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# A Simple Scoring System for Quick, Accurate, and Reliable Early Diagnosis of Hand, Foot, and Mouth Disease

Authors' Contribution:  
Study Design A  
Data Collection B  
Statistical Analysis C  
Data Interpretation D  
Manuscript Preparation E  
Literature Search F  
Funds Collection G

ACDE 1 **Shaoxing Chen**  
BF 1 **Kaihong Yi**  
DE 1 **Xiaojun Chen**  
AG 2 **Liping Li\***  
ADG 3 **Xuerui Tan\***

1 Department of Community Monitoring, First Affiliated Hospital of Shantou University Medical College, Shantou, Guangdong, P.R. China  
2 Injury Prevention Research Center, Shantou University Medical College, Shantou, Guangdong, P.R. China  
3 Department of Cardiology, First Affiliated Hospital of Shantou University Medical College, Shantou, Guangdong, P.R. China

\* Contributed equally to this work

**Corresponding Authors:** Xuerui Tan, e-mail: tanxuerui@vip.sina.com, Liping Li, e-mail: lpli@stu.edu.cn

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**Background:** To construct an accurate, reliable, and simple scoring system of improving HFMD diagnosis.





**Material/Methods:** Based on the following 3 steps, a simple scoring diagnostic system was built: (1) we selected basic markers (age and sex), markers recommended in HFMD diagnosis guidelines, and significant biomarkers among severity groups found in a large dataset; (2) we used positive constituent ratio for determining scores of each marker; and (3) we applied receiving operating curve in an external dataset to determine the optimal cut-off score.

**Results:** The selected markers were sex, age, fever, skin rashes, nervous system disorder, respiratory system disorder, digestive system disorder and cardiopulmonary complications, C-reactive-protein, White Blood Cell, Creatinine Kinase, Creatinine Kinase Isoenzyme, Gamma-Glutamyl Transpeptidase, Albumin, Globulin, Albumin/Globulin Ratio, Sodium, Chloride, Calcium, and Glucose. A simple scoring system with 3.9684 as the lower cut-off was constructed. The AUC was 0.918 (95% CI: 0.874-0.963,  $P < 0.01$ ). The sensitivity, specificity, and Youden Index, which were based on the validation dataset of 200 subjects (80 cases, 120 non-cases with skin rashes or fever), were 0.95, 0.90, and 0.85, respectively.

**Conclusions:** This simple scoring system is an effective method to diagnose HFMD.

**MeSH Keywords:** **Epidemiologic Study Characteristics as Topic • Hand, Foot and Mouth Disease • Research Design**

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## Background

Hand, foot, and mouth disease (HFMD) is an acute infectious disease caused by Enterovirus A, of which the most commonly reported genotypes are Coxsackievirus A6, A10, and A16 (CV-A6, CV-A10, CV-A16) and Enterovirus A71 (EV-A71) [1]. HFMD commonly occurs in preschool children. In China, children under 3 years old are at the highest risk of HFMD [2]. The main clinical manifestations of HFMD are maculopapular rash and vesicles on the hands, feet, mouth, or buttocks. Some rare HFMD cases have other symptoms that are related to EV-A71, such as viral meningitis, encephalitis, encephalomyelitis, pulmonary edema, and circulatory disorder. The primary causes of death are brainstem encephalitis and neurogenic pulmonary edema.

Since its original identification in New Zealand in 1957, HFMD has been frequently reported worldwide [3]. Additionally, it has become a public health priority in China before its first large-scale epidemic in Fuyang City, Anhui Province in 2008 [4,5]. During the epidemic season, kindergarten, nursery, and family can be infected by intra-familial and intra-community transmissions [6]. The enterovirus of HFMD transmits rapidly, is strongly contagious, and is a main cause of asymptomatic infection; thus, the disease can be potential dissemination and controlling its transmission is challenging [7].

The clinical manifestations of HFMD are diverse and nonspecific. Therefore, it is difficult to perform an accurate tentative diagnosis in clinical practice. During the epidemic season, preschool children who have rashes on hands, feet, mouth, or buttocks, and who are with or without fever, are suspected to have HFMD according to the diagnostic criteria of HFMD [1]. Although the main symptoms of HFMD are skin rashes with or without fever, it can be easily confused with other types of pyretic or exanthematous diseases, such as measles, urticaria, rubella, exanthema subitum, varicella, scarlatina, herpangina, and herpes simplex. Laboratory examinations are usually used to confirm suspected HFMD cases, but the detection rates are highly dependent on factors such as quality of specimens and incubation periods [8]. As a result, the diagnosis of HFMD can be misdiagnosed or missed, which delays delivery of optimal treatments.

With advances in the development and application of mathematical methods, quantitative methods, and statistical methods in medicine, analysis methods such as diagnostic scale, discrimination model, regression analysis, and system evaluation are being integrated into epidemiological and clinical research. Scoring systems have been applied to evaluate the risk of mortality in intensive care units (ICU) and are integrated into the intensive care process. Scoring systems allow between- and within-individual comparisons over time and also provide useful information for comparing the severity of illness in patients who were enrolled into clinical trials [9].

Two mortality risk scoring systems – Pediatric Risk of Mortality (PRISM) and the Pediatric Index of Mortality (PIM) scores – are widely applied in pediatrics [10,11]. A number of studies, mainly in developed countries, as well as in a few resource-restricted settings, have validated PIM and its updated version, PIM2 scores [12,13]. Previous studies in China have established scoring systems based on the internationally applied scoring systems (PIM or PCIS) and the clinical manifestation of HFMD, to assess the severity of HFMD [14,15]. To improve HFMD disease management by detecting disease at early stages, our study developed a simple scoring system to quickly and accurately diagnose HFMD, especially where laboratory-confirmed tests are not available.

## Material and Methods

### Ethics statement

The present study was approved by the Ethics Committee of the First Affiliated Hospital of Shantou University Medical College. The methods carried out were accordance with the approved guidelines as well as the guidelines for the protection of human subjects. Due to the retrospective nature of the study, informed consent was not required.

