Viral infection in community acquired pneumonia patients with fever: a prospective observational study

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Background: Patients with community acquired pneumonia (CAP) caused by viruses can develop severe complications, which result in hospitalization and death. The purpose of this study was to analyse the aetiology, incidence, clinical characteristics, and outcomes of CAP patients with fever during non-pandemics, and then to provide theoretical basis for accurate diagnosis and treatment in CAP patients.

Methods: An enrolment system was established for monitoring the CAP patients with fever. Multiplex polymerase chain reaction (mPCR) kits were used to detect 10 viruses [influenza A and B, adenovirus (ADV), respiratory syncytial virus (RSV) A and B, picornavirus, parainfluenza virus (PIV), coronavirus, human metapneumovirus (HMPV), and bocavirus]. Data on age, gender, underlying diseases, complications, laboratory indexes, and outcomes were collected by physicians.

Results: This prospective study included 320 patients with fever. Among them, 23.4% were viral-positive by mPCR, with influenza virus most prominent followed by picornavirus. Strong variation in seasonal distribution was shown in viral infections, with peak months from December to February. Patients with influenza infection were likely to be taken to emergency rooms and have respiratory failure with higher creatinine kinase levels and lower white blood cell counts. Streptococcus pneumoniae followed by haemophilus influenzae were the most common bacteria in viral co-infections, which accounted for one third of virus-positive patients. Viral CAP and mixed CAP were not independent factors for death. In addition, lactate dehydrogenase (LDH) >246 IU/L [odds ratio (OR) =7.06, 95% confidence interval (CI): 2.15–23.2, P=0.001], and serum calcium <2.18 mmol/L (OR =6.67, 95% CI: 1.42–31.3, P=0.016) were associated with death

Conclusions: Viruses play an important role in CAP patients with fever, a systematic clinical, radiological and biological analysis of these patients can contribute to effective therapy that may prevent the development of CAP and improve the outcomes. The present work showed an elaborate analysis evidence of viral infection among fever CAP inpatients.

Keywords: Community acquired pneumonia (CAP); fever; multiplex polymerase chain reaction (mPCR); viruses

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Introduction

Community acquired pneumonia (CAP) is a common disease, with a reported mortality rate of 7.3% in Asia, based on a large prospective multicentre study (1). In Germany, the mortality rate of adult CAP was 8.6%, outpatients and inpatients had the mortality rate of 0.8% and 12.2% respectively (2). The plague of influenza A (H1N1) in 2009 caused serious viral pneumonia with high mortality rates in a large part of the world. In addition, severe hypoxemic acute respiratory failure and acute respiratory distress syndrome (ARDS) were strongly associated with influenza A (H1N1) in 2009 (3,4). The influenza outbreak was so severe that extracorporeal membrane oxygenation (ECMO) was used to decrease death (5,6).

A multiplex polymerase chain reaction (mPCR) method was a quick and effective diagnostic strategy (7,8) used by many clinical researchers and led to higher positive rates of pathogens associated with pneumonia. Compared with traditional cell culture, antigen assays, and serology, mPCR has improved accuracy and efficiency for the viral diagnosis (9). Many studies have focused on the incidence and aetiology of CAP patients seen at emergency departments and the intensive care unit (ICU) (10,11), and analysed the clinical characteristics and outcomes of viral CAP. To our knowledge, this is a prospective study on the pathogens of CAP inpatients with a fever using an established enrolment system. Most CAP patients suffer from fever, which is typical for many diseases, but there are few studies on virus-associated CAP patients with fever in non-epidemic seasons. In this study, we examined the incidence of viral pneumonia in CAP inpatients with fever and the associated risk factors or indicative laboratory indexes for viral infection and outcomes, such as death during hospitalization

Methods

Patients

This prospective study was conducted through our enrolment system: (I) feverish adult (body temperature ≥37.5 °C, aged ≥18 years inpatients in Shanghai pulmonary hospital from January 2015 to October 2016 were enrolled in this system; (II) chest X-rays or CT scans were conducted as soon as all the fever patients were admitted to the hospital. A new infiltrate on the chest radiograph of these fever patients was diagnosed as CAP, according to the criteria of the British Thoracic Society (12);

