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Research Article

T Lymphocytes and Cytokines: Earlier time to peak indicates better prognosis in COVID-19 patients?

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Abstract

Aims: To explore the changes of cytokines and T lymphocyte subgroups in COVID-19 patients, to find the turning point to severe clinical course of patients, to timely intervene, and to reduce the severity.

Methods: Thirty-nine patients with COVID-19 admitted to Third People's Hospital of Kunming from January 31, 2020 to March 2, 2020 were divided into 3 groups: mild, common and severe/critical according to 'Clinical Protocol for COVID-19, 8th Edition' (National Health Commission of People's Republic of China). The changes of serum T lymphocyte subgroups and cytokines level during hospitalization were monitored and statistically analyzed. The difference was considered statistically significant when $p < 0.05$.

Results: 39 patients with COVID-19 were 41.51±19.03 years old, 18 of them were male (46.2%). 8 patients (20.5%) were classified as mild group, 10 patients (25.6%) were classified as common group, 21 patients (53.8%) were classified as severe/critical group. The hospitalization days were 17.51±6.19 days, and the nucleic acid seroconversion days from positive to negative were 11.41±6.77 days. Comparison of T lymphocyte subgroups in patients with different clinical classifications showed that T cell count and CD4/CD8⁺T cell counts were different among different clinical classifications ($p < 0.05$). In pairwise comparison, the count of lymphocyte subgroups in severe/critical group patients was significantly lower than that in mild and common groups. Intra-group Comparison showed that, IL-2, IL-8, IL-12p70 and IL-5 examinations had significant statistical differences at different examination time points, but the effects of the above indicators on clinical classification would not change with time. According to develop trends, lymphocytes increased in patients of three clinical groups after treatment, the increase trends were obvious especially in severe/critical group patients. The OR value calculated by Logistics model showed that the probability of severe/critical disease was 1.276 times higher for each day increase of T cell time to peak (95%CI: 1.039-1.569), $\chi^2 = 5.379$, $p = 0.02$. The probability of severe/critical disease was 1.408 times (95%CI: 1.066-1.859) with each day increase of IL1 β time to peak, $\chi^2 = 5.824$, $p = 0.016$. Independent sample *t* test showed that in CD4⁺T cells, IL-1 β , IL-2, IL-8, IL-12p70, IL-17A, IFN- γ , TNF- α and other detection indicators, the time to peak of the above indicators in the fast seroconversion group was significantly shorter than that in slow seroconversion group.

Conclusions: The early peak, timely decrease of cytokines and T cell counts in COVID-19 patients has a positive effect on prognosis. Cytokines and T cell counts should be closely monitored in COVID-19 patients.



Since December 2019, a novel coronavirus was first discovered in Wuhan, China, new Coronavirus Disease (COVID-19) has been a worldwide disease. Up to March, there have been more than 128 million confirmed cases of COVID-19 globally, the mortality rate is about 2.2% [1]. At the beginning of the disease, acute respiratory symptoms are commonly seen, while some patients would progress to Acute Respiratory Distress Syndrome (ARDS), in some of them, respiratory failure may appear. With the development of COVID-19 immunological research, many researchers have elaborated on the immunological characteristics of COVID-19. Huang, et al. compared the clinical manifestations and laboratory indicators of COVID-19 ICU patients and non-ICU patients, and found that the serum levels of serum cytokines such as IL-2, IL-7, IL-10 and TNF- α in ICU patients were significantly higher than those in non-ICU patients. The serum levels of CD4⁺T cells, CD8⁺T cells and Natural Killer (NK) cells in critical patients were lower than those in mild patients, suggesting that critical patients may have significant immune cell depletion [1]. In a variety of viral diseases, it has been found that the imbalance of cytokines and the decrease or exhaustion of T cell count play important roles in the severity of the disease [2]. Above studies, however, were merely descriptive studies on cytokines and lymphocytes of COVID-19 patients. Few involved the time point of transferring from mild disease to severe disease. Our study aims to explore the relationship between cytokines and T lymphocyte subgroups, to find the turning point of mild to severe clinical course, to block the severity of disease as early as possible, to intervene in time, and to explore the clinical possibility of reducing severe clinical course.

