hypoxia, such as GCs <8 after ROSC. The patients were not stratified by age, sex, and occurrence of comorbidities or mild systemic ischemic hypoxia. Second, since this was a preliminary observational study, we observed only early changes. A dynamic observation for a longer duration would be helpful to understand the significance of GR expression in evolving immunity during the clinical course of CA after ROSC. Third, the samples used in this study were from the clinical laboratory; thus, plasma total cortisol and ACTH in the samples were at a risk of degradation before we collected the samples. Finally, we did not discuss the changes in and roles of GR isoforms, free cortisol, and corticosteroid-binding globulin. Therefore, future studies on these aspects are warranted to better understand the immunosuppressive effects of ROSC among patients who experienced CA. In conclusion, this study revealed that GR expression, cell counts, and ratios rapidly decreased, whereas plasma total cortisol levels increased, in the early period

307 after ROSC among CA patients. These findings may provide important information

- 3 4	308	about GC sensitivity and immunosuppressive status in these patients. In addition, this
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6 7 8	309	study provides a new perspective for clinical targeted treatment using GCs and high-
9 10	310	quality prognosis in CA patients.
11 12 13	311	
14 15	312	Acknowledgements: We thank all researchers who participated in this study and all
16 17 18	313	colleagues in the emergency department who provided support.
19 20 21	314	Ethics Approval: This study was approved by the Medical Ethics Committee of
22 23	315	Beijing Chaoyang Hospital (2013-KE-1). Patient consent to participate was obtained
24 25 26	316	prior to enrolment in this study.
27 28	317	Contributorship statement: CL designed the study and reviewed the manuscript.
29 30 31	318	YNY searched the literature and contributed to the experimental studies, data
32 33 34	319	analysis, and writing of the manuscript. ZRT, CCH, and LA collected and analyzed
35 36	320	data. JBL and MRX helped with the statistical analyses. All authors have read and
37 38 39	321	approved the final manuscript.
40 41	322	Competing interests: All authors declare no competing interest associated with this
42 43 44	323	project.
45 46	324	Funding: This research received no specific grant from any funding agency in the
47 48 49	325	public, commercial or not-for-profit sectors.
50 51	326	Data sharing statement: All data relevant to the study are included in the article or
52 53 54	327	uploaded as supplementary information. Due to privacy and ethical concerns, data can
55 56 57	328	not be shared.
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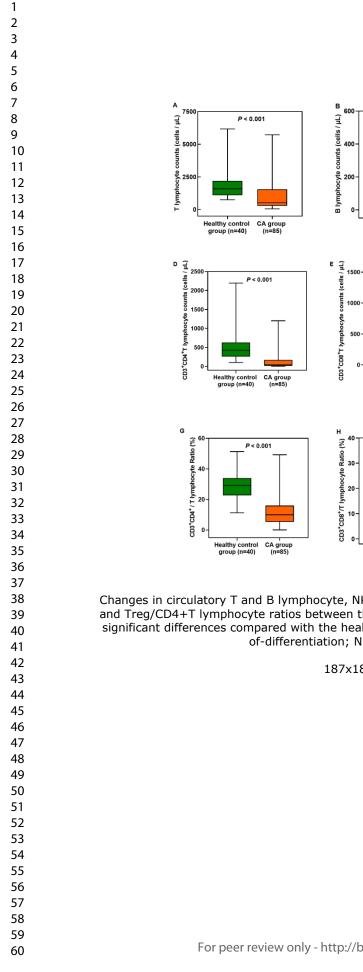
414 **Figure legends**

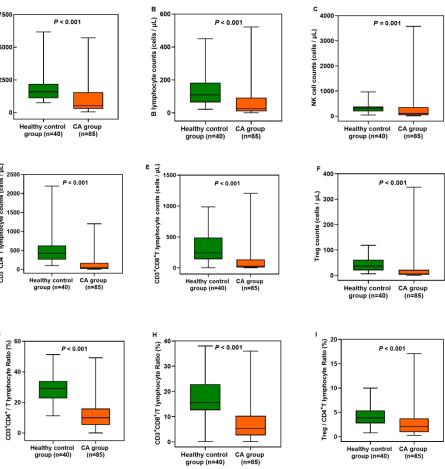
Fig. 1. Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts, and
CD3⁺CD4⁺/T, CD3⁺CD8⁺/T, and Treg/CD4⁺T lymphocyte ratios between the healthy
control group and CA group. The CA group showed significant differences compared
with the healthy control group (P<0.001). CA, cardiac arrest; CD, cluster-of-
differentiation; NK, natural killer; Treg, regulatory T.

Fig. 2. Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells
in the healthy control group and CA group. The CA group showed significant
differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD,
cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC,
return of spontaneous circulation; Treg, regulatory T.

Fig. 3. (A, B) Plasma total cortisol and ACTH levels (the natural logarithmic
conversion values after adding 1) after ROSC in the healthy control group and CA
group. (C, D) Plasma total cortisol and ACTH levels in survivors and non-survivors
after ROSC. The CA group showed significant differences compared with the healthy
control group (P<0.05). ACTH, adrenocorticotrophic hormone; CA, cardiac arrest;
ROSC, return of spontaneous circulation.

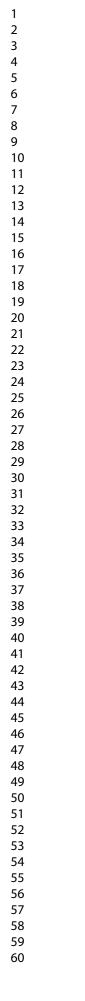
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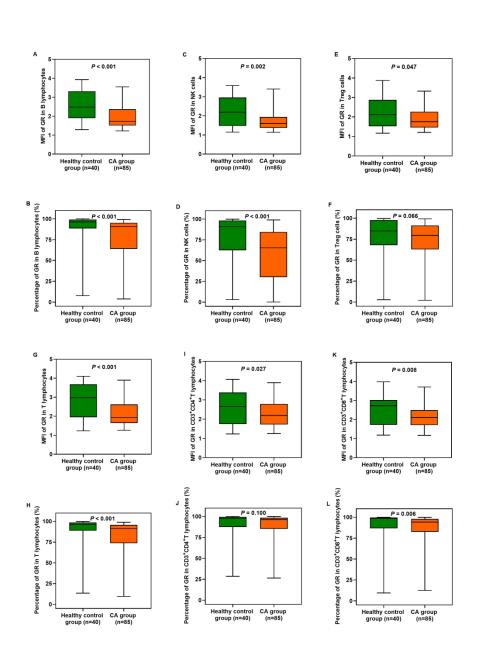




Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts, and CD3+CD4+/T, CD3+CD8+/T, and Treg/CD4+T lymphocyte ratios between the healthy control group and CA group. The CA group showed significant differences compared with the healthy control group (P<0.001). CA, cardiac arrest; CD, cluster-of-differentiation; NK, natural killer; Treg, regulatory T.

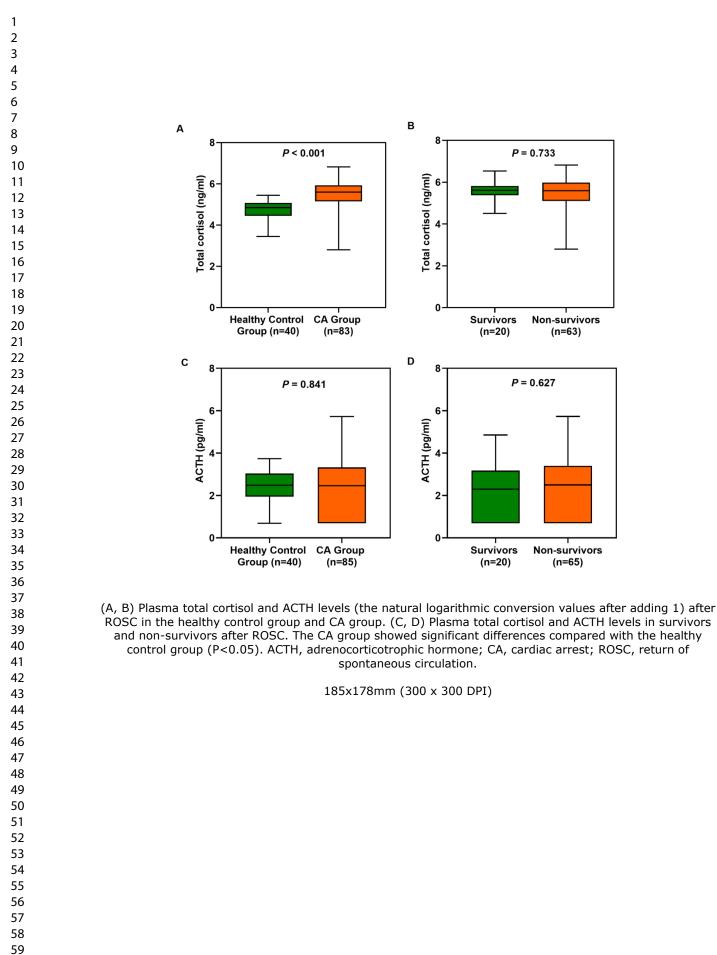
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Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells in the healthy control group and CA group. The CA group showed significant differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD, cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC, return of spontaneous circulation; Treg, regulatory T.

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Electronic supplemental material

Expression of glucocorticoid receptors early after the return of spontaneous circulation in patients who

experienced cardiac arrest: A prospective observational study

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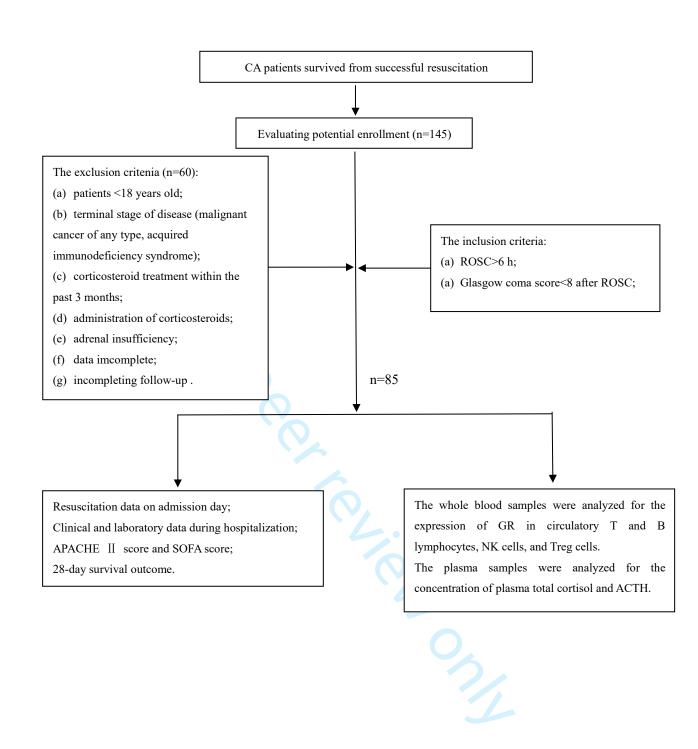
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Supplemental Figure 1. The flow chart of the study.

Abbreviations: CA, cardiac arrest; ROSC, return of spontaneous circulation; APACHE II, acute physiology and chronic health evaluation II; SOFA, sequential organ failure assessment; GR, glucocorticoid receptor; Treg, regulatory T; ACTH, adrenocorticotrophic hormone.

Antigen	Catalog Number	Fluorescein Conjugate	Source
CD3	558117	Pacific Blue	BD Pharmingen ^a
CD4	555347	PE	BD Pharmingen
CD4	560345	Horizon V450	BD Pharmingen
CD8	557746	PE-Cy7	BD Pharmingen
CD19	557835	PE-Cy7	BD Pharmingen
CD16	558122	Pacific Blue	BD Pharmingen
CD56	557747	PE-Cy7	BD Pharmingen
CD25	557741	PE-Cy7	BD Pharmingen
CD127	557938	PE	BD Pharmingen
GR	MCA2469F	FITC	Bio-Rad ^b
Mouse IgG1 Isotype	MCA928F	FITC	Bio-Rad
Mouse IgG1,к Isotype	557872	PE-Cy7	BD Pharmingen
Mouse IgG1,к Isotype	554680	РЕ	BD Pharmingen
Mouse IgG1,к Isotype	558120	Pacific Blue	BD Pharmingen
^a BD Pharmingen, San Di	ego, USA; ^b Bio-Rad Abl	D Serotec, Oxford, UK.	
Abbreviations: CD, clust	er-of-differentiation; PE,	phycoerythrin; FITC, fluores	cein isothiocyanate; GR, g

	Survivors (n=20)	Non-survivors (n=65)	<i>P</i> -value
Age (years), median [IQR]	59.0 (53.3, 72.8)	66.0 (59.0, 75.5)	0.070
Male/Female (n)	12/8	46/19	0.366
Cardiac arrest cause (n, %)			
Cardiac	10 (50.0%)	24 (36.9%)	0.297
Non-Cardiac	10 (50.0%)	41 (63.1%)	0.297
Initial resuscitation			
Time to ROSC (min), median [IQR]	15.0 (7.3, 26.0)	20.0 (15.0, 30.0)	0.032
Adrenaline (mg), median [IQR]	1.0 (0.0, 3.0)	2.0 (0.0, 5.0)	0.091
Initial rhythm VF/VT, n (%)	11 (55.0%)	19 (29.2%)	0.035
MAP (mmHg), median [IQR]	89.9 (70.5, 104.9)	70.7 (50.0, 93.5)	0.033
White cell count (×10 ⁹ /L), median [IQR]	12.40 (6.98, 18.76)	13.80 (11.67, 18.20)	0.286
Lactate (mmol/L), median [IQR]	3.50 (1.33, 7.05)	7.50 (3.80, 11.20)	0.008
APACHE II score, mean±SD	27.8±6.6	34.4±5.6	< 0.001
SOFA score, median [IQR]	9.0 (7.3, 11.8)	12.0 (9.0, 15.0)	0.011

Supplemental Table 2. Characteristics of CA survivors and non-survivors on admission.

Data are presented as mean±SD or interquartile range (IQR) as appropriate. The *P*-value represents comparison between groups. Abbreviations: ROSC: return of spontaneous circulation; VF: ventricular fibrillation; VT: ventricular tachycardia; MAP: mean arterial pressure; APACHE II: acute physiology and chronic health evaluation; SOFA: sequential organ failure assessment.

Supplemental Table 3. The flow cytometry results of cell counts and ratios of healthy control group and successful resuscitation group

	Healthy Control	Successful	Z-value	P-value
	Group (n=40)	Resuscitation Group		
		(n=85)		
T lymphocyte count (cells /µL)	1586.0 (1101.5, 2192.5)	514.0 (287.5, 1555.0)	-4.515	< 0.001
NK cell count (/µL)	311.5 (191.0, 378.8)	101.0 (36.0, 351.5)	-3.332	0.001
B lymphocyte count (/μL)	109.3 (63.7, 183.3)	25.7 (9.4, 92.3)	-5.076	<0.001
Treg count (/µL)	0.259 (0.095, 0.516)	0.233 (0.135, 0.488)	-5.518	<0.001
Treg / CD4 ⁺ T lymphocyte Ratio	0.039 (0.028, 0.054)	0.021 (0.010, 0.038)	-4.418	<0.001
CD3 ⁺ CD4 ⁺ T lymphocyte count (/ μ L)	421.7 (258.6, 627.4)	38.9 (17.6, 168.3)	-6.256	<0.001
CD3 ⁺ CD4 ⁺ / T lymphocyte Ratio	0.292 (0.227, 0.340)	0.100 (0.054, 0.160)	-7.066	<0.001
CD3 ⁺ CD8 ⁺ T lymphocyte count (/ μ L)	241.1 (139.5, 488.6)	26.3 (7.2, 135.9)	-5.287	<0.001
CD3 ⁺ CD8 ⁺ / T lymphocyte Ratio	0.157 (0.126, 0.229)	0.053 (0.026, 0.104)	-5.719	< 0.001

All the data in Supplemental table 3 are represented as the median [IQR]; IQR: Interquartile Range; CD: cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

Supplemental Table 4. The flow cytometry results of cell counts and ratios of the CA patients on admission based on 28-day survival

	Survivors (n=20)	Non-survivors	Z-value	<i>P</i> -value
		(n=65)		
T lymphocyte count (/µL)	502.0 (353.8, 1199.8)	514.0 (282.5, 1891.0)	-0.186	0.852
NK cell count (/µL)	167.0 (29.8, 309.3)	100.0 (36.0, 404.0)	-0.218	0.828
B lymphocyte count (/µL)	38.6 (15.7, 103.5)	19.2 (7.1, 65.7)	-0.632	0.527
Tregs count (/µL)	0.318 (0.145, 0.552)	0.212 (0.128, 0.479)	-0.611	0.396
Treg / CD4 ⁺ T lymphocyte Ratio	0.025 (0.009, 0.043)	0.021 (0.010, 0.034)	-0.498	0.619
CD3 ⁺ CD4 ⁺ T lymphocyte count (/µL)	55.1 (32.4, 228.0)	38.0 (16.0, 168.1)	-0.850	0.396
CD3 ⁺ CD4 ⁺ / T lymphocyte Ratio	0.118 (0.070, 0.236)	0.097 (0.049, 0.142)	-1.565	0.118
CD3 ⁺ CD8 ⁺ T lymphocyte count (/µL)	25.4 (12.5, 96.2)	26.3 (6.3, 138.8)	-0.021	0.983
CD3 ⁺ CD8 ⁺ / T lymphocyte Ratio	0.054 (0.033, 0.104)	0.053 (0.025, 0.104)	-0.187	0.852

All the data in Supplemental table 4 are represented as the median [IQR]; IQR: Interquartile Range; CD:

cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

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59 60 Supplemental Table 5. The flow cytometry results of GR expression in the CA group and successful resuscitation group.

