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Retrospective Study

Coxsackievirus A6 was the most common enterovirus serotype causing hand, foot, and mouth disease in Shiyan City, central China

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Abstract**BACKGROUND**

Hand, foot, and mouth disease (HFMD) has become one of the most common infectious diseases in China. Before 2016, the primary causal serotypes were enterovirus A71 (EV-A71) and coxsackievirus A16 (CV-A16). Following the introduction of EV-A71 vaccines in China since 2016, the situation could change. CV-A6 has recently replaced EV-A71 and CV-A16 in some areas of China. However, the epidemiological characteristics of central China remain unknown.

AIM

To investigate the clinical symptoms and pathogen spectrum of HFMD in Shiyan City, central China, in recent years.

METHODS

The epidemiological, clinical, and laboratory data from HFMD cases reported to the Shiyan Center for Disease Control and Prevention between January 2016 and

December 2020 were analyzed. 196 throat swab specimens were collected from hospitalized HFMD patients between January 2018 and December 2020. To detect and genotype enteroviruses, real-time reverse transcription-polymerase chain reaction and sequencing of the 5'-untranslated region were used. In Shiyang, 168 laboratory-confirmed HFMD cases were studied using a logistic regression model to determine the effect of predominant enterovirus serotypes. Based on the logistic regression model, the least absolute shrinkage and selection operator model was used to analyze the correlation between CV-A6 infection and various clinical characteristics in HFMD patients in Shiyang.

RESULTS

From 2016 to 2020, 35840 HFMD cases were reported in Shiyang. The number of cases decreased by 48.4% from 2016 to 2017. Approximately 1.58-fold increases were found in 2018 and 2019 when compared to the previous year, respectively. In 2020, a decrease of about 85.5% was reported when compared to 2019. The most common serotypes shifted from EV-A71 and CV-A16 (about 60%-80% in 2016 and 2018) to others (more than 80.0% in 2017, 2019, and 2020). EV-A71 lost its dominance in 2017 in Shiyang. Among 196 confirmed HFMD cases, 85.7% tested positive for enterovirus, with CV-A6 being the most common serotype (121/168, 72.0%). The positive rates for CV-A16 and CV-A10 were 4.8% and 3.0%, respectively. There was no EV-A71 discovered. Infection with CV-A6 was linked to fever, myocardial damage, increased creatine kinase MB isoenzyme, and lactate dehydrogenase levels.

CONCLUSION

CV-A6 was the most common enterovirus serotype in Shiyang City, replacing EV-A71 and CV-A16 as the HFMD pathogen. Developing vaccines against CV-A6 or multiple pathogens, as well as rising CV-A6 surveillance, will help prevent HFMD in central China.

Key Words: Hand, foot, and mouth disease; Coxsackievirus A6; Enterovirus; Prevalence serotype; Fever; Myocardial damage

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Core Tip: Hand, foot, and mouth disease (HFMD) has become China's most common infectious disease since 2008. Enterovirus A71 (EV-A71) and coxsackievirus A16 (CV-A16) were the primary causal serotypes. However, the dominant serotypes shifted recently to other enteroviruses. Nevertheless, there has been no report on the prevalence of the pathogen spectrum in Hubei, central China, since 2017. Our findings revealed that CV-A6 was the most common HFMD pathogen in Shiyang City, displacing EV-A71 and CV-A16. Fever and myocardial damage were more common in HFMD caused by CV-A6.

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INTRODUCTION

Hand, foot, and mouth disease (HFMD) is one of the most widespread infectious diseases in the world. It disproportionately affects children under the age of five[1]. The Chinese Ministry of Health has designated HFMD as a category III notifiable infectious disease since 2008. The Chinese Center for Disease Control and Prevention (China CDC) has documented more than 20 million cases of HFMD from 2008 to 2018. The clinical presentation of HFMD is characterized by low-grade fever, a maculopapular or papulovesicular rash on the palms and soles of the feet, and severe oral ulcerations[2]. The illness is self-limiting, and the majority of children recover in a week without sequelae; however, a small proportion of children who experience neurologic or cardiopulmonary complications could be lethal[2].

As ever reported, the pathogens that caused HFMD are enterovirus-A (EV-A) and enterovirus-B (EV-B) species of the *Enterovirus* genus of the family *Picornaviridae*[3]. There are at least 23 serotypes, including nine coxsackievirus-A (CV-A) serotypes, enterovirus-A71 (EV-A71), six CV-B serotypes, six enteric cytopathogenic human orphan (ECHO) virus serotypes, and EV-B84[3]. The most common serotypes are CV-A16 and EV-A71, particularly in Asia-Pacific[4,5]. Furthermore, EV-A71 is also

responsible for the most severe and fatal cases[6,7].

