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Performances of Clinical Characteristics and Radiological Findings in Identifying COVID- 19 From Suspected Cases

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Research Article

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Abstract

Background: To identify effective factors and establish a model to distinguish COVID-19 patients from suspected cases.

Methods: The clinical characteristics, laboratory results and initial chest CT findings of suspected COVID-19 patients in 3 institutions were retrospectively reviewed. Univariate and multivariate logistic regression were performed to identify significant features. A nomogram was constructed, with calibration validated internally and externally.

Results: 239 patients from 2 institutions were enrolled in the primary cohort including 157 COVID-19 and 82 non-COVID-19 patients. 11 features were included for multivariate logistic regression analysis after LASSO selection. We found that the COVID-19 group are more likely to have fever (OR, 4.22), contact history (OR, 284.73), lower WBC count (OR, 0.63), left lower lobe involvement (OR, 9.42), multifocal lesions (OR, 8.98), pleual thickening (OR, 5.59), peripheral distribution (OR, 0.09), and less mediastinal lymphadenopathy (OR, 0.037). The nomogram developed accordingly for clinical practice showed satisfactory internal and external validation.

Conclusions: In conclusion, fever, contact history, decreased WBC count, left lower lobe involvement, pleural thickening, multifocal lesions, peripheral distribution and absence of mediastinal lymphadenopathy are able to distinguish COVID-19 patients from other suspected patients. The corresponding nomogram is a useful tool in clinical practice.

1. Introduction:

In December 2019, a few pneumonia cases of unknown etiology were reported in Wuhan, Hubei Province, China.(1) The disease, now named ©coronavirus disease 2019 (COVID-19) then spread at a striking speed worldwide. The causative organism was identified as a novel coronavirus named severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) due to the ©phylogenetic similarity to SARS-CoV.(2) As of Oct 15th, there were a total of 238,940,176 cumulative cases and 4,882,066 cumulative deaths worldwide. COVID-19 was declared as a public health emergency of international concern (PHEIC) by the World Health Organization (WHO) as early as January 30, 2020.(3)(4)

The confirmation of COVID-19 relies on 0the positive result of the nucleic acid amplification test (NAAT) of the respiratory tract or blood specimens using the 0real-time reverse transcriptase-polymerase chain reaction (RT-PCR) tests.(5) However, the limitations of RT-PCR tests include: 1) The severity and progression of the disease cannot be quantitatively judged. 2) They have long turnaround times, especially in less developed regions; 3) They require certified laboratories, expensive equipments and trained technicians. (6, 7)

Chest CT scan is relatively easy to perform with fast diagnosis high in sensitivity, thus is considered an ideal primary tool for COVID-19 detection.(8–10) The Diagnosis and Treatment Program of COVID-19 (trail version 7)(11, 12) fomulated by the National Health Commission of China has summarized the typical CT manifestations of COVID-19 as follows and incorporated it in the diagnosis criteria: multiple small patchy shadows and interstitial changes, notably at the peripheral zone, at the early stage. As the disease progresses, multiple ground-glass opacities (GGO) and infiltration occur bilaterally, and consolidation is found in severe cases. Pleural effusion is rarely seen.

Patients with above-said CT manifestations are suspected as COVID-19 infectors therefore need further examinations. Before the RT-PCR result is available, the patient needs isolation, but the quarantine of the patients may lead to a waste of medical resources and a possible delay of essential treatment. Hence, effective and convenient methods to better distinguish COVID-19 patients are needed.

The aim of our study is to identify the useful clinical, laboratory and radiographic features that are able to distinguish COVID-19 patients from other suspected cases and generate a nomogram as a useful tool for clinical practice.

2. Materials And Methods:

The schematic workflow is depicted in Figure 1.

2.1. Patient cohort:

Data were de-identified to guarantee the patients' confidentiality. From Jan 21 to Mar 5, 2020, patients admitted to a hospital in Anhui province, China and our institution in Shanghai, China who met the following requirements were enrolled as the primary cohort in our study: 1) Patients with chest CT manifestations suggested by the Diagnosis and Treatment Program of COVID-19 (trail version 7) (12) that had a suspicion of COVID-19. 2) Patients that took laboratory examination at admission. 3) Patients with the diagnosis of COVID-19 ruled out or confirmed by the RT-PCR. Exclusion criteria included: 1) Patients who were hospitalized before (n=4). 2) Significant motion artefacts in CT images (n=12). 3) Patients lacking essential data (n=21). The epidemiological history, the symptoms, the laboratory test results and the imaging features of their first CT scan after onset were recorded.

From Feb 6 to Mar 13, 2020, an independent cohort of CT-suspected patients from another institution in Anhui Province was prospectively studied, using the same inclusion and exclusion criteria. These patients formed the validation cohort.

