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Serum cytokines and clinical features in patients with fever and thrombocytopenia syndrome



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ABSTRACT

Purpose: To explore the clinical, microbiological and immunological features of patients with fever and thrombocytopenia.

Methods: Patients with unexplained fever and thrombocytopenia were enrolled. Viruses were detected using real-time PCR, and bacteria were measured by culturing methods. Serum cytokines, platelet antibody IgG (PA-IgG) and *Helicobacter pylori* (HP) were detected using ELISA.

Results: Pathogens were detected in 74.68% of patients, which included single fungal/viral/bacterial infection and multiple infection. The pathogens could not be unidentified in 25.32% of cases. Cytokines including Interleukin (IL)-6, IL-10, interferon- γ (IFN- γ), platelet activating factor (PAF) and PA-IgG were significantly higher in patients as compared to healthy controls ($P < .01$ or $P < .05$). Principal component analyses extracted four groups of parameters that have a strong positive predicting value, revealing that disease status evaluation would be more accurate if we combined the platelet parameters and inflammatory biomarkers. While event-free survival (EFS) that indicates the time of platelet elevated after therapy was the highest in patients with single bacterial or fungal infection, EFS was affected by the levels of cytokines and PA-IgG.

Conclusions: Differences in immune function may be the main factors affecting the prognosis of patients with fever and thrombocytopenia, while treatment based on precise etiological diagnosis is important for therapeutic efficacy.

1. Introduction

Fever and thrombocytopenia are frequent symptoms in patients with viral infection. Severe fever with thrombocytopenia syndrome (SFTS), caused by a novel bunyavirus, was initially identified in 2009. SFTS produces a broad spectrum of clinical manifestations that range from acute self-limited febrile illness to severe disease, with a reported fatality rate varying between 12% and 30% [1]. Due to its poor prognosis and higher mortality, SFTS has quickly become an emerging infectious disease that has aroused extensive surveillance in public health. Currently, SFTS, together with meningitis syndrome and respiratory syndrome, are the primary three syndromes that are monitored in clinic. Unlike other inflammatory disease, the etiology of fever and thrombocytopenia is quite complicated; thrombocytopenia may

arise due to a lack of platelet production, retention of platelets (PLTs) in the spleen, platelet destruction, or increased use and dilution of PLTs. In clinic, most thrombocytopenia patients are diagnosed as primary immune thrombocytopenia (ITP) that is defined as a peripheral blood platelet count $< 100 \times 10^9/l$ in the absence of any obvious cause of thrombocytopenia [2]. Because the etiology of ITP is so complicated that physicians seldom relies on its etiology to formulate treatment plans, then, it is difficult to explain why some patients have poor therapeutic efficacy.

Fever is a common feature and consequence for ITP patients with viral infection. Evidence regarding the cross-reactivity of virus-specific antibodies with normal platelet antigens was firstly obtained in patients developing ITP after contracting human immune deficiency virus [3] and varicella-zoster virus infection [4]. Since then, it has been

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discovered that thrombocytopenia and viral infections occur in succession. Thrombocytopenia as a major clinical sign is found in some viral infectious diseases including Zika virus, HIV-1, hepatitis B/hepatitis C virus, dengue virus, Epstein-Barr virus, B19 virus, and novel bunyavirus [1,5–12]. A previous study showed that patients with influenza virus A/B or parainfluenza virus 1 infections tend to have a lower PLT count than patients who did not have respiratory viral infection, while patients with human metapneumovirus, parainfluenza virus 2, coronavirus 229, or adenovirus infections did not show significant changes in the PLT count [13]. In some patients, thrombocytopenia caused by viral infection was transient without serious bleeding and spontaneously resolved within a few weeks without treatment [14]; however, other patients have a severe bleeding syndrome and worse prognosis. The mechanism remains to be elucidated.

Thrombocytopenia is also a common complication during or after bacterial infections. Direct interaction between platelets and bacteria may contribute to thrombocytopenia [15]. *Helicobacter pylori* (HP) has proven to be the most important bacterium causing ITP. Additionally, it was observed that salmonella survived and replicated in macrophages that spread to the lymphatic vessels, spleen, and bone marrow, and thrombocytopenia may be associated with hemophagocytic histiocytes in the bone marrow [16]. Fecal microbial transplantation may also lead to thrombocytopenia during inflammation [17]. *Staphylococcus aureus* activates platelets through several surface-expressed proteins, and binds to adhered platelets to cause platelet activation and aggregation, resulting in thrombocytopenia [18]. Unfortunately, the role of bacterial infection in thrombocytopenia is often clinically ignored.

A viral/bacterial antigen may be recognized as a similar platelet antigen, a process termed molecular mimicry, which then gives rise to cross-reactive anti-platelet autoantibodies [19,20]. Platelets with bound autoantibodies are subsequently recognized by phagocytes bearing Fc γ receptors (Fc γ Rs), which results in enhanced antibody-mediated platelet phagocytosis and destruction primarily in the spleen [21]. In patients with ITP autoantibodies targeting platelet surface glycoproteins, primarily GPIIb-IIIa and GPIb-IX, lead to Fc-dependent clearance via macrophages [22]. Platelets express Fc γ RIIA, an IgG receptor used to identify immune complexes (ICs). Fc γ RIIA is involved in the initiation of strong effector functions in leukocytes, including cytokine release, antibody-dependent cell-mediated pathogen killing, and ICs internalization, which is critical for immune and inflammatory responses. Virus or bacteria can activate platelets via Fc γ RIIA that plays an important role in the interaction of platelet and pathogen infection, and the polymorphism of Fc γ RIIA-131 directly affects the susceptibility and severity of infection [23]. However, the relationship among the platelet autoantibodies, cytokines, and other parameters associated infection is still not clear.