(III) nasopharyngeal swab and sputum samples were collected from enrolled CAP patients after admission to the hospital; (IV) mPCR kits were used to detect viruses in both nasopharyngeal swabs and sputum samples. During the study period, samples of swab and sputum were done for mPCR within 72 hours after collection; (V) clinical data were systematically collected by physicians and analysed using statistical software. The CURB-65 score consists of 5 points: confusion, BUN >7 mmol/L, respiratory rate >30 breaths/min, systolic blood pressure <90 mmHg and/or diastolic blood pressure ≤60 mmHg, and age ≥65 yrs (13). This project was approved by the ethical committee of Shanghai Pulmonary Hospital.

Samples and detection method

After admission to the hospital, we collected all the nasopharyngeal swab and sputum samples from the patients for viral detection. Samples of swab and sputum were done for mPCR within 72 hours after collection. Viral nucleic acids were purified from each nasopharyngeal swab or sputum specimen using the Axyprep Body Fluid Viral DNA/RNA Miniprep Kit, following the manufacturer's instructions. The resultant DNA/RNA was eluted with 60 µL elution buffer and stored at −80 °C until testing. Ten kinds of viral DNA/RNA [influenza A and B, adenovirus (ADV), respiratory syncytial virus (RSV) A and B, picornavirus, parainfluenza virus (PIV), coronavirus, human metapneumovirus (HMPV), and bocavirus] were detected using the one step primeScript RT-PCR kit (Perfect Realtime, DRR064; Takara) according to the manufacturer's instructions. The RT-PCR assay conditions were: 42 °C for 5 min, 95 °C for 10 s, 40 cycles of 95 °C for 3 s and 60 °C for 25 s with the CFX96 real-time PCR detection system (Bio-Rad). Blood, bronchoalveolar lavage fluid, and sputum samples were collected to obtain general information on the bacteria, which were cultured by the laboratory clinicians of our hospital.

Microbiological diagnosis

Virus infection confirmed by positive RT-PCR on nasopharyngeal swab or sputum specimen. Bacterial infection was identified by one of the following sample culture positive: (I) bronchoalveolar lavage fluid (threshold 10^4 CFU/mL); (II) blood culture; (III) sputum culture (considering >25 neutrophils and <10 epithelial cells as a valid specimen, threshold 10^5 CFU/mL).

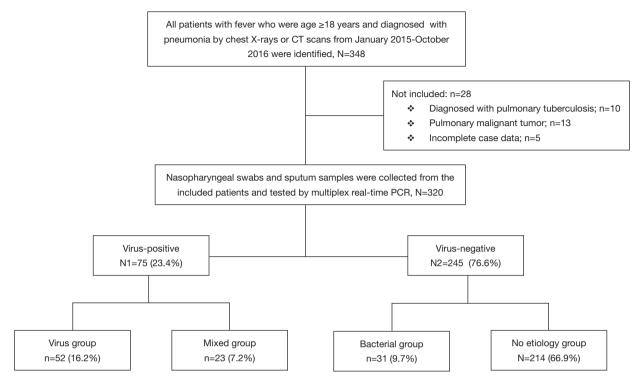


Figure 1 Flow chart of recruitment of adult fever community-acquired pneumonia and aetiology detection of by multiplex PCR. PCR, polymerase chain reaction.

Statistical analysis

Clinical data, including demographic variables such as age, gender, clinical symptoms, underlying diseases and complications, severity score assessed by CURB65, laboratory indexes, and outcomes were collected by our physicians. Statistical analyses were performed in SPSS (version 24.0). Dichotomous variables were analysed with chi-squared or Fisher's exact test. The Mann-Whitney U test or Kruskal-Wallis method was used to analyse numerical data. Risk factors associated with severe situations, death during hospitalization, analysed by multiple logistic regression. Logistic regression analysis was also used to examine the risk factors associated with influenza A viral infection. A P value ≤0.05 was considered statistically significant.