The laboratory tests were carried out in the outpatient department or in the wards on admission, mostly on the same day when CT scan was done. Collected laboratory indices included the white blood cell (WBC) count, lymphocyte count, lactate dehydrogenase (LDH), C reactive protein (CRP), procalcitonin (PCT), alanine aminotransferase (ALT), aspartate aminotransferase (AST).

2.2. CT protocol

105 patients from Huashan Hospital Affiliated to Fudan University were imaged with 1.5mm-thickness with a 256-slice spiral CT scanner (Philips). 134 patients from Fuyang No.2 People

 $sHos\pi talwereima \ge dwith1mm - thick \ne sswitha64 - sectionCTsca \cap er(Aquilion64, ToshibaMedicalSystems).$ 59 $patientsomBozhouPeop \le$ s Hospital in the validation cohort were imaged with 5mm-thickness with a 64-section CT scanner (Siemens Somantom Sensation).

2.3. CT manifestation analysis

All imaging data were analyzed with consensus by two experienced radiologists (12 and 7 years of experience). 23 features were collected as listed below: a) The involved pulmonary lobes including five features: right upper, right middle, right lower, left upper, left lower lobes; b) Main distribution of lesions including anterior and posterior part of lungs; c) The location of lesions that is set as dummy variables: peripheral, central or both; d) The extent of the lesions that is set as dummy variables: unifocal, multifocal and diffuse. e) An extent score was semi-quantitatively calculated. Both lungs were divided into upper (above tracheal carina), lower (below inferior pulmonary vein) and middle (in between) zones, and involved percentage in each zone was scored: 0, 0%; 1, < 25%; 2, 25% - 49%; 3, 50% - 74%; 4, > 75%, and they added up to the extent score(range 0-24). f) The existence of opacification set as dummy variables included GGO, mixed (mainly GGO), mixed (mainly consolidation) and consolidation; g) The shape of the lesions, including nodular, linear, patchy and large patchy; h) The halo sign; i) The reversed halo sign ; j) Reticulated changes; k) The existence of vascular enlargement; l) The existence of air bronchogram; m) Bronchiectasis; n) Pleural thickening; o) Pleural traction; p) Pleural effusion; q) Mediastinal lymphadenopathy (the maximal axial diameter >1 cm). The description of the radiological features followed the definition compiled by the Fleischner Society.(13) Five 1cm² regions of interests (ROI) were drawn in the liver and spleen parenchymal to obtain the mean CT values of liver and spleen. Liver spleen ratio (LS ratio) was calculated as CT_{liver}/CT_{spleen} to indicate the relative density.

2.4. Feature selection

The clinical (8), laboratory (7) and CT features (23) were analyzed altogether, but with the limited sample size, a total of 38 features would lead to overfitting in multivariate analysis. Thus, the least absolute shrinkage and selection operator (LASSO) method was adopted to select the most relevant features. This method is able to shrink the coefficients and diminish some to zero, thus can be used for feature reduction and selection. The R software and the "glmnet" package (version 3.6.0; R foundation for Statistical computing) were used.

2.5. Statistical analysis

All statistical analyses were executed with R software. The Shapiro-Wilk test was used to evaluate the distribution type and Bartlett's test was used to evaluate the homogeneity of variance. Normally distributed data were displayed as mean \pm standard deviation. Non-normally distributed data and ordinal data were displayed as median (inter-quartile range). Categorical variables were summarized as counts and percentages. Both univariate and multivariate logistic regression were analyzed to demonstrate the correlation of the features with COVID-19 diagnosis. The regression coefficient (β) was calculated using the odds ratio (OR). The model was estimated as follows:

$$\beta = \log(\text{OR})$$

$$\text{logitP} = \beta_1 \chi_1 + \beta_2 \chi_2 + \dots + \beta_i \chi_i$$

A nomogram was established. The calibration ability was internally assessed with the bootstraping method and the Hosmer-Lemeshow test (HL test) was performed to test the goodness of fit.

For the external validation of the nomogram, the prediction value of each case were calculated according to the nomogram and compared with the observed diagnosis. The accuracy was validated by correctly predicted case proportion, and the HL goodness-of-fit test. A *P*-value of <0.05 was defined as statistical significance.

2.6. IRB approval

This multi-center retrospective study was approved by the institutional review board (IRB) and the requirement of written informed consent was waived.