Materials and methods

Patients

The study was approved by the ethics committee of the Third People's Hospital of Kunming, following the principles of Helsinki declaration. 39 patients admitted in Third People's Hospital of Kunming with COVID-19 were collected since January 31st to March 2nd retrospectively. Informed consent were signed by all the 39 patients or their guardians.

Clinical classifications of disease severity

All the patients were diagnosed according to "Clinical Protocol for COVID-19, 8th Edition", National Health Commission of People's Republic of China [3]. They were classified as: (1) Mild: mild clinical symptoms, no pneumonia in imaging; (2) Common: fever, respiratory symptoms, imaging findings of pneumonia; (3) Severe: any of the following: (1) Shortness of breath, respiratory rate (RR) \geq 30 times/min; (2) in resting state, oxygen saturation \leq 93%; (3) PaO₂/FiO₂ \leq 300 mmHg (1 mmHg = 0.133 kPa); High altitudes areas (over 1000 m) should correct for PaO₂/FiO₂ according to the following formula: PaO₂/FiO₂ \times [760/atmospheric pressure (mmHg)]. 44 progressive exacerbation of clinical symptoms and pulmonary imaging showing significant progression of > 50% within 24- 48 hours. (4) Critical: One of the following conditions: A. Respiratory failure requiring mechanical ventilation; B. Shock; C. complicated with other organ failures required for ICU monitoring and treatments.

Data collection

The basic data of 39 COVID-19 patients, including age, gender, diagnosis, days of hospital stay, and days of nucleic acid seroconversion (from positive to negative) were collected. T cell subgroups and cytokine values were dynamically monitored at different time points during patients' hospitalization. All the laboratory tests were carried out by Clinical Laboratory of the Third People's Hospital of Kunming.

Test methods

BD company FACSCANTO II flow cytometer was used for flow fluorescence detection. The T-lymphocyte subgroups detection reagent was the BD Mult-Test IMK Kit by BD Company, and the cytokine detection reagent was the 12 Cytokine Detection Kits provided by Jiangxi Saiji Biotechnology Co., Ltd. Bio-Rad Real-time fluorescence quantitative PCR was used for 2019 novel coronavirus (2019-nCoV) RNA detection. The reagent was 2019-nCoV nucleic acid detection kit by Shanghai BioGerm Medical Technology Co., Ltd.

Statistic analysis

GraphPad Prism 8.02 was used for plotting, and IBM SPSS Data Editor Version 23.0 was used for statistical analysis. Variables of normal distribution are represented by mean \pm standard deviation, the variables of non-normal distribution are represented by median, and the classified variables are represented by frequency and rate (%). Repeated measurement ANOVA was used. If the repeated measurement ANOVA did not satisfy the spherical symmetry test, the correction results of Greenhouse-Geisser were used. The difference was considered statistically significant when $p < 0.05$.

Results

Basic characteristics of patients

Among the 39 patients with COVID-19 in our hospital, most of them were female, with the age ranging from 3-79 years old. There were 8 cases of mild type, 10 cases of common type and 21 cases of severe and critical type. Days of nucleic acid seroconversion is the days from admission to the time of nucleic acid negative collected by throat swab (Table 1).

Results of sphericity test

Results of Mauchly's Test of Sphericity for all indicators in the COVID-19 patient's laboratory (Table 2). The results of Mauchly's Test of Sphericity showed that $p > 0.05$, which

Table 1: Basic Characteristics of Patients.

Gender (male%)		18 (46.2%)
Age (years old)		3-79 (41.51 \pm 19.03)
Clinical Classifications	Mild Group	8(20.5%)
	Common Group	10(25.6%)
	Severe and Critical Group	21(53.8%)
Days in Hospital		6-31(17.51 \pm 6.19)
Days of Nucleic Acid Seroconversion		1-24(11.41 \pm 6.77)



satisfied the hypothesis of spherical symmetry. When $p < 0.05$ dissatisfied with the hypothesis of spherical symmetry, Greenhouse-Geisser's correction results were used.

Results of analysis of repeated measures ANOVA for different clinical classifications of patients' laboratory indicators.