	Healthy Control	Successful	Z-value	<i>P</i> -value
	Group (n=40)	Resuscitation Group		
		(n=85)		
Percentage of GR on B lymphocytes	0.963 (0.885, 0.992)	0.896 (0.605, 0.949)	-3.742	<0.001
MFI of GR on B lymphocytes	2.48 (1.91, 3.31)	1.73 (1.50, 2.37)	-3.980	< 0.001
Percentage of GR on T lymphocytes	0.964 (0.889, 0.986)	0.900 (0.703, 0.955)	-3.755	< 0.001
MFI of GR on T lymphocytes	2.98(1.95, 3.68)	1.92 (1.36, 1.99)	-3.853	<0.001
Percentage of GR on NK cells	0.907 (0.624, 0.983)	0.611 (0.306, 0.840)	-3.792	< 0.001
MFI of GR on NK cells	2.19 (1.48, 2.96)	1.60 (1.36, 1.99)	-3.171	0.002
Percentage of GR on Treg cells	0.848 (0.680, 0.978)	0.784 (0.589, 0.911)	-1.837	0.066
MFI of GR on Treg cells	2.12 (1.53, 2.88)	1.76 (1.44, 2.30)	-1.990	0.047
Percentage of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	0.980 (0.874, 0.996)	0.957 (0.824, 0.985)	-2.204	0.100
MFI of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	2.65 (1.75, 3.38)	2.17 (1.70, 2.92)	-1.646	0.027
Percentage of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	0.986 (0.868, 0.996)	0.938 (0.823, 0.979)	-2.758	0.006
MFI of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	2.73 (1.73, 3.02)	2.10 (1.68, 2.54)	-2.668	0.008

All the data in Supplemental table 5 are represented as the median [IQR]. Abbreviations: IQR, interquartile Range; CD, cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, Glucocorticoid receptor; MFI, mean fluorescence intensity.

	Survivors (n=20)	Non-survivors (n=65)	Z-value	<i>P</i> -value
Percentage of GR on B lymphocytes	0.904 (0.595, 0.976)	0.906 (0.657, 0.946)	-0.787	0.431
MFI of GR on B lymphocytes	1.92 (1.52, 2.54)	1.72 (1.51, 2.31)	-0.881	0.378
Percentage of GR on T lymphocytes	0.899 (0.778, 0.969)	0.913 (0.692, 0.951)	-1.057	0.291
MFI of GR on T lymphocytes	2.05 (1.67, 2.83)	1.91 (1.64, 2.46)	-1.031	0.303
Percentage of GR on NK cells	0.717 (0.292, 0.886)	0.556 (0.302, 0.823)	-0.756	0.449
MFI of GR on NK cells	1.54 (1.37, 2.09)	1.61 (1.34, 1.87)	-0.565	0.572
Percentage of GR on Tregs	0.780 (0.667, 0.849)	0.799 (0.576, 0.923)	-0.440	0.660
MFI of GR on Tregs	1.61 (1.48, 2.30)	1.77 (1.45, 2.27)	-0.005	0.996
Percentage of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	0.975 (0.876, 0.985)	0.957 (0.845, 0.987)	-0.617	0.538
MFI of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	2.08 (1.72, 3.35)	2.22 (1.71, 2.69)	-0.865	0.387
Percentage of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	0.963 (0.816, 0.977)	0.938 (0.834, 0.980)	-0.254	0.800
MFI of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	2.08 (1.68, 3.10)	2.11(1.71, 2.46)	-0.653	0.514

All the data in Supplemental table 6 are represented as the median [IQR]. Abbreviations: IQR, Interquartile Range; CD, Cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, glucocorticoid receptor; MFI, mean fluorescence intensity.

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STROBE Statement-checklist of items that should be included in reports of observational studies

	Item No	Recommendation	Page No
Title and abstract	1	(<i>a</i>) Indicate the study's design with a commonly used term in the title or the abstract	2
		(<i>b</i>) Provide in the abstract an informative and balanced summary of what was done and what was found	3
Introduction			1
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	3-5
Objectives	3	State specific objectives, including any prespecified hypotheses	5
Methods			1
Study design	4	Present key elements of study design early in the paper	Supplemental Figure 1
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection	5-9, Supplemental Figure 1
Participants	6	 (a) Cohort study—Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up Case-control study—Give the eligibility criteria, and the sources and methods of case ascertainment and control selection. Give the rationale for the choice of cases and controls Cross-sectional study—Give the eligibility criteria, and the sources and methods of selection of participants 	5,6,8,9
		(b) Cohort study—For matched studies, give matching criteria and number of exposed and unexposed Case-control study—For matched studies, give matching criteria and the number of controls per case	5
Variables	7	Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable	5, 6, 8, Supplemental Figure 1
Data sources/ measurement	8*	For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group	6-8
Bias	9	Describe any efforts to address potential sources of bias	6-8
Study size	10	Explain how the study size was arrived at	Supplemental Figure 1
Quantitative variables	11	Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why	6-8
Statistical methods	12	(<i>a</i>) Describe all statistical methods, including those used to control for confounding	8, 11

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(b) Describe any methods used to examine subgroups and interactions	N/A
(c) Explain how missing data were addressed	8, 11
(<i>d</i>) <i>Cohort study</i> —If applicable, explain how loss to follow-up was addressed	11
<i>Case-control study</i> —If applicable, explain how matching of cases and controls was addressed	
Cross-sectional study—If applicable, describe analytical	
methods taking account of sampling strategy	
(<u>e</u>) Describe any sensitivity analyses	

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Participants	13	(a) Report numbers of individuals at each stage of study—eg numbers	9,
1	*	potentially eligible, examined for eligibility, confirmed eligible, included in	Supplei
		the study, completing follow-up, and analysed	l Figure
		(b) Give reasons for non-participation at each stage	11,
			Suppler
			1 Figure
		(c) Consider use of a flow diagram	Suppler
			l Figure
Descriptive	14	(a) Give characteristics of study participants (eg demographic, clinical,	9
data	*	social) and information on exposures and potential confounders	
		(b) Indicate number of participants with missing data for each variable of	9-11
		interest	
		(c) Cohort study—Summarise follow-up time (eg, average and total amount)	8
Outcome data	15	<i>Cohort study</i> —Report numbers of outcome events or summary measures	9-11
	*	over time	
		Case-control study-Report numbers in each exposure category, or summary	
		measures of exposure	
		Cross-sectional study—Report numbers of outcome events or summary	
		measures	
Main results	16	(a) Give unadjusted estimates and, if applicable, confounder-adjusted	9-11,
		estimates and their precision (eg, 95% confidence interval). Make clear	Electro
		which confounders were adjusted for and why they were included	supplen
			materia
		(b) Report category boundaries when continuous variables were categorized	
		(c) If relevant, consider translating estimates of relative risk into absolute	
		risk for a meaningful time period	
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and	N/A
		sensitivity analyses	
Discussion			1
Key results	18	Summarise key results with reference to study objectives	12
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias	15
		or imprecision. Discuss both direction and magnitude of any potential bias	
Interpretation	20	Give a cautious overall interpretation of results considering objectives,	12-14
		limitations, multiplicity of analyses, results from similar studies, and other	
		relevant evidence	
Generalisabilit	21	Discuss the generalisability (external validity) of the study results	15
y Other informat	ion		I
Funding	10n 22	Give the source of funding and the role of the funders for the present study	16
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*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies.

Note: An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely

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available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.

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Glucocorticoid receptor expression in patients with cardiac arrest in the early period after the return of spontaneous circulation: A prospective observational study

Journal:	BMJ Open
Manuscript ID	bmjopen-2021-060246.R1
Article Type:	Original research
Date Submitted by the Author:	24-May-2022
Complete List of Authors:	Yu, Yanan; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Tang, Ziren; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Xie, Miaorong; Capital Medical University Affiliated Beijing Friendship Hospital, Department of Emergency Medicine Li, Jiabao; Capital Medical University Affiliated Beijing Friendship Hospital, Department of Critical Care Hang, Chen-Chen; Beijing Chao-Yang Hospital, Emergency Medicine An, Le; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Li, Chunsheng; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine
Primary Subject Heading :	Emergency medicine
Secondary Subject Heading:	Intensive care
Keywords:	ACCIDENT & EMERGENCY MEDICINE, INTENSIVE & CRITICAL CARE, Adult intensive & critical care < INTENSIVE & CRITICAL CARE

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1	Glucocorticoid receptor expression in patients with cardiac arrest in the early
2	period after the return of spontaneous circulation: A prospective observational
3	study
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16	
17	Keywords: Cardiac arrest, glucocorticoid receptor, immunosuppression, cortisol
18	Word count of the main text: 3,335 words
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23 Abstract

Objectives: Rapid changes in glucocorticoid (GC) levels and adrenal insufficiency are related to the development of post-cardiac arrest (CA) syndrome. However, GC receptor (GR) expression changes have not been studied. Hence, this study aimed to investigate the association of early changes in GR expression and prognosis and immune response in patients who experienced CA.

- 29 **Design:** Prospective observational study.
- 30 Setting: Emergency department.

Participants: Patients (85) in the early period of return of spontaneous circulation
(ROSC) after CA were admitted between October 2018 and October 2019. After a
physical examination, age- and sex-matched healthy individuals (40) were recruited for
the control group.

Primary and secondary outcome measures: GR expression and cell counts of
circulatory T and B lymphocytes, natural killer cells, and regulatory T (Treg) cells were
assessed. Plasma total cortisol and adrenocorticotrophic hormone (ACTH) levels were
also tested.

Results: All cell counts were lower, and plasma total cortisol levels were higher (P<0.001) in patients who experienced CA than in the healthy control group. GR expression in Treg cells and CD3⁺CD4⁺ T lymphocytes were not significantly different, but the mean fluorescence intensity and GR expression in other cells were lower in patients who experienced CA (P<0.05) than in the healthy control group. ACTH levels were not different. There were no significant differences between survivors and non-

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45	survivors.
46	Conclusions: This study revealed that GR expression and cell counts rapidly decreased,
47	whereas plasma total cortisol levels increased in the early period after ROSC among
48	patients who experienced CA. Our findings provide important information about GR
49	level and function, and immunosuppressive status in these patients. Assessing GR
50	expression in CA patients may help screening for those who are more sensitive to
51	glucocorticoid therapy.
52	
53	Strengths and limitations of this study
54	1. The study design will be single-center, prospective.
55	2. This is the first study to evaluate the GR expression in the early period following
56	ROSC among CA patients.
57	3. Only CA patients in the early period following ROSC will be included, limiting the
58	generalisability of the results.
59	4. Decreased GR expression may affect the sensitivity of CA patients to GCs.
60	5. Decreased GR expression may affect potential immune consequences of CA
61	patients.
62	
63	Introduction
64	Cardiac arrest (CA) is a significant health problem globally; about 356,500 people

experience medical emergencies due to CA in the United States, and over 544,000

66 people die from sudden CA in China annually. [1, 2] The systemic ischemia-reperfusion

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67	response in patients who have experienced CA can present as post-cardiac arrest
68	syndrome (PCAS) or systematic inflammatory response syndrome (SIRS), which
69	increases the risk of multiple organ failure and infection and affects the inflammatory
70	response and prognosis of patients after the return of spontaneous circulation (ROSC).
71	[3-6]
72	CA is the most intense among acute stress events, which seriously affect the pituitary
73	and adrenal axis function. [7] Studies have shown that abnormal cortisol levels and
74	relative adrenocortical insufficiency after ROSC in patients who experienced CA are
75	related to their prognosis. [8-11] However, the clinical application of glucocorticoids
76	(GCs) is controversial. In the 2015 International Cardiopulmonary Resuscitation
77	Guidelines, the routine use of GCs is not recommended for the resuscitation of patients
78	with in-hospital or out-of-hospital CA. [12] Recent clinical studies have shown that
79	early administration of corticosteroids after CA can improve the success rate of ROSC,
80	nervous system functional outcome, and prognosis, which is speculated to be related to
81	its influence on hemodynamics, and SIRS response, and other mechanisms. [12-17]
82	Therefore, the role of GCs in the occurrence and development of PCAS needs to be
83	studied further.
81	GCs combine with intracellular GC recentors (GRs) to evert anti-inflammatory and

GCs combine with intracellular GC receptors (GRs) to exert anti-inflammatory and immunosuppressive effects and reduce the production and the release of inflammatory cytokines. [18, 19] The affinity of GRs to GCs in circulating monocytes is decreased in patients with acquired immunodeficiency syndrome. [20] The expression of GR alpha and beta in peripheral polymorphonuclear cells is decreased in patients with critical

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3 4	89	illness, [21] pediatric septic shock, and high serum cortisol levels. [22] However, no
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6 7	90	study has reported the GR expression after ROSC in patients who experienced CA.
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9 10	91	Previous studies have found that the counts of circulating B and T lymphocytes,
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12	92	regulatory T (Treg) cells, and monocytes and expression of human leukocyte antigen
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15	93	DR (HLA-DR) on circulatory monocytes and B and T lymphocytes are reduced. [23,
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17 18	94	24] Hence, this study aimed to investigate the relationship between GR expression and
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20	95	immune alteration in the early period after ROSC in patients who experienced CA by
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22	96	observing GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells,
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25 26	97	their cell counts, and total plasma cortisol and adrenocorticotrophic hormone (ACTH)
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28	98	levels.
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32	100	Materials and methods
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2015 International Cardiopulmonary Resuscitation Consensus. [13] After a physical
examination, age- and sex-matched healthy individuals were recruited for the control
group.