Many questions regarding the pathophysiology, mechanism, diagnosis, natural history, and management of ITP remain to be resolved [24,25]. The obscure etiology and mechanisms of thrombocytopenia greatly affect its therapeutic efficacy. First-line therapy for ITP consists of corticosteroids with or without intravenous immunoglobulin [2], and some patients respond to this while others do not. Nowadays, there are no valuable biomarkers to predict the therapeutic efficacy. Etiologic diagnosis of thrombocytopenia may be beneficial for treatment, to some ITP caused by viral infections, accurate etiologic diagnosis can reduce the amount of antibiotics used. Unfortunately, the diagnosis of ITP is still based on exclusion [26]. The relationships among the etiology of fever and thrombocytopenia, the intrinsic immune mechanism, and therapeutic efficacy remain to be elucidated. To clarify the above issues, patients with unexplained fever and thrombocytopenia were enrolled in this study. The associations among the etiology, platelet autoantibody, cytokine expression, and therapeutic efficacy were investigated with the aim of determining valuable parameters that mirror the pathogenesis and predict therapeutic efficacy of thrombocytopenia.

2. Material and methods

2.1. Patients and sample collection

During the period of January 2016 to June 2017, patients with unexplained fever (body temperature > 37.6 °C) and thrombocytopenia (platelet count < 100 × 10⁹/l) were enrolled in our study. Patients with solid tumors, leukemia, liver disease, or upper and lower gastrointestinal or other traumatic bleeding were excluded. Ten healthy volunteers were selected as the controls. The study was approved by the Ethics Committee of the First Affiliated Hospital of Shantou University Medical College, and informed consent was obtained from all patients. All samples were collected at the onset of disease and before treatment.

2.2. Real-time PCR detection of viruses

The nucleic acid from the serum samples was extracted using a viral DNA/RNA magnetic bead extraction kit (TIANLONG, China) and NP968 nucleic acid extraction instrument (TIANLONG, China). A single-tube TaqMan® real-time reverse transcription polymerase chain reaction (PCR) strategy (AgPath-ID™ One-Step RT-PCR kit from Applied Biosystems, USA) was used to detect viruses. Bunyavirus was detected using the severe fever with thrombocytopenia syndrome virus (SFTSV) Bunyavirus Real Time RT-PCR Kit (Liferiver, China), and human parvovirus B19 was detected using the Human Parvovirus B19 Nucleic Acid Detection Kit (Daangene, China). PCR amplification was performed using a 7500 Real-Time PCR System (Applied Biosystems, USA), according to the manufacturer's instructions.

2.3. Fungus/bacteria detection

For bacteria identification, the blood and sputum specimens of patients were cultured according to National Standard Protocols of China. Specimens were inoculated in blood agar, eosin methylene blue agar, and chocolate agar and incubated for 24–48 h at 37 °C. Colony identification was undertaken using a VITEK 2 Compact system (Biomérieux, France). *Helicobacter pylori* was detected using a commercial kit (Human Anti-*Helicobacter Pylori* Antibody (IgG) ELISA kit (Cusabio, China)).

2.4. Detection of PA-IgG and cytokines in serum

The PA-IgG and six cytokines PAF, interleukin (IL)-6, IL-10, IL-17A, IL-12/23P40, and IFN- γ in serum samples were detected using a Human platelet-associated IgG antibody (PA-IgG) ELISA kit (Shanghai Jingkang, China), Human Platelet Activating Factor ELISA Kit (Dakewe, China) and Human IL-6/IL-10/IL-17A/IL-12/23P40/IFN- γ precoated ELISA Kit (CUSABIO, China), according to the manufacturer's instructions. The absorbance values of six cytokines were measured using an Infinite M200 Pro microplate reader (Tecan, Switzerland) and then converted to concentration values according to the reagent instructions. The sensitivity of PA-IgG, PAF, IL-6, IL-10, IL-17A, IL-12/23P40, and IFN- γ detection was 0.1 μ g/l, 1.17 ng/ml, 2 pg/ml, 5 pg/ml, 31.3 pg/ml, 15 pg/ml, and 5 pg/ml, respectively.

2.5. Statistical analyses

Statistical analyses were performed using SPSS 19.0 software. Continuous variables with normal distributions were used mean \pm standard deviation (SD) to describe and compare using multiple variance of multiple samples. Medians and interquartile ranges (IQRs) were used to describe non-normal distribution, with comparisons based on the Mann-Whitney *U* test between any two groups. The correlations between serum cytokine and other parameters were obtained using the non-parametric rank-based Spearman's correlation coefficient. Principal component analysis (PCA) was performed to obtain the significant

Table 1
Patient's general characteristics and pathogen detection results.

	Cases	Percentage
Total	79	
Age		
0–14	4	5.06%
≥15	75	94.94%
Gender		
Male	48	60.76%
Female	31	39.24%
Temperature (°C)		
37.6–37.9	11	13.92%
38.0–38.9	44	55.70%
39.0–39.9	16	20.25%
≥40.0	8	10.13%
Pathogen infection		
Single virus		
PIV2	3	3.80%
229E	2	2.53%
Flu A	2	2.53%
Single fungus	5	6.33%
Single bacteria		
<i>Helicobacter pylori</i>	24	30.38%
<i>Acinetobacter baumannii</i>	3	3.80%
<i>Acinetobacter baumannii</i> + HP	3	3.80%
<i>Haemophilus influenzae</i>	1	1.27%
<i>Serratia marcescens</i> + HP	1	1.27%
Multiple pathogens		
229E + HP	6	7.59%
PIV2 + HP	3	3.80%
PIV2 + Fungus	1	1.27%
229E + Fungus + HP	1	1.27%
Flu A + Fungus + HP	1	1.27%
PIV2 + <i>Klebsiella pneumoniae</i>	1	1.27%
PIV2 + <i>Acinetobacter baumannii</i> + HP	1	1.27%
Flu A + <i>Escherichia coli</i> + HP	1	1.27%
Unidentified	20	25.32%