However, since the first inactivated EV-A71 whole virus vaccine was launched in China in December 2015, the prevalence of EV-A71 has decreased dramatically[8,9]. A CV-A6-associated HFMD outbreak was reported in Finland in 2008[10]. Since then, CV-A6 has emerged as one of the pathogens responsible for the prevalence of HFMD in America, Europe, Africa, and Asia[11-14]. Between 2009 and 2017, several Chinese provinces reported sporadic HFMD cases caused by CV-A6[15-18]. Furthermore, the CV-A6 epidemic is becoming more widespread[14]. However, etiological data on HFMD in China after 2017 is scarce. There was even no report from Central China after 2017. Nonetheless, monitoring the circulating pathogens following the cumulative coverage of the EV-A71 vaccine is critical because it will help determine how to best implement a preventative strategy.

In this study, we focused on changes in the pathogen spectrum in Shiyan City, Hubei Province, between 2016 and 2020. We also analyzed the relationship between the primary pathogens and the clinical features of HFMD.

MATERIALS AND METHODS

Data collection

We obtained the surveillance data of HFMD in Shiyan city from 2016 to 2020 from Shiyan CDC. Since 2008, all clinically diagnosed and laboratory-confirmed HFMD cases have been reported to the local CDC within 24 h. The diagnostic criteria were based on the People's Republic of China Hygiene Industry Standard for Hand, Foot, and Mouth Disease Diagnosis (WS 588-2018)[19].

Patients and specimen collection

From 2018 to 2020, 196 hospitalized HFMD patients were enrolled in this study. The demographics of the patients, clinical data from their hospitalization, and throat swab specimens were all collected. The study was approved by the Institutional Review Board of the Taihe Hospital, Affiliated Hospital of Hubei University of Medicine (2017KS035). Participants' legal guardians signed informed consent forms.

Viral RNA extraction and real-time RT-PCR

Viral RNAs were extracted from the throat swab specimens using the Tianlong automatic nucleic acid extractor (NP968, Tianlong Science and Technology, Xi'an, China) and its corollary automatic nucleic acid extraction kit according to the manufacturer's instructions. In brief, 200 μ L specimens were used. The extracted RNAs were eluted in 100 μ L elution buffer and stored at -70 °C until used.

Real-time reverse transcription-polymerase chain reaction (RT-PCR) was performed to identify the enteroviruses in clinical samples using EV-A71 specific, CV-A16 specific, and pan-enterovirus real-time RT-PCR Kits (Tianlong Science and Technology).

5'-untranslated region amplification and sequencing for enterovirus serotypes identification

Enterovirus-positive samples were amplified for 5'-untranslated region (5'-UTR) and sequenced to identify the serotypes. The amplification was performed by RT-PCR using the 5'-UTR primers of the enteroviruses as described previously[20]. The primer sequences were as follows: EV-F: AYCYYTGTRCGCCTGTTTT; and EV-R: CCCAAAGTGTCCGGTCCGC. The products of RT-PCR were sequenced using the EV-R primer by Biotecan (Shanghai, China). To identify the serotypes, the detected sequences were compared to reference strains in GenBank from the National Center for Biotechnology Information.

Statistical analyses

Categorical variables were summarized as frequencies and proportions using descriptive statistics. Chi-square and Fisher's exact tests were used to examine the relationships between categorical variables. Continuous variables were expressed as mean \pm SD. For normally distributed data, the means of continuous variables were compared using independent group *t*-tests, and for non-normally distributed data, the Mann-Whitney test. To determine the relationship between viral pathogens and HFMD characteristics, a logistic regression model was used. The least absolute shrinkage and selection operator (LASSO) method, which is suitable for analyzing high-dimensional data, was used to select the most significant pathogen-related characteristics[21]. In the forward stepwise logistic regression model, features with non-zero coefficients (automatically removing unnecessary/uninfluential covariates in the LASSO regression model) were set[22]. All the statistical analyses were performed using R software version 4.1.2 (R Foundation for Statistical Computing, Vienna, Austria) and GraphPad Prism 7 (GraphPad Software, La Jolla, California, United States).