114 Data collection

Data collection was performed according to the 2004 guidelines of the Utstein Style 115 template. [26] We collected data on demographics, resuscitation (initial heart rhythm, 116 ROSC time, and cumulative adrenaline [epinephrine] dose, and laboratory findings 117 routine blood cell counts, blood gas analysis, and blood biochemical tests performed > 118 119 6 h and < 24 h after ROSC). Acute Physiology and Chronic Health Evaluation (APACHE) II and the Sequential Organ Failure Assessment (SOFA) were used to 120 determine disease severity. Residual blood samples from routine clinical tests or 121 122 physical health examinations in the morning were collected, maintained at 4 °C during transport and storage, and used to determine GR expression in circulatory T and B 123 lymphocytes, NK cells, and Treg cells and their cell counts. The plasma was maintained 124 125 at -80 °C during storage and used to determine total cortisol and ACTH levels. During follow-up, 28-day survival data were also collected. Supplemental Figure 1 shows the 126 workflow of this study. 127

128 Outcome measures

The primary outcomes of this study were GR expression and cell counts of T and B
cells, NK cells, and Treg cells, measured by flow cytometry. Venous blood samples
collected in ethylenediaminetetraacetic acid tubes, then used to measure GR expression
in T and B lymphocytes, NK cells, and Treg cells. Briefly, a 100-μL peripheral blood

133	sample was stained for 20 min with surface antibodies (CD3, CD4, CD8, CD19, CD16,
134	CD56, CD25, and CD127) in a dark place. Erythrocytes were lysed for 15 min, and the
135	debris was washed away. Before intracellular GR staining, surface-stained cells were
136	fixed and permeabilized using the BD Transcription Factor Buffer Set (BD
137	Pharmingen, San Diego, USA, Catalogue No. 562574). Monoclonal antibodies and
138	their isotype controls were all purchased from BD Biosciences (San Jose, CA, USA).
139	Details of all antibodies are shown in Supplemental Table 1. According to the
140	manufacturer's recommendations, all antibodies and their isotype controls were used at
141	a concentration of 1 μ L per 100 μ L of whole blood. Samples were measured using the
142	Gallios flow cytometer (Beckman Coulter, Brea, CA, USA) and analyzed using Gallios
143	Software version 1.0 (Beckman Coulter). The flow cytometer was periodically
144	calibrated by an engineer. Cells were stained for 20 min; thresholds were defined using
145	the manufacturer's recommended isotype controls. Representative plots and gating
146	strategy from a single sample are shown in Supplemental Figure 2. T cells were gated
147	by CD3 ⁺ CD4 ⁺ or CD3 ⁺ CD8 ⁺ , B cells were gated by CD3 ⁻ CD19 ⁺ , NK cells were gated
148	by CD16 ⁺ CD56 ⁺ , and Tregs were gated by CD4 ⁺ CD25 ^{high} CD127 ^{low} . At least 10,000
149	events were collected in the lymphocyte cell gate for each sample. Results are expressed
150	as percentages and mean fluorescence intensity (MFI) values.
151	Absolute CD3 ⁺ and CD4 ⁺ lymphocyte, NK cell, and Treg cell counts were obtained
152	using Flow-Count fluorospheres (Beckman Coulter, Catalogue No. 7547053),
153	according to the manufacturer's instructions. B, CD3+CD4+T, CD3+CD8+T, and Treg

154 cell counts were calculated by their percentages in CD3⁺ or CD4⁺ lymphocytes

155	multiplied by CD3+	or CD4 ⁺ lymphocyte counts.
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The secondary outcomes of this study were plasma total cortisol and ACTH levels after ROSC. Venous blood samples were collected in heparin anticoagulant tubes, centrifuged 10 min at 3000 rpm, and then stored at -80 °C. Plasma total cortisol (IMMULITE 2000 Cortisol, L2KCO2, UK) and ACTH (IMMULITE 2000 ACTH, L2KAC2, UK) levels were assayed using a chemiluminescent immunoassay on a Siemens automated analyzer (IMMULITE 2000 XPi; Siemens Healthcare Diagnostics, Erlangen, Germany). The equipment and reagents were calibrated by engineers before use. The lower detection limit of total cortisol was 2.00 ng/mL, and that of ACTH was 5.00 pg/mL.

165 Statistical analyses

Data analysis was used in SPSS version 22.0 (IBM Corp., Armonk, NY, USA) and sample size calculation in PASS15.0 software (NCSS, LLC, Kaysville, UT, USA). For normally distributed data, continuous variables are expressed as means with standard deviations. Since the data for total cortisol and ACTH levels had a skewed distribution, we compared our results with the natural logarithmic conversion values after adding 1 (ln [total cortisol+ 1], ln [ACTH+ 1]). Measurement data with a skewed distribution are expressed as medians (25th and 75th percentiles). The Mann-Whitney U test was used to compare variables between groups. The qualitative parameters in the 2 \times 2 contingency table were used for analysis. All statistical tests were two-tailed, and a P-value of <0.05 was considered statistically significant.

176 Follow-up

> Patients were classified into survivor and non-survivor groups according to the 28-day survival endpoint. Those with all-cause mortality within the follow-up period were considered non-survivors. If data were lost, the corresponding candidate was excluded. Patient and public involvement Patients and/or the public were not involved in the design, or conduct, or reporting, or dissemination plans of this research. Results Patient characteristics 40 healthy individuals and 85 patients who experienced CA were analyzed. The demographics and clinical characteristics of both groups are shown in Table 1. In this study, acute cardiac and brain events were the main causes of CA, with those in the latter category emanating from strokes. Other causes of CA included poisoning

> (including carbon monoxide poisoning) and hypokalemia. Sex and age were not
> significantly different between the CA and healthy control groups. The comparisons of
> clinical characteristics of the survivor and non-survivor groups based on 28-day
> survival are shown in Supplemental Table 2. The APACHE II and SOFA scores were
> significantly different between the CA and healthy control groups (P<0.001 for all) and

survivor and non-survivor groups (P<0.001 and P=0.011, respectively).

Table 1. Patient Characteristics at Admission

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		Healthy Control	Successful Resuscitatio
	Characteristics	Group (n=40)	Group (n=85)
-	Age (years), median [IQR]	64.0 (54.3, 69.8)	65.0 (55.0, 74.0)
	Male/Female (n)	23/17	58/27
	Previous medical history, n (%)		
	Hypertension	5 (12.5%)	38 (44.7%)
	Diabetes	3 (7.5%)	27 (31.8%)
	Coronary heart disease	2 (5.0%)	29 (34.1%)
	Chronic lung disease	1 (2.5%)	9 (10.6%)
	Chronic kidney disease	0	9 (10.6%)
	Cardiac arrest cause (n, %)		
	Cardiac		34 (40.0%)
	Respiratory		20 (23.5%)
	Cerebral		23 (27.1%)
	Others		7 (8.2%)
	Unknow		1 (1.2%)
	Initial resuscitation		
	Time to ROSC (min), median [IQR]		20.0 (10.0, 30.0)
	Adrenaline (mg), median [IQR]		2.0 (0.0, 5.0)
	Initial rhythm VF/VT, n (%)		30 (35.3%)
	MAP (mmHg), median [IQR]	95.7 (86.0, 103.2)	74.3 (56.2, 97.2)
	White cell count (×10 ⁹ /L), median [IQR]	5.81 (4.85, 6.53)	13.56 (10.84, 18.29)
	APACHE II score, mean±SD	0	32.9±6.5
	SOFA score, median [IQR]	0	11.5 (8.5, 14.0)
	28-day mortality, n (%)	0	65 (76.5%)
	28-day CPC 1–2, n (%)		14 (16.5%)
-	Abbreviations: IQR: interquartile rang		

200 APACHE II: acute physiology and chronic health evaluation; SOFA: sequential

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3 4	201	organ failure assessment; SD: standard deviation; CPC: cerebral performance
5 6	202	category.
7 8 9	203	Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts after
10 11	204	ROSC
12 13 14	205	The T and B lymphocyte, NK cell, and Treg cell counts were significantly lower after
15 16 17	206	ROSC in patients who experienced CA than in healthy controls (P<0.001 for all).
18 19	207	Additionally, the CD3+CD4+/T lymphocyte, CD3+CD8+/T lymphocyte, and Treg
20 21 22	208	cell/CD4 ⁺ T lymphocyte ratios were significantly lower after ROSC in patients who
23 24	209	experienced CA than in healthy controls (P<0.001 for all) (Fig. 1; Supplemental Table
25 26 27	210	3). However, there were no significant differences in these cell counts and ratios
28 29 30	211	between survivors (n=20) and non-survivors (n=65) (P>0.05 for all) (Supplemental
31 32	212	Table 4).
33 34 35	213	GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells after
36 37	214	ROSC
38 39 40	215	The MFI and percentages of GR expression in B and T lymphocytes, NK cells, and
41 42	216	CD3 ⁺ CD8 ⁺ T lymphocytes were significantly lower after ROSC in patients who
43 44 45	217	experienced CA than in healthy individuals (P<0.01 for all) (Fig. 2A–D, G, H, K, L).
46 47 48	218	There were also significant reductions in the MFI in Treg cells and CD3 ⁺ CD4 ⁺ T
49 50	219	lymphocytes (P<0.05 for all) (Fig. 2E, I) but not in the percentages of GR expression
51 52 53	220	(P>0.05 for all) (Fig. 2F, J; Supplemental Table 5). However, there were no significant
54 55	221	differences in the MFI and percentages of GR expression in these cells between
56 57 58	222	survivors and non-survivors (P>0.05 for all) (Supplemental Table 6).
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223 Changes in plasma total cortisol and ACTH levels after RC	ROSC
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We measured the plasma total cortisol and ACTH levels of the 40 healthy individuals and 85 patients who experienced CA (two samples were excluded because their total cortisol levels were not measured). Plasma total cortisol levels were significantly higher in patients who experienced CA than in healthy controls (P<0.001), but ACTH levels were not (Fig. 3A, C). No significant differences in ln (total cortisol+1) and ln (ACTH+1) values were observed between survivors and non-survivors (P>0.05 for all) (Fig. 3B, D).

232 Discussion

In this study, we examined the levels of GR expression and plasma corticosteroids in patients with CA in the early period after ROSC. We found that GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells, cell counts and ratios in patients with CA was significantly lower compared to that in controls. Furthermore, plasma total cortisol levels in patients with CA were significantly higher compared to the controls.

The ischemia-reperfusion response initiates an acute inflammatory response that contributes to post-resuscitation shock after CA.[27] The immune response of patients who experience CA is impaired, and the systemic inflammatory response increases. [6, 242 28] The T and B lymphocyte, NK cell, and Treg cell counts and CD3⁺CD4⁺/T, 243 CD3⁺CD8⁺/T, and Treg cell/CD4⁺ T lymphocyte ratios were significantly reduced after 244 ROSC. NK cells, which are special innate immune cells with cytotoxic functions

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245	similar to CD3 ⁺ CD8 ⁺ T lymphocytes, mainly distinguish infected and stressed cells
246	from healthy cells and eliminate intracellular infection and dysfunctional cells. [29, 30]
247	T lymphocytes are also crucial because they function as adaptive immune cells to
248	control and eliminate the infection. [29] Moreover, B and T lymphocytes mediate
249	humoral and cellular immunity, respectively. This study was performed earlier and
250	involved a more comprehensive assessment of the immune system of patients who
251	experienced CA. Our findings more substantially supported the rapid emergence of
252	immune dysfunction in these patients after ROSC than in previous reports.
253	The binding of GCs to GR inside different peripheral blood mononuclear cells
254	(PBMC) leads to changes in the ability of cells to regulate apoptosis, proliferation, and
255	activity, and GC-GR complexes limit the transcription (trans-repression) of
256	inflammatory genes, including those encoding for proinflammatory cytokines.[31, 32]
257	This study is the first to explore GR expression in circulating immune cells in patients
258	who experienced CA after ROSC. We observed that GR expression in B and T
259	lymphocytes, NK cells, and CD3+CD8+ T lymphocytes decreased significantly in
260	patients who experienced CA, whereas the percentage of GR ⁺ Treg cells and
261	CD3 ⁺ CD4 ⁺ T lymphocytes decreased slightly. Moreover, we observed a more
262	significant decrease in the MFI of GR expression in Treg cells and CD3 ⁺ CD4 ⁺ T
263	lymphocytes but not in the percentage of GR expression. Previous studies have found
264	decreased expression of GRs in peripheral polymorphonuclear cells in critically ill
265	patients, [21] and antagonism to GRs aggravates viral and bacterial infections. [33]
266	GCs induced upon infections help to maintain homeostasis and mitigate the life-

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threatening impact of sepsis on the host.[31] Although studies have reported that the use of GCs during and after CPR seems to confer benefits concerning ROSC rates and long-term survival, the evidence is scant. [13,18,34,35] Since cortisol signaling is mediated by GRs, we hypothesized that the differential responses of CA patients to GC may be related to their levels of GR expression. This study suggests that the decrease in intracellular GR expression in patients who experienced CA is one of the causes of GC resistance due to insufficient binding of GRs and GCs, GC insensitivity, and the inability of GCs to exert anti-inflammatory and immunosuppressive effects effectively. These findings may also explain why different results regarding the clinical application of GCs have been reported previously. Furthermore, it is vital to measure GR levels as sufficient expression of GR is essential for mediating adequate GC effects during and after CPR.

We also found that the total plasma cortisol levels were significantly higher in patients who experienced CA, but ACTH levels were not. High levels of inflammatory cytokines inhibit ACTH release. [18] During critical illness, the body does not sufficiently metabolize cortisol. [36] In addition, the continuous increase in plasma cortisol levels may trigger the negative feedback pathway of the hypothalamic-pituitary-adrenal axis, inhibiting the release of ACTH and cortisol and eventually leading to adrenal insufficiency [37]. These factors may explain the opposite trends of plasma ACTH and cortisol levels in the patients included in this study and who experienced CA. Notably, this result suggests that low GR expression levels are not matched by high plasma total cortisol levels in patients who experienced CA. The

dissociation between low GR expression and high cortisol implies an abnormal stress response. [38] Although systemic cortisol levels may be high, its availability is low during cardiac arrest. Previous studies have found that GC use during resuscitation may benefit patients who experience CA. [13-16] Possible reasons for this response may be that large doses of GCs given to CA patients may stimulate the function of GRs, or that GR expression or GC sensitivity was better in some patients. The probability of systemic inflammatory response and immunosuppression may also have been reduced in some CA patients. This study did not provide data on plasma GC levels and GR expression in a group of patients who were administered GCs and successfully resuscitated; therefore, further studies are required.

300 Limitations

Our study has several limitations. First, to assess changes, we only enrolled patients who experienced CA and had signs of systemic ischemic hypoxia, such as GCS <8 after ROSC. The patients were not stratified by age, sex, the occurrence of comorbidities, or mild systemic ischemic hypoxia. Second, since this was a preliminary observational study, we observed only early changes. A more relevant control group and dynamic observations obtained over a longer duration would be helpful to understand the significance of GR expression in evolving immunity during the clinical course of CA after ROSC. Third, the samples used in this study were from clinical laboratories; thus, plasma total cortisol and ACTH in the samples were at risk of degradation before we collected the samples. Finally, we did not discuss the changes in and roles of GR

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3 4 5	311	isoforms, free cortisol, and corticosteroid-binding globulin. Therefore, future studies
6 7	312	on these aspects are warranted to better understand the immunosuppressive effects of
8 9 10	313	ROSC among patients who experienced CA.
11 12	314	In conclusion, this study revealed that GR expression, cell counts and ratios rapidly
13 14 15	315	decreased, whereas plasma total cortisol levels increased, in the early period after
16 17 18	316	ROSC among CA patients. These findings may provide important information about
19 20	317	GR expression levels and function, and immunosuppressive status in these patients. The
21 22 23	318	assessment of GR expression in CA patients may help screening for those who are more
24 25	319	sensitive to glucocorticoid therapy.
26 27 28	320	
29 30 31	321	Acknowledgments: We thank all the patients and their families who were enrolled in
32 33	322	this study and colleagues from the emergency department who provided support. And
34 35 36	323	we are grateful for the efforts of the staff for ongoing resuscitation in hospitals.
37 38	324	Contributorship statement: CL designed the study and reviewed the manuscript.
39 40 41	325	YNY searched the literature and contributed to the experimental studies, data analysis,
42 43 44	326	and manuscript writing. ZRT, CCH, and LA collected and analyzed data. JBL and MRX
45 46	327	helped with the statistical analyses. All authors have read and approved the final
47 48 49	328	manuscript.
50 51	329	Competing interests: All authors declare no competing interest associated with this
52 53 54	330	project.
55 56 57	331	Funding: This research received no specific grant from any funding agency in public,
57 58 59	332	commercial or not-for-profit sectors.

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3 4	333	Provenance and peer review: Not commissioned; externally peer-reviewed.
5 6	334	Data sharing statement: All data relevant to the study are included in the article or
7 8	554	Data sharing statement. All data relevant to the study are mended in the attere of
9 10	335	uploaded as supplementary information. Due to privacy and ethical concerns, data can
11 12 13	336	not be shared.
14 15	337	
16 17 18	338	Ethics statements
19 20 21	339	Patient consent for publication: Not applicable.
22 23	340	Ethics approval: This study was approved by the Medical Ethics Committee of Beijing
24 25 26	341	Chaoyang Hospital (2013-KE-1). After successful resuscitation, informed consent was
27 28 29	342	obtained from the families of the patients to enroll them in the study.
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465 Figure legends

Fig. 1. Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts,
CD3⁺CD4⁺/T, CD3⁺CD8⁺/T, and Treg/CD4⁺T lymphocyte ratios between the healthy
control group and CA group. The CA group showed significant differences compared
with the healthy control group (P<0.001). CA, cardiac arrest; CD, cluster-of-
differentiation; NK, natural killer; Treg, regulatory T.

Fig. 2. Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells
in the healthy control group and CA group. The CA group showed significant
differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD,
cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC,
return of spontaneous circulation; Treg, regulatory T.