Results

The 320 patients enrolled in our study were divided into four categories (viral group, mixed group, bacterial group, and no pathogen group) (*Figure 1*) according to the causative agent of pneumonia, and we also classified

all the patients into two sub-groups (viral-positive and viral-negative) based on viral infection or not. The characteristics of clinical symptoms, complications, and outcomes according to the aetiology diagnosis are shown in Tables 1,S1. In this study, 64.7% were male with a median age of 60 years (IQR, 53-68 years). There was no statistically significant difference among the four categories of underlying diseases. Approximately one third of patients in the mixed-group had a CURB65 ≥2 score that was higher than the other three groups. The median hospitalization span was 9 days (IQR, 7.0-13.0 days). When examining outcomes according to the pathogens (Table 1), the mixedgroup had the highest incidences of death in the hospital. Laboratory indexes are shown in Table 1. As to the two subgroups (viral-positive and viral-negative), there were 71 (22.2%) and 28 (8.8%) patients with respiratory failure and ARDS, respectively, and the virus-positive group had a larger population for these two complications than the virus-negative group (Table S1). Virus-positive patients had higher serum lactic dehydrogenase and creatine kinase (CK) compared with virus-negative patients (Table S1).

The most common virus detected in CAP patients with fever was influenza A, followed by picornavirus. The

Table 1 Clinical characteristics, complications, laboratory indexes, and outcomes of 320 CAP patients with fever, according to the aetiology diagnosis