Abbreviations: 229E, coronaviruses 229E; PIV2, parainfluenza viruses 2; Flu A, influenza viruses types A; HP, *Helicobacter pylori*.

principal components that most accurately reflect the status of thrombocytopenia. Receiver operating characteristic (ROC) curve analysis was used to determine and distinguish the usefulness of biomarkers. EFS was calculated from the date of complete recovery (platelet count return to normal range) until the first event (any cause of recurrence or death) or until the last follow-up. Survival curves were calculated using

Table 2
The clinical data among different pathogens infected with fever and thrombocytopenia patients.

	Single virus	Single f/b	Multiple pathogens	Unidentified
Number of cases	7	37	15	20
Improved case after healing	3	19	5	8
Deaths cases	1	2	2	4
Gender (M/F)	5/2	20/17	9/6	14/6
Age	57.14 ± 16.24	59.22 ± 19.58	55.53 ± 20.51	54.85 ± 23.10
Temperature (°C)	38.81 ± 0.78	38.64 ± 0.81	38.85 ± 0.97	38.45 ± 0.51
WBC (×10 ⁹ /l)	7.43(5.99–9.16)	9.81(5.25–11.29)	9.58(8.46–13.20)	7.07(3.79–11.83)
LY (%)	8.64(2.96–19.2)	9.28(4.10–14.01)*	13.10(5.55–15.32)	18.37(6.35–26.33)
N (%)	84.97(71.52–95.22)	84.28(71.07–92.58)	82.70(76.21–86.84)	70.72(59.18–87.19)
RBC (×10 ¹² /l)	3.54 ± 0.98	3.54 ± 0.80	3.87 ± 0.80	3.77 ± 1.06
HB (g/l)	98.29 ± 23.82	105.65 ± 23.16	118.00 ± 23.47	113.05 ± 29.61
PLT (×10 ⁹ /l)	75.86 ± 14.98▲	64.57 ± 21.12	52.27 ± 22.78	61.85 ± 26.10
MPV (fl)	9.50 ± 0.86	12.44 ± 2.69	10.36 ± 1.42	9.79 ± 1.69
PDW (fl)	17.57 ± 0.89	17.07 ± 2.20	16.19 ± 2.33	16.77 ± 1.70
CRP (mg/l)	181.4(107.00–196.00)	93.40(28.30–138.50)	128.44(110.00–140.00)	75.70(13.25–109.55)
PCT (ng/ml)	1.72(0.54–14.45)	6.41(2.56–195.46)	7.75(0.61–61.87)	1.09(0.23–8.27)
ALT (U/l)	23.0(19.00–62.00)	29.00(15.50–142.00)	50.00(24.00–436.00)	56.50(26.50–118.50)
AST (U/l)	83.00(28.00–129.50)	66.00(23.00–235.00)	59.00(34.00–789.00)	83.50(33.25–133.50)

Abbreviations: f/b, fungus/bacteria; M, Male; F, Female; WBC, white blood cell; LY, lymphocyte; N, neutrophil; RBC, red blood cell; Hb, hemoglobin; PLT, platelet; MPV, mean platelet volume; PDW, platelet distribution width; CRP, C-reaction protein; PCT, procalcitonin; ALT, alanine aminotransferase; AST, aspartate aminotransferase.

* $P < 0.05$ when the single f/b group is compared with unidentified group.

▲ $P < 0.05$ when the single virus group is compared with multiple pathogens group.

the Kaplan–Meier method, and survival was compared using Mantel's log-rank test. Thrombocytopenia with or without improvement were analyzed using binary logistic regression. The level of statistical significance was set at 5%.

3. Results

3.1. The basic clinical characteristics of the patients

In Table 1, the 79 patients included 48 males and 31 females. Four cases occurred in children with a mean age (years) of 7.75 (1–14 years), and 75 cases occurred in adults with a mean age of 59.87 (17–93 years). Various body temperatures (T) were obtained from the patients, including 11 patients with normal T (37.6–37.9 °C), 44 (55.7%) patients with moderate T (38.0–38.9 °C), and 24 cases with high T (≥39.0 °C). Pathogen detection showed that a single virus infection was found in 7 patients (8.86%), which mainly consisted of three types of viruses: 229E, PIV2, and Flu A. Single fungus infection was found in 5 cases, and single bacterial infection was found in 24 cases of *Helicobacter pylori* (HP), 3 cases of *Acinetobacter baumannii*, 3 cases of *Acinetobacter baumannii* concurrent with HP, 1 case of *Haemophilus influenzae* and 1 case of *Serratia marcescens* concurrent with HP. Multiple-infection was found in 15 patients (18.99%) (shown in Table 1). The pathogens could not be measured in 20 cases (25.32%). Of these 79 patients, 52 patients treated with antimicrobials, and most used antibiotics were quinolones, followed by cephalosporins and beta-lactams. Few patients were treated with anti-viral drugs.

We then divided all patients into four groups according to the pathogens detected (shown in Table 2): single viral group (7 cases), single fungal/bacterial (f/b) group (37 cases), multiple pathogens group (15 cases), and unidentified pathogen group (20 cases). Males were more than females, and average age ranged from 54 to 60 years in each group.