I. Intergroup comparison: Analysis of variance ($p < 0.05$) suggested that there were differences in T cell count and CD4/CD8⁺T cell count among different clinical classifications (Table 3). Further pairwise comparison results are shown in Table 4.

T cell count is significantly lower in severe/critical group compared with in mild ($p = 0.033$) and common group ($p = 0.001$);

CD4⁺T cell count is significantly lower in severe/critical group compared with in common group ($p = 0.003$);

CD8⁺T cell count is significantly lower in severe/critical group compared with in mild ($p = 0.003$) and common group ($p = 0.003$);

It can be seen that the total number of T lymphocyte, CD4⁺T and CD8⁺T cell counts in severe/critical patients were remarkable lower than those in mild and normal patients.

II. Intra-group Comparison: Time for IL-2, IL-8, IL-12p70 and IL-5 $p < 0.05$, Time*Classification $p > 0.05$. It suggested that IL-2, IL-8, IL-12p70 and IL-5 examinations had significant statistical differences at different examination time points, but the effects of the above indicators on classification would not change with time.

Development trends for CD3+ Cell /CD4+/CD8+ T Cell in different clinical classifications

The horizontal axis shows the times of examinations

Table 2: Mauchly's Test of Sphericity Results of Patients' Laboratory Indicators.

Indicators	Mauchly's W	Appro Chi-square	df	sig.	Greenhouse-Geisser
T Cell	0.606	14.376	5	0.013	0.798
CD4 ⁺ T Cell	0.727	9.176	5	0.102	0.827
CD8 ⁺ T Cell	0.661	11.898	5	0.036	0.822
IL-1	0.073	20.200	5	0.001	0.638
IL-2	0.421	6.687	5	0.248	0.720
IL-4	0.547	4.665	5	0.461	0.785
IL-5	0.407	6.935	5	0.229	0.672
IL-6	0.079	19.605	5	0.002	0.430
IL-8	0.126	15.991	5	0.007	0.658
IL-10	0.658	3.227	5	0.667	0.810
IL-12p70	0.427	6.579	5	0.257	0.741
IL-17A	0.291	9.527	5	0.092	0.633
IFN- γ	0.361	7.868	5	0.167	0.744
IFN- α	0.000	62.175	5	0.000	0.339
TNF- α	0.271	10.078	5	0.075	0.575

during the patients' hospitalization, and the vertical axis shows the levels of T lymphocyte, CD4⁺ and CD8⁺T cell count. As can be seen from Figures 1,2, T cell count in mild group and common group patients increased slowly over time, while those in severe/critical group increased significantly over time, especially in the third and fourth examinations compared with the second examination. CD4⁺T cell count in mild group and common group patients increased slowly with time, and the trend was relatively gentle. CD4⁺T cell count in severe/critical patients increased significantly with time, and the elevation was more obvious in the third and fourth examinations than in the second examination. CD8⁺T cell count also increased significantly in severe/critical patients compared with mild and common patients during treatment.

Results of analysis of single factor for different clinical classifications

The T lymphocyte subgroups and the time to peak of cytokines collected were analyzed by one-way ANOVA according to the clinical classification, as shown in Table 5. At the same time, multiple linear tests were conducted for all indicators in the table with $p < 0.05$, and finally, the days of nucleic acid seroconversion, the time to peak of T lymphocytes (CD 3⁺ cells), the time to peak of IL1 β and the time to peak of IL17A were included in the logistics regression statistics (Table 6).

Logistics analysis results showed that parallel test was satisfied ($\chi^2 = 7.837$ $p = 0.098$); There is proportional advantage hypothesis; Deviance goodness-of-fit test showed that the model could fit well ($\chi^2 = 45.127$, $p = 0.991$). This model was superior to the model with constant only ($\chi^2 = 45.127$, $p < 0.001$).

As can be seen from Table 6, the time to peak of T cells (CD3⁺ cells) was $p = 0.02$ and the time to peak of IL-1 β was $p = 0.016$, which were statistically significant and could be included in the logistics model. It is suggested that, in mild and common group vs severe/critical group, T cell time to peak and IL-1 β time to peak had influence on clinical classifications. Meanwhile, the OR value showed that the probability of progressing to severe/critical disease was 1.276 times (95%CI: 1.039-1.569) with the increase of T cell time to peak ($\chi^2 = 5.379$, $p = 0.02$). The probability of progressing to severe/critical disease was 1.408 times (95%CI: 1.066-1.859) with each day increase of IL1 β time to peak ($\chi^2 = 5.824$, $p = 0.016$).