### Training dataset

Data on subjects were retrospectively collected from medical records of all the clinically diagnosed HFMD pediatric inpatients, aged at 14 years old or younger, who visited the pediatric department of the First Affiliated Hospital of Shantou University Medical College between January 2012 and December 2014. The diagnostic criteria of HFMD were based on the HFMD diagnosis and treatment guideline (2010 edition) of the Ministry of Health, China [1]: (1) HFMD emerge mainly in summer and autumn in preschool children, especially in infants; (2) Patients have fever accompanied with rash on hands, feet, mouth, or buttocks. Some patients may have no fever; (3) In a minority of severe patients who have atypical rash and are difficult to diagnose, clinical diagnosis primarily depends on etiologic or serologic test results; and (4) It is not recommended to diagnose HFMD for patients without rash.

HFMD cases were categorized into 3 groups (mild, severe, and critical) [1]. Patients with any neurological complications (e.g., aseptic meningitis, encephalitis, encephalomyelitis, and acute flaccid paralysis or autonomic nervous system dysregulation) were categorized as severe cases. Patients with any 1 of the following 3 groups of clinical manifestations were categorized as critical cases: (1) frequent cramp, coma, and cerebral hernia; (2) cardiopulmonary complications (e.g., dyspnea, cyanosis, frothy sputum, cardiopulmonary edema, and pulmonary rale;

and (3) shock or other dysfunctions in the circulatory system. Probable or confirmed HFMD patients that were neither critical nor severe were categorized as mild cases.

All data were collected from medical records of the patients' first physical examination and blood test on the first day of hospital admission, including: (1) sex and age; (2) clinical symptoms such as skin rashes, fever, nervous system, respiratory system, digestive system, and cardiopulmonary complications; and (3) biomarkers such as HBDH, LDH, LDH1, AST, ALT, GGT, ALP, CHE, MAO, AFU, TP, ALB, GLB, Tbil, Dbil, Ibil, A/G, K<sup>+</sup>, Na<sup>+</sup>, Cl<sup>-</sup>, Ca<sup>2+</sup>, CO<sub>2</sub>CP, BUN, Cr, CRP, WBC, CK, CK-MB, and GLU. The diagnosis of clinical symptoms was based on the clinical diagnosis of their first physical examination. The measurements of biomarkers were based on the blood samples of their first laboratory test since admission. In addition, patients were considered to have skin rashes if they had maculopapular or vesicular rash on locations such as hands, feet, mouth, buttocks, elbows, trunk, or face. Patients were considered to have a nervous system disorder if they had altered mental status, somnolence, skittishness, headache, vomiting, delirium or coma, extremity tremor, myoclonus, nystagmus, ataxia, oculomotor disorder, atony or acute flaccid paralysis, convulsions, startle, seizures, or neck resistance.

### Development of the scoring system

A simple scoring system for HFMD diagnosis was developed in the following steps: (1) We selected relevant markers including basic markers (age and sex), markers listed in HFMD clinical guidelines of China, and biochemical markers that were significantly different among severity groups in the training dataset; (2) We estimated the score of each marker by using positive constituent ratio to construct the simple scoring system; and (3) We determined the optimal cut-off of the simple scoring system in the validation dataset.

### Markers selection through data analysis of training dataset

All biochemical markers in the training dataset were common biochemical markers used in clinical practice.

Analysis of variance (ANOVA) was used to analyze the association between HFMD and biochemical markers. When a biochemical marker was found to be statistically different among the 3 groups, multiple comparisons were used for pairwise comparison of the mean difference. Bonferroni corrections were applied to adjust for multiple comparisons. In the cases of homogenous variance, Fisher's least significant difference (LSD) and Student-Newman-Keuls (SNK) tests were used for pairwise multiple comparison; in the cases of heterogeneous variance, Dunnett analysis (T3) was used for pairwise multiple

comparison. For mean square analysis of all groups, all 2-sided  $P < 0.01$  were considered statistically significant; for pairwise multiple comparison, all 2-side  $P < 0.05$  were considered statistically significant. Markers that were significantly different in 2 of the 3 groups were selected as markers of the simple scoring system.

### Assign score to markers

The score of each marker was estimated using positive constituent ratio. The age and sex of the 1404 cases were detailed and accurately recorded; hence, their scores were established based on 1404 cases of the sample. The other 18 selected markers were completely recorded in 985 cases; therefore, their scores were established based on 985 cases of the sample.

Age and sex were categorical markers. Of the 1404 cases of this study, 915 (65.17%) were male and 489 (34.83%) were female, which means if the patient was male, his sex score would be 0.6517; if the patient was female, the sex score would be 0.3483. Similarly, the score of age group 0–3 years old, 4–6 years old, and 7–14 years old were 0.7479, 0.2321, and 0.020, respectively.

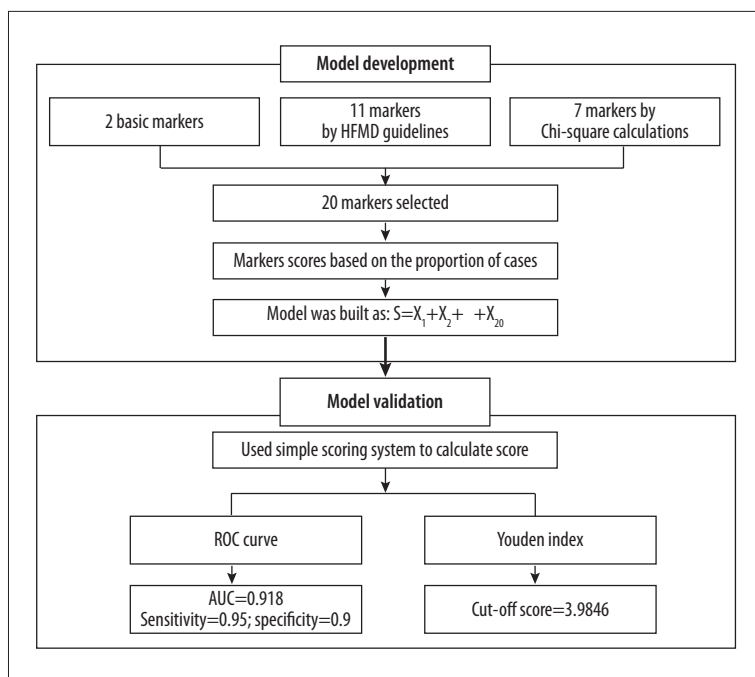
The score of 6 clinical manifestations (fever, skin rashes, nervous system disorder, respiratory system disorder, digestive system disorder, and cardiopulmonary complications) were separately obtained by calculating the proportion of cases with positive test results. For example, if 882 of the 985 cases had fever, then the percentage of fever was  $(882/985) \times 100\% = 89.54\%$ ; which means the score of fever in those pediatric patients who had fever was 0.8954, and the score of fever for pediatric patient who did not have fever was 0. The calculation of score for the remaining second type of markers followed the same rules.