Parameters	All patients (N=320)	Viral group (n=52, 16.2%)	Mixed group (n=23, 7.2%)	Bacterial group (n=31, 9.7%)	No aetiology group (n=214, 66.9%)	P value
Age, y	60 [53–68]	58 [47–68]	61 [56–72]	61 [54–72]	60 [52–67]	0.4
Male sex	207 (64.7)	33 (63.5)	16 (69.6)	19 (61.3)	139 (65.0)	0.94
Smoking history	120 (37.5)	14 (26.9)	10 (43.5)	8 (25.8)	88 (41.1)	0.12
Underlying disease						
COPD	52 (16.3)	9 (17.3)	3 (13.0)	5 (16.1)	35 (16.4)	0.99
Asthma	15 (4.7)	1 (1.9)	1 (4.3)	2 (6.5)	11 (5.1)	0.74
Bronchiectasis	68 (21.3)	10 (19.2)	4 (17.4)	7 (22.6)	47 (22.0)	0.94
Diabetes mellitus	38 (11.9)	2 (3.8)	2 (8.7)	7 (22.6)	27 (12.6)	0.07
Interstitial lung disease	35 (10.9)	2 (3.8)	4 (17.4)	3 (9.7)	26 (12.1)	0.22
Heart disease	30 (9.4)	3 (5.8)	5 (21.7)	4 (12.9)	18 (8.4)	0.13
Immunodeficiency	27 (8.4)	5 (9.6)	3 (13.0)	2 (6.5)	17 (7.9)	0.74
Renal disease	48 (15.0)	9 (17.3)	2 (8.7)	6 (19.4)	31 (14.5)	0.7
Antibiotics treatment outside	168 (52.5)	30 (57.7)	10 (43.4)	17 (54.8)	111 (51.9)	0.67
CURB65 ≥2	34 (10.6)	4 (7.7)	7 (30.4)	4 (12.9)	19 (8.9)	0.025*
Clinical symptoms						
Cough	295 (92.2)	45 (86.5)	23 (100.0)	27 (87.1)	200 (93.5)	0.1
Chills	30 (9.4)	7 (13.5)	1 (4.3)	2 (6.5)	20 (9.3)	0.63
Dyspnoea	169 (52.8)	31 (59.6)	16 (69.6)	16 (51.6)	106 (49.5)	0.22
Myalgia	11 (3.4)	5 (9.6)	1 (4.3)	0	5 (2.3)	0.06
Complications						
Respiratory failure	71 (22.2)	19 (36.5)	7 (30.4)	8 (25.8)	37 (17.3)	0.02*
ARDS	28 (8.8)	8 (15.4)	5 (21.7)	3 (9.7)	12 (5.6)	0.01*
Thrombocytopenia	16 (5.0)	4 (7.7)	3 (13.0)	3 (9.7)	6 (2.8)	0.02*
Pleural effusion	102 (31.9)	15 (28.8)	7 (30.4)	16 (51.6)	64 (29.9)	0.1
Laboratory indexes						
PCT, ng/mL	0.05 (0.02-0.13)	0.04 (0.03-0.1)	0.03 (0.02-0.08)	0.11 (0.04–0.2)	0.04 (0.02-0.1)	0.04*
ESR, mm/h	34 (14.0–59.0)	32 (15.5–53.5)	33 (10.0–65.0)	61.5 (30.3–77.3)	33.5 (13.3–56.0)	0.06
CRP, mg/L	19.2 (2.0-75.0)	11.9 (3.0–45.3)	10.3 (2.0–72.0)	42 (9.1–93.9)	8.7 (1.9–57.8)	0.02*
WBC <4×10 ⁹ /L	21 (6.6)	7 (13.5)	4 (17.4)	1 (3.2)	9 (4.2)	0.01*
IL-6, pg/mL	20.1 (8.0-52.0)	22.5 (9.0-68.6)	11 (8.0–46.5)	39.5 (9.3–72.5)	19 (7.3–43.5)	0.64
TNF-α, U/mL	35.8 (21.0–55.0)	33 (15.8–9.1)	35 (10.0–104.0)	30.7 (19.4–56.7)	37 (21.1–51.7)	0.97
Serum potassium, mmol/L	4 (3.6–4.3)	3.9 (3.6-4.2)	3.8 (3.5–4.1)	4 (3.4–4.5)	4 (3.7–4.3)	0.4
Serum calcium, <2.18 mmol/L	151 (47.2)	31 (59.6)	11 (47.8)	17 (54.8)	92 (43.0)	0.14
LDH >246, IU/L	98 (30.6)	23 (44.2)	11 (47.8)	10 (32.3)	54 (25.2)	0.007*
CK >294, IU/L	30 (9.4)	8 (15.4)	7 (30.4)	4 (12.9)	11 (5.1)	<0.001*
PLT <100×10 ⁹ /L	12 (3.8)	5 (9.6)	1 (4.3)	3 (9.7)	3 (1.4)	0.005*

Table 1 (continued)

Table 1 (continued)

(/						
Parameters	All patients (N=320)	Viral group (n=52, 16.2%)	Mixed group (n=23, 7.2%)	Bacterial group (n=31, 9.7%)	No aetiology group (n=214, 66.9%)	P value
Outcomes						
Emergency treatment	52 (16.3)	16 (30.8)	5 (21.7)	7 (22.6)	24 (11.2)	0.004*
ICU	9 (2.8)	3 (5.8)	1 (4.3)	2 (6.5)	3 (1.4)	0.15
Invasive mechanical ventilation	9 (2.8)	1 (1.9)	3 (13.0)	4 (12.9)	1 (0.5)	<0.001*
Length of stay, d	9 (7.0–13.0)	11 (8.0–14.0)	8 (6.5–13.5)	10.5 (7.0–13.5)	8 (7.0–12.0)	0.03*
Death during hospitalization	21 (6.6)	0	6 (26.1)	4 (12.9)	11 (5.1)	<0.001*

Data are presented as the median (first through third quartiles) or number (%). P values refer to the comparison among viral community acquired pneumonia (CAP), bacterial CAP, mixed CAP, and no pathogen CAP. *, P<0.05. COPD, chronic obstructive pulmonary disease; CURB65, confusion of new onset, blood urea nitrogen >7 mmol/L, respiratory rate ≥30 breaths per minute, blood pressure <90 mmHg systolic or ≤60 mmHg diastolic, age ≥65 years; ARDS, acute respiratory distress syndrome; PCT, procalcitonin; ESR, erythrocyte sedimentation rate; CRP, C-reactive protein; LDH, lactate dehydrogenase; CK, creatine kinase; PLT, platelets; ICU, intensive care unit.