RESULTS

Shiyang City experienced a fluctuant decrease in the annual number of HFMD cases from 2016 to 2020

From 2016 to 2020, the Shiyang CDC surveillance system received 35840 HFMD cases, with 1690 (4.7%) laboratory-confirmed cases, 4 (0.01%) severe cases, and 2 (0.06%) fatal cases. In 2017 and 2020, the temporal distribution showed two low ebbs. The yearly number of HFMD decreased by 48.4% in 2017 ($n = 5405$) compared to 2016 ($n = 10473$), following 1.6-fold increases in 2018 ($n = 8513$) and 2019 ($n = 9997$). In 2020 ($n = 1452$), the incidence of HFMD fell by 85.5% compared to 2019 (Table 1 and Figure 1A). When compared to 2016 (308.63/100000), the annual incidence rate of HFMD decreased by 48.8% in 2017 (158.05/100000), then increased by 1.6-fold and 1.2-fold in 2018 (247.78/100000) and 2019 (292.48/100000) respectively. In 2020, the incidence of HFMD reduced by 85.4% (42.73/100000) compared to 2019 (Table 1 and Figure 1B), which could be attributed to the COVID-19 epidemic and China's isolation policy.

EV-A71 lost its dominance since the year 2017 in Shiyang

Since 2017, the proportion of EV-A71-associated HFMD cases among 1690 laboratory-confirmed HFMD cases in Shiyang has decreased significantly. In 2016, 45.5% of EV-A71-positive cases (167/367) were confirmed; however, this proportion dropped to 2.5% (9/363) in 2017 and nearly zero between 2018 and 2020 (Table 1 and Figure 1C). According to these findings, EV-A71 was the dominant pathogen of HFMD in Shiyang in 2016, but it lost the advantage between 2017 and 2020. The China Food and Drug Administration approved the EV-A71 whole virus vaccine in December 2015[8]. In Shiyang city, there were 8234 inoculated people in 2016, 28064 in 2017, 45948 in 2018, 45109 in 2019, and 50337 in 2020, with the number steadily increasing year after year (Figure 1C). These findings suggest that EV-A71 vaccine inoculation may be linked to the recent decrease in EV-A71 prevalence.

The CV-A16 was another common pathogen in China[5]. Its prevalence in Shiyang fluctuates between 2016 and 2020. In 2016, 34.1% of cases (125/367) tested positive for CV-A16. However, the ratio in 2017 was 11.8% (43/363), 61.0% (316/518) in 2018, 15.1% (56/370) in 2019, and 2.8% (2/72) in 2020 (Table 1 and Figure 1C). These findings indicate that CV-A16 was also a dominant HFMD pathogen in Shiyang in 2016, but that its dominance waned sharply in 2018.

In contrast to EV-A71 and CV-A16, the prevalence of other enteroviruses tends to rise in Shiyang from 2016 to 2020. Other enteroviruses ratios were 20.4% (75/367) in 2016, 85.7% (311/363) in 2017, 39.0% (202/518) in 2018, 84.6% (313/370) in 2019, and 97.2% (70/72) in 2020. 971 laboratory-confirmed cases were linked to other enteroviruses between 2016 and 2020, accounting for 57.5% of all cases (Table 1 and Figure 1C). These results suggest that other enteroviruses will gradually replace EV-A71 and CV-A16 as the most common HFMD pathogens in Shiyang. The specific prevailing serotype, however, remained unknown.

CV-A6 was the pathogen that had recently emerged in Shiyang

From 2018 to 2020, we collected 196 throat swab samples from hospitalized HFMD patients to investigate the newly prevalent enterovirus serotype (s) in Shiyang. Using EV-A71 specific, CV-A16 specific, and pan-enterovirus real-time RT-PCR Kits, 168 enterovirus-positive specimens were identified (Figure 2A). According to the 5'-UTR sequences of 168 enterovirus-positive samples, 121 (72.0%) were CV-A6 positive, 8 (4.8%) were CV-A16 positive, and 5 (3.0%) were CV-A10. Other enteroviruses were found in 34 of the samples (20.2%) (Figure 2B). From these results, CV-A6 was the newly prevalent serotype contributing to HFMD in Shiyang between 2018 and 2020.