The scores of the 12 laboratory parameters were obtained by calculating the proportion of abnormal cases in the whole sample. For example, if among the 985 cases, 356 CRP values were abnormal, then the proportion of abnormal CRP was  $(356/985) \times 100\% = 36.14\%$ , which means the score of CRP for pediatric patient who had abnormal CRP value was 0.3614, and the score of CRP for pediatric patient who had normal CRP value was 0. The calculation of score for the remaining third type of markers followed the same principle.

The formula of the scoring system was  $S = X_1 + X_2 + X_3 + \dots + X_{20}$ , where  $X_1, X_2, X_3, \dots + X_{20}$  were the score of each marker, and  $S$  was the HFMD diagnosis score.

### Validation dataset

Confirmed HFMD cases by IgM antibody assays whose age were 14 years old or younger were retrospectively and randomly



**Figure 1.** The flowchart of creating a simple score system. HFMD – hand, foot and mouth disease; ROC – receiver operating curve;  $X_1, X_2, \dots, X_{20}$  – the score of the 20 markers;  $S$  – score of hand, foot and mouth disease.

sampled from the inpatients in the Pediatric Department of the First Affiliated Hospital of Shantou University Medical College in 2015. To assess the performance of our scoring system in distinguishing HFMD cases from other diseases such as measles, urticaria, rubella, exanthema subitum that share similar symptoms as HFMD, non-cases in the validation dataset were chosen among patients of other diseases (e.g., measles and rubella) that showed symptoms of skin rashes and with or without fever. All non-cases were age 14 years or younger, with fever or skin rashes, and were confirmed to be other diseases without HFMD.

A total of 200 participants, including 80 confirmed HFMD cases and 120 non-cases, were enrolled into the validation dataset. Among 80 cases, the serotypes were EV-A71 (32 cases), CV-A16 (22 cases) and cross-contamination of both viruses (26 cases). The types of non-cases were measles (9 cases), rubella (12 cases), exanthema subitum (8 cases), and other similar diseases (91 cases).

### EVs detection

Blood samples were clinically collected from serum when patients visited the clinic or at admission. EV71-IgM and CV-A16-IgM antibody in serum and cerebrospinal fluid specimens were detected using immune colloidal gold technique, according to the manufacturer's instructions. EV71-IgM and CV-A16-IgM antibody detection kits (Wantai Biological Pharmacy Enterprise Co., Ltd., Beijing, China) were used for the detection.

### Discrimination analysis

The diagnostic accuracy of the simple scoring system was assessed in 3 steps: (1) We calculated the score of all subjects in the validation dataset; (2) We calculated the sensitivity, specificity, Youden Index (YI) and AUC; and (3) We determined the optimal cut-off score that corresponded to the largest YI. The outcomes (case, non-case) were masked, and then the score of all subjects in the validation dataset was calculated using the simple scoring system. The sensitivity and 1-specificity of all cut-off values of ROC curve were exported to a spreadsheet to obtain the optimal cut-off score that corresponded to the largest YI (Supplementary Table 1). The scoring system was performed with complete data and missing data at different levels of severity to assess the robustness of its sensitivity.

The flowchart is shown in Figure 1.

Examples of how the scoring system calculates the scores of patients are shown in Figure 2. All statistical analyses were performed using IBM SPSS version 21 (SPSS, Inc., Chicago, IL, USA).

## Results

### Subject characteristics of training dataset

A total of 1404 cases (1104 mild, 252 severe, and 48 critical) were recruited into the training dataset (Table 1). The frequency (percentage) of male and female were 915 (65.2%) and 489 (34.8%), respectively. The age of cases ranged from 0 to 14