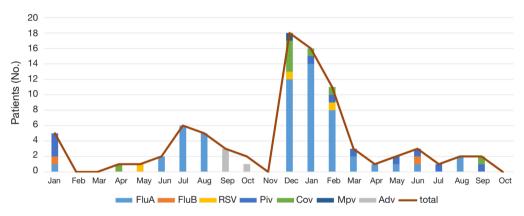


Figure 2 Seasonal distribution of detected viruses in nasopharyngeal swabs or sputum samples by multiplex PCR in 320 patients enrolled in the management system from January 2015 to October 2016. PCR, polymerase chain reaction.

epidemic seasons are autumn and winter and viral detection peaks were in December and February (*Figure 2*). The distribution of the highest body temperature during the whole course, according to the pathogen group, is shown in *Figure S1*. The viral-group and mixed group were more likely to have a higher fever (T >39 °C) than the no pathogen group. The mPCR positive results for nasopharyngeal swab samples and sputum samples are shown in *Table 2*. The causes of bacterial co-infections in the mixed group are shown in *Table S2*. Streptococcus pneumoniae followed by haemophilus influenzae were the most common bacteria in co-infections.

Influenza virus was critical for viral pneumonia in our study. In addition, CK >294 IU/L, WBC <4×10°/L, emergency admission, and respiratory failure were

independent factors associated with influenza infection in multivariate logistic regression analysis (*Table 3*). In this study, we also analysed two risk factors for outcomes of CAP patients with fever, i.e., death. Viral CAP and mixed CAP were not independent factors for death. In addition, lactate dehydrogenase (LDH) >246 IU/L, and serum calcium <2.18 mmol were associated with death (*Table 4*).

Discussion

The purpose of this study was to analyse the aetiology, incidence, clinical characteristics, and outcomes of CAP patients with fever during non-pandemics in one pulmonary hospital. This study revealed there was a viral aetiology in 75 (23.4%) of 320 fever patients diagnosed with CAP. This

Table 2 Detection rates of all the viruses by multiplex real-time PCR from nasopharyngeal swabs and sputum samples

Virus	All	Nasopharyngeal swab samples (%)	Sputum samples (%)
			,
Cases, No.	320	320	230
Total virus, No.	84	45	66
Influenza	57 (17.8)	31 (9.7)	43 (18.7)
Influenza A	55 (17.2)	30 (9.4)	41 (17.8)
Influenza B	2 (0.6)	1 (0.3)	2 (0.9)
Adenovirus	4 (1.3)	2 (0.6)	3 (1.3)
Respiratory syncytial virus	3 (0.9)	1 (0.3)	3 (1.3)
Picornavirus	10 (3.1)	6 (1.9)	9 (3.9)
Coronavirus	8 (2.5)	5 (1.6)	6 (2.6)
Metapneumovirus	2 (0.6)	0 (0)	2 (0.9)

Data are presented as number (%).

Table 3 Factors associated with influenza CAP by multivariate logistic analysis

Variables	OR	95% CI	P value
CK >294 IU/L	2.53	1.02-6.29	0.046*
WBC <4×10 ⁹ /L	4.02	1.48–10.9	0.006*
Emergency	2.56	1.22-5.38	0.01*
Respiratory failure	2.37	1.21-4.67	0.01*

^{*,} P<0.05. CAP, community-acquired pneumonia; CK, creatine kinase; WBC, white blood cells; respiratory failure, arterial partial pressure of oxygen (PaO₂) <60 mmHg or percutaneous oxygen saturation (SpO₂) <90% without oxygen therapy; OR, odds ratio; 95% CI, 95% confidence interval.