Clinical characteristics of CV-A6-associated HFMD in Shiyang from 2018 to 2020

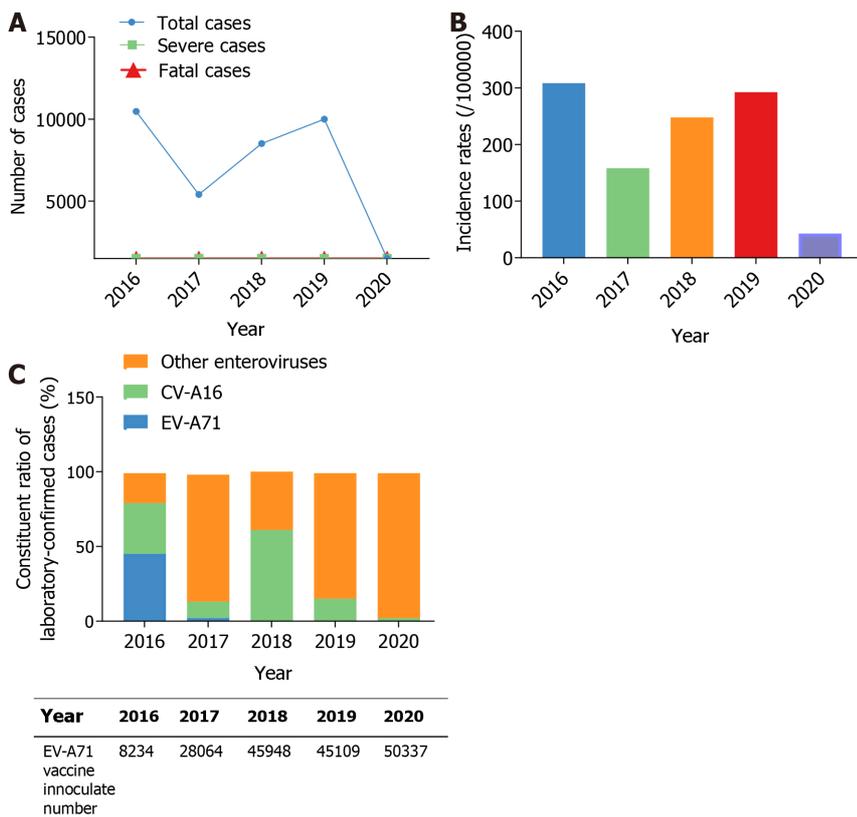
As shown in Table 2 and Figure 3A, the male-to-female ratio among the 168 enterovirus-positive cases was 1.67:1. 91 cases (54.2%) were under one year old, and no case was older than six years old. All patients had oral mucosal or skin rashes on the hand, foot, crissum, or hip. However, no significant differences in gender, age, or rashes were found between CV-A6 and other enteroviruses (CV-A16, CV-A10, and others). Fever was present in 95.9% of CV-A6-positive HFMD patients and 83.0% of CV-A6-negative HFMD patients. Fever levels differed significantly between HFMD with CV-A6 infection and HFMD without CV-A6 infection, according to univariate and multivariate logistic regression analyses (Table 2 and Figure 3B). A subsequent study using the LASSO regression model discovered that CV-A6 infection was more likely to result in fever ($P = 0.023 < 0.05$, Table 3).

There were no severe or fatal cases among the 168 enterovirus-positive cases. However, some patients had respiratory inflammation (such as bronchopneumonia, bronchitis, other upper respiratory tract diseases, or abnormal chest X-ray), and vomited. Nonetheless, there were no significant differences in these clinical manifestations between CV-A6 and other enteroviruses (CV-A16, CV-A10, and others) (Table 2). Myocardial damage, on the other hand, was significantly more common in CV-A6 infected patients than in other enterovirus infections (presenting as polyhedrosis, flustered, electrocardiographic abnormality, increased CK, CK-MB, and LDH). Myocardial damage was present in approximately 25.6% (31/121) of CV-A6-positive HFMD cases, whereas it was present in 8.5% (4/47) of CV-A6-negative HFMD patients (Table 2, Figure 3C). Furthermore, the LASSO regression model revealed that

Table 1 Incidence, severe illness, mortality, and etiological results of hand, foot, and mouth disease cases in 2016-2020 of Shiyang, central China, *n* (%)

Characteristics	Year				
	2016	2017	2018	2019	2020
Total cases	10473	5405	8513	9997	1452
Incidence rates (/100000)	308.63	158.05	247.78	292.48	42.73
Laboratory-detected cases	660	530	700	530	146
Laboratory-confirmed cases	367	363	518	370	72
Severe cases	1	0	2	1	0
Fatal cases	2	0	0	0	0
Etiological results					
EV-A71	167 (45.5)	9 (2.5)	0 (0.0)	1 (0.3)	0 (0.0)
CV-A16	125 (34.1)	43 (11.8)	316 (61.0)	56 (15.1)	2 (2.8)
Other enteroviruses	75 (20.4)	311 (85.7)	202 (39.0)	313 (84.6)	70 (97.2)

HFMD: Hand, foot, and mouth disease; EV-A71: Enterovirus A71; CV-A16: Coxsackievirus A16.