Comparison of the differences between the time to peak of cytokines and T lymphocyte subgroups

Independent sample t test was used to analyze whether there were differences in the time to peak, when cytokines value and T cell count reached the maximum between the fast nucleic acid seroconversion group and slow seroconversion group (Table 7). It can be seen that in CD4⁺T cells, IL-1 β , IL-2, IL-8, IL-12p70, IL-17A, IFN- γ , TNF- α and other detection indicators, the time to peak of the above indicators in the fast seroconversion group was significantly shorter than that in slow seroconversion group, that is, faster time to peak of the above indicators related to faster recovery.



Table 3: Results of Analysis of Repeated Measures ANOVA for Different Classifications of COVID-19 Patients Laboratory Indicators.

Indicators	Source	Type III sum of Squares	df	Mean Squares	F	sig.
CD3+ cell (#/ul)	Classification	5428471.702	2	2714235.851	7.789	0.002
	Time	431339.213	2.395	180106.572	2.698	0.064
	Time*Classification	647274.241	4.790	135135.343	2.024	0.088
CD4+T cell (#/ul)	Classification	928441.455	2	464220.727	8.797	0.001
	Time	35284.493	3	11761.498	1.479	0.226
	Time*Classification	104564.728	6	17427.455	2.191	0.051
CD8+T cell (#/ul)	Classification	1778700.931	2	889350.466	5.140	0.012
	Time	199738.030	3	66579.343	2.705	0.050
	Time*Classification	237864.583	6	39644.097	1.610	0.153
IL-1	Classification	9.132	1	9.132	0.186	0.677
	Time	18.892	1.913	9.874	3.066	0.074
	Time*Classification	5.276	1.913	2.757	0.856	0.438
IL-2	Classification	0.712	1	.712	0.135	0.721
	Time	1.699	3	0.566	4.142	0.015
	Time*Classification	1.144	3	0.381	2.788	0.060
IL-4	Classification	8.835	1	8.835	2.643	0.138
	Time	0.731	3	0.244	1.303	0.294
	Time*Classification	1.707	2.355	0.725	3.040	0.062
IL-5	Classification	0.412	1	0.412	0.514	0.492
	Time	0.694	3	0.231	3.069	0.045
	Time*Classification	0.169	3	0.056	0.746	0.534
IL-6	Classification	0.287	1	0.287	0.016	0.902
	Time	0.885	1.291	0.686	0.055	0.875
	Time*Classification	37.388	1.291	28.969	2.339	0.150
IL-8	Classification	64.060	1	64.060	0.142	0.715
	Time	6961.616	1.973	3528.994	4.087	0.035
	Time*Classification	437.890	1.973	221.976	0.257	0.773
IL-10	Classification	0.847	1	0.847	0.033	0.861
	Time	3.465	3	1.155	0.936	0.437
	Time*Classification	0.397	3	0.132	0.107	0.955
IL-12p70	Classification	44.217	1	44.217	2.836	0.126
	Time	14.976	3	4.992	6.458	0.002
	Time*Classification	5.293	3	1.764	2.282	0.102
IL-17A	Classification	1312.225	1	1312.225	0.524	0.487
	Time	1345.865	3	448.622	1.907	0.152
	Time*Classification	394.612	3	131.537	0.559	0.647
IFN-γ	Classification	0.276	1	0.276	0.001	0.979
	Time	131.474	3	43.825	1.535	0.228
	Time*Classification	53.051	3	17.684	0.619	0.609
IFN-α	Classification	500.336	1	500.336	0.450	0.519
	Time	1501.757	1.016	1477.983	0.855	0.381
	Time*Classification	2114.547	1.016	2081.071	1.204	0.302
TNF-α	Classification	9.419	1	9.419	0.213	0.656
	Time	6.449	3	2.150	2.663	0.068
	Time*Classification	6.956	3	2.319	2.872	0.055

Table 4: Multiple Comparison Results of CD3+ cell /CD4+/CD8+ T Cell Counts of Different Classifications.