Table 4 Risk factors for death in fever CAP patients

Variables	OR	95% CI	P value
Pathogen			
No pathogen	Ref	_	
Viral	0.19	0.02-1.62	0.13
Mixed group	3.72	0.86-16.1	0.08
Bacterial	2.43	0.63-0.36	0.19
CURB65 >2	3.13	0.98-9.96	0.053
LDH >246 IU/L	7.06	2.15-23.2	0.001*
Serum calcium <2.18 mmol/L	6.67	1.42–31.3	0.016*

^{*,} P<0.05. CAP, community-acquired pneumonia; Ref, reference; OR, odds ratio; 95% CI, 95% confidence interval.

is in line with other studies, where the rates of viral infection in CAP patients varied from 23-56% (10,11,14,15). The differences in aetiologies of CAP patients between studies is likely due to the different test methods, the diversity of the aetiologies detected, and the distinctions of different regions and populations. During the observational period in our study, the viral infection peak months were from December to February. The predominant virus was influenza A followed by picornavirus, and streptococcus was the most common bacteria in viral co-infection cases, which is consistent with the results from other studies (16). de Roux et al. reported 65% of mixed pneumococcal infections were associated with a respiratory virus in hospitalized patients, mainly the influenza virus (17). Another study suggested that pneumococcus is critical for the development of pneumonia associated with influenza virus A, RSV, PIV, and ADV (18).

In our study, most viral-positive patients, with or without bacterial co-infection, had high temperatures. This finding is in contrast to the common opinion that high temperatures frequently occur in patients with bacterial or mixed infections. This discrepancy may be attributed to the difficulty in identifying a bacterial infection because of the use of antibiotics before testing. Although, this result also indicated that fever cannot be an empirical indicator for antibiotic treatment because a bacterial infection may have been misdiagnosed before any diagnostic tests.

In our study, nasopharyngeal swabs and sputum samples were collected for mPCR. Only 230 patients had sputum

samples, which had high detection rates for viruses, suggesting there is a lower viral load in nasopharyngeal samples from CAP patients (19). Furthermore, this also suggested that sensitivity of viral pathogen identification could be enhanced by simultaneous examination both NPS and sputum samples rather than only NPS samples (20). Influenza virus is the most common virus in most reports; although, the presence of myalgia and neutropenia were associated with influenza pneumonia in the study by Jennings et al. (21). Furthermore, we identified that having influenza pneumonia increased the odds of emergency room visits and respiratory failure; while, previous reports revealed that patients admitted with influenza who have pneumonia are more likely to be admitted to the ICU or die (22,23). The more accurate the detection of influenza infection, the more correct the use of antivirals will be and unnecessary antibiotics use will be reduced among patients. In our study, high levels of CK were also associated with influenza pneumonia; therefore, CK may be a predictive marker in influenza infection. The combined use of biologic markers and viral testing may correctly identify patients and safely limit antibiotic exposure.

Due to pre-hospitalization antibiotic usage in the majority of subjects, the rates of bacterial and viral-bacterial detection may be underestimated in our study. Interesting, in our research, only one case had bacteremia which was diagnosed with positive blood culture, which was lower than other studies (10,24,25). Approximately one third of subjects with viral pneumonia had bacterial co-infection. This rate was lower than found in other studies (10), but we observed mild-to-moderate patients; while, other studies focused on severe CAP patients, which may have led to differences. In our study, we found that the co-infection group had the highest death rate (21.6%) compared with the other three groups, but co-infection was not an independent risk factor for death during hospitalization. The lower mortality in our mild-to-moderate subjects should be re-inspected in a much larger study group. The mixed group was also identified as an independent risk factor for complicated courses (death and invasive mechanical ventilation) in ICU samples (10). This trend may be explained by the finding that influenza virus (predominant virus in co-infection with bacterium) neuraminidase could contribute to excess mortality in bacterial pneumonia (26). Possible explanations for mixed CAP progressing into severe cases include epithelial damage, changes in airway function, up-regulation of receptors, and changes in the innate immune response (27). Previous study reports that patients with co-infection were

much more likely to have a pneumonia severity index (PSI) score of IV or V and a longer length of stay (10). The influenza outbreak was so severe that ECMO was used to decrease death. Influenza plays a dominant role in co-infection, which was so severe that needed effective means to decrease death. The Influenza patients diagnosed with severity score CURB 65 and detection method mPCR can be admitted to ICU sooner and start treatment like ECMO and Tamiflu sooner with increased survival (5,6).