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Figure 1 Yearly distribution of hand, foot, and mouth disease cases from 2016 to 2020, in Shiyang, central China. A: Yearly distribution of hand, foot, and mouth disease (HFMD) cases, severe cases, and fatal cases; B: Yearly incidence rates of HFMD; C: Yearly distribution of laboratory-confirmed HFMD cases and enterovirus A71 vaccine inoculation numbers. CV-A16: Coxsackievirus A16; EV-A71: Enterovirus A71.

myocardial damage ($P = 0.037 < 0.05$) was significantly associated with CV-A6 infection. Additionally, elevated CK-MB ($P = 0.015 < 0.05$) and LDH ($P = 0.001$), two myocardial damage markers, were strongly linked with CV-A6 infection (Table 3), implying that myocardial damage is more frequent in HFMD caused by CV-A6.

Table 2 Logistic regression analysis on the effect of predominant enterovirus serotypes in 168 laboratory-confirmed hand, foot, and mouth disease cases in Shiyang, central China, *n* (%)

	HFMD pathogens				Univariate analysis	Multivariate analysis	
	CV-A6 (<i>n</i> = 121)	CV-A16 (<i>n</i> = 8)	CV-A10 (<i>n</i> = 5)	Other enteroviruses (<i>n</i> = 34)	<i>P</i> value	Coefficient (SD)	<i>P</i> value
Gender					1	-0.224 (0.451)	0.619
Male (<i>n</i> = 105)	76 (62.8)	5 (62.5)	4 (80.0)	20 (58.8)			
Female (<i>n</i> = 63)	45 (37.2)	3 (37.5)	1 (20.0)	14 (41.2)			
Age (yr)					0.794		
≤ 1 (<i>n</i> = 91)	65 (53.7)	4 (50.0)	3 (60.0)	19 (55.9)			
≤ 2 (<i>n</i> = 49)	37 (30.6)	2 (25.0)	2 (40.0)	8 (23.5)		0.131 (0.497)	0.792
≤ 3 (<i>n</i> = 18)	13 (10.7)	0 (0.0)	0 (0.0)	5 (14.7)		0.236 (0.717)	0.742
≤ 6 (<i>n</i> = 10)	6 (5.0)	2 (25.0)	0 (0.0)	2 (5.9)		-0.371 (0.785)	0.637
Fever					0.013 ^a	2.124 (0.747)	0.004 ^b
Yes	116 (95.9)	5 (62.5)	5 (100.0)	29 (85.3)			
No	5 (4.1)	3 (37.5)	0 (0.0)	5 (14.7)			
Clinical manifestations							
Skin or oral mucosal rashes	121 (100.0)	8 (100.0)	5 (100.0)	34 (100.0)			
Respiratory system syndromes ¹	24 (19.8)	3 (37.5)	1 (20.0)	3 (8.8)	0.603	0.811 (0.589)	0.168
Myocardial damage	31 (25.6)	1 (12.5)	1 (20.0)	2 (5.9)	0.025 ^a	1.040 (0.678)	0.125
Vomit	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	0.623	-15.923 (882.744)	0.986
Laboratory findings, <i>n</i> (%), mean ± SD)							
WBC increased	69 (56.6, 14.07 ± 3.42)	4 (50.0, 19.82 ± 10.74)	5 (100.0, 20.68 ± 7.67)	15 (44.1, 13.88 ± 4.38)	0.042 ^a	-0.058 (0.050)	0.247
CRP increased	79 (64.8, 24.64 ± 26.90)	5 (62.5, 25.28 ± 26.64)	5 (100.0, 47.68 ± 29.52)	15 (44.1, 25.97 ± 43.35)	0.431	-0.007 (0.006)	0.271
CK-MB increased	56 (45.9, 33.77 ± 13.78)	2 (25.0, 30.00 ± 4.24)	4 (80.0, 28.00 ± 6.48)	17 (50.0, 30.94 ± 7.04)	0.262	0.045 (0.027)	0.097
LDH increased	98 (80.3, 313.08 ± 59.37)	6 (75.0, 298.17 ± 38.97)	5 (100.0, 275.00 ± 24.81)	27 (79.4, 334.52 ± 59.64)	0.487	-0.009 (0.004)	0.043 ^a
ALT increased	20 (16.4)	1 (12.5)	1 (20.0)	3 (8.8)	0.333	-0.575 (1.328)	0.665
AST increased	29 (23.8)	1 (12.5)	1 (20.0)	3 (8.8)	0.086	1.778 (1.196)	0.137
Radiographic evaluation							
Abnormal chest X-ray ²	48 (39.3)	4 (50.0)	1 (20.0)	8 (23.5)	0.267	0.275 (0.449)	0.540
Abnormal electrocardiogram ³	24 (19.7)	1 (12.5)	1 (20.0)	5 (14.7)	0.389	0.236 (0.566)	0.677

¹Respiratory system syndromes were defined as the presence of at least one of the followings: Bronchopneumonia, bronchitis, or other upper respiratory tract diseases.