In Table 2, the improvement rate for the single f/b infectious group was 51.35% (19/37), the single viral group was 42.86% (3/7), the unidentified pathogen group was 40.00% (8/20), and the multiple infectious group was 33.33% (5/15). In the comparison of laboratory parameters, the white blood cell (WBC) counts in the single f/b group and multiple infectious group were higher than those in the single viral and unidentified pathogen group, and the highest lymphocyte count

was found in the unidentified pathogen group ($P < .05$). The mean PLT count of the four groups was $< 76.0 \times 10^9/L$, and the mean PLT count of the multiple pathogens group was the lowest ($52.27 \times 10^9/L$), which was significantly lower than that in the single virus group ($P < .05$). The mean platelet volume (MPV) count was the highest in the single f/b group (mean = 12.44 fl), which was higher than that in the other three groups. The mean C-reactive protein (CRP) in thrombocytopenia groups was higher than the normal value (CRP < 0.05 mg/l), similarly, the median PCT in the four groups were higher than the clinical limit of detection (PCT < 0.1 ng/ml), with the median value of the multiple pathogens group being the highest (7.75) (Table 2). The clinical parameters of patients with or without HP infection were also analyzed. The median LY (%) in HP-positive patients were lower than those in negative patients ($P < .05$). The median PAF concentration was 2.5 times higher in HP-positive patients than that in the negative group.

3.2. The serum PA-IgG and cytokines in different groups

Among the six cytokines measured in this study, there were no significant differences for serum IL-12/23P40 and IL-17A among five groups ($P > .05$) (Fig. 1). However, we found that serum levels of IL-6, IL-10, and IFN- γ , as well as PAF were significantly higher in all patient groups as compared to those in healthy controls ($P < .01$). Compared to the unidentified pathogen group, IL-6 was significantly higher in the single viral group with a mean concentration of 153.03 pg/ml ($P < .05$). The highest level of IL-10 was found in the unidentified pathogen group (501.67 pg/ml). The IFN- γ concentration (582.59 pg/ml) was the highest in the single viral group among the five groups. The highest PAF value was found in the single f/b group (400.02 ng/ml). Similarly, the concentration of PA-IgG of healthy people was lower than those of fever and thrombocytopenia patients, and there was a significant difference when compared with three of patients' groups (healthy vs. single virus: $P < .01$; healthy vs. single f/b: $P < .05$; healthy vs. unidentified: $P < .01$) (Fig. 1).

3.3. The correlations between cytokines and platelet parameters in patients with fever and thrombocytopenia

Inner associations were also analyzed between cytokines and the platelet parameters. In this study, the PLT quantity was associated with platelet distribution width (PDW) ($r = 0.231$, $P < .05$) and IL-10 ($r = 0.331$, $P < .01$). A correlation exists between IL-6 with IL-10 ($r = 0.458$, $P < .01$), with PAF ($r = 0.370$, $P < .01$), with CRP ($r = 0.667$, $P < .01$), and with PCT ($r = 0.440$, $P < .01$). The correlations were found between IL-10 with IL-12/23P40 ($r = 0.298$, $P < .01$), IFN- γ ($r = 0.446$, $P < .01$), PAF ($r = 0.257$, $P < .05$) and PCT ($r = 0.379$, $P < .01$). Additionally, IL-12/23P40 was correlated with IFN- γ ($r = 0.447$, $P < .01$), and PCT was correlated with CRP ($r = 0.410$, $P < .01$). No correlations were tested between IL-17A and other cytokines (Table 3).

3.4. Principal component analysis (PCA)

PCA method was exploited to determine whether the parameters profile reflects disease status in patients with fever and thrombocytopenia. PCA extracted four important principal components with eigenvalues > 1 that explained 58.71% of the total variance of the data set. It can be seen from component 1 that IL-6, CRP, and PCT have a strong positive loading. Component 2 includes IL-10, IFN- γ , and PLT, where PLT exert a strong negative loading. Similarly, component 3 included IL-12/23P40 and PDW, component 4 included PA-IgG, IL-17A, PAF and MPV, while PDW, PAF and MPV contributed strong negative loadings (Table 4). Our results showed that cytokine loading is larger than other parameters in reflecting disease status (component 1: IL-6 = 0.873 $>$ CRP = 0.767/PCT = 0.758; component 2: IL-

10 = 0.674/IFN- γ = 0.714 $>$ PLT = -0.615; component 3: IL-12/23P40 = 0.783 $>$ PDW = -0.678; component 4: PA-IgG = 0.811 $>$ IL-17A = 0.659/PAF = -0.458 $>$ MPV = -0.302). Therefore, to patients with fever and thrombocytopenia, the disease status evaluation would be more accurate if we combined the platelet parameters and inflammatory biomarkers.

3.5. Risk evaluation in patients with/without improvement after treatment

The factors that significantly influenced the prognosis of patients were also analyzed, these were age, PLT quantities, PCT, and IL-17A. Among them, PLT quantities, PCT, and IL-17A were risk factors (OR > 1.0), and age was a protective factor (OR < 1.0) (Table 5). We then used Kaplan-Meier survival curves to further assess the suitability of six cytokines and PA-IgG as prognostic factors and to compare the prognosis of different types of pathogen infections. We conducted a follow-up of the patient from hospital admittance to discharge, and survival information was collected. The levels of cytokines and PA-IgG were considered as a stratification variable for survival time analysis. Our data showed that patients with single f/b infection have the highest event-free survival (EFS), and patients with single viral infection had the lowest. EFS in patients with lower levels of IL-6, IL-10, IL-17A, IFN- γ and PA-IgG tended to be longer than in patients with higher levels. However, EFS in patients with lower levels of IL-12/23P40 and PAF tended to be shorter when compared to higher-level patients. We observed no significant difference between the higher and lower groups, and the p -values of the K-M curves were all > 0.05 , probably due to the small sample size (Fig. 2).