There were some limitations in our study. The single centre design restricted the observed populations. We used cultures instead of mPCR for bacterial detection. The mPCR method can detect dead as well as viable bacteria (14); therefore, the culture method would have lower positive rates of bacterial and co-infection. Moreover, we underestimated the role of atypical pathogens such as *C. pneumonia* and *M. pneumonia* that would manifest fever in most cases. The low mortality rate could impair the accuracy of the independent risk factors in the multiple logistic regression analysis.

In conclusion, we found strong variations in the virus distribution in non-pandemic years with relatively high viral rates in the CAP population with fever. It is likely that ARDS and respiratory failure are associated with viral infection, even though there are no definite predictors for causative virus in CAP. The role of co-infection in CAP is unclear. It was not an independent risk factor for death but LDH >246 IU/L [odds ratio (OR) =7.06, 95% confidence interval (CI): 2.15-23.2, P=0.001], and serum calcium <2.18 mmol/L (OR =6.67, 95% CI: 1.42-31.3, P=0.016) were associated with death. The implications of positive testing can offer the physicians accurate information with early treatment of influenza, and then to prevent progression to CAP and respiratory failure. More studies on the treatment of viral CAP are necessary because effective antiviral therapy may prevent the development of CAP and improve the outcomes.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The study was approved by the ethical committee of Shanghai Pulmonary Hospital (K17-113) and written informed consent was obtained from all patients.

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Supplementary

Table S1 Clinical characteristics, complications, laboratory indexes, and outcomes of all 320 patients with CAP and fever and according to the virus-positive and virus-negative groups

Parameters	All patients (n=320)	Virus-positive group (n=75)	Virus-negative group (n=245)	P value
Age, y	60 [53–68]	60 [49–69]	60 [53–67]	0.67
Male sex	207 (64.7)	49 (65.3)	158 (64.5)	1
Smoking history	120 (37.5)	24 (32.0)	96 (39.2)	0.28
Underlying disease				
COPD	52 (16.3)	12 (16.0)	40 (16.3)	1
Asthma	15 (4.7)	2 (2.7)	13 (5.3)	0.53
Bronchiectasis	68 (21.3)	14 (18.7)	54 (22.0)	0.63
Diabetes mellitus	38 (11.9)	4 (5.3)	34 (13.9)	0.06
Interstitial lung disease	35 (10.9)	6 (8.0)	29 (11.8)	0.4
Heart disease	30 (9.4)	8 (10.7)	22 (9.0)	0.82
Immunodeficiency	27 (8.4)	8 (10.7)	19 (7.8)	0.48
Renal disease	48 (15.0)	11 (14.7)	37 (15.1)	1
Antibiotics treatment outside	168 (52.5)	40 (53.3)	128 (52.2)	0.62
CURB65 ≥2	34 (10.6)	11 (14.7)	23 (9.4)	0.2
Clinical symptoms				
Cough	295 (92.2)	68 (90.7)	227 (92.7)	0.62
Chills	30 (9.4)	8 (10.7)	22 (9.0)	0.82
Dyspnoea	169 (52.8)	47 (62.7)	122 (49.8)	0.06
Myalgia	11 (3.4)	6 (8.0)	5 (2.0)	0.02*
Complications				
Respiratory failure	71 (22.2)	26 (34.7)	45 (18.4)	0.004*
ARDS	28 (8.8)	13 (17.3)	15 (6.1)	0.005*
Thrombocytopenia	16 (5.0)	7 (9.3)	9 (3.7)	0.07
Pleural effusion	102 (31.9)	22 (29.3)	80 (32.7)	0.67
Laboratory indexes				
PCT, ng/mL	0.05 (0.02-0.13)	0.04 (0.02-0.08)	0.04 (0.02-0.1)	0.42
ESR, mm/h	34 (14.0–59.0)	32.5 (13.0–54.8)	35.5 (14.0–60.0)	0.47
CRP, mg/L	19.2 (2.5–75.2)	11.6 (2.6–53.3)	11 (1.9–72.5)	0.64
WBC <4×10 ⁹ /L	21 (6.6)	11 (14.7)	10 (4.1)	0.003*
IL-6, pg/mL	20.1 (7.95–52.0)	18.7 (9.2–62.4)	20.2 (7.6–51.0)	0.83
TNF-α, U/mL	35.8 (21.1–54.8)	33 (15.0–64.7)	36 (21.1–51.8)	0.81
Serum potassium, mmol/L	4 (3.6–4.3)	3.8 (3.6–4.15)	4 (3.6–4.3)	0.12
Serum calcium <2.18 mmol/L	151 (47.2)	42 (56.0)	109 (44.5)	0.06
LDH >246, IU/L	98 (30.6)	34 (45.3)	64 (26.1)	0.001*
CK >294, IU/L	30 (9.4)	15 (20.8)	15 (6.1)	0.001*
PLT <100, 4×10 ⁹ /L	12 (3.8)	6 (8.1)	6 (2.4)	0.04*
Outcomes				
Emergency treatment	52 (16.3)	21 (28.0)	31 (12.7)	0.002*
ICU	9 (2.8)	4 (5.3)	5 (2.0)	0.22
Invasive mechanical ventilation	9 (2.8)	4 (5.3)	5 (2.0)	0.22
Length of stay, d	9 [7–13]	10 [7–14]	8 [7–12]	0.1
Death during hospitalization	21 (6.6)	6 (8.0)	15 (6.1)	0.2