Fig. 3. (A, B) Plasma total cortisol and ACTH levels (the natural logarithmic
conversion values after adding 1) after ROSC in the healthy control group and CA
group. (C, D) Plasma total cortisol and ACTH levels in survivors and non-survivors
after ROSC. The CA group showed significant differences compared with the healthy
control group (P<0.05). ACTH, adrenocorticotrophic hormone; CA, cardiac arrest;
ROSC, return of spontaneous circulation.

P = 0.001

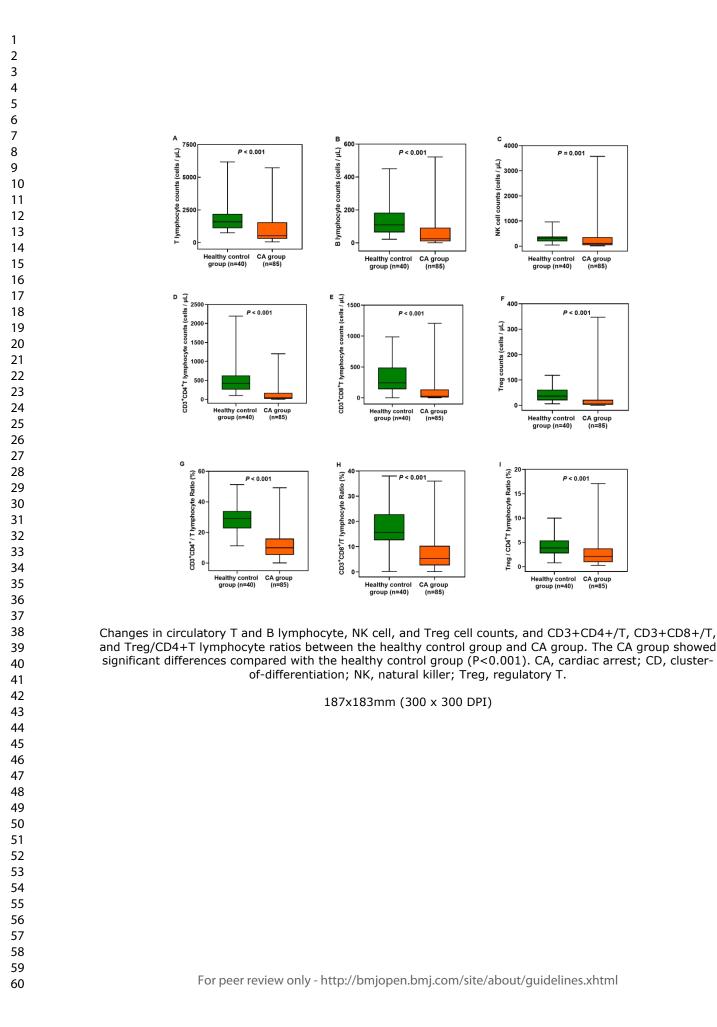
CA group (n=85)

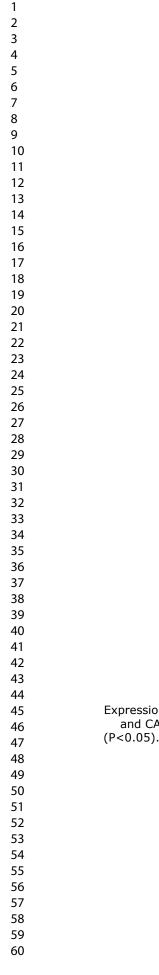
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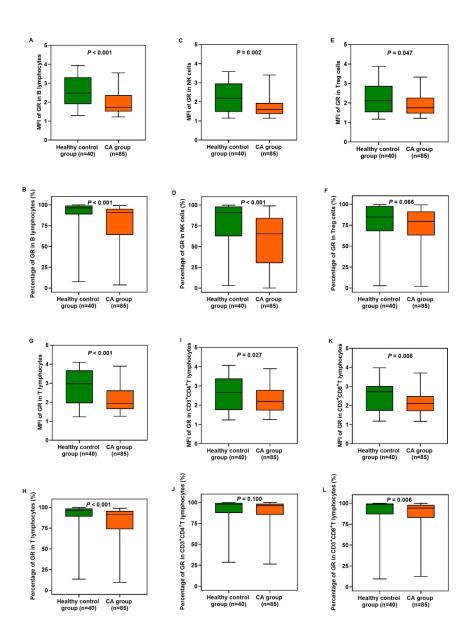
CA group (n=85)

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P < 0.001

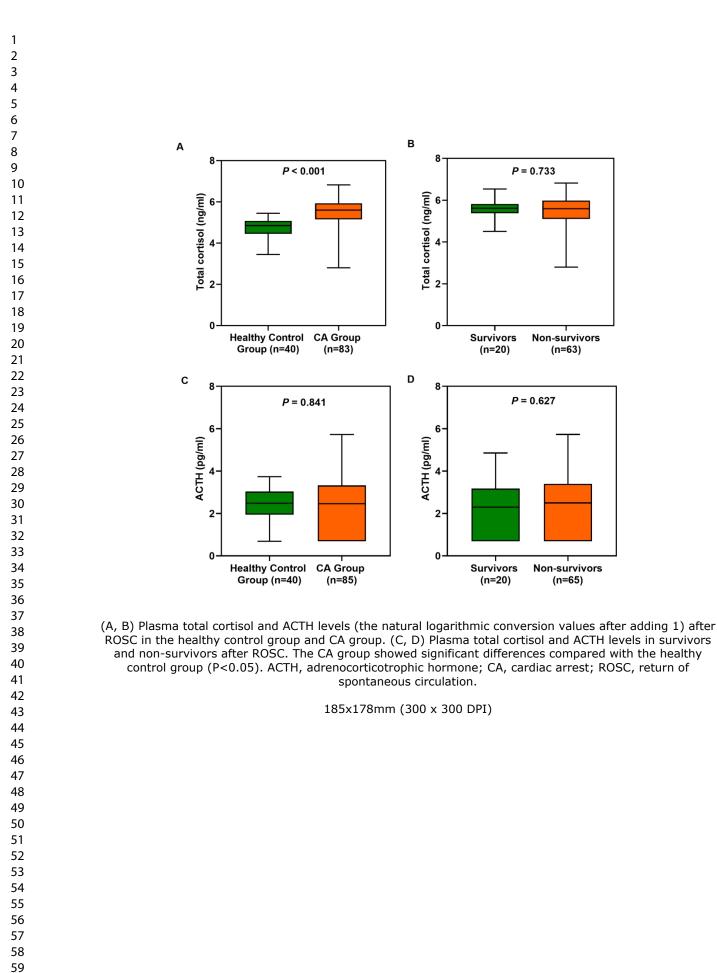






Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells in the healthy control group and CA group. The CA group showed significant differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD, cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC, return of spontaneous circulation; Treg, regulatory T.

199x256mm (300 x 300 DPI)



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Electronic supplemental material

Glucocorticoid receptor expression in patients with cardiac arrest in the early period after the return of

spontaneous circulation: A prospective observational study

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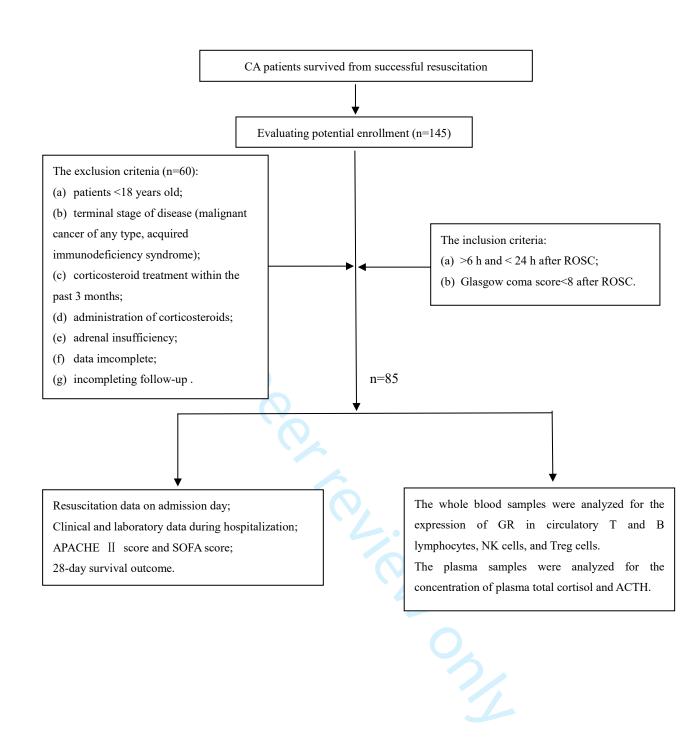
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Contents

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- Supplemental Table 1
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Supplemental Figure 1. The flow chart of the study.

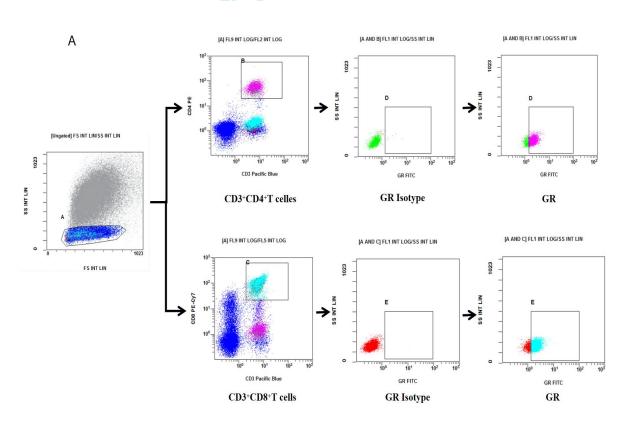
Abbreviations: CA, cardiac arrest; ROSC, return of spontaneous circulation; APACHE II, acute physiology and chronic health evaluation II; SOFA, sequential organ failure assessment; GR, glucocorticoid receptor; Treg, regulatory T; ACTH, adrenocorticotrophic hormone.

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Supplemental Figure 2. Representative plots and gating strategies for analyzing glucocorticoid receptor (GR) in the whole blood.

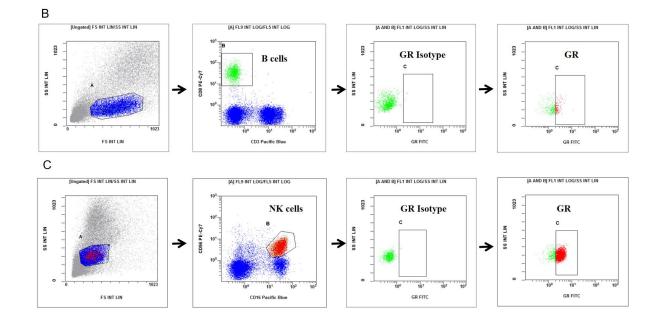
GR expression levels were determined on T cells, B cells, NK cells, and T regulatory (Treg) cells. Single cells were gated from all cellular events (FSC/SSC gate). B cells were identified as CD3⁻CD19⁺ cells. NK cells were identified as CD16⁺56⁺ cells. T cells were identified as CD3⁺CD4⁺ T cells and CD3⁺CD8⁺ T cells. Treg cells were identified as CD4⁺CD25^{high}CD127^{low}.

A. Expression of GR on T cells



B. Expression of GR on B cells

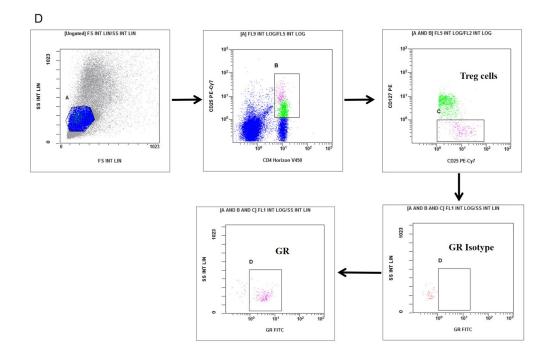
C. Expression of GR on NK cells



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D. Expression of GR on Treg cells



Source

BD Pharmingen^a

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

BD Pharmingen

Bio-Rad^b

Bio-Rad

Antigen	Catalog Number	Fluorescein
CD3	558117	Pacific Blue
CD4	555347	PE
CD4	560345	Horizon V4
CD8	557746	PE-Cy7
CD19	557835	PE-Cy7
CD16	558122	Pacific Blue
CD56	557747	PE-Cy7
CD25	557741	PE-Cy7
CD127	557938	РЕ
GR	MCA2469F	FITC
Mouse IgG1 Isotype	MCA928F	FITC
Mouse IgG1,к Isotype	557872	PE-Cy7
Mouse IgG1,к Isotype	554680	PE
Mouse IgG1,к Isotype	558120	Pacific Blue

acific Blue **BD** Pharmingen otec, Oxford, UK. coerythrin; FITC, fluorescein isothiocyanate; GR, glucocorticoid

receptor; Ig: immunoglobulin.

	Survivors (n=20)	Non-survivors (n=65)
Age (years), median [IQR]	59.0 (53.3, 72.8)	66.0 (59.0, 75.5)
Male/Female (n)	12/8	46/19
Cardiac arrest cause (n, %)		
Cardiac	10 (50.0%)	24 (36.9%)
Non-Cardiac	10 (50.0%)	41 (63.1%)
Initial resuscitation		
Time to ROSC (min), median [IQR]	15.0 (7.3, 26.0)	20.0 (15.0, 30.0)
Adrenaline (mg), median [IQR]	1.0 (0.0, 3.0)	2.0 (0.0, 5.0)
Initial rhythm VF/VT, n (%)	11 (55.0%)	19 (29.2%)
MAP (mmHg), median [IQR]	89.9 (70.5, 104.9)	70.7 (50.0, 93.5)
White cell count (×10 ⁹ /L), median [IQR]	12.40 (6.98, 18.76)	13.80 (11.67, 18.20)
Lactate (mmol/L), median [IQR]	3.50 (1.33, 7.05)	7.50 (3.80, 11.20)
APACHE II score, mean±SD	27.8±6.6	34.4±5.6
SOFA score, median [IQR]	9.0 (7.3, 11.8)	12.0 (9.0, 15.0)

Supplemental Table 2. Characteristics of CA survivors and non-survivors on admission.

Data are presented as mean±SD or interquartile range (IQR) as appropriate. Abbreviations: ROSC: return of spontaneous circulation; VF: ventricular fibrillation; VT: ventricular tachycardia; MAP: mean arterial pressure; APACHE II: acute physiology and chronic health evaluation; SOFA: sequential organ failure assessment.