	Common Group			Severe/Critical Group		
	CD3+	CD4+	CD8+	CD3+	CD4+	CD8+
Mild Group	87.281	204.28	47.38	360.21*	71.76	156.29*
Common Group				477.50*	277.04*	156.29*

* p<0.05

Discussion

In the study, T cell count of COVID-19 patients with various clinical classifications were analyzed, and it was found that all T cell subgroups in severe/critical patients were lower than those in mild or common patients, which was consistent with the conclusions of previous other studies. Moreover, it is worth mentioning that our study is the first one to describe the

immunological characteristics of COVID-19 patients in time dimension.

With the deepening of immunological studies on COVID-19 patients, it is the key point for the treatment to explore the changes of cytokines and T lymphocytes, timely find the changing point of severe clinical course of disease and timely intervene. As one of the important cells of the acquired immune system, T lymphocytes play important roles in all infectious diseases. T cell count is positively correlated with the disease prognosis [4]. As cytotoxic T cells, CD8+T cells have a direct killing effect on virus-infected cells, and a significant decrease in CD8+T cell levels can be seen in patients with COVID-19 [5,6]. Decrease in T cell count can be observed in the early stage of COVID-19 disease, and the decrease in T cell count is more

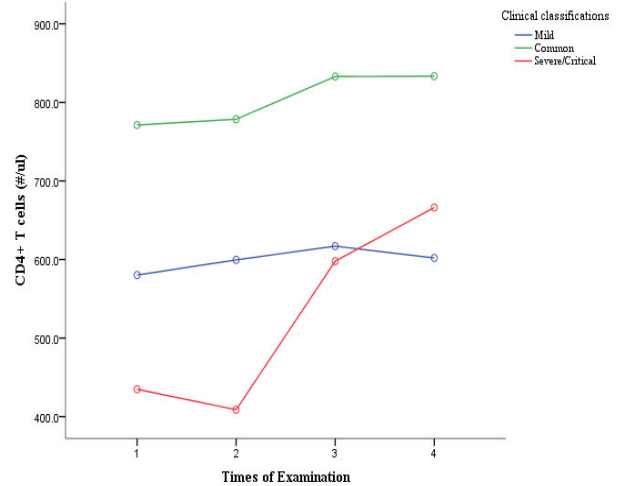
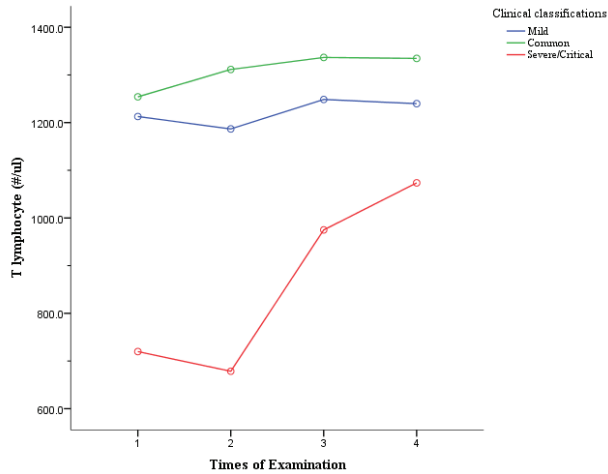


Figure 1: CD3⁺ (left) and CD4⁺ (right) T Cell Development Trends of Patients with Different Clinical Classifications of COVID-19.

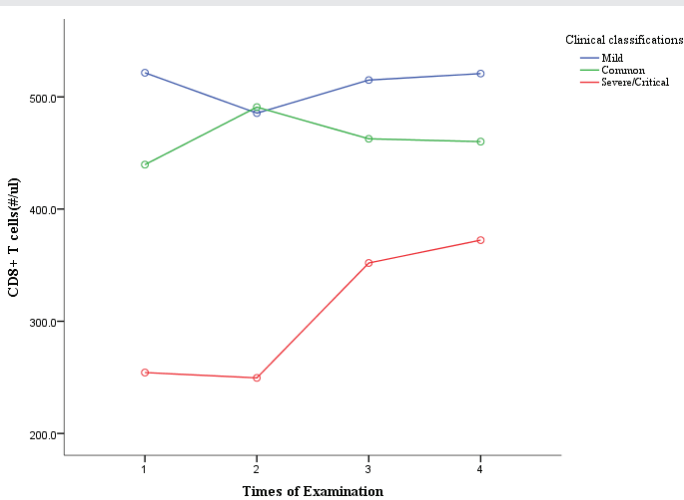


Figure 2: CD8⁺ T Cell Development Trends of Patients with Different Types of COVID-19.