Simple scoring system	Patient A	Patient B
<p><b>Markers and score</b></p> <p>1. Gender (male: 0.6517; female: 0.3483)                  2. Age (0~3: 0.7479; 4~6: 0.2321; 7~14: 0.020)                  3. Fever (Yes: 0.8954; No: 0)                  4. Skin rashes (Yes: 0.9817; No: 0)                  5. Nervous system disorder (Yes:0.2376; No:0)                  6. Respiratory system disorder (Yes: 0.2985; No: 0)                  7. Digestive system disorder (Yes: 0.1046; No: 0)                  8. Cardiopulmonary complications (Yes: 0.0061; No: 0)                  9. CRP (A: 0.3614; No: 0)                  10. WBC (A: 0.4964; N: 0)                  11. CK (A: 0.1533; N: 0)                  12. CK-MB (A: 0.1797; N: 0)                  13. GGT (A: 0.0061; N: 0)                  14. ALB (A: 0.2274; N:0)                  15. GLB (A: 0.3563; N:0)                  16. A/G (A: 0.2650; N:0)                  17. Na<sup>+</sup> (A: 0.3299; N: 0)                  18. Cl<sup>-</sup> (A: 0.1848; N:0)                  19. Ca<sup>2+</sup> (A: 0.1848; N:0)                  20. GLU (A: 0.2782; N:0)</p>	<p>1. male: 0.6517                  2. 4~6: 0.2321                  3. Yes: 0.8954                  4. Yes: 0.9817                  5. Yes: 0.2376                  6. Yes: 0.2985                  7. No: 0                  8. No: 0                  9. No: 0                  10. A: 0.4964                  11. N: 0                  12. N: 0                  13. N: 0                  14. N: 0                  15. N: 0                  16. N: 0                  17. A: 0.3299                  18. A: 0.1848                  19. N: 0                  20. N: 0</p>	<p>1. male: 0.6517                  2. 0~3: 0.7479                  3. No: 0                  4. Yes: 0.9817                  5. No: 0                  6. No: 0                  7. No: 0                  8. No: 0                  9. A: 0.3614                  10. A: 0.4964                  11. N: 0                  12. N: 0                  13. N: 0                  14. N: 0                  15. N: 0                  16. N: 0                  17. N: 0                  18. A: 0.1848                  19. N: 0                  20. A: 0.2782</p>
<p><b>Calculation</b></p> $S = X_1 + X_2 + \dots + X_{20}$	$S = 0.6517 + 0.2321 + 0.8954 + 0.9817 + 0.2376 + 0.2985 + 0.4964 + 0.3299 + 0.1848 = 4.3081$	$S = 0.6517 + 0.7479 + 0.9817 + 0.3614 + 0.4964 + 0.1848 + 0.2782 = 3.7021$
<p><b>Diagnosis</b></p> <p>Cut-off score = 3.9684</p>	<p>Score = 4.3081 &gt; 3.9684                  ↓                  HFMD case</p>	<p>Score = 3.7021 &lt; 3.9684                  ↓                  Non-HFMD case</p>

**Figure 2.** Examples of using the simple scoring system to diagnose HFMD cases. CRP – C-reactive protein; WBC – white blood cell; CK – creatinine kinase; CK-MB – creatinine kinase isoenzyme; GGT – gamma-glutamyl transpeptidase; ALB – albumin; GLB – globulin; A/G – albumin/globulin ratio; Na<sup>+</sup> – natrium ion; Cl<sup>-</sup> – chloride ion; Ca<sup>2+</sup> – calcium ion; GLU – glucose; HFMD – hand, foot and mouth disease; X<sub>1</sub>, X<sub>2</sub>, ..., X<sub>20</sub> – the score of the 20 markers; S – score of hand, foot, and mouth disease.

years old. Subjects who were 3 years old or younger had the highest incidence, which were 1050 cases (74.8%); subjects who were 4–6 years old had the second highest incidence, which were 326 cases (23.2%); and subjects who were 7–14 years old were the least, which were 28 cases (2.0%). There were 799 cases (56.9%) from rural areas and 605 cases (43.1%) from urban areas. In total, 152 cases (10.8%) had a history of HFMD.

### Subject characteristics of validation dataset

A total of 200 subjects (80 subjects with HFMD, 120 subjects without HFMD) were randomly selected into the validation dataset (Table 1). Among HFMD cases, the frequency of males and females were 45 (56.3%) and 35 (43.7%), respectively; their mean age was 1.56 years old, ranging from 0 to 14 years old, in which 58 (72.5%) were under 3 years old; 71 subjects (88.8%) were mild cases and 9 subjects (11.2%) were severe cases. Among subjects without HFMD, the frequency of males and females were 72 cases (60.0%) and 48 cases (40.0%), respectively; their mean age was 1.48 years old, ranging from 0 to 14 years old.

### Markers selection

Summary statistics of biochemical markers that were significantly different among the mild, severe, and critical cases in the training dataset are shown in Table 2. White Blood Cell (WBC), Creatinine Kinase (CK), Creatinine Kinase Isoenzyme (CK-MB), Glucose (GLU), C-Reactive Protein (CRP), Gamma-Glutamyl Transpeptidase (GGT), Albumin (ALB), Globulin (GLB), Albumin/Globulin ratio (A/G), Natrium Ion (Na<sup>+</sup>), Chloride Ion (Cl<sup>-</sup>), and Calcium Ion (Ca<sup>2+</sup>) were significantly different among the groups (mild, severe, and critical) with  $P < 0.01$ . Those biochemical markers were included as markers of the simple scoring system. A total of 20 markers were included in the simple scoring system, including basic markers (age and sex), markers selected from the HFMD diagnosis and treatment guidelines of China and markers that were found to be significantly different among the groups in the training dataset.

The assigned scores of 20 markers are shown in Table 3, they were sex (male: 0.6517; female: 0.3483), age (0–3 years old: 0.7479; 4–6 years old: 0.2321; 7–14 years old: 0.020),

**Table 1.** Subject characteristics of training and validation dataset.

Dataset	Training dataset (n=1404)	Validation dataset (n=200)	
		HFMD cases (n=80)	Non-cases (n=120)
Sex (n and%)			
Male	915 (65.2)	45 (56.3)	72 (60.0)
Female	489 (34.8)	35 (43.7)	48 (40.0)
Age (n and%)			
0–3 y	1050 (74.8)	58 (72.5)	107 (89.2)
4–6 y	326 (23.2)	21 (26.3)	12 (10.0)
7–14 y	28 (2.0)	1 (1.2)	1 (0.8)
Clinical manifestations (n and%)	Based on 985 cases		
Fever	882 (89.5)	71 (88.8)	68 (56.7)
Skin rashes	967 (98.2)	80 (100.0)	63 (52.5)
Nervous system disorder	234 (23.8)	47 (58.8)	32 (26.7)
Respiratory system disorder	294 (29.9)	25 (31.3)	23 (19.2)
Digestive system disorder	103 (10.5)	17 (21.3)	9 (7.5)
Cardiopulmonary complications	6 (0.6)	3 (3.8)	1 (0.8)

fever (0.8954), skin rashes (0.9817), nervous system disorder (0.2376), respiratory system disorder (0.2985), digestive system disorder (0.1046), cardiopulmonary complications (0.0061), C-Reactive Protein (CRP, 0.3614), WBC (0.4964), Creatinine Kinase (CK, 0.1533), Creatinine Kinase Isoenzyme (CK-MB, 0.1797), GGT (0.0061), ALB (0.2274), GLB (0.3563), A/G (0.2650), Na<sup>+</sup> (0.3299), Cl<sup>-</sup> (0.1848), Ca<sup>2+</sup> (0.1858), and Glucose (GLU, 0.2782).