4. Discussion

Platelets participate in the interaction between pathogens and host defense [27], but their function in host defense against infection has received much less attention [28]. Pathogenic microbiome may cause thrombocytopenia by autoimmunity mechanisms, mainly through molecules or epitopes that mimic autoantigens, the immunological recognition of previously unexposed auto-antigens, or via bystanders that activate their own responses to lymphocytes and super-antigen-mediated polyclonal T cell activation with exaggerated cytokine release [29]. Studies have shown that an imbalance of T helper cell (Th) and regulatory T cell (Treg) cytokines may mediate the pathogenesis of thrombocytopenia, which is closely related to the expression of Th1 cytokines (i.e., IFN- γ , IL-2), Th2 cytokines (i.e., IL-4, IL-10), Th17 cytokines (IL-17), and Treg cytokines (TGF- β 1) [30]. Consistent with previous studies, up-regulated cytokines were found in our patients with fever and thrombocytopenia, especially IL-6, IL-10, and IFN- γ , indicating that cytokine disorders may be involved in the pathogenesis of newly diagnosed ITPs [31,32]. Notably, IL-6 in thrombocytopenia patients was significantly higher as compared to healthy controls, and IL-6 in the viral group was the highest among all groups. Moreover, our results revealed that IL-6, together with CRP and PCT, has a strong positive loading in mirroring the pathogenesis of thrombocytopenia. PAF is a potent and versatile mediator of pro-inflammation secreted by WBCs [33], which triggers platelet aggregation and activation. Monocytes that encounter dengue virus begin to generate PAF [34], and this leads to enhanced apoptosis of platelets and accelerates platelets clearance in secondary dengue infection [35]. The PAF concentration in our patients with fever and thrombocytopenia was significantly higher as compared to the healthy controls, revealing that PAF may involve in the platelet's apoptosis and clearance in infectious thrombocytopenia. The dose-dependent effect of PAF may induce the release of cytokines and affect the host immune response. A previous study showed that PAF can stimulate mononuclear cells of peripheral blood leading to the release of TNF- α and IL-6 in viral infection [36]. PAF itself or combined with LPS can promote platelet activation and recruitment and worsen endotoxin-induced injury [37,38]. Therefore, PAF and cytokines

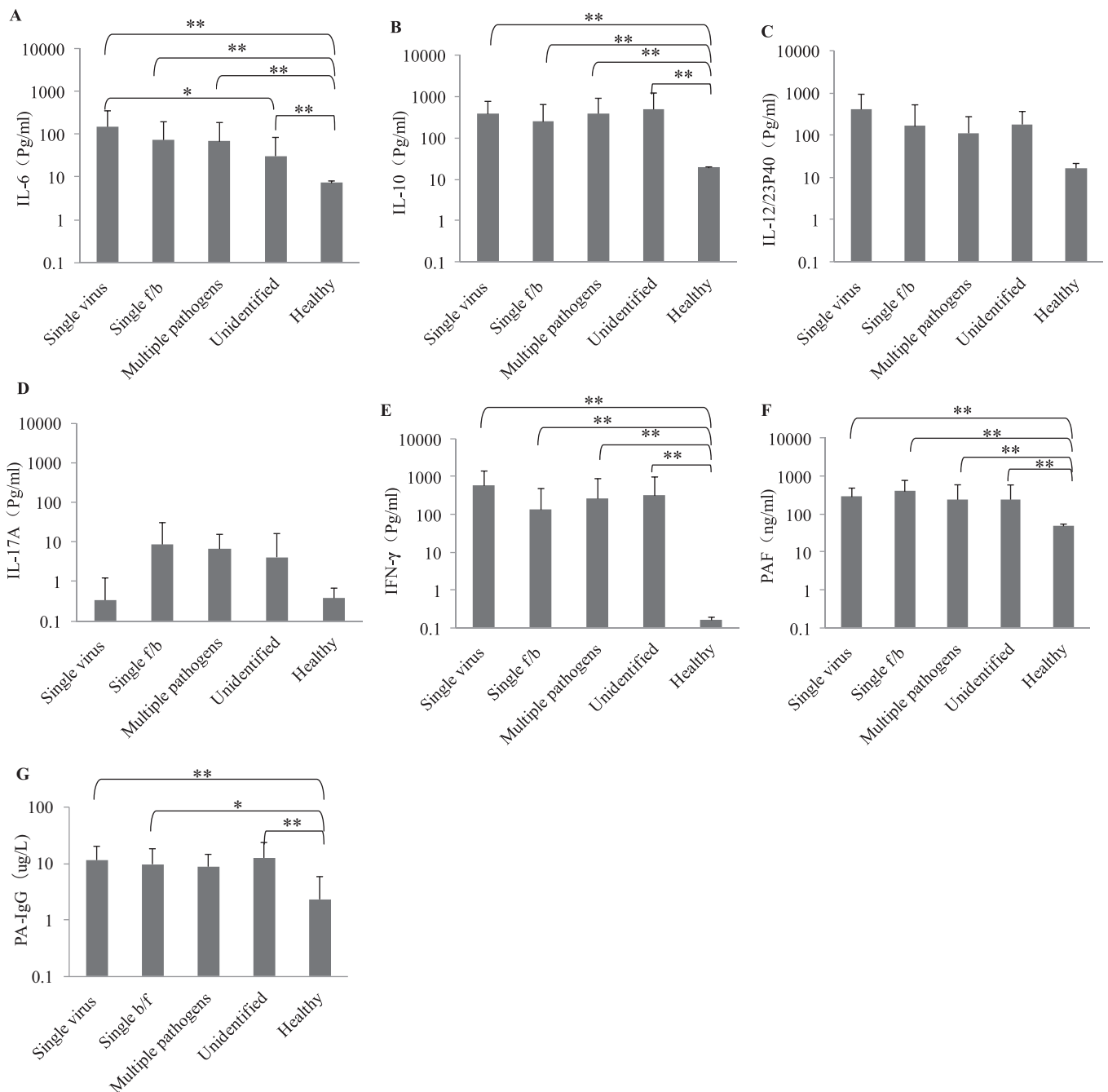


Fig. 1. The serum cytokines and PA-IgG levels in different groups and comparison (Single virus: $n = 7$; Single f/b: $n = 37$; Multiple pathogens: $n = 15$; Unidentified: $n = 20$; Healthy: $n = 10$). The bars represent the mean of each group of data, and the vertical line represents the error line and the end is the standard deviation value. * and ** indicate the statistical significant at $P < .05$ and $P < 0.01$, respectively.