significant in patients with severe disease than in patients with mild disease. In some critical patients, even, depletion of T lymphocyte may occur. As can be seen in other study, decrease in CD4⁺T cell counts is also common in severely ill patients [7]. We observed that, patients with severe/critical disease had significantly lower T cell, CD4 and CD8⁺T cell counts than patients with mild and common disease, which was consistent with the characteristics of patients with COVID-19 described by other researchers. According to develop trends described in our study, however, T lymphocytes, CD4⁺ T and CD8⁺ T cell counts have not decreased dramatically. Lymphocytes increased in patients of all three clinical groups after treatment, the increase trends were significant especially in severe/critical group patients. A possible explanation is, although there is a certain proportion of severe/critical patients, all patients got good prognosis after hospitalization. Again, the T cell count and its changing trend probably determine the progression and outcome of the patient's disease.

In our study, it was suggested that the possibility of severe/

critical patients was significantly increased with prolonged time to peak of T cell count and IL1 β value. At the same time, the faster CD4, IL-1 β , IL-2, IL-8, IL-12p70, IL-17A, IFN- γ , TNF- α and other indicators reached the peak, the faster nucleic acid negatively seroconvert. Previous immunology studies on COVID-19 infection suggested that the virus produced two cytokine spikes after attacking the body. In the first stage, the natural immune system of the body activates type I interferon response and causes a small peak of cytokines. The second was a cytokines storm that dramatically worsened the symptoms [8]. Type I interferon response is an important mechanism of the innate immune system to protect the body against virus. It produces IL-1 β , TNF- α , IL-6 and regulates the immune response of T cells to prevent the further amplification of the cytotoxic reaction [9]. In the process of respiratory virus

Table 5: Results of Analysis of Single Factor for Different Types of COVID-19 Laboratory Indicators.

	M	SD	F	sig.
Days in hospital	9.692	6.9364	11.697	<0.001
Nucleic acid seroconversion days	10.846	6.8577	7.033	0.003
CD3 ⁺ T cell time to peak	9.462	6.6207	5.520	0.008
CD4 ⁺ Tcell time to peak	16.895	7.0779	5.284	0.010
CD8 ⁺ Tcell time to peak	5.579	5.3858	4.136	0.024
IL-1 β time to peak	5.000	5.9729	12.396	<0.001
IL-2 time to peak	5.132	5.2152	1.383	0.264
IL-4 time to peak	4.237	4.3086	0.968	0.390
IL-5 time to peak	4.947	5.6614	0.767	0.472
IL-6 time to peak	4.237	4.4505	0.404	0.671
IL-8 time to peak	5.342	5.3790	2.017	0.148
IL-10 time to peak	6.184	6.1065	0.507	0.607
IL-12p70 time to peak	4.763	4.5704	1.446	0.249
IL-17A time to peak	3.579	2.9097	4.449	0.019
IFN- γ time to peak	5.263	5.7595	1.816	0.178
IFN- α time to peak	9.692	6.9364	1.454	0.247
TNF- α time to peak	10.846	6.8577	1.374	0.266

**Table 6:** Estimated Values of Logistics Parameters.

Parameters	B	Std. Error	Wald	sig.	Exp(B)	95% Confidence Interval for Exp(B)	
						Lower Bound	Upper Bound
Mild Group	5.080	1.8770	0.007	0.007	0.007	4.059	6364.650
Common Group	7.308	2.1162	0.001	0.001	0.001	23.582	94463.486
Nucleic acid turns cloudy days	-0.084	0.1164	0.523	0.470	0.470	0.732	1.155
CD3+T cell Peak time	0.244	0.1052	5.379	0.020	0.020	1.039	1.569
IL-1 β Peak time	0.342	0.1418	5.824	0.016	0.016	1.066	1.859
IL-17A Peak time	0.164	0.1278	1.644	0.200	0.200	0.917	1.513

Table 7: Differences in the Time to Peak of each Indicator between the Fast and Slow nucleic acid Seroconversion Groups.