The score of all cases without missing data (985 cases) ranged from 1.65 to 6.07 of 419 cases that had missing data; 364 cases were mild, 37 cases were severe, and 18 cases were critical.

### Discrimination analysis of ROC curve

Figure 3 reveals the AUC of the proposed HFMD diagnosis scoring system. The results showed that the simple scoring system had high diagnostic accuracy for HFMD [20] (sensitivity=0.95, specificity=0.90, AUC=0.918, 95% CI= 0.874–0.963,  $P<0.01$ ). The optimal cut-off point of the simple scoring system that corresponded to the largest YI was 3.9684. The largest YI (sensitivity + specificity – 1) was 0.850. There were 4 (5%) false-negative cases, which were all mild cases. There were 12 (10%) false-positive cases, which were severe pneumonia, Kawasaki disease, urticaria, and herpangina. At prevalence=40% (80 HFMD cases out of 200 subjects), the positive and negative predictive

values of the scoring system were 86.4% and 96.4%, respectively. Therefore, at prevalence=40%, the probability of being cases among subjects who tested positive was 86.4%, and the probability being non-cases among subjects who tested negative was 96.4%.

### Sensitivity analysis

Using the scoring system in cases with completed data, 3 (10.0%) of the 29 critical cases, 23 (11.0%) of the 213 severe cases, and 222 (30.0%) of the 743 mild cases were detected as non-cases. Using the scoring system in cases with missing data, 134 (36.8%) of the 364 mild cases, 11 (29.7%) of the 37 severe cases, and 9 (50%) of the critical cases were detected as non-cases (Table 4).

### Discussion

HFMD is a major public health concern in China, seriously threatening the health of children. Despite the long history of HFMD, China still lacks effective prevention and treatment measures. Therefore, in this study, a simple scoring system was developed to diagnose HFMD quickly, accurately, and reliably at the early stage to prevent disease progression by implementing timely disease management, especially in the medical centers

**Table 2.** Mean comparison of biochemistry parameters.

Biochemical markers	Mild (n=1104)	Severe (n=252)	Critical (n=48)	P
HBDH	258±81.2	265±73.3	277±92.8	0.195
LDH	348±102	360±102	371.5±150	0.114
LDH1	91.5±36.3	92.5±31.9	106±48.2	0.034
AST	40.5±15.7	43.3±42.2	45.8±22.1	0.115
ALT	19.7±22.5	25.0±65.2	20.6±12.2	0.112
GGT	11.0±7.09 <sup>+</sup>	11.8±8.08 <sup>+</sup>	14.7±17.0 <sup>#*</sup>	<b>0.006</b>
ALP	229±74.4	238±127	242±72.6	0.247
CHE	8.57±1.95	8.65±1.95	7.98±1.75	0.147
MAO	8.70±1.94	5.40±3.56	4.98±3.13	0.384
AFU	22.3±7.20	22.0±8.16	23.3±6.59	0.606
TP	65.9±6.12	67.0±6.47	67.7±7.27	0.016
ALB	42.7±3.78 <sup>+</sup>	42.8±3.56 <sup>+</sup>	40.1±4.67 <sup>#*</sup>	<b>&lt;0.000</b>
GLB	23.3±5.34 <sup>*+</sup>	24.2±5.74 <sup>#+</sup>	27.6±7.89 <sup>#*</sup>	<b>&lt;0.000</b>
Tbil	8.94±5.05	8.48±4.73	8.25±4.54	0.332
Dbil	1.95±1.54	1.85±1.15	1.68±0.882	0.341
Ibil	7.02±4.46	6.63±4.08	6.58±4.20	0.401
A/G	1.94±0.631 <sup>+</sup>	1.88±0.527 <sup>+</sup>	1.58±0.492 <sup>#*</sup>	<b>0.001</b>
K <sup>+</sup>	4.33±0.535	4.37±0.509	4.41±0.696	0.441
Na <sup>+</sup>	137±3.63 <sup>*+</sup>	136±3.73 <sup>#</sup>	134.6±4.40 <sup>#</sup>	<b>&lt;0.000</b>
Cl <sup>-</sup>	102±3.34 <sup>*</sup>	101±3.77 <sup>#</sup>	100.6±4.20	<b>&lt;0.000</b>
Ca <sup>2+</sup>	2.42±0.171 <sup>+</sup>	2.40±0.174	2.32±0.238 <sup>#</sup>	<b>0.001</b>
CO <sub>2</sub> CP	19.0±2.98	18.9±3.10	19.1±3.63	0.855
BUN	3.29±1.29	3.13±1.14	3.03±1.37	0.103
Cr	39.2±15.8	37.2±15.0	42.8±16.8	0.049

Data are summarised as mean ±SD. # Compared with Mild Group  $P < 0.05$ , \* Compared with Severe Group, + Compared with Critical Group. HBDH –  $\alpha$ -hydroxybutyrate acid; LDH – lactate dehydrogenase; LDH1 – isoenzyme; AST – aspartate aminotransferase; ALT – alanine transaminase; GGT – gamma-glutamyl transpeptidase; ALP – alkaline phosphate; CHE – cholinesterase; MAO – monoamine oxidase; AFU –  $\alpha$ -L-fucosidase; TP – total protein; ALB – albumin; GLB – globulin; Tbil – total bilirubin; Dbil – direct bilirubin; Ibil – indirect bilirubin; A/G – albumin/globulin ratio; K<sup>+</sup> – potassium ion; Na<sup>+</sup> – sodium ion; Cl<sup>-</sup> – chloride ion; Ca<sup>2+</sup> – calcium ion; CO<sub>2</sub>CP – carbon dioxide containing power; BUN – blood urea nitrogen; Cr – creatinine.

without laboratory-confirmed tests. Our major findings were as follows: (1) This scoring system demonstrated a good discrimination in HFMD cases, with an AUC of 0.918, based on 20 clinical variables collected at the time of admission. (2) All 20 markers of this scoring system can be easily obtained and are comprehensive, including 2 basic markers, 11 markers selected by guidelines of HFMD diagnosis and treatment, and

7 laboratory parameters selected by the chi-square statistic. (3) The score of each marker is determined by calculating its positive constituent ratio, which is simple and objective and can be estimated by its weight. (4) We provided a framework and step-by-step guidelines of the simple scoring system, which can be applied to diagnose other diseases by re-adjusting the markers and scores.