Data are presented as the median (first through third quartiles) or number (%). P values refer to the comparison between virus-positive and virus-negative fever patients diagnosed with CAP. *, P<0.05. COPD, chronic obstructive pulmonary disease; CURB65, confusion of new onset, blood urea nitrogen >7 mmol/L, respiratory rate \geq 30 breaths per minute, blood pressure <90 mmHg systolic or \leq 60 mmHg diastolic, age \geq 65 years; ARDS, acute respiratory distress syndrome; PCT, procalcitonin; ESR, erythrocyte sedimentation rate; CRP, C-reactive protein; LDH, lactate dehydrogenase; CK, creatine kinase; PLT, platelets; ICU, intensive care unit.

Table S2 Causative agents of viral co-infections in 320 adults with fever diagnosed with community-acquired pneumonia

lever diagnosed with community-acquired pheumoma				
Pathogens	Subjects (n)			
Influenza A				
Streptococcus pneumoniae	4			
Haemophilus influenzae	5			
Staphylococcus aureus	1			
Other staphylococci	3			
Other streptococci	1			
Pseudomonas aeruginosa	2			
Enterobacterium	1			
Acinetobacter baumannii + streptococcus pneumoniae	1			
Adenovirus				
Xanthomonas maltophilia	1			
Picornavirus				
Enterobacter aerogenes	1			
Staphylococcus	1			
Coronavirus				
Acinetobacter baumannii	1			
Klebsiella pneumoniae	1			
Total	23			

Staphylococcus epidermidis, staphylococcus saprophyticus and staphylococcus haemolyticus are considered into the other staph. And streptococcus pyogenes belongs to the other staph.

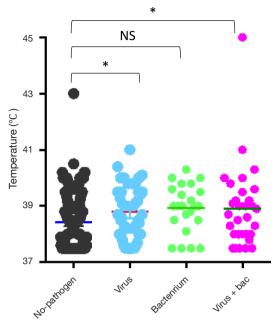


Figure S1 Distribution of the high temperature during the course of CAP according to the aetiology. *, P<0.05. CAP, community-acquired pneumonia.