potentiate each other's effects in vivo to modulate the immune response [39]. The strong associations were found between PAF and IL-6, between PAF and IL-10, between PAF and CRP, revealing the PAF, interacting with IL-6 and IL-10, are jointly responsible for thrombocytopenia [25].

Both antibody-mediated and/or T cell-mediated platelet destruction are key processes of thrombocytopenia [40]. Through antibody-induced immune-mediated phagocytosis and complement activation, alloantibodies destroy platelets and eliminate them from the circulation [41]. In as many as 30% to 40% of the patients, no detectable antibodies can be found, however, studies have shown that antibodies can cause thrombocytopenia even if the platelet-associated antibody levels in humans are lower than the detection limits of conventional methods.

Therefore, the detection of the relevant antibodies is the primary condition for diagnosis and treatment [42] as the presence of auto-antibodies targeting GPIb-IX is an effective predictor for refractoriness to steroid or IVIG therapy [43]. In this study, only 2 patients were negative in the PA-IgG test in 79 selected patients, indicating that antibody-mediated clearance maybe the main reason for our patients with fever and thrombocytopenia.

Although an informal connection between *H. pylori* infection and ITP has been suggested by various clinical studies, there are studies that have demonstrated that *H. pylori* is not associated with thrombocytopenia [44]. The curative effects of anti-*H. pylori* therapy on improved blood platelet quantity in chronic immune thrombocytopenia is inconsistent [45,46]. In this study, *H. pylori* has caused the highest

Table 3
The spearman correlation coefficients of six cytokines in patients with fever and thrombocytopenia.

	IL-6 (pg/ml)	IL-10 (pg/ml)	IL-12/23P40 (pg/ml)	IL-17A (pg/ml)	IFN- γ (pg/ml)	PAF (ng/ml)	PLT ($\times 10^9/l$)	MPV (fl)	PDW (fl)	CRP (mg/l)	PCT (ng/ml)
IL-6	-										
IL-10	$r = 0.458^{**}$ $P = .000$	-									
IL-12/23P40	$r = 0.009$ $P = .937$	$r = 0.298^{**}$ $P = .008$	-								
IL-17A	$r = -0.018$ $P = .875$	$r = 0.167$ $P = .142$	$r = 0.116$ $P = .310$	-							
IFN- γ	$r = 0.104$ $P = .360$	$r = 0.446^{**}$ $P = .000$	$r = 0.447^{**}$ $P = .000$	$r = 0.092$ $P = .419$	-						
PAF	$r = 0.370^{**}$ $P = .001$	$r = 0.257^*$ $P = .022$	$r = 0.062$ $P = .589$	$r = -0.09$ $P = .433$	$r = -0.059$ $P = .604$	-					
PLT	$r = -0.040$ $P = .726$	$r = -0.331^{**}$ $P = .003$	$r = 0.212$ $P = .061$	$r = 0.049$ $P = .668$	$r = -0.065$ $P = .571$	$r = 0.125$ $P = .271$	-				
MPV	$r = -0.129$ $P = .258$	$r = 0.280$ $P = .807$	$r = -0.137$ $P = .229$	$r = 0.012$ $P = .915$	$r = 0.019$ $P = .869$	$r = 0.024$ $P = .831$	$r = -0.179$ $P = .115$	-			
PDW	$r = 0.080$ $P = .486$	$r = -0.007$ $P = .952$	$r = -0.077$ $P = .500$	$r = -0.024$ $P = .836$	$r = -0.057$ $P = .617$	$r = 0.156$ $P = .171$	$r = -0.231^*$ $P = .040$	$r = -0.058$ $P = .614$	-		
CRP	$r = 0.667^{**}$ $P = .000$	$r = 0.148$ $P = .193$	$r = -0.083$ $P = .465$	$r = 0.028$ $P = .807$	$r = -0.057$ $P = .807$	$r = 0.256$ $P = .023^*$	$r = -0.031$ $P = .787$	$r = -0.134$ $P = .239$	$r = 0.203$ $P = .072$	-	
PCT	$r = 0.440^{**}$ $P = .000$	$r = 0.379^{**}$ $P = .001$	$r = -0.017$ $P = .880$	$r = -0.195$ $P = .086$	$r = 0.104$ $P = .363$	$r = 0.185$ $P = .103$	$r = -0.206$ $P = .069$	$r = -0.118$ $P = .301$	$r = 0.046$ $P = .687$	$r = 0.410^{**}$ $P = .000$	-

The level of statistical significance (P) was set at 5%.

* and ** indicate the statistically significant spearman correlation at the 0.05 level and 0.01 level, respectively.

Table 4
Loading scores of variables on significant PCA that combined components to predict cytokine profile, PA-IgG and clinical indicators in thrombocytopenia, and receiver operating characteristic curve analysis of components.