3. Results:

3.1 Clinical information:

In the primary cohort, 239 patients with COVID-19 (134 males and 105 females) were included in this study with an average age of 46.31 \pm 15.90 years old. 28.87% of the patients had a direct contact with confirmed COVID-19 patients before the onset or had travelled/lived in the Hubei Province. 17.57% of the patients had indirect contact. Most common symptoms the patients presented were fever (70.29%), cough (44.35%), and chest distress (11.30%). Some patients had digestive symptoms such as diarrhea 2.09%) and anorexia (2.09%). The median interval between the onset and the date of CT scan was 8 (range 1-22) days (Table 1). 157 patients were confirmed as COVID-19 by RT-PCR and were allocated to the COVID-19 group. They were put in quarantine and treated with the antiviral therapy based on the evolving recommendations.(12) The other 82 patients had negative RT-PCR results. They were eventually diagnosed as other viral pneumonia, bacterial infection, or other respiratory conditions. Clinical information of two groups were compared using univariate analysis (Table 3). COVID-19 patients were found to be younger (*P* = 0.037), more likely to have fever (*P* = 0.001) or cough (*P* < 0.001), and more likely to have contact history (*P* < 0.001).

Table 1

Clinical Characteristics	Primary cohort	Validation cohort	Pvalue
	(n = 239)	(n = 59)	
Age, mean ± SD	46.30±15.90	45.64±16.57	0.614
Gender			
Male	134 (56.07%)	31 (52.54%)	0.733
Female	105 (43.93%)	28 (47.46%)	
Epidemiological history			
Direct contact	69 (19.12%)	19 (32.20%)	0.546
Indirect contact	42 (48.53%)	13 (22.03%)	
None contact	128 (32.35%)	27 (45.76%)	
Symptom			
Fever	168 (70.29%)	47 (79.66%)	0.202
Cough	106 (44.35%)	31 (52.54%)	0.097
Chest distress	27 (11.30%)	6 (10.17%)	0.988
Diarrhea	5 (2.09%)	5 (8.47%)	0.042*
Anorexia	5 (2.09%)	1 (1.69%)	1.000
Laboratory Test, median (inter-quartile range)			
WBC, median (range), ×10 ⁹ /L	5.28 (4.30-10.44)	5.96 (3.91-6.00)	0.101
Lymphocyte count, median (range), ×10 ⁹ /L	1.19 (0.90-1.63)	1.21 (0.85-1.44)	0.746
LDH, median (range), U/L	233.00 (193.00-271.40)	234 (199-290)	0.158
CRP, median (range), mg/L	14.80 (4.8-42.93)	25.90 (3.7-30.30)	0.038*
PCT, median (range), ng/mL	0.05 (0-0.19)	0.04 (0.02-0.06)	0.743
ALT, median (range), U/L	30.00 (20.00-51.50)	29.90 (17.30-37.70)	0.558
AST, median (range), U/L	28.00 (21.00-46.75)	28.00 (20.40-34.70)	0.450
Abbreviations: WBC: White blood cell count: I DH: Lactate	debudrogenase: CPD: C-reactive protein: DC	T: Procelcitonin: ALT: Alenine emi	notransferaça: AST.

Abbreviations: WBC: White blood cell count; LDH: Lactate dehydrogenase; CRP: C-reactive protein; PCT: Procalcitonin; ALT: Alanine aminotransferase; AST: Aspartate aminotransferase

3.2 Laboratory Tests

Compared with the non-COVID-19 group, COVID-19 group showed lower WBC (P < 0.001) and lymphocyte count(P = 0.002), lower levels of PCT(P = 0.002). (Table 3)

3.3 Chest CT imaging findings

Imaging characteristics were assessed and compared between two groups (Table 2, 3). Regarding the location and the distribution of the lesions, COVID-19 patients were found to be more located in posterior part of the lungs (P < 0.001) compared with non-COVID-19 patients. They had more involvement in every lobe of the lung (P < 0.05) due to more multifocal distribution (P < 0.001). Besides, they are more likely to have specific signs including reticular changes (P = 0.04), vascular enlargement (P < 0.001), air bronchogram (P = 0.043), and pleural thickening (P < 0.001). They are less likely to show pleural effusion (OR = 0.16, P = 0.007) or mediastinal lymphadenopathy (P < 0.001). Other parameters were not significantly different.