²Abnormal chest X-ray: unilateral or bilateral enhanced lung texture.

³abnormal electrocardiogram: sinus arrhythmia, or sinus tachycardia.

^a*P* < 0.05.

^b*P* < 0.01. HFMD: Hand, foot, and mouth disease; CV-A6: Coxsackievirus A6; CV-A16: Coxsackievirus A16; CV-A10: Coxsackievirus A10; SD: Standard deviation; WBC: White blood cell; CRP: C-reactive protein; CK-MB: Creatine kinase MB isoenzyme; LDH: Lactate dehydrogenase; ALT: Alanine transaminase; AST: Aspartate aminotransferase.

DISCUSSION

In 1957, HFMD was initially reported in New Zealand and Canada[23,24]. Then it spread all over the world, including the United States, Bulgaria, Hungary, and a slew of Asia countries[25-33]. For a long time, little attention was paid to this disease due to its mild clinical manifestations and self-limiting nature. Nonetheless, over the last two decades, an increasing number of outbreaks with atypical manifestations and expanding pathogen spectrums have been reported worldwide, attracting public attention[3,34,35]. Because of the widespread prevalence of HFMD in China, the Chinese Ministry of Health has also designated it as a category III notifiable infectious disease since 2008. The main pathogens in the pathogen spectrums were CV-A16 and EV-A71[36]. The disease caused by CV-A16 was typically mild, with classical rashes. EV-A71, on the other hand, can cause severe neurological manifestations such as polio-like paralysis, fatal encephalitis, and cardiopulmonary complications in addition to rashes[37]. As a result, developing effective pathogen-specific vaccines is a wise decision.

Since 2016, three inactivated, whole EV-A71 vaccines have been launched in China[38]. And the EV-A71 vaccines were proved to be effective in a recent epidemiologic survey in Guangxi Province (Southern China). The incidence and severity of EV-A71 cases have both decreased significantly following vaccination with EV-A71 vaccines from 2017 to 2019[9]. A report from Nanchang (Southeastern China) found no EV-A71 after a 2-year implementation of the EV-A71 vaccination[39]. Our findings also showed the efficacy of EV-A71 vaccines. It was discovered that, as the number of inoculated people in Shiyang (Central China) increased year after year, the incidence of HFMD decreased sharply in 2017 (Table 1 and Figure 1A-B). Furthermore, EV-A71 prevalence in Shiyang decreased sharply in 2017 and will almost completely disappear between 2018 and 2020 (Table 1 and Figure 1C), indicating that EV-A71 vaccine inoculation may be beneficial. However, the HFMD incidence increased again in Shiyang in 2018 and 2019, indicating that, while EV-A71 was under control, other pathogens may be contributing to the increased HFMD cases. In fact, the EV-A71 vaccine provided no cross-protection against other serotypes. Then, which pathogen(s) were responsible for the rise in HFMD cases in Shiyang between 2018 and 2020?

In reality, a shift in HFMD etiology has been reported in recent years. CV-A6, CV-A10, and CV-A12 have emerged as important HFMD pathogens, except for CV-A16 and EV-A71[40-42]. HFMD caused by CV-A6 has been reported in countries around the world, including Finland, the United States, Japan, Spain, *etc*[10,11,43,44]. CV-A6 was also a significant pathogen contributing to HFMD in China. Between 2010 and 2012, CV-A6 surpassed CV-A16 as the second most common serotype in southern China (Guangdong Province)[45,46]. In 2013, CV-A6 replaced EV-A71 and CV-A16 as the primary pathogen of HFMD in northern (Beijing, Tianjin), northeastern (Jilin), eastern (Taizhou), and middle China (Chongqing)[47-51]. In 2015, CV-A6 outbreaks were reported in northern (Beijing) and middle (Chongqing) China[52,53]. During 2018-2019, CV-A6 was found to be the most common cause of HFMD in eastern (Shanghai), northern (Beijing, 2018, and 2020), southeastern (Nanchang, 2019), and southern (Kunming, 2018) China[39,54-56]. In our study, CV-A6 surpassed EV-A71 and CV-A16 as the leading cause of HFMD in central China (Shiyang) between 2018 and 2020 (Table 2 and Figure 2), demonstrating that CV-A6 has become the most common serotype recently in China. Intriguingly, CV-A6 nonetheless overtook CV-A16 and EV-A71 as the primary causal agent in some regions in 2013 despite the absence of EV-A71 vaccines. The reason remained unknown.