Supplemental Table 3. The flow cytometry results of cell counts and ratios of the healthy control group and successful resuscitation group

	Healthy Control	Successful	Z-value	<i>P</i> -value
	Group (n=40)	Resuscitation Group		
		(n=85)		
T lymphocyte count (cells /µL)	1586.0 (1101.5, 2192.5)	514.0 (287.5, 1555.0)	-4.515	<0.001
NK cell count (/µL)	311.5 (191.0, 378.8)	101.0 (36.0, 351.5)	-3.332	0.001
B lymphocyte count (/μL)	109.3 (63.7, 183.3)	25.7 (9.4, 92.3)	-5.076	<0.001
Treg count (/µL)	0.259 (0.095, 0.516)	0.233 (0.135, 0.488)	-5.518	<0.001
Treg / CD4 ⁺ T lymphocyte Ratio	0.039 (0.028, 0.054)	0.021 (0.010, 0.038)	-4.418	<0.001
$CD3^+CD4^+ T$ lymphocyte count (/µL)	421.7 (258.6, 627.4)	38.9 (17.6, 168.3)	-6.256	<0.001
CD3 ⁺ CD4 ⁺ / T lymphocyte Ratio	0.292 (0.227, 0.340)	0.100 (0.054, 0.160)	-7.066	<0.001
CD3 ⁺ CD8 ⁺ T lymphocyte count (/µL)	241.1 (139.5, 488.6)	26.3 (7.2, 135.9)	-5.287	<0.001
CD3 ⁺ CD8 ⁺ / T lymphocyte Ratio	0.157 (0.126, 0.229)	0.053 (0.026, 0.104)	-5.719	< 0.001

All the data in Supplemental table 3 are represented as the median [IQR]; IQR: Interquartile Range; CD: cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

Supplemental Table 4. The flow cytometry results of cell counts and ratios of the CA patients on admission based on

Survivors (n=20)	Non-survivors	Z-value	<i>P</i> -value
	(n=65)		
502.0 (353.8, 1199.8)	514.0 (282.5, 1891.0)	-0.186	0.852
167.0 (29.8, 309.3)	100.0 (36.0, 404.0)	-0.218	0.828
38.6 (15.7, 103.5)	19.2 (7.1, 65.7)	-0.632	0.527
0.318 (0.145, 0.552)	0.212 (0.128, 0.479)	-0.611	0.396
0.025 (0.009, 0.043)	0.021 (0.010, 0.034)	-0.498	0.619
55.1 (32.4, 228.0)	38.0 (16.0, 168.1)	-0.850	0.396
0.118 (0.070, 0.236)	0.097 (0.049, 0.142)	-1.565	0.118
25.4 (12.5, 96.2)	26.3 (6.3, 138.8)	-0.021	0.983
0.054 (0.033, 0.104)	0.053 (0.025, 0.104)	-0.187	0.852
	502.0 (353.8, 1199.8) 167.0 (29.8, 309.3) 38.6 (15.7, 103.5) 0.318 (0.145, 0.552) 0.025 (0.009, 0.043) 55.1 (32.4, 228.0) 0.118 (0.070, 0.236) 25.4 (12.5, 96.2)	(n=65) 502.0 (353.8, 1199.8) 514.0 (282.5, 1891.0) 167.0 (29.8, 309.3) 100.0 (36.0, 404.0) 38.6 (15.7, 103.5) 19.2 (7.1, 65.7) 0.318 (0.145, 0.552) 0.212 (0.128, 0.479) 0.025 (0.009, 0.043) 0.021 (0.010, 0.034) 55.1 (32.4, 228.0) 38.0 (16.0, 168.1) 0.118 (0.070, 0.236) 0.097 (0.049, 0.142) 25.4 (12.5, 96.2) 26.3 (6.3, 138.8)	(n=65)502.0 (353.8, 1199.8)514.0 (282.5, 1891.0)-0.186167.0 (29.8, 309.3)100.0 (36.0, 404.0)-0.21838.6 (15.7, 103.5)19.2 (7.1, 65.7)-0.6320.318 (0.145, 0.552)0.212 (0.128, 0.479)-0.6110.025 (0.009, 0.043)0.021 (0.010, 0.034)-0.49855.1 (32.4, 228.0)38.0 (16.0, 168.1)-0.8500.118 (0.070, 0.236)0.097 (0.049, 0.142)-1.56525.4 (12.5, 96.2)26.3 (6.3, 138.8)-0.021

All the data in Supplemental table 4 are represented as the median [IQR]; IQR: Interquartile Range; CD: cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

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59 60 Supplemental Table 5. The flow cytometry results of GR expression in the CA group and successful resuscitation group.

	Healthy Control	Successful	Z-value	<i>P</i> -value
	Group (n=40)	Resuscitation Group		
		(n=85)		
Percentage of GR on B lymphocytes	0.963 (0.885, 0.992)	0.896 (0.605, 0.949)	-3.742	<0.001
MFI of GR on B lymphocytes	2.48 (1.91, 3.31)	1.73 (1.50, 2.37)	-3.980	< 0.001
Percentage of GR on T lymphocytes	0.964 (0.889, 0.986)	0.900 (0.703, 0.955)	-3.755	< 0.001
MFI of GR on T lymphocytes	2.98(1.95, 3.68)	1.92 (1.36, 1.99)	-3.853	< 0.001
Percentage of GR on NK cells	0.907 (0.624, 0.983)	0.611 (0.306, 0.840)	-3.792	< 0.001
MFI of GR on NK cells	2.19 (1.48, 2.96)	1.60 (1.36, 1.99)	-3.171	0.002
Percentage of GR on Treg cells	0.848 (0.680, 0.978)	0.784 (0.589, 0.911)	-1.837	0.066
MFI of GR on Treg cells	2.12 (1.53, 2.88)	1.76 (1.44, 2.30)	-1.990	0.047
Percentage of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	0.980 (0.874, 0.996)	0.957 (0.824, 0.985)	-2.204	0.100
MFI of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	2.65 (1.75, 3.38)	2.17 (1.70, 2.92)	-1.646	0.027
Percentage of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	0.986 (0.868, 0.996)	0.938 (0.823, 0.979)	-2.758	0.006
MFI of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	2.73 (1.73, 3.02)	2.10 (1.68, 2.54)	-2.668	0.008

All the data in Supplemental table 5 are represented as the median [IQR]. Abbreviations: IQR, interquartile range; CD, cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, Glucocorticoid receptor; MFI, mean fluorescence intensity.

	Survivors	Non-survivors	Z-value	P-value
	(n=20)	(n=65)		
Percentage of GR on B lymphocytes	0.904 (0.595, 0.976)	0.906 (0.657, 0.946)	-0.787	0.431
MFI of GR on B lymphocytes	1.92 (1.52, 2.54)	1.72 (1.51, 2.31)	-0.881	0.378
Percentage of GR on T lymphocytes	0.899 (0.778, 0.969)	0.913 (0.692, 0.951)	-1.057	0.291
MFI of GR on T lymphocytes	2.05 (1.67, 2.83)	1.91 (1.64, 2.46)	-1.031	0.303
Percentage of GR on NK cells	0.717 (0.292, 0.886)	0.556 (0.302, 0.823)	-0.756	0.449
MFI of GR on NK cells	1.54 (1.37, 2.09)	1.61 (1.34, 1.87)	-0.565	0.572
Percentage of GR on Tregs	0.780 (0.667, 0.849)	0.799 (0.576, 0.923)	-0.440	0.660
MFI of GR on Tregs	1.61 (1.48, 2.30)	1.77 (1.45, 2.27)	-0.005	0.996
Percentage of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	0.975 (0.876, 0.985)	0.957 (0.845, 0.987)	-0.617	0.538
MFI of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	2.08 (1.72, 3.35)	2.22 (1.71, 2.69)	-0.865	0.387
Percentage of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	0.963 (0.816, 0.977)	0.938 (0.834, 0.980)	-0.254	0.800
MFI of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	2.08 (1.68, 3.10)	2.11(1.71, 2.46)	-0.653	0.514

All the data in Supplemental table 6 are represented as the median [IQR]. Abbreviations: IQR, Interquartile Range; CD, Cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, glucocorticoid receptor; MFI, mean fluorescence intensity.

Supplemental Table 6. The flow cytometry results of GR expression in the survivors and non-survivors.

STROBE Statement—checklist of items that should be included in reports of observational studies

	Item No	Recommendation	Page No
Title and abstract	1	(a) Indicate the study's design with a commonly used term in	1
		the title or the abstract	
		(b) Provide in the abstract an informative and balanced	3
		summary of what was done and what was found	
Introduction			
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	3-5
Objectives	3	State specific objectives, including any prespecified hypotheses	5
Methods		51	1
Study design	4	Present key elements of study design early in the paper	Supplemental Figure 1
Setting	5	Describe the setting, locations, and relevant dates, including	5-9,
-		periods of recruitment, exposure, follow-up, and data	Supplemental
		collection	Figure 1
Participants	6	(a) Cohort study—Give the eligibility criteria, and the sources	5,6,8,9
		and methods of selection of participants. Describe methods of	
		follow-up	
		Case-control study—Give the eligibility criteria, and the	
		sources and methods of case ascertainment and control	
		selection. Give the rationale for the choice of cases and	
		controls	
		Cross-sectional study—Give the eligibility criteria, and the	
		sources and methods of selection of participants	
		(b) Cohort study—For matched studies, give matching criteria	5
		and number of exposed and unexposed	
		Case-control study—For matched studies, give matching	
		criteria and the number of controls per case	
Variables	7	Clearly define all outcomes, exposures, predictors, potential	5, 6, 8,
		confounders, and effect modifiers. Give diagnostic criteria, if	Supplemental
		applicable	Figure 1
Data sources/	8*	For each variable of interest, give sources of data and details	6, 8
measurement		of methods of assessment (measurement). Describe	., -
		comparability of assessment methods if there is more than one	
		group	
Bias	9	Describe any efforts to address potential sources of bias	6-8
Study size	10	Explain how the study size was arrived at	Supplemental
<u>,</u>	-		Figure 1
Quantitative variables	11	Explain how quantitative variables were handled in the	6-8
		analyses. If applicable, describe which groupings were chosen	
		and why	
Statistical methods	12	(<i>a</i>) Describe all statistical methods, including those used to control for confounding	8, 11

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(b) Describe any methods used to examine subgroups and interactions	N/A
(c) Explain how missing data were addressed	8, 11
(<i>d</i>) <i>Cohort study</i> —If applicable, explain how loss to follow-up was addressed	11
<i>Case-control study</i> —If applicable, explain how matching of cases and controls was addressed	
Cross-sectional study—If applicable, describe analytical	
methods taking account of sampling strategy	
(e) Describe any sensitivity analyses	

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Participants	13	(a) Report numbers of individuals at each stage of study—eg numbers	9,
	*	potentially eligible, examined for eligibility, confirmed eligible, included in	Supplem
		the study, completing follow-up, and analysed	1 Figure 1
		(b) Give reasons for non-participation at each stage	9,12,
			Suppleme
			l Figure 1
		(c) Consider use of a flow diagram	Suppleme
			1 Figure 1
Descriptive	14	(a) Give characteristics of study participants (eg demographic, clinical,	9
data	*	social) and information on exposures and potential confounders	
		(b) Indicate number of participants with missing data for each variable of	Suppleme
		interest	1 Figure 1
		(c) <i>Cohort study</i> —Summarise follow-up time (eg, average and total amount)	8
Outcome data	15	Cohort study—Report numbers of outcome events or summary measures	9-12
	*	over time	
		Case-control study-Report numbers in each exposure category, or summary	
		measures of exposure	
		Cross-sectional study—Report numbers of outcome events or summary	
		measures	
Main results	16	(a) Give unadjusted estimates and, if applicable, confounder-adjusted	9-12,
		estimates and their precision (eg, 95% confidence interval). Make clear	Electronic
		which confounders were adjusted for and why they were included	suppleme
			material
		(b) Report category boundaries when continuous variables were categorized	
		(c) If relevant, consider translating estimates of relative risk into absolute	
		risk for a meaningful time period	
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and	N/A
		sensitivity analyses	
Discussion			
Key results	18	Summarise key results with reference to study objectives	12
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias	15
		or imprecision. Discuss both direction and magnitude of any potential bias	
Interpretation	20	Give a cautious overall interpretation of results considering objectives,	12-14
		limitations, multiplicity of analyses, results from similar studies, and other	
		relevant evidence	
Generalisabilit	21	Discuss the generalisability (external validity) of the study results	15
у			
Other informat	ion		
Funding	22	Give the source of funding and the role of the funders for the present study	17
		and, if applicable, for the original study on which the present article is based	

*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies.

Note: An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely

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available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.

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Glucocorticoid receptor expression in patients with cardiac arrest in the early period after the return of spontaneous circulation: A prospective observational single-center study

Journal:	BMJ Open
	lined of the line
Manuscript ID	bmjopen-2021-060246.R2
Article Type:	Original research
Date Submitted by the Author:	10-Aug-2022
Complete List of Authors:	Yu, Yanan; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Tang, Ziren; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Xie, Miaorong; Capital Medical University Affiliated Beijing Friendship Hospital, Department of Emergency Medicine Li, Jiabao; Capital Medical University Affiliated Beijing Friendship Hospital, Department of Critical Care Hang, Chen-Chen; Beijing Chao-Yang Hospital, Emergency Medicine An, Le; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine Li, Chunsheng; Beijing Chao-Yang Hospital Capital Medical University, Department of Emergency Medicine
Primary Subject Heading :	Emergency medicine
Secondary Subject Heading:	Intensive care
Keywords:	ACCIDENT & EMERGENCY MEDICINE, INTENSIVE & CRITICAL CARE, Adult intensive & critical care < INTENSIVE & CRITICAL CARE

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1	Glucocorticoid receptor expression in patients with cardiac arrest in the early
2	period after the return of spontaneous circulation: A prospective observational
3	single-center study
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16	
17	Keywords: Cardiac arrest, glucocorticoid receptor, immunosuppression, cortisol
18	Word count of the main text: 3,437 words
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23 Abstract

Objectives: Rapid changes in glucocorticoid (GC) levels and adrenal insufficiency are related to the development of post-cardiac arrest (CA) syndrome. However, GC receptor (GR) expression changes have not been studied. Hence, this study aimed to investigate the association of early changes in GR expression and prognosis and immune response in patients who experienced CA.

- 29 **Design:** Prospective observational study.
- 30 Setting: Emergency department.

Participants: Patients (85) in the early period of return of spontaneous circulation
(ROSC) after CA were admitted between October 2018 and October 2019. After a
physical examination, age- and sex-matched healthy individuals (40) were recruited for
the control group.

Primary and secondary outcome measures: GR expression and cell counts of
circulatory T and B lymphocytes, natural killer cells, and regulatory T (Treg) cells were
assessed. Plasma total cortisol and adrenocorticotrophic hormone (ACTH) levels were
also tested.

Results: All cell counts were lower, and plasma total cortisol levels were higher (P<0.001) in patients who experienced CA than in the healthy control group. GR expression in Treg cells and CD3⁺CD4⁺ T lymphocytes were not significantly different, but the mean fluorescence intensity and GR expression in other cells were lower in patients who experienced CA (P<0.05) than in the healthy control group. ACTH levels were not different. There were no significant differences between survivors and non-

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45	survivors.
46	Conclusions: This study revealed that GR expression and cell counts rapidly decreased,
47	whereas plasma total cortisol levels increased in the early period after ROSC among
48	patients who experienced CA. Our findings provide important information about GR
49	level and function, and immunosuppressive status in these patients. Assessing GR
50	expression in CA patients may help screening for those who are more sensitive to
51	glucocorticoid therapy.
52	
53	Strengths and limitations of this study
54	1. The study was designed as single-center, prospective study.
55	2. This is the first study to evaluate the GR expression in the early period following
56	ROSC among CA patients.
57	3. We only studied the GR expression of CA patients in the early period following
58	ROSC; therefore, our results cannot be extrapolated to time points beyond 24 hours.
59	4. Decreased GR expression may affect the sensitivity of CA patients to GCs.
60	5. Decreased GR expression may affect potential immune consequences of CA
61	patients.
62	
63	Introduction
64	Cardiac arrest (CA) is a significant health problem globally; about 356,500 people

- experience medical emergencies due to CA in the United States, and over 544,000
- people die from sudden CA in China annually. [1, 2] The systemic ischemia-reperfusion 66

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67	response in patients who have experienced CA can present as post-cardiac arrest
68	syndrome (PCAS) or systematic inflammatory response syndrome (SIRS), which
69	increases the risk of multiple organ failure and infection and affects the inflammatory
70	response and prognosis of patients after the return of spontaneous circulation (ROSC).
71	[3-6]
72	CA is the most intense among acute stress events, which seriously affect the pituitary
73	and adrenal axis function. [7] Studies have shown that abnormal cortisol levels and
74	relative adrenocortical insufficiency after ROSC in patients who experienced CA are
75	related to their prognosis. [8-11] However, the clinical application of glucocorticoids
76	(GCs) is controversial. In the 2015 International Cardiopulmonary Resuscitation
77	Guidelines, the routine use of GCs is not recommended for the resuscitation of patients
78	with in-hospital or out-of-hospital CA. [12] Recent clinical studies have shown that
79	early administration of corticosteroids after CA can improve the success rate of ROSC,
80	nervous system functional outcome, and prognosis, which is speculated to be related to
81	its influence on hemodynamics, and SIRS response, and other mechanisms. [12-17]
82	Therefore, the role of GCs in the occurrence and development of PCAS needs to be
83	studied further.
81	GCs combine with intracellular GC recentors (GRs) to evert anti-inflammatory and

GCs combine with intracellular GC receptors (GRs) to exert anti-inflammatory and immunosuppressive effects and reduce the production and the release of inflammatory cytokines. [18, 19] The affinity of GRs to GCs in circulating monocytes is decreased in patients with acquired immunodeficiency syndrome. [20] The expression of GR alpha and beta in peripheral polymorphonuclear cells is decreased in patients with critical

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3 4	89	illness, [21] pediatric septic shock, and high serum cortisol levels. [22] However, no
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6 7	90	study has reported the GR expression after ROSC in patients who experienced CA.
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9 10	91	Previous studies have found that the counts of circulating B and T lymphocytes,
11		
12	92	regulatory T (Treg) cells, and monocytes and expression of human leukocyte antigen
13 14		
15	93	DR (HLA-DR) on circulatory monocytes and B and T lymphocytes are reduced. [23,
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17 18	94	24] Hence, this study aimed to investigate the relationship between GR expression and
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20	95	immune alteration in the early period after ROSC in patients who experienced CA by
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22	96	observing GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells,
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25 26	97	their cell counts, and total plasma cortisol and adrenocorticotrophic hormone (ACTH)
20	20	
28	98	levels.
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32	100	Materials and methods
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35 36	101	Study participants
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35 36 37 38 39 40	102	Study participants This was an observational study conducted in the Emergency Department (ED).
35 36 37 38 39 40 41		Study participants
35 36 37 38 39 40	102 103	Study participants This was an observational study conducted in the Emergency Department (ED). According to the 2015 International Cardiopulmonary Resuscitation Guidelines, [25]
35 36 37 38 39 40 41 42 43 44	102	Study participants This was an observational study conducted in the Emergency Department (ED).
35 36 37 38 39 40 41 42 43 44 45	102 103 104	Study participants This was an observational study conducted in the Emergency Department (ED). According to the 2015 International Cardiopulmonary Resuscitation Guidelines, [25] we enrolled patients in the early ROSC period after CA (both in-hospital and out-of-
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2015 International Cardiopulmonary Resuscitation Consensus. [13] After a physical
examination, age- and sex-matched healthy individuals were recruited for the control
group.