**Table 3.** Score of markers.

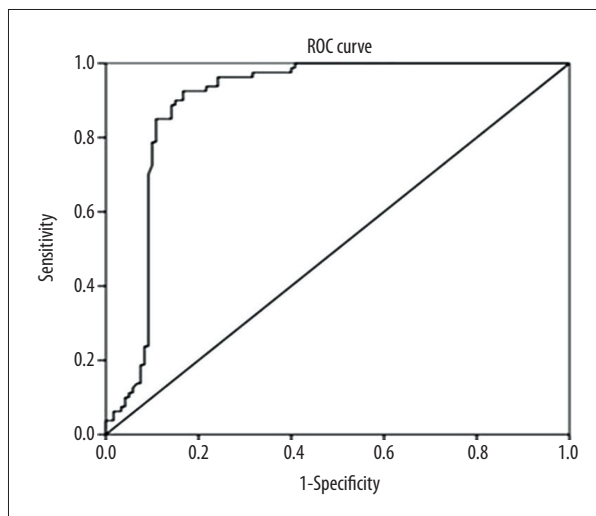
Markers	Category	Frequency	Score
Sex	Male	915	0.652
	Female	489	0.348
Age	0–3 years	1050	0.748
	4–6 years	326	0.232
	7–14 years	28	0.020
Fever	Symptomatic	882	0.895
	Asymptomatic		0
Skin rashes	Symptomatic	967	0.982
	Asymptomatic		0
Nervous system disorder	Symptomatic	234	0.238
	Asymptomatic		0
Respiratory system disorder	Symptomatic	294	0.299
	Asymptomatic		0
Digestive system disorder	Symptomatic	103	0.105
	Asymptomatic		0
Cardiopulmonary complications	Symptomatic	6	0.0061
	Asymptomatic		0
CRP	Abnormal	356	0.361
	Normal		0
WBC	Abnormal	489	0.496
	Normal		0
CK	Abnormal	151	0.153
	Normal		0
CK-MB	Abnormal	177	0.180
	Normal		0
GGT	Abnormal	6	0.0061
	Normal		0
ALB	Abnormal	224	0.227
	Normal		0
GLB	Abnormal	351	0.356
	Normal		0
A/G	Abnormal	261	0.265
	Normal		0
Na <sup>+</sup>	Abnormal	325	0.330
	Normal		0



**Table 3 continued.** Score of markers.

Markers	Category	Frequency	Score
Cl <sup>-</sup>	Abnormal	182	0.185
	Normal		0
Ca <sup>2+</sup>	Abnormal	183	0.186
	Normal		0
GLU	Abnormal	274	0.278
	Normal		0

(I) age and sex, score=(/1404); (II) 6 clinical manifestation, score= (/985); (III) 12 lab parameters, score= (/985), where n=the number of abnormal cases. CRP – c-reactive protein; WBC – white blood cell; CK – creatinine kinase; CK-MB – creatinine kinase isoenzyme; GGT – gamma-glutamyl transpeptidase; ALB – albumin; GLB – globulin; A/G – albumin/globulin ratio; Na<sup>+</sup> – natrium ion; Cl<sup>-</sup> – chloride ion; Ca<sup>2+</sup> – calcium ion; GLU – glucose.



**Figure 3.** The Receiver Operating Characteristic (ROC) curve of the simple scoring system. The curve represents the receiver operating characteristics of the simple scoring system. The blank area bounded by the curve and straight line is the area under the curve of ROC.

Our validation results showed that the simple scoring system had high diagnostic accuracy for HFMD (sensitivity=0.95, specificity=0.90, AUC=0.918, the positive predictive value=86.4%, the negative predictive value=96.4%). A previous report presented a mortality risk score model comprising 4 laboratory parameters with good discrimination (AUC >0.9) [16]; however, the model can only discriminate children with high mortality risk from the severe HFMD cases. Another study developed a prediction system for identification of the severe HFMD based on 14 variables with an AUC of 0.916 [17], but this system can only discriminate the severe HFMD cases from the mild HFMD cases. Compared to these studies previous [16,17], our study had a different purpose – to discriminate HFMD cases from non-HFMD cases. In addition, the present study exhibited superior performance in sensitivity, specificity, and AUC. One possibility is that our system covers more parameters than other studies, including clinical features, laboratory indicators, and demographic variables.

Our scoring system included a total of 12 biochemical markers: CRP, WBC, CK, CK-MB, GLU, GGT, ALB, GLB, A/G, Na<sup>+</sup>, Cl<sup>-</sup>, and Ca<sup>2+</sup>.

**Table 4.** Sensitivity and false negative of the simple scoring system in HFMD cases with complete and missing data by severity levels.

HFMD severity	Cases with complete data				Cases with missing data			
	Sensitivity		False-negative		Sensitivity		False-negative	
Mild	521/743	(70%)	222/743	(30%)	134/364	(36.8%)	230/364	(63.2%)
Severe	190/213	(89%)	23/213	(11%)	11/37	(29.7%)	26/37	(71.3%)
Critical	26/29	(90%)	3/29	(10%)	9/18	(50%)	9/18	(50%)
Total	736/984	(75%)	248/984	(25%)	154/419	(36.8%)	265/419	(63.2%)

The significant differences in severity among different cases suggest that these markers could play a role in the disease progression of HFMD. Addressed by multiple studies, some of the biochemical markers change with the severity of HFMD [18–20], and the results can be inconsistent. Previous studies [21,22] reported that GLU or WBC is the marker for HFMD disease progression, rather than the risk factors for HFMD complications. Moreover, the CRP and CK-MB levels are correlated with HFMD severity [23]. Furthermore, Cl<sup>-</sup> is an independent risk factor, along with suitable combinations of other risk factors, and can be useful for the detection of severe cases [24]. A recent study [17] found that GLU, platelet, percentage of lymphocytes, LDH, ALP, CK, CK-MB, Cr, uric acid, Cl<sup>-</sup>, and ALT are important independent risk factors for severe HFMD. In addition to the factors mentioned in previous studies, our study indicated that other factors can also be used to predict HFMD, such as GGT ALB, GLB, A/G, Na<sup>+</sup>, and Ca<sup>2+</sup>.