It has been reported that CV-A6 can cause severe HFMD in children and atypical HFMD in adults[57-59]. Clinical manifestations involved vesiculobullous exanthema, palm and sole desquamation, and onychomadesis[14,60]. A study of 8 confirmed HFMD children found severe clinical features such as 100% fever (higher than 39 °C), 87.5% herpangina, 100% meningitis, and 25% myocardial damage caused by CV-A6[57]. In our collected samples, we found that more than half of the cases occurred in boys and children under the age of one year. No patient was older than the age of six (Table 2). As a result, the prevention of infant HFMD should be prioritized. Because we collected samples from the pediatric department, no adult patients were found. In addition, we showed that CV-A6 was more likely to cause fever (95.9% *vs* 83%) and myocardial damage (25.6% *vs* 8.5%) (Table 3 and Figure 3), which was consistent with the previous study[57]. However, no onychomadesis was observed in our study. It might be due to different CV-A6 genotypes. The limitation is that genotypes of prevailing CV-A6 in Shiyang and other regions were not identified. Further study can be done to determine the association between different CV-A6 genotypes and clinical features.

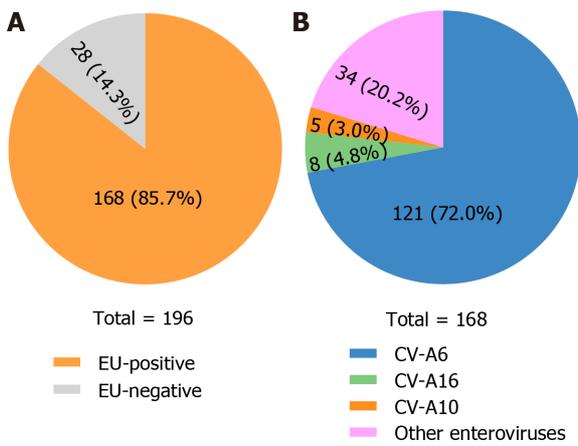
The median R0 (interquartile range) of the major HFMD serotypes was calculated in a recent study. Among EV-A71, CV-A16, and CV-A6, CV-A6 had the highest R0 (5.94, and 25.80 after adjusting for seroprevalences), implying that CV-A6 may be the major challenge in the prevention and control of HFMD in China[61]. As a result, developing vaccines against CV-A6 or multiple pathogens will help prevent HFMD. Recently, an inactivated CV-A6 vaccine candidate based on Vero cells was shown to be effective in mouse models[62]. However, more clinical trials were required to provide evidence of its efficacy. A recent study[63] used cryo-electron microscopy to identify the virion structure of CV-A6, laying the groundwork for the development of novel vaccines and drugs. In addition, multiple enterovirus serotypes, including CV-A6, should be included in long-term HFMD surveillance.

Table 3 The least absolute shrinkage and selection operator regression analysis of the correlation between coxsackievirus A6 infection and various characteristics in hand, foot, and mouth disease patients in Shiyang, China

Characteristic	CV-A6 infection	
	Coefficient (SE)	P value
Temperature	-8.880e-01 (3.899e-01)	0.023 ^a
Myocardial damage	1.224e+00 (5.894e-01)	0.037 ^a
CK-MB increased	7.187e-02 (2.978e-02)	0.015 ^a
LDH increased	-1.355e-02 (4.137e-03)	0.001 ^b

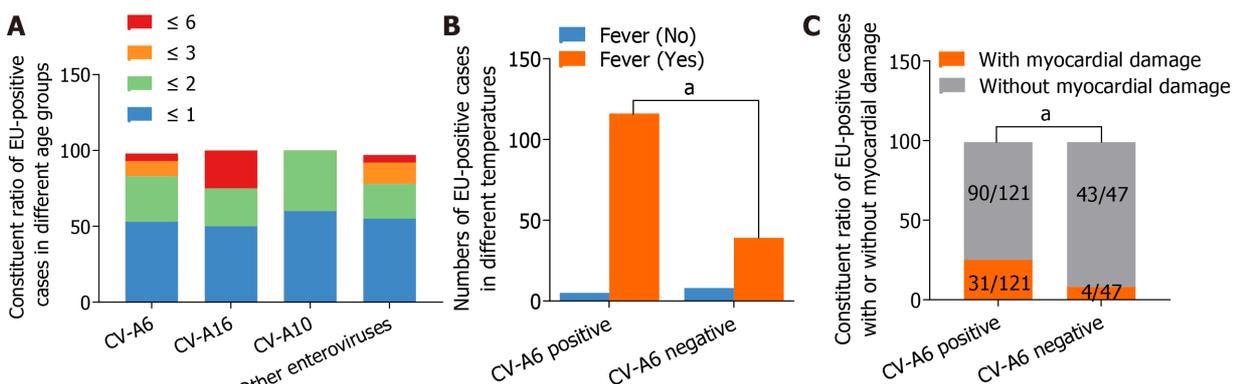
^aP < 0.05.