Variables	Components			
	1	2	3	4
PA-IgG ($\mu g/l$)	0.044	0.039	-0.016	0.811
IL-6 (pg/ml)	0.873	-0.013	0.012	0.025
IL-10 (pg/ml)	0.462	0.674	0.076	-0.024
IL-12/23P40 (pg/ml)	-0.010	0.250	0.783	0.011
IL-17A (pg/ml)	-0.143	-0.041	-0.009	0.659
IFN- γ (pg/ml)	-0.125	0.714	0.118	0.030
PAF (ng/ml)	0.421	0.058	0.194	-0.458
PLT ($\times 10^9/l$)	0.032	-0.615	0.521	0.055
MPV (fl)	-0.168	-0.028	-0.205	-0.302
PDW (fl)	0.158	0.111	-0.678	0.028
CRP (mg/l)	0.767	-0.147	-0.124	-0.053
PCT (ng/ml)	0.758	0.187	-0.118	-0.034
Receiver operating characteristic curve analysis				
AUC	0.521	0.358	0.505	0.564
P	0.745	0.031	0.937	0.333
Cut off value	0.150	-0.252	0.199	0.282
Sensitivity (%)	40	65.7	97.1	60
Specificity (%)	75	9.1	22.7	68.2

Abbreviations: AUC, areas under curve.

morbidity among bacteria inducing thrombocytopenia. There may be a cross-reaction between *H. pylori* cytotoxin-related gene (CagA) antibodies and platelet antigens, causing platelet-accelerated clearance. In *H. pylori*-positive patients, significantly higher levels of IL-6, IL-10, and PAF were found when compared with those in the *H. pylori*-negative group. Because of the involvement of cytokines, the pathogenesis of *H. pylori*-induced thrombocytopenia is more complex than initially believed. Unfortunately, the mechanism of *H. pylori*-induced thrombocytopenia is obscure. Revealing the interaction between *H. pylori* and thrombocytopenia will greatly expand the choices of therapeutic options.

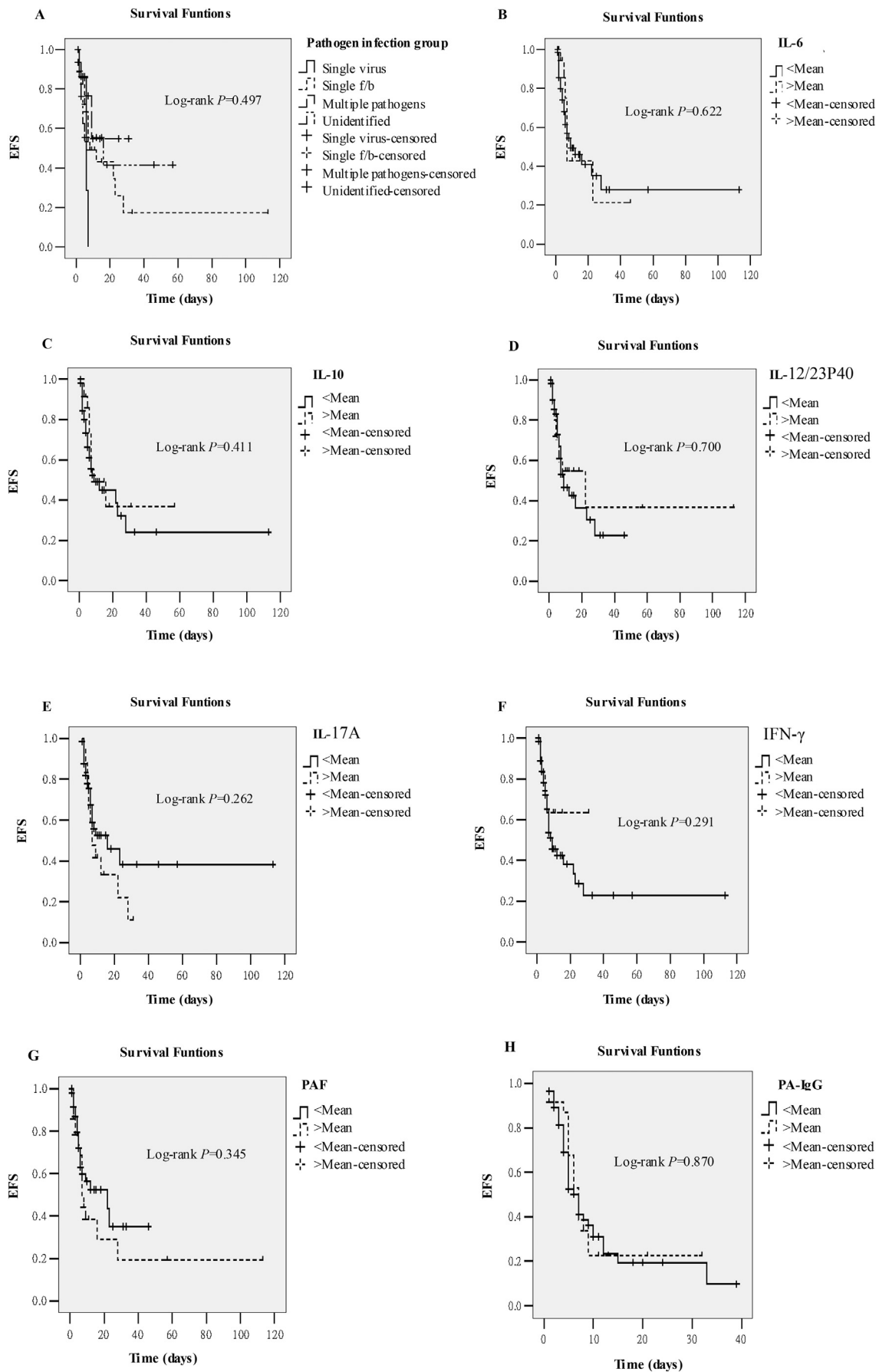
In this study, Kaplan-Meier survival curves were used to further assesses the prognosis of infectious thrombocytopenia. Our data showed that best therapeutic efficacy occurred in the single f/b

Table 5
Binary logistic regression analysis of risk factors in patients with or without improvement.