Indicators	Fast group (n=19)	Slow group (n=20)	t	sig.
CD3 ⁺ cell	7.68 \pm 4.75	11.6 \pm 8.185	-1.838	0.076
CD4 ⁺ T cell	8.37 \pm 4.81	13.2 \pm 7.757	-2.35	0.025*
CD8 ⁺ T cell	8.89 \pm 5.216	10 \pm 7.827	-0.521	0.606
IL-1 β	3.39 \pm 2.81	6.9 \pm 5.803	-2.41	0.023*
IL-2	3.72 \pm 2.927	7.25 \pm 6.528	-2.185	0.038*
IL-4	3.39 \pm 2.789	6.45 \pm 7.605	-1.679	0.106
IL-5	3.72 \pm 3.045	6.4 \pm 6.411	-1.67	0.106
IL-6	3.72 \pm 2.761	4.7 \pm 5.371	-0.694	0.492
IL-8	2.78 \pm 1.987	6.9 \pm 7.1	-2.49	0.021*
IL-10	3.06 \pm 2.014	5.3 \pm 5.695	-1.652	0.112
IL-12p70	3.28 \pm 2.296	7.2 \pm 6.63	-2.485	0.02*
IL-17A	3.89 \pm 3.179	8.25 \pm 7.355	-2.413	0.023*
IFN- γ	2.94 \pm 1.83	6.4 \pm 5.633	-2.596	0.016*
IFN- α	3.33 \pm 2.473	3.8 \pm 3.302	-0.489	0.628
TNF- α	3 \pm 2.249	7.3 \pm 7.131	-2.559	0.018*

infection, type I interferon response protects alveolar epithelial cells, reduces inflammatory exudation, and ultimately improves prognosis and reducing mortality rates. The timing and intensity determine the prognosis of viral infection [10,11]. Channappanavar et al. infected mice with SARS-CoV and found that delayed type I interferon response resulted in an increase in proinflammatory mononuclear macrophages, leading to increased inflammatory exudation of mouse lung tissue and intensified immune response, ultimately leading to increased mortality in mice [12]. These adverse outcomes were also observed in mice infected with MERS-CoV with delayed or absent type I interferon response [13]. However, studies on the timing of immune response in clinical patients are still lacking at present. Our study is the first to show that the early onset and timely termination of the immune response has positive effects on the prognosis of disease and the speed of 2019-nCoV nucleic acid seroconversion. Patients with the above immune response may be cured before the cytokine storm occurs in the body.

There are some deficiencies in our study. Due to the lack of death cases, no in-vivo immune response was observed in the patients who were finally dead. And as a single-center retrospective study, the sample size and follow-up time is short. In the future, multicenter, prospective study is needed.

Longer follow-up period is also required to clarify the timing and strength of COVID-19 patients' immune functions, so that we accurately grasp the turning point of disease, and give timely treatment. Moreover, with the application of Next Generation Sequencing (NGS), researchers have found that the B cell receptor cluster correlates with the SARS-CoV-2 antibodies. Clonality and skewing of TCR repertoires were associated with interferon type I responses, and early CD4⁺ and CD8⁺ T cell activation [14]. This might be a new direction of our research. In a future study, NGS may be used to analyze TCR and BCR in above three groups of patients to find out whether there are differences among them. That would be momentous in the coming era of vaccine studies.

Conclusions

The early peak, timely decrease of cytokines values and T cell counts in COVID-19 patients has a positive effect on prognosis. Dynamic monitoring of T cell counts and quantitative changes of cytokines is a simple and easy method to evaluate disease severity in clinical practice for COVID-19 patients. By closely monitoring the time and quantity changes of the above indicators, the disease turning point can be found in time, which is convenient for clinicians intervene and improve the prognosis of patients timely.

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