114 Data collection

Data collection was performed according to the 2004 guidelines of the Utstein Style 115 template. [26] We collected data on demographics, resuscitation (initial heart rhythm, 116 ROSC time, and cumulative adrenaline [epinephrine] dose, and laboratory findings 117 routine blood cell counts, blood gas analysis, and blood biochemical tests performed > 118 119 6 h and < 24 h after ROSC). Acute Physiology and Chronic Health Evaluation (APACHE) II and the Sequential Organ Failure Assessment (SOFA) were used to 120 determine disease severity. Residual blood samples from routine clinical tests or 121 122 physical health examinations in the morning were collected, maintained at 4 °C during transport and storage, and used to determine GR expression in circulatory T and B 123 lymphocytes, NK cells, and Treg cells and their cell counts. The plasma was maintained 124 125 at -80 °C during storage and used to determine total cortisol and ACTH levels. During follow-up, 28-day survival data were also collected. Supplemental Figure 1 shows the 126 workflow of this study. 127

128 Outcome measures

The primary outcomes of this study were GR expression and cell counts of T and B
cells, NK cells, and Treg cells, measured by flow cytometry. Venous blood samples
collected in ethylenediaminetetraacetic acid tubes, then used to measure GR expression
in T and B lymphocytes, NK cells, and Treg cells. Briefly, a 100-μL peripheral blood

13	33	sample was stained for 20 min with surface antibodies (CD3, CD4, CD8, CD19, CD16,
13	34	CD56, CD25, and CD127) in a dark place. Erythrocytes were lysed for 15 min, and the
13	35	debris was washed away. Before staining of the intracellular GR antibody and its
13	36	isotype control (Bio-Rad AbD Serotec, Oxford, UK), surface-stained cells were fixed
13	37	and permeabilized using the BD Transcription Factor Buffer Set (BD Pharmingen, San
13	38	Diego, USA, Catalogue No. 562574). Monoclonal antibodies and their isotype controls
13	39	were all purchased from BD Biosciences (San Jose, CA, USA). Details of all antibodies
14	40	are shown in Supplemental Table 1. According to the manufacturer's recommendations,
14	41	all antibodies and their isotype controls were used at a concentration of 1 μ L per 100
14	42	μ L of whole blood. Samples were measured using the Gallios flow cytometer (Beckman
14	43	Coulter, Brea, CA, USA) and analyzed using Gallios Software version 1.0 (Beckman
14	14	Coulter). The flow cytometer was periodically calibrated by an engineer. Cells were
14	45	stained for 20 min; thresholds were defined using the manufacturer's recommended
14	46	isotype controls. Representative plots and gating strategy from a single sample are
14	47	shown in Supplemental Figure 2. T cells were gated by CD3+CD4+ or CD3+CD8+, B
14	48	cells were gated by CD3 ⁻ CD19 ⁺ , NK cells were gated by CD16 ⁺ CD56 ⁺ , and Tregs were
14	19	gated by CD4+CD25 ^{high} CD127 ^{low} . At least 10,000 events were collected in the
15	50	lymphocyte cell gate for each sample. Results are expressed as percentages and mean
15	51	fluorescence intensity (MFI) values.
15	52	Absolute CD3 ⁺ and CD4 ⁺ lymphocyte, NK cell, and Treg cell counts were obtained

Absolute CD3⁺ and CD4⁺ lymphocyte, NK cell, and Treg cell counts were obtained
 using Flow-Count fluorospheres (Beckman Coulter, Catalogue No. 7547053),
 according to the manufacturer's instructions. B, CD3⁺CD4⁺T, CD3⁺CD8⁺T, and Treg

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155 cell counts were calculated by their percentages in CD3⁺ or CD4⁺ lymphocytes
156 multiplied by CD3⁺ or CD4⁺ lymphocyte counts.

The secondary outcomes of this study were plasma total cortisol and ACTH levels after ROSC. Venous blood samples were collected in heparin anticoagulant tubes, centrifuged 10 min at 3000 rpm, and then stored at -80 °C. Plasma total cortisol (IMMULITE 2000 Cortisol, L2KCO2, UK) and ACTH (IMMULITE 2000 ACTH, L2KAC2, UK) levels were assayed using a chemiluminescent immunoassay on a Siemens automated analyzer (IMMULITE 2000 XPi; Siemens Healthcare Diagnostics, Erlangen, Germany). The equipment and reagents were calibrated by engineers before use. The lower detection limit of total cortisol was 2.00 ng/mL, and that of ACTH was 5.00 pg/mL.

166 Sample size calculation and statistical analysis

The sample size was calculated using the PASS15.0 software (NCSS, LLC, Kaysville, UT, USA) and the non-parametric test method. The median GR expression was 0.93 and 0.80 in the healthy and CA groups, respectively, and the interquartile spacing was 0.1 and 0.3. According to the ratio of 1:2 between the two groups, with a test level of 0.05 and a confidence interval of 0.90, a total of 105 samples were required, comprising at least 35 in the healthy group and 70 in the CA group. The number of people included in the two groups in this study was 40 and 85, respectively, which met our research requirements. Data analysis was used in SPSS version 22.0 (IBM Corp., Armonk, NY, USA). For normally distributed data, continuous variables are expressed as means with standard deviations. Since the data for total cortisol and ACTH levels

177	had a skewed distribution, we compared our results with the natural logarithmic
178	conversion values after adding 1 (ln [total cortisol+ 1], ln [ACTH+ 1]). Measurement
179	data with a skewed distribution are expressed as medians (25th and 75th percentiles).
180	The Mann-Whitney U test was used to compare variables between groups. The
181	qualitative parameters in the 2 \times 2 contingency table were used for analysis. All
182	statistical tests were two-tailed, and a P-value of <0.05 was considered statistically
183	significant.
184	Follow-up
185	Patients were classified into survivor and non-survivor groups according to the 28-
186	day survival endpoint. Those with all-cause mortality within the follow-up period were
187	considered non-survivors. If data were lost, the corresponding candidate was excluded.
188	Patient and public involvement
189	Patients and/or the public were not involved in the design, or conduct, or reporting,
190	or dissemination plans of this research.
191	
192	Results
193	Patient characteristics
194	40 healthy individuals and 85 patients who experienced CA were analyzed. The
195	demographics and clinical characteristics of both groups are shown in Table 1. In this
196	study, acute cardiac and brain events were the main causes of CA, with those in the
197	latter category emanating from strokes. Other causes of CA included poisoning
198	(including carbon monoxide poisoning) and hypokalemia. Sex and age were not

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significantly different between the CA and healthy control groups. The comparisons of clinical characteristics of the survivor and non-survivor groups based on 28-day survival are shown in Supplemental Table 2. The APACHE II and SOFA scores were significantly different between the CA and healthy control groups (P<0.001 for all) and survivor and non-survivor groups (P<0.001 and P=0.011, respectively).

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205 **Table 1.** Patient Characteristics at Admission

	Healthy Control	Successful Resuscitation
Characteristics	Group (n=40)	Group (n=85)
Age (years), median [IQR]	64.0 (54.3, 69.8)	65.0 (55.0, 74.0)
Male/Female (n)	23/17	58/27
Previous medical history, n (%)		
Hypertension	5 (12.5%)	38 (44.7%)
Diabetes	3 (7.5%)	27 (31.8%)
Coronary heart disease	2 (5.0%)	29 (34.1%)
Chronic lung disease	1 (2.5%)	9 (10.6%)
Chronic kidney disease	0	9 (10.6%)
Cardiac arrest cause (n, %)		
Cardiac		34 (40.0%)
Respiratory		20 (23.5%)
Cerebral		23 (27.1%)
Others		7 (8.2%)
Unknow		1 (1.2%)
Initial resuscitation		
Time to ROSC (min), median [IQR]		20.0 (10.0, 30.0)
Adrenaline (mg), median [IQR]		2.0 (0.0, 5.0)
Initial rhythm VF/VT, n (%)		30 (35.3%)

	MAP (mmHg), median [IQR]	95.7 (86.0, 103.2)	74.3 (56.2, 97.2)
	White cell count (×10 ⁹ /L), median [IQR]	5.81 (4.85, 6.53)	13.56 (10.84, 18.29)
	APACHE II score, mean±SD	0	32.9±6.5
	SOFA score, median [IQR]	0	11.5 (8.5, 14.0)
	28-day mortality, n (%)		65 (76.5%)
	28-day CPC 1–2, n (%)		14 (16.5%)
206	Abbreviations: IQR: interquartile range; ROSC: return of spontaneous circulation;		
207	VF: ventricular fibrillation; VT: ventricular tachycardia; MAP: mean arterial pressure;		
208	APACHE II: acute physiology and chronic health evaluation; SOFA: sequential		
209	organ failure assessment; SD: standard deviation; CPC: cerebral performance		
210	category.		
211	Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts after		
212	ROSC		
212	KUSC		
213	The T and B lymphocyte, NK cell, and	nd Treg cell counts	were significantly lower after
214	ROSC in patients who experienced (CA than in health	y controls (P<0.001 for all).
215	Additionally, the CD3+CD4+/T lymp	phocyte, CD3+CD	8+/T lymphocyte, and Treg
216	cell/CD4 ⁺ T lymphocyte ratios were s	significantly lower	after ROSC in patients who
217	experienced CA than in healthy controls (P<0.001 for all) (Fig. 1; Supplemental Table		
218	3). However, there were no significant differences in these cell counts and ratios		
219	between survivors (n=20) and non-su	nrvivors (n=65) (P	>0.05 for all) (Supplemental
220	Table 4).		
221	GR expression in circulatory T and	B lymphocytes, N	K cells, and Treg cells after
222	ROSC		
223	The MFI and percentages of GR exp	pression in B and T	Γ lymphocytes, NK cells, and

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CD3⁺CD8⁺ T lymphocytes were significantly lower after ROSC in patients who
experienced CA than in healthy individuals (P<0.01 for all) (Fig. 2A–D, G, H, K, L).
There were also significant reductions in the MFI in Treg cells and CD3⁺CD4⁺ T
lymphocytes (P<0.05 for all) (Fig. 2E, I) but not in the percentages of GR expression
(P>0.05 for all) (Fig. 2F, J; Supplemental Table 5). However, there were no significant
differences in the MFI and percentages of GR expression in these cells between
survivors and non-survivors (P>0.05 for all) (Supplemental Table 6).

231 Changes in plasma total cortisol and ACTH levels after ROSC

We measured the plasma total cortisol and ACTH levels of the 40 healthy individuals and 85 patients who experienced CA (two samples were excluded because their total cortisol levels were not measured). Plasma total cortisol levels were significantly higher in patients who experienced CA than in healthy controls (P<0.001), but ACTH levels were not (Fig. 3A, C). No significant differences in ln (total cortisol+1) and ln (ACTH+1) values were observed between survivors and non-survivors (P>0.05 for all) (Fig. 3B, D).

240 Discussion

In this study, we examined the levels of GR expression and plasma corticosteroids in patients with CA in the early period after ROSC. We found that GR expression in circulatory T and B lymphocytes, NK cells, and Treg cells, cell counts and ratios in patients with CA was significantly lower compared to that in controls. Furthermore, plasma total cortisol levels in patients with CA were significantly higher compared to

the controls.

The ischemia-reperfusion response initiates an acute inflammatory response that contributes to post-resuscitation shock after CA.[27] The immune response of patients who experience CA is impaired, and the systemic inflammatory response increases. [6, 28] The T and B lymphocyte, NK cell, and Treg cell counts and CD3⁺CD4⁺/T, CD3⁺CD8⁺/T, and Treg cell/CD4⁺ T lymphocyte ratios were significantly reduced after ROSC. NK cells, which are special innate immune cells with cytotoxic functions similar to CD3⁺CD8⁺ T lymphocytes, mainly distinguish infected and stressed cells from healthy cells and eliminate intracellular infection and dysfunctional cells. [29, 30] T lymphocytes are also crucial because they function as adaptive immune cells to control and eliminate the infection. [29] Moreover, B and T lymphocytes mediate humoral and cellular immunity, respectively. This study was performed earlier and involved a more comprehensive assessment of the immune system of patients who experienced CA. Our findings more substantially supported the rapid emergence of immune dysfunction in these patients after ROSC than in previous reports. The binding of GCs to GR inside different peripheral blood mononuclear cells (PBMC) leads to changes in the ability of cells to regulate apoptosis, proliferation, and activity, and GC-GR complexes limit the transcription (trans-repression) of

inflammatory genes, including those encoding for proinflammatory cytokines.[31, 32]
This study is the first to explore GR expression in circulating immune cells in patients
who experienced CA after ROSC. We observed that GR expression in B and T

267 lymphocytes, NK cells, and CD3⁺CD8⁺ T lymphocytes decreased significantly in

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268	patients who experienced CA, whereas the percentage of GR ⁺ Treg cells and
269	CD3+CD4+ T lymphocytes decreased slightly. Moreover, we observed a more
270	significant decrease in the MFI of GR expression in Treg cells and CD3 ⁺ CD4 ⁺ T
271	lymphocytes but not in the percentage of GR expression. Previous studies have found
272	decreased expression of GRs in peripheral polymorphonuclear cells in critically ill
273	patients, [21] and antagonism to GRs aggravates viral and bacterial infections. [33]
274	GCs induced upon infections help to maintain homeostasis and mitigate the life-
275	threatening impact of sepsis on the host.[31] Although studies have reported that the
276	use of GCs during and after CPR seems to confer benefits concerning ROSC rates and
277	long-term survival, the evidence is scant. [13,18,34,35] Since cortisol signaling is
278	mediated by GRs, we hypothesized that the differential responses of CA patients to GC
279	may be related to their levels of GR expression. This study suggests that the decrease
280	in intracellular GR expression in patients who experienced CA is one of the causes of
281	GC resistance due to insufficient binding of GRs and GCs, GC insensitivity, and the
282	inability of GCs to exert anti-inflammatory and immunosuppressive effects effectively.
283	These findings may also explain why different results regarding the clinical application
284	of GCs have been reported previously. Furthermore, it is vital to measure GR levels as
285	sufficient expression of GR is essential for mediating adequate GC effects during and
286	after CPR.

We also found that the total plasma cortisol levels were significantly higher in patients who experienced CA, but ACTH levels were not. High levels of inflammatory cytokines inhibit ACTH release. [18] During critical illness, the body does not

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290	sufficiently metabolize cortisol. [36] In addition, the continuous increase in plasma
291	cortisol levels may trigger the negative feedback pathway of the hypothalamic-
292	pituitary-adrenal axis, inhibiting the release of ACTH and cortisol and eventually
293	leading to adrenal insufficiency [37]. These factors may explain the opposite trends of
294	plasma ACTH and cortisol levels in the patients included in this study and who
295	experienced CA. Notably, this result suggests that low GR expression levels are not
296	matched by high plasma total cortisol levels in patients who experienced CA. The
297	dissociation between low GR expression and high cortisol implies an abnormal stress
298	response. [38] Previous studies have reported that GR-action was clearly suppressed
299	throughout critical illness; GR resistance could not be overcome by further increasing
300	glucocorticoid availability.[21,39,40] Adequate GR levels and function are also
301	required for normal GC function, which may explain differences in the responsiveness
302	of cardiac arrest patients to exogenous steroid administration or endogenous cortisol
303	secretion. Thus, actual GR levels cannot be reflected by measuring total cortisol levels
304	alone. Therefore, the GR level should be considered when applying personalized GC
305	therapy. The determination of GR expression might help to screen those who might
306	respond better to glucocorticoid prescription.