Factors	P value	OR	95.0% C.I. for OR	
			Lower	Upper
Age	0.029	0.956	0.918	0.995
T	0.504	1.458	0.483	4.404
WBC	0.483	1.063	0.896	1.261
LY	0.055	1.087	0.998	1.183
N	0.067	1.072	0.995	1.154
RBC	0.558	2.588	0.108	62.253
HB	0.337	0.949	0.853	1.056
PLT	0.005	1.081	1.024	1.141
MPV	0.472	1.108	0.838	1.467
PDW	0.060	1.483	0.984	2.237
CRP	0.591	1.003	0.993	1.013
PCT	0.040	1.031	1.001	1.061
ALT	0.741	0.999	0.991	1.006
IL-6	0.180	0.992	0.981	1.004
IL-10	0.702	1.000	0.998	1.003
IL-12/23P40	0.075	0.998	0.995	1.000
IL-17A	0.040	1.101	1.004	1.207
IFN- γ	0.583	1.000	0.998	1.001
PAF	0.092	1.002	1.000	1.004
PA-IgG	0.923	1.004	0.918	1.099

OR, Odds ratio.

infectious group, and patients in this group had the highest EFS. The worst therapeutic efficacy was found in the multiple infectious group. Platelet-virus interaction can occur via a variety of receptors expressed on platelets, which are mainly mediated by toll like receptors (TLRs) [47]. Accordingly, different pathogens may interact with its special TLRs on platelets. It is postulated that pathogens recognized by TLRs increase platelet destruction, which is multiplied in the presence of anti-platelet antibodies due to the synergy with TLRs, and then increases R-Fc-mediated phagocytosis [48]. The TLRs activate different signaling pathways and affect cells producing inflammatory cytokines [49]. Therefore, the cytokine production may vary due to the different types of microbiomes. As cytokines involved in immune function adjustment and modulation of the anti-platelet antibody response by B



(caption on next page)

Fig. 2. Kaplan-Meier survival curves were used to assess the prognosis of different types of pathogen infection groups and the applicability of six serum cytokines and PA-IgG as prognostic factors. (A) The EFS for the single virus group was 7 days shorter than for the multiple pathogens group (EFS = 31 days), followed by the unidentified group for 57 days. The maximum EFS in single f/b group was 113 days; (B) The mean of the IL-6 is 69.90 pg/ml, EFS in patients with IL-6 > 69.90 pg/ml was shorter than the patients with IL-6 < 69.90 pg/ml. (C) The mean of the IL-10 is 353.80 pg/ml, EFS in patients with IL-10 > 353.80 pg/ml was shorter than the patients with IL-10 < 353.80 pg/ml. (D) The mean of the IL-12/23P40 is 182.50 pg/ml, EFS in patients with IL-12/23P40 > 182.50 pg/ml was longer than the patients with IL-12/23P40 < 182.50 pg/ml. (E) The mean of the IL-17A is 6.33 pg/ml, EFS in patients with IL-17A > 6.33 pg/ml was shorter than the patients with IL-17A < 6.33 pg/ml. (F) The mean of the IFN- γ is 246.07 pg/ml EFS in patients with IFN- γ > 246.07 pg/ml was shorter than the patients with IFN- γ < 246.07 pg/ml. (G) The mean of the PAF is 321.80 ng/ml, EFS in patients with PAF > 321.80 ng/ml was longer than the patients with PAF < 321.80 ng/ml. (H) The mean of the PA-IgG is 11.164 μ g/l, EFS in patients with PA-IgG > 11.164 μ g/l was shorter than the patients with PA-IgG < 11.164 μ g/l.

cells [50,51], they may closely related to the platelet destruction. Therefore, we used the levels of cytokines as a stratification variable for EFS, the results showed that cytokine play a significant role in EFS. Patients with lower levels of IL-6, IL-10, IL-17A, and IFN- γ tended to have longer EFS than in patients with higher levels of cytokines. Previous study revealed that IL-22 is elevated in active phases of disease, and decreases in patients responsive to dexamethasone [52]. These results indicate that cytokines could reflect the disease status and the therapeutic effect.

The etiology of thrombocytopenia with a viral/bacterial infection may also be due to non-immunological mechanisms. The virus-induced pro-inflammatory environment itself often leads to further platelet activation in viremic patients [53]. Viruses can trigger a decrease in platelet production by infection of megakaryocytes and hematopoietic stem cells to decrease platelet formation; additionally, virus-infected platelets can expose P-selectin or phosphatidylserine, promoting their phagocytosis by macrophages [35]. Another viable mechanism is the formation of mixed platelet-leukocyte aggregates in adenovirus infection [54] that rapidly reduce platelet counts [55].

5. Conclusions

The etiology, immunologic and clinical features of patients with fever and thrombocytopenia is quite complicated, and multiple factors including type of pathogens, cytokines and autoantibody participate in the platelet's clearance. Finding reliable biomarkers for diagnosis and prognosis prediction are urgently needed for effective therapy.

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Authors' contributions

Huanzhu Chen and Chun Lin are responsible for the collection of case data, sample collection and testing; Ping He, Zhiqiang Fan, Wenjun Yu and Manxiong Cao are responsible for data collection, collation and statistics; Changwen Ke, Xiaoyang Jiao are responsible for the design of the experiment and the writing of the manuscript.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Ethics approval and informed consent

This study was approved by the Ethics Committee of Shantou University Medical College and informed consents were obtained from the patients.

Consent for publication

We have read and agree to the final draft before submission.

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