^bP < 0.01. CV-A6: Coxsackievirus A6; SE: Standard error; CK-MB: Creatine kinase MB isoenzyme; LDH: Lactate dehydrogenase.



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Figure 2 Enterovirus serotypes of 196 hospitalized hand, foot, and mouth disease patients enrolled in this study from 2018 to 2020 in Shiyang, central China. A: Enterovirus A71 specific, coxsackievirus A16 specific, and pan-enterovirus real-time reverse transcription-polymerase chain reaction Kits were used to identify enterovirus-positive specimens; B: The 5'-untranslated region sequences identified enterovirus serotypes of 168 enterovirus-positive samples. EU-positive: Enterovirus-positive; CV: Coxsackievirus.



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Figure 3 Clinical characteristics of hand, foot, and mouth disease in Shiyang from 2018 to 2020. A: The constituent ratio of different enterovirus serotypes associated with hand, foot, and mouth disease cases in different age groups; B: Numbers of enterovirus-positive cases with or without fever; C: The constituent ratio of enterovirus-positive cases with or without myocardial damage. ^aP < 0.05. EU-positive: Enterovirus-positive; CV: Coxsackievirus.

CONCLUSION

In conclusion, CV-A6 is the most common enterovirus serotype in Shiyang City, replacing EV-A71 and CV-A16 as the HFMD pathogen. Fever and myocardial damage are more common in CV-A6-caused HFMD. CV-A6 prevalence surveillance should be prioritized.

ARTICLE HIGHLIGHTS

Research background

Before 2016, the main causative serotypes of hand, foot, and mouth disease (HFMD) were coxsackievirus A16 (CV-A16) and enterovirus A71 (EV-A71). Some regions in China have shown that CV-A6 has recently supplanted EV-A71 and CV-A16 due to the introduction of EV-A71 vaccines. Finding out Central China's epidemiological characteristics is equally important.

Research motivation

By investigating the clinical symptoms and pathogen spectrum of HFMD in Shiyang City, central China, after 2016, it will provide critical proof for future knowledge of the etiology of HFMD.

Research objectives

To investigate the epidemiological and etiological features of HFMD in Shiyang City, which may provide information on the disease's recent prevalence in central China.

Research methods

The Shiyang Center for Disease Control and Prevention HFMD data from 2016 to 2020 were examined. Between 2018 and 2020, throat swab specimens were collected from hospitalized HFMD patients in Shiyang. To detect and genotype enteroviruses, real-time reverse transcription-polymerase chain reaction and sequencing of the 5'-untranslated region were used. The effect of predominant enterovirus serotypes and the correlation between CV-A6 infection and various clinical characteristics in Shiyang HFMD patients were studied using a logistic regression model and the least absolute shrinkage and selection operator model.

Research results

The frequency of HFMD dramatically dropped in 2017 as the number of immunized persons in Shiyang grew year after year. But the prevalence of HFMD rose once more in 2018 and 2019. Between 2018 and 2020, CV-A6 superseded EV-A71 and CV-A16 as the main cause of HFMD in Shiyang. CV-A6 was associated with an increased risk of fever and myocardial damage. The limitation is that the dominant CV-A6 genotypes in Shiyang and other regions were not identified. More research is needed to determine the relationship between different CV-A6 genotypes and clinical features.

Research conclusions

The most common enterovirus serotype in Shiyang City was CV-A6, which replaced EV-A71 and CV-A16 as the HFMD pathogen.

Research perspectives

The prevention of HFMD in central China will be aided by the creation of vaccines against CV-A6 or multiple infections as well as increased CV-A6 surveillance.

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FOOTNOTES

Author contributions: Li JF, Zhang CJ, and Li YW contributed equally to this study; Li JF, Zhang CJ, and Li YW designed and performed the research and co-wrote the paper; Li C, Wang SS, Zhang SC, Jiang Y, Luo XB, and Liao XJ collected the clinical specimens and the corresponding characteristics; Wu SX, and Lin L performed the serotypes identification of clinical samples; all authors have read and approved the final manuscript.

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Informed consent statement: Patients were not required to give informed consent to the study because the analysis used anonymous clinical data that were obtained after each patient agreed to treatment by written consent.

Conflict-of-interest statement: All authors hereby declare that no conflict of interest exists.

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