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308 Limitations

Our study has several limitations. First, to assess changes, we only enrolled patients
who experienced CA and had signs of systemic ischemic hypoxia, such as GCS <8 after
ROSC. The patients were not stratified by age, sex, the occurrence of comorbidities, or

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mild systemic ischemic hypoxia. Second, since this was a preliminary observational study, we observed only early changes. A more relevant control group and dynamic observations obtained over a longer duration would be helpful to understand the significance of GR expression in evolving immunity during the clinical course of CA after ROSC. Third, the samples used in this study were from clinical laboratories; thus, plasma total cortisol and ACTH in the samples were at risk of degradation before we collected the samples. Finally, we did not discuss the changes in and roles of GR isoforms, free cortisol, and corticosteroid-binding globulin. Therefore, future studies on these aspects are warranted to better understand the immunosuppressive effects of ROSC among patients who experienced CA. In conclusion, this study revealed that GR expression, cell counts and ratios rapidly decreased, whereas plasma total cortisol levels increased, in the early period after ROSC among CA patients. These findings may provide important information about

325 GR expression levels and function, and immunosuppressive status in these patients.

Acknowledgments: We thank all the patients and their families who were enrolled in
 this study and colleagues from the emergency department who provided support. And
 we are grateful for the efforts of the staff for ongoing resuscitation in hospitals.

Contributorship statement: CL designed the study and reviewed the manuscript.
YNY searched the literature and contributed to the experimental studies, data analysis,
and manuscript writing. ZRT, CCH, and LA collected and analyzed data. JBL and MRX

333	helped with the statistical analyses. All authors have read and approved the final
334	manuscript.
335	Competing interests: All authors declare no competing interest associated with this
336	project.
337	Funding: This research received no specific grant from any funding agency in public,
338	commercial or not-for-profit sectors.
339	Provenance and peer review: Not commissioned; externally peer-reviewed.
340	Data sharing statement: All data relevant to the study are included in the article or
341	uploaded as supplementary information. Due to privacy and ethical concerns, data can
342	not be shared.
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344	Ethics statements
345	Patient consent for publication: Not applicable.
346	Ethics approval: This study was approved by the Medical Ethics Committee of Beijing
347	Chaoyang Hospital (2013-KE-1). Because CA is sudden and life-threatening, the
348	consent was usually obtained orally from relatives or bystanders and in writing with
349	some delay from relatives or bystanders after successful resuscitation.
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480 Figure legends

Fig. 1. Changes in circulatory T and B lymphocyte, NK cell, and Treg cell counts,
CD3⁺CD4⁺/T, CD3⁺CD8⁺/T, and Treg/CD4⁺T lymphocyte ratios between the healthy
control group and CA group. The CA group showed significant differences compared
with the healthy control group (P<0.001). CA, cardiac arrest; CD, cluster-of-
differentiation; NK, natural killer; Treg, regulatory T.

Fig. 2. Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells
in the healthy control group and CA group. The CA group showed significant
differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD,
cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC,
return of spontaneous circulation; Treg, regulatory T.

Fig. 3. (A, B) Plasma total cortisol and ACTH levels (the natural logarithmic
conversion values after adding 1) after ROSC in the healthy control group and CA
group. (C, D) Plasma total cortisol and ACTH levels in survivors and non-survivors
after ROSC. The CA group showed significant differences compared with the healthy
control group (P<0.05). ACTH, adrenocorticotrophic hormone; CA, cardiac arrest;
ROSC, return of spontaneous circulation.

P = 0.001

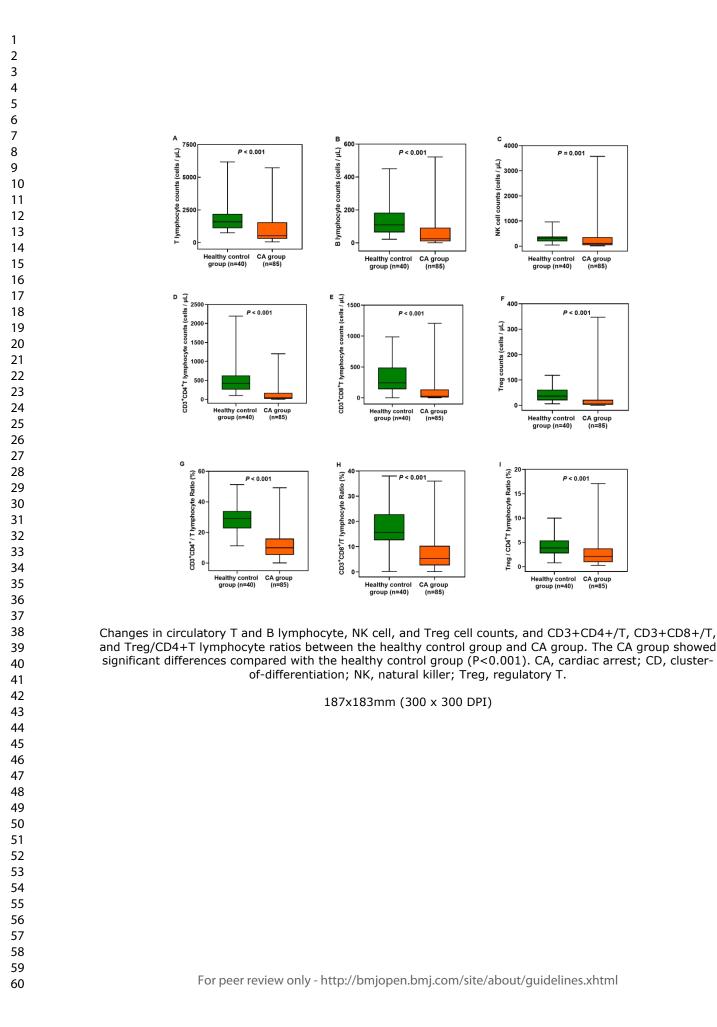
CA group (n=85)

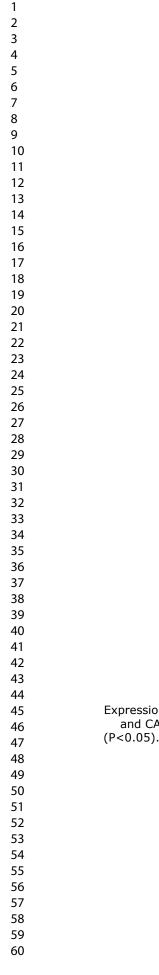
CA group (n=85)

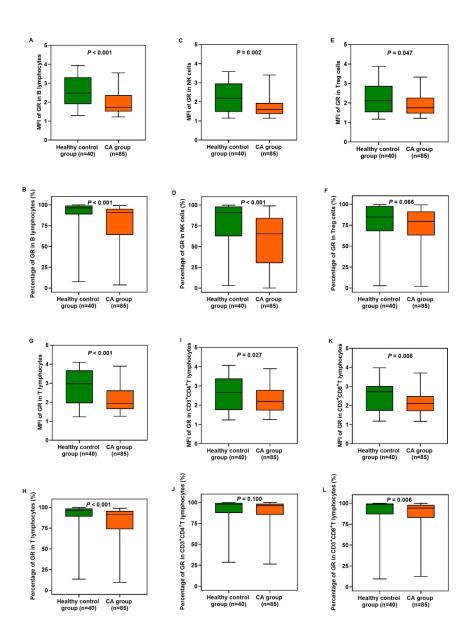
CA group (n=85)

P < 0.001

P < 0.001

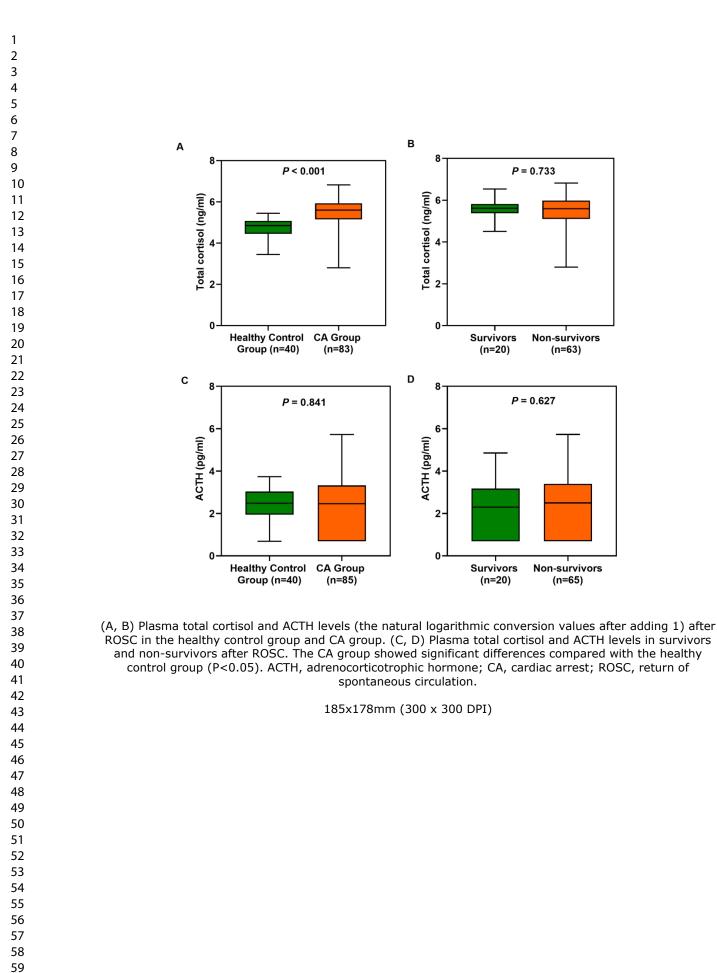






Expression of GRs in circulatory T and B lymphocytes, NK cells, and Treg cells in the healthy control group and CA group. The CA group showed significant differences compared with the healthy control group (P<0.05). CA, cardiac arrest; CD, cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; ROSC, return of spontaneous circulation; Treg, regulatory T.

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Electronic supplemental material

Glucocorticoid receptor expression in patients with cardiac arrest in the early period after the return of spontaneous circulation: A prospective observational single-center study

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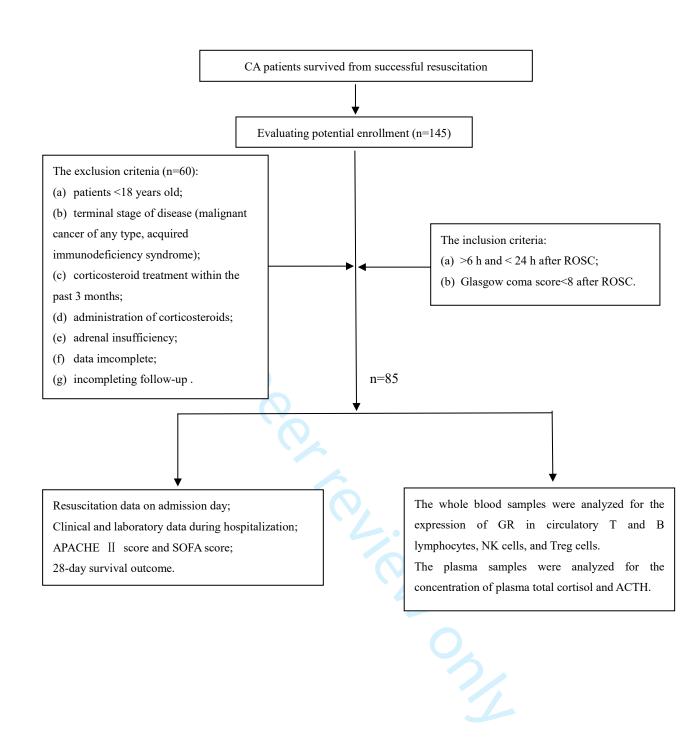
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Supplemental Figure 1. The flow chart of the study.

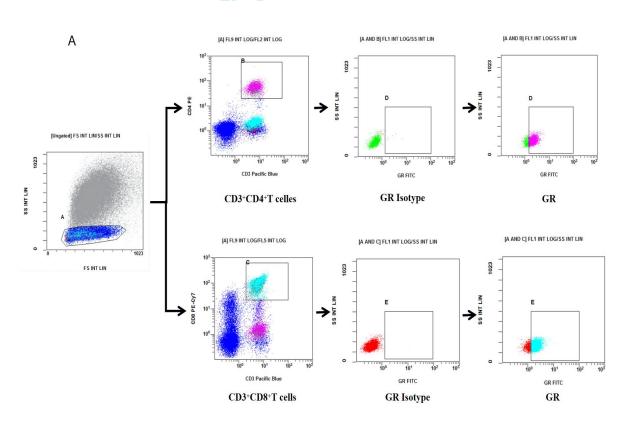
Abbreviations: CA, cardiac arrest; ROSC, return of spontaneous circulation; APACHE II, acute physiology and chronic health evaluation II; SOFA, sequential organ failure assessment; GR, glucocorticoid receptor; Treg, regulatory T; ACTH, adrenocorticotrophic hormone.

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Supplemental Figure 2. Representative plots and gating strategies for analyzing glucocorticoid receptor (GR) in the whole blood.

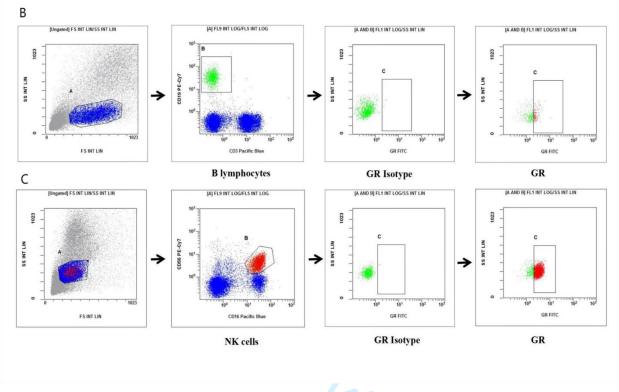
GR expression levels were determined on T cells, B cells, NK cells, and T regulatory (Treg) cells. Single cells were gated from all cellular events (FSC/SSC gate). B cells were identified as CD3⁻CD19⁺ cells. NK cells were identified as CD16⁺56⁺ cells. T cells were identified as CD3⁺CD4⁺ T cells and CD3⁺CD8⁺ T cells. Treg cells were identified as CD4⁺CD25^{high}CD127^{low}.

A. Expression of GR on T cells



B. Expression of GR on B cells

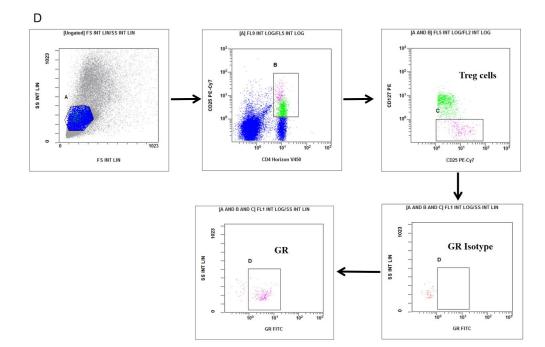
C. Expression of GR on NK cells



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D. Expression of GR on Treg cells





Source

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Bio-Rad^b

Bio-Rad

Antigen	Catalog Number	Fluorescein
CD3	558117	Pacific Blue
CD4	555347	PE
CD4	560345	Horizon V4
CD8	557746	PE-Cy7
CD19	557835	PE-Cy7
CD16	558122	Pacific Blue
CD56	557747	PE-Cy7
CD25	557741	PE-Cy7
CD127	557938	РЕ
GR	MCA2469F	FITC
Mouse IgG1 Isotype	MCA928F	FITC
Mouse IgG1,к Isotype	557872	PE-Cy7
Mouse IgG1,к Isotype	554680	PE
Mouse IgG1,к Isotype	558120	Pacific Blue

acific Blue **BD** Pharmingen otec, Oxford, UK. coerythrin; FITC, fluorescein isothiocyanate; GR, glucocorticoid

receptor; Ig: immunoglobulin.