The criterion gold standard diagnosis for HFMD is the serological test. However, this diagnosis requires special equipment, and is time-consuming and difficult to perform in general medical centers or for every patient. In these settings, our study provides a simple and alternative method for HFMD diagnosis. All the markers in our scoring system can be collected within 1 h after hospital admission. In addition, we can get the marker scores as long as the parameter results are within the normal range; if they are outside the normal range, one does not need to know the specific value of the laboratory parameters. This is one of the important differences in our simple scoring system compared to other scoring systems [25,26].

The application of our scoring system can be extended. Although this system was built based on inpatient data, it can also be applied to outpatients. Because the markers included in this system are common and easy to obtain, most outpatients also need to undergo these physical examinations and laboratory texts. Therefore, the scoring system for the outpatients could

be developed by refining the assigned score. Since we have established a relatively simple diagnostic system, and because external data validation has proved its high performance for HFMD diagnosis, our diagnostic method can also be applied to other disease diagnostic systems.

In this study, the limitations include: first, this was a single-center study, which could be population-specific. Second, the sensitivity and false-negative results were poor in some HFMD cases with missing data by severity levels. These findings indicate that our scoring system may not be suitable when only some of the 20 parameters are available. Fortunately, these 20 parameters are available in most medical institutions in China. Third, although the results of the comparison of diagnostic values across different levels of severity indicated that the sensitivities of the critical and severe cases were greater than that of the mild cases, it cannot distinguish the severity of HFMD. Due to these limitations, this scoring system should be further improved by refining the parameters in prospective and multi-center clinical samples.

### Conclusions

In summary, we developed an alternative simple scoring system for HFMD diagnosis with high accuracy and reliability. This simple scoring system is recommended for use in similar clinical settings for early HFMD diagnosis or can be further updated or developed for use in different clinical settings by using the framework and step-by-step guidance provided in this study.

We acknowledge Jianping Xiong and Suihong Qiu for establishing the database.

### Conflict of interest

None.

### Supplementary Table

Supplementary Table 1. Cut-off scores of ROC curve.

Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI
-.141300	1.000	1.000	0.000	0.000	2.962550	1.000	.417	0.583	0.583	4.016550	.688	.083	0.917	0.604
.871200	1.000	.992	0.008	0.008	2.984350	1.000	.408	0.592	0.592	4.019000	.675	.083	0.917	0.592
1.066650	1.000	.983	0.017	0.017	2.989450	.988	.408	0.592	0.579	4.021900	.663	.083	0.917	0.579
1.324550	1.000	.975	0.025	0.025	3.033900	.988	.400	0.600	0.588	4.023300	.650	.083	0.917	0.567
1.480850	1.000	.958	0.042	0.042	3.079450	.988	.392	0.608	0.596	4.024000	.638	.083	0.917	0.554
1.594550	1.000	.950	0.050	0.050	3.085800	.988	.383	0.617	0.604	4.026550	.625	.083	0.917	0.542
1.642600	1.000	.933	0.067	0.067	3.094500	.988	.375	0.625	0.613	4.029100	.600	.083	0.917	0.517

Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI
1.699600	1.000	.925	0.075	0.075	3.107200	.988	.367	0.633	0.621	4.029750	.588	.083	0.917	0.504
1.750550	1.000	.917	0.083	0.083	3.118400	.988	.358	0.642	0.629	4.031000	.575	.083	0.917	0.492
1.760550	1.000	.908	0.092	0.092	3.137150	.988	.350	0.650	0.638	4.031900	.563	.083	0.917	0.479
1.770000	1.000	.900	0.100	0.100	3.167550	.988	.342	0.658	0.646	4.032650	.538	.083	0.917	0.454
1.792850	1.000	.883	0.117	0.117	3.194100	.988	.333	0.667	0.654	4.034550	.525	.083	0.917	0.442
1.832000	1.000	.875	0.125	0.125	3.205850	.988	.325	0.675	0.663	4.036600	.513	.083	0.917	0.429
1.876650	1.000	.867	0.133	0.133	3.212300	.988	.317	0.683	0.671	4.037700	.500	.083	0.917	0.417
1.900200	1.000	.858	0.142	0.142	3.226800	.988	.308	0.692	0.679	4.038650	.488	.083	0.917	0.404
1.914050	1.000	.850	0.150	0.150	3.239000	.988	.300	0.700	0.688	4.039700	.475	.083	0.917	0.392
1.938850	1.000	.842	0.158	0.158	3.263950	.988	.292	0.708	0.696	4.041300	.463	.083	0.917	0.379
2.016050	1.000	.833	0.167	0.167	3.290850	.988	.283	0.717	0.704	4.043300	.450	.083	0.917	0.367
2.079350	1.000	.817	0.183	0.183	3.312150	.988	.275	0.725	0.713	4.044800	.438	.083	0.917	0.354
2.082750	1.000	.808	0.192	0.192	3.338600	.988	.267	0.733	0.721	4.045800	.413	.083	0.917	0.329
2.107250	1.000	.800	0.200	0.200	3.353850	.988	.258	0.742	0.729	4.046600	.400	.083	0.917	0.317
2.150550	1.000	.792	0.208	0.208	3.363450	.988	.250	0.750	0.738	4.048600	.388	.083	0.917	0.304
2.217100	1.000	.783	0.217	0.217	3.419400	.988	.242	0.758	0.746	4.050150	.375	.083	0.917	0.292
2.266350	1.000	.775	0.225	0.225	3.502600	.975	.242	0.758	0.733	4.050700	.363	.083	0.917	0.279
2.272700	1.000	.767	0.233	0.233	3.539500	.975	.233	0.767	0.742	4.051400	.350	.083	0.917	0.267
2.283150	1.000	.758	0.242	0.242	3.562550	.975	.225	0.775	0.750	4.052100	.338	.083	0.917	0.254
2.323500	1.000	.750	0.250	0.250	3.598600	.975	.217	0.783	0.758	4.053650	.325	.083	0.917	0.242
2.356650	1.000	.733	0.267	0.267	3.624850	.975	.208	0.792	0.767	4.055000	.313	.083	0.917	0.229
2.369800	1.000	.725	0.275	0.275	3.636000	.975	.200	0.800	0.775	4.057450	.300	.083	0.917	0.217
2.391450	1.000	.717	0.283	0.283	3.646100	.975	.192	0.808	0.783	4.060500	.300	.075	0.925	0.225
2.402950	1.000	.708	0.292	0.292	3.661550	.975	.183	0.817	0.792	4.062550	.288	.075	0.925	0.213
2.412300	1.000	.700	0.300	0.300	3.673300	.975	.175	0.825	0.800	4.063800	.275	.075	0.925	0.200
2.419450	1.000	.692	0.308	0.308	3.686300	.975	.167	0.833	0.808	4.064100	.263	.075	0.925	0.188
2.438700	1.000	.683	0.317	0.317	3.722300	.975	.158	0.842	0.817	4.064350	.250	.075	0.925	0.175
2.469800	1.000	.675	0.325	0.325	3.761350	.975	.150	0.850	0.825	4.064950	.238	.075	0.925	0.163
2.490100	1.000	.667	0.333	0.333	3.776550	.963	.150	0.850	0.813	4.065650	.238	.067	0.933	0.171
2.516850	1.000	.658	0.342	0.342	3.786400	.963	.142	0.858	0.821	4.067150	.225	.067	0.933	0.158
2.540200	1.000	.650	0.350	0.350	3.806900	.963	.133	0.867	0.829	4.077100	.213	.067	0.933	0.146
2.546800	1.000	.642	0.358	0.358	3.835850	.963	.125	0.875	0.838	4.091350	.200	.067	0.933	0.133
2.553800	1.000	.633	0.367	0.367	3.874750	.950	.125	0.875	0.825	4.110800	.188	.067	0.933	0.121
2.559400	1.000	.625	0.375	0.375	3.913650	.950	.117	0.883	0.833	4.127150	.175	.067	0.933	0.108
2.576300	1.000	.617	0.383	0.383	3.945600	.950	.108	0.892	0.842	4.132050	.175	.058	0.942	0.117
2.613000	1.000	.608	0.392	0.392	3.968400	.950	.100	0.900	0.850	4.139000	.163	.050	0.950	0.113
2.634650	1.000	.600	0.400	0.400	3.982850	.938	.100	0.900	0.838	4.148700	.150	.050	0.950	0.100
2.647700	1.000	.592	0.408	0.408	3.992250	.925	.100	0.900	0.825	4.154250	.150	.042	0.958	0.108
2.685250	1.000	.583	0.417	0.417	3.994750	.913	.100	0.900	0.813	4.156950	.138	.042	0.958	0.096
2.714850	1.000	.575	0.425	0.425	3.997250	.900	.100	0.900	0.800	4.177450	.125	.042	0.958	0.083
2.719650	1.000	.567	0.433	0.433	3.998100	.888	.100	0.900	0.788	4.210250	.113	.042	0.958	0.071
2.732650	1.000	.558	0.442	0.442	3.999400	.875	.100	0.900	0.775	4.233500	.100	.042	0.958	0.058

Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI	Cut-off score	Sensitivity	1-Specificity	Specificity	YI
2.758350	1.000	.550	0.450	0.450	4.002150	.875	.092	0.908	0.783	4.244150	.088	.042	0.958	0.046
2.779850	1.000	.542	0.458	0.458	4.004200	.863	.092	0.908	0.771	4.247300	.088	.033	0.967	0.054
2.789450	1.000	.533	0.467	0.467	4.005000	.850	.092	0.908	0.758	4.268100	.075	.033	0.967	0.042
2.791300	1.000	.533	0.467	0.467	4.005700	.838	.092	0.908	0.746	4.298400	.075	.025	0.975	0.050
2.804650	1.000	.517	0.483	0.483	4.006200	.825	.092	0.908	0.733	4.369000	.075	.017	0.983	0.058
2.824500	1.000	.508	0.492	0.492	4.007700	.813	.092	0.908	0.721	4.492850	.063	.017	0.983	0.046
2.840250	1.000	.500	0.500	0.500	4.009250	.800	.092	0.908	0.708	4.708050	.050	.017	0.983	0.033
2.849900	1.000	.475	0.525	0.525	4.009600	.800	.092	0.908	0.708	4.899100	.038	.017	0.983	0.021
2.863100	1.000	.467	0.533	0.533	4.009700	.775	.083	0.917	0.692	4.966100	.038	.008	0.992	0.029
2.879800	1.000	.458	0.542	0.542	4.010200	.763	.083	0.917	0.679	5.176100	.038	0.000	1.000	0.038
2.898900	1.000	.450	0.550	0.550	4.010750	.738	.083	0.917	0.654	5.483750	.025	0.000	1.000	0.025
2.916200	1.000	.442	0.558	0.558	4.010900	.738	.083	0.917	0.654	5.837050	.013	0.000	1.000	0.013
2.931450	1.000	.433	0.567	0.567	4.013500	.700	.083	0.917	0.617	7.065500	0.000	0.000	1.000	0.000

Minimum cut-off score was the result of smallest observed test score subtracting 1; Maximum cut-off score was the result of largest observed test score adding 1. All other cut-off values were the average of 2 adjacent test scores.

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