	Survivors (n=20)	Non-survivors (n=65)
Age (years), median [IQR]	59.0 (53.3, 72.8)	66.0 (59.0, 75.5)
Male/Female (n)	12/8	46/19
Cardiac arrest cause (n, %)		
Cardiac	10 (50.0%)	24 (36.9%)
Non-Cardiac	10 (50.0%)	41 (63.1%)
Initial resuscitation		
Time to ROSC (min), median [IQR]	15.0 (7.3, 26.0)	20.0 (15.0, 30.0)
Adrenaline (mg), median [IQR]	1.0 (0.0, 3.0)	2.0 (0.0, 5.0)
Initial rhythm VF/VT, n (%)	11 (55.0%)	19 (29.2%)
MAP (mmHg), median [IQR]	89.9 (70.5, 104.9)	70.7 (50.0, 93.5)
White cell count (×10 ⁹ /L), median [IQR]	12.40 (6.98, 18.76)	13.80 (11.67, 18.20)
Lactate (mmol/L), median [IQR]	3.50 (1.33, 7.05)	7.50 (3.80, 11.20)
APACHE II score, mean±SD	27.8±6.6	34.4±5.6
SOFA score, median [IQR]	9.0 (7.3, 11.8)	12.0 (9.0, 15.0)

Supplemental Table 2. Characteristics of CA survivors and non-survivors on admission.

Data are presented as mean±SD or interquartile range (IQR) as appropriate. Abbreviations: ROSC: return of spontaneous circulation; VF: ventricular fibrillation; VT: ventricular tachycardia; MAP: mean arterial pressure; APACHE II: acute physiology and chronic health evaluation; SOFA: sequential organ failure assessment.

Supplemental Table 3. The flow cytometry results of cell counts and ratios of the healthy control group and successful resuscitation group

	Healthy Control	Successful	Z-value	<i>P</i> -value
	Group (n=40)	Resuscitation Group		
		(n=85)		
T lymphocyte count (cells /µL)	1586.0 (1101.5, 2192.5)	514.0 (287.5, 1555.0)	-4.515	<0.001
NK cell count (/µL)	311.5 (191.0, 378.8)	101.0 (36.0, 351.5)	-3.332	0.001
B lymphocyte count (/μL)	109.3 (63.7, 183.3)	25.7 (9.4, 92.3)	-5.076	<0.001
Treg count (/µL)	0.259 (0.095, 0.516)	0.233 (0.135, 0.488)	-5.518	<0.001
Treg / CD4 ⁺ T lymphocyte Ratio	0.039 (0.028, 0.054)	0.021 (0.010, 0.038)	-4.418	<0.001
$CD3^+CD4^+ T$ lymphocyte count (/µL)	421.7 (258.6, 627.4)	38.9 (17.6, 168.3)	-6.256	<0.001
CD3 ⁺ CD4 ⁺ / T lymphocyte Ratio	0.292 (0.227, 0.340)	0.100 (0.054, 0.160)	-7.066	<0.001
CD3 ⁺ CD8 ⁺ T lymphocyte count (/µL)	241.1 (139.5, 488.6)	26.3 (7.2, 135.9)	-5.287	<0.001
CD3 ⁺ CD8 ⁺ / T lymphocyte Ratio	0.157 (0.126, 0.229)	0.053 (0.026, 0.104)	-5.719	< 0.001

All the data in Supplemental table 3 are represented as the median [IQR]; IQR: Interquartile Range; CD: cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

Supplemental Table 4. The flow cytometry results of cell counts and ratios of the CA patients on admission based on

Survivors (n=20)	Non-survivors	Z-value	<i>P</i> -value
	(n=65)		
502.0 (353.8, 1199.8)	514.0 (282.5, 1891.0)	-0.186	0.852
167.0 (29.8, 309.3)	100.0 (36.0, 404.0)	-0.218	0.828
38.6 (15.7, 103.5)	19.2 (7.1, 65.7)	-0.632	0.527
0.318 (0.145, 0.552)	0.212 (0.128, 0.479)	-0.611	0.396
0.025 (0.009, 0.043)	0.021 (0.010, 0.034)	-0.498	0.619
55.1 (32.4, 228.0)	38.0 (16.0, 168.1)	-0.850	0.396
0.118 (0.070, 0.236)	0.097 (0.049, 0.142)	-1.565	0.118
25.4 (12.5, 96.2)	26.3 (6.3, 138.8)	-0.021	0.983
0.054 (0.033, 0.104)	0.053 (0.025, 0.104)	-0.187	0.852
	167.0 (29.8, 309.3) 38.6 (15.7, 103.5) 0.318 (0.145, 0.552) 0.025 (0.009, 0.043) 55.1 (32.4, 228.0) 0.118 (0.070, 0.236) 25.4 (12.5, 96.2)	502.0 (353.8, 1199.8) 514.0 (282.5, 1891.0) 167.0 (29.8, 309.3) 100.0 (36.0, 404.0) 38.6 (15.7, 103.5) 19.2 (7.1, 65.7) 0.318 (0.145, 0.552) 0.212 (0.128, 0.479) 0.025 (0.009, 0.043) 0.021 (0.010, 0.034) 55.1 (32.4, 228.0) 38.0 (16.0, 168.1) 0.118 (0.070, 0.236) 0.097 (0.049, 0.142) 25.4 (12.5, 96.2) 26.3 (6.3, 138.8)	502.0 (353.8, 1199.8)514.0 (282.5, 1891.0)-0.186167.0 (29.8, 309.3)100.0 (36.0, 404.0)-0.21838.6 (15.7, 103.5)19.2 (7.1, 65.7)-0.6320.318 (0.145, 0.552)0.212 (0.128, 0.479)-0.6110.025 (0.009, 0.043)0.021 (0.010, 0.034)-0.49855.1 (32.4, 228.0)38.0 (16.0, 168.1)-0.8500.118 (0.070, 0.236)0.097 (0.049, 0.142)-1.56525.4 (12.5, 96.2)26.3 (6.3, 138.8)-0.021

All the data in Supplemental table 4 are represented as the median [IQR]; IQR: Interquartile Range; CD: cluster-of-differentiation; GR, glucocorticoid receptor; NK, natural killer; Treg, regulatory T.

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59 60 Supplemental Table 5. The flow cytometry results of GR expression in the CA group and successful resuscitation group.

Healthy Control	Successful	Z-value	<i>P</i> -value
Group (n=40)	Resuscitation Group		
	(n=85)		
0.963 (0.885, 0.992)	0.896 (0.605, 0.949)	-3.742	<0.001
2.48 (1.91, 3.31)	1.73 (1.50, 2.37)	-3.980	< 0.001
0.964 (0.889, 0.986)	0.900 (0.703, 0.955)	-3.755	< 0.001
2.98(1.95, 3.68)	1.92 (1.36, 1.99)	-3.853	< 0.001
0.907 (0.624, 0.983)	0.611 (0.306, 0.840)	-3.792	< 0.001
2.19 (1.48, 2.96)	1.60 (1.36, 1.99)	-3.171	0.002
0.848 (0.680, 0.978)	0.784 (0.589, 0.911)	-1.837	0.066
2.12 (1.53, 2.88)	1.76 (1.44, 2.30)	-1.990	0.047
0.980 (0.874, 0.996)	0.957 (0.824, 0.985)	-2.204	0.100
2.65 (1.75, 3.38)	2.17 (1.70, 2.92)	-1.646	0.027
0.986 (0.868, 0.996)	0.938 (0.823, 0.979)	-2.758	0.006
2.73 (1.73, 3.02)	2.10 (1.68, 2.54)	-2.668	0.008
	Group (n=40) 0.963 (0.885, 0.992) 2.48 (1.91, 3.31) 0.964 (0.889, 0.986) 2.98(1.95, 3.68) 0.907 (0.624, 0.983) 2.19 (1.48, 2.96) 0.848 (0.680, 0.978) 2.12 (1.53, 2.88) 0.980 (0.874, 0.996) 2.65 (1.75, 3.38) 0.986 (0.868, 0.996)	Group (n=40) Resuscitation Group 0.963 (0.885, 0.992) 0.896 (0.605, 0.949) 0.963 (0.885, 0.992) 0.896 (0.605, 0.949) 2.48 (1.91, 3.31) 1.73 (1.50, 2.37) 0.964 (0.889, 0.986) 0.900 (0.703, 0.955) 2.98(1.95, 3.68) 1.92 (1.36, 1.99) 0.907 (0.624, 0.983) 0.611 (0.306, 0.840) 0.907 (0.624, 0.983) 0.611 (0.306, 0.840) 2.19 (1.48, 2.96) 1.60 (1.36, 1.99) 0.848 (0.680, 0.978) 0.784 (0.589, 0.911) 2.12 (1.53, 2.88) 1.76 (1.44, 2.30) 0.980 (0.874, 0.996) 0.957 (0.824, 0.985) 2.65 (1.75, 3.38) 2.17 (1.70, 2.92) 0.986 (0.868, 0.996) 0.938 (0.823, 0.979)	Group (n=40) Resuscitation Group (n=85) 0.963 (0.885, 0.992) 0.896 (0.605, 0.949) -3.742 2.48 (1.91, 3.31) 1.73 (1.50, 2.37) -3.980 0.964 (0.889, 0.986) 0.900 (0.703, 0.955) -3.755 2.98(1.95, 3.68) 1.92 (1.36, 1.99) -3.853 0.907 (0.624, 0.983) 0.611 (0.306, 0.840) -3.792 2.19 (1.48, 2.96) 1.60 (1.36, 1.99) -3.171 0.848 (0.680, 0.978) 0.784 (0.589, 0.911) -1.837 2.12 (1.53, 2.88) 1.76 (1.44, 2.30) -1.990 0.980 (0.874, 0.996) 0.957 (0.824, 0.985) -2.204 2.65 (1.75, 3.38) 2.17 (1.70, 2.92) -1.646 0.986 (0.868, 0.996) 0.938 (0.823, 0.979) -2.758

All the data in Supplemental table 5 are represented as the median [IQR]. Abbreviations: IQR, interquartile range; CD, cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, Glucocorticoid receptor; MFI, mean fluorescence intensity.

	Survivors	Non-survivors	Z-value	P-value
	(n=20)	(n=65)		
Percentage of GR on B lymphocytes	0.904 (0.595, 0.976)	0.906 (0.657, 0.946)	-0.787	0.431
MFI of GR on B lymphocytes	1.92 (1.52, 2.54)	1.72 (1.51, 2.31)	-0.881	0.378
Percentage of GR on T lymphocytes	0.899 (0.778, 0.969)	0.913 (0.692, 0.951)	-1.057	0.291
MFI of GR on T lymphocytes	2.05 (1.67, 2.83)	1.91 (1.64, 2.46)	-1.031	0.303
Percentage of GR on NK cells	0.717 (0.292, 0.886)	0.556 (0.302, 0.823)	-0.756	0.449
MFI of GR on NK cells	1.54 (1.37, 2.09)	1.61 (1.34, 1.87)	-0.565	0.572
Percentage of GR on Tregs	0.780 (0.667, 0.849)	0.799 (0.576, 0.923)	-0.440	0.660
MFI of GR on Tregs	1.61 (1.48, 2.30)	1.77 (1.45, 2.27)	-0.005	0.996
Percentage of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	0.975 (0.876, 0.985)	0.957 (0.845, 0.987)	-0.617	0.538
MFI of GR on CD3 ⁺ CD4 ⁺ T lymphocytes	2.08 (1.72, 3.35)	2.22 (1.71, 2.69)	-0.865	0.387
Percentage of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	0.963 (0.816, 0.977)	0.938 (0.834, 0.980)	-0.254	0.800
MFI of GR on CD3 ⁺ CD8 ⁺ T lymphocytes	2.08 (1.68, 3.10)	2.11(1.71, 2.46)	-0.653	0.514

All the data in Supplemental table 6 are represented as the median [IQR]. Abbreviations: IQR, Interquartile Range; CD, Cluster-of-differentiation; NK, natural killer; Treg, regulatory T; GR, glucocorticoid receptor; MFI, mean fluorescence intensity.

Supplemental Table 6. The flow cytometry results of GR expression in the survivors and non-survivors.

STROBE Statement—checklist of items that should be included in reports of observational studies

	Item No	Recommendation	Page No
Title and abstract	1	(a) Indicate the study's design with a commonly used term in	1
		the title or the abstract	
		(b) Provide in the abstract an informative and balanced	3
		summary of what was done and what was found	
Introduction			
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	3-5
Objectives	3	State specific objectives, including any prespecified hypotheses	5
Methods			1
Study design	4	Present key elements of study design early in the paper	Supplemental Figure 1
Setting	5	Describe the setting, locations, and relevant dates, including	5-9,
-		periods of recruitment, exposure, follow-up, and data	Supplemental
		collection	Figure 1
Participants	6	(a) Cohort study—Give the eligibility criteria, and the sources	5,6,8,9
		and methods of selection of participants. Describe methods of	
		follow-up	
		Case-control study—Give the eligibility criteria, and the	
		sources and methods of case ascertainment and control	
		selection. Give the rationale for the choice of cases and	
		controls	
		Cross-sectional study—Give the eligibility criteria, and the	
		sources and methods of selection of participants	
		(b) Cohort study—For matched studies, give matching criteria	5
		and number of exposed and unexposed	
		Case-control study—For matched studies, give matching	
		criteria and the number of controls per case	
Variables	7	Clearly define all outcomes, exposures, predictors, potential	5, 6, 8,
		confounders, and effect modifiers. Give diagnostic criteria, if	Supplemental
		applicable	Figure 1
Data sources/	8*	For each variable of interest, give sources of data and details	6, 8
measurement	-	of methods of assessment (measurement). Describe	
		comparability of assessment methods if there is more than one	
		group	
Bias	9	Describe any efforts to address potential sources of bias	6-8
Study size	10	Explain how the study size was arrived at	Supplemental
<u>,</u>	-	· · · · · · · · · · · · · · · · · · ·	Figure 1
Quantitative variables	11	Explain how quantitative variables were handled in the	6-8
		analyses. If applicable, describe which groupings were chosen	
		and why	
Statistical methods	12	(<i>a</i>) Describe all statistical methods, including those used to control for confounding	8, 11

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(b) Describe any methods used to examine subgroups and interactions	N/A
(c) Explain how missing data were addressed	8, 11
(<i>d</i>) <i>Cohort study</i> —If applicable, explain how loss to follow-up was addressed	11
<i>Case-control study</i> —If applicable, explain how matching of cases and controls was addressed	
Cross-sectional study—If applicable, describe analytical	
methods taking account of sampling strategy	
(\underline{e}) Describe any sensitivity analyses	

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Participants	13	(a) Report numbers of individuals at each stage of study—eg numbers	9,
	*	potentially eligible, examined for eligibility, confirmed eligible, included in	Supplem
		the study, completing follow-up, and analysed	1 Figure 1
		(b) Give reasons for non-participation at each stage	9,12,
			Suppleme
			l Figure 1
		(c) Consider use of a flow diagram	Suppleme
			1 Figure 1
Descriptive	14	(a) Give characteristics of study participants (eg demographic, clinical,	9
data	*	social) and information on exposures and potential confounders	
		(b) Indicate number of participants with missing data for each variable of	Suppleme
		interest	1 Figure 1
		(c) <i>Cohort study</i> —Summarise follow-up time (eg, average and total amount)	8
Outcome data	15	Cohort study—Report numbers of outcome events or summary measures	9-12
	*	over time	
		Case-control study-Report numbers in each exposure category, or summary	
		measures of exposure	
		Cross-sectional study—Report numbers of outcome events or summary	
		measures	
Main results	16	(a) Give unadjusted estimates and, if applicable, confounder-adjusted	9-12,
		estimates and their precision (eg, 95% confidence interval). Make clear	Electronic
		which confounders were adjusted for and why they were included	suppleme
			material
		(b) Report category boundaries when continuous variables were categorized	
		(c) If relevant, consider translating estimates of relative risk into absolute	
		risk for a meaningful time period	
Other analyses	17	Report other analyses done-eg analyses of subgroups and interactions, and	N/A
		sensitivity analyses	
Discussion			
Key results	18	Summarise key results with reference to study objectives	12
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias	15
		or imprecision. Discuss both direction and magnitude of any potential bias	
Interpretation	20	Give a cautious overall interpretation of results considering objectives,	12-14
		limitations, multiplicity of analyses, results from similar studies, and other	
		relevant evidence	
Generalisabilit	21	Discuss the generalisability (external validity) of the study results	15
у			
Other informat	ion		
Funding	22	Give the source of funding and the role of the funders for the present study	17
		and, if applicable, for the original study on which the present article is based	

*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies.

Note: An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely

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available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.

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