Imaging manifestations on chest CT of the primary and validation cohort							
Imaging manifestation	Primary cohort	Validation cohort	<i>P</i> value				
	(n = 239)	(n = 59)					
Involved lobes:							
Right Upper Lobe	144 (60.25%)	39 (66.1%)	0.498				
Right Middle Lobe	129 (53.97%)	32 (54.24%)	1.000				
Right Lower Lobe	179 (74.9%)	39 (66.1%)	0.230				
Left Upper Lobe	143 (59.83%)	40 (67.8%)	0.329				
Left Lower Lobe	176 (73.64%)	44 (74.58%)	1.000				
Main distribution:							
Anterior Part of Lungs	44 (18.41%)	18 (30.51%)	0.061				
Posterior Part of Lungs	168 (70.29%)	40 (67.8%)	0.847				
Location of lesions:							
Peripheral	158 (66.11%)	33 (55.93%)	0.191				
Central	16 (6.69%)	2 (3.39%)	0.516				
Both	65 (27.2%)	24(40.68%)	0.482				
Extent of lesions:							
Unifocal	58 (24.27%)	16 (27.12%)	0.775				
Multi-focal	141 (59%)	26 (44.07%)	0.055				
Diffuse	40 (16.74%)	17 (28.82%)	0.971				
Extent score:	4 (2-5)	5 (3-7)	0.057				
Density of lesions:							
GGO	77 (32.22%)	11(18.64%)					
Mixed (Mainly GGO)	98 (41.00%)	27 (45.76%)	0.606				
Mixed (Mainly Consolidation)	57 (23.85%)	20 (33.9%)	0.158				
Consolidation	7 (2.93%)	1(1.69%)	0.940				
Shape of lesions:							
Nodular	1 (0.42%)	1 (1.69%)	0.853				
Linear	5 (2.09%)	3 (5.08%)	0.410				
Patchy	161 (67.6%)	41 (69.49%)	0.875				
Large patchy	72 (30.13%)	14 (23.73%)					
Halo sign	67 (28.03%)	22 (37.29%)	0.218				
Reverse halo sign	11 (4.60%)	2 (3.39%)	0.958				
Reticulation	61 (25.52%)	11 (18.64%)	0.349				
Air bronchogram	85 (35.56%)	26 (44.07%)	0.289				
Bronchiectasis	25 (10.46%)	2 (3.39%)	0.150				
Vascular enlargement	82 (34.31%)	21 (35.59%)	0.974				
Pleural thickening	101 (42.26%)	27 (45.76%)	0.734				
Pleural traction	60 (25.10%)	15 (25.42%)	1.000				
Pleural effusion	12 (5.02%)	6 (10.17%)	0.237				
Mediastinal Lymphadenopathy	23 (9.62%)	7 (11.86%)	0.787				
Liver-spleen CT value ratio	1.17 (1.05-1.27)	1.19 (1.07-1.37)	0.278				
Abbreviations: GGO: Ground-alass	sopacities						

Table 2

Univariate logistic regression analys	sis of features for differentiating COVID-19 p	patients and non-COVID patients in Primary	cohort

Features	Non-COVID	p-19	COVID-19	Coefficient		OR	<i>P</i> value
	(n=82)		(n=157)				
Clinical characteristics							
Age, mean ± SD	49.29±17.4	19	44.75±14.82	-0.02		0.98	0.037*
Gender, male/female	50/32		84/73	-0.31		0.74	0.270
Epidemiological history#							
Direct contact	1 (1.22%)		68 (43.31%)	4.13		61.89	<0.001*
Indirect contact	3 (3.66%)		39 (24.84%)	2.16		8.70	<0.001*
None contact	78 (95.12%)	50 (31.85%)	-3.73		0.02	<0.001*
Symptom							
Fever	42 (51.22%)))	126 (80.25%)	1.35		3.87	<0.001*
Cough	24 (29.27%)))	82 (52.23%)	0.47		1.60	0.084
Chest distress	9 (10.98%)		18 (11.46%)	0.05		1.05	0.910
Diarrhea	1 (1.22%)		4 (2.55%)	0.75		2.12	0.505
Anorexia	1 (1.22%)		5 (2.55%)	0.75		2.12	0.505
Laboratory Test, median (range)							
WBC, ×10 ⁹ /L	8	3.72±4.15	5.068±1.80		-0.54	0.58	<0.001*
Lymphocyte count, ×10 ⁹ /L	1	.42±0.68	1.18±0.47		-0.77	0.46	0.002*
LDH, U/L	2	231.78±109.50	250.66±72.02		0.003	1.00	0.114
CRP, mg/L	3	31.08±40.56	23.06±29.40		-0.01	0.99	0.089
PCT, ng/mL	0).91±4.28	0.07±0.13		-3.56	0.03	0.002*
ALT, U/L	4	17.80±32.60	38.51±61.19		-0.003	1.00	0.226
AST, U/L	4	4.95±40.05	34.38±43.01		-0.01	0.99	0.091
Imaging manifestation							
Involved lobes:							
Right Upper Lobe	3	35 (42.68%)	109 (69.43%)		1.12	3.05	<0.001*
Right Middle Lobe	3	86 (43.90%)	93 (59.24%)		0.62	1.86	0.025*
Right Lower Lobe	4	18 (58.54%)	131 (83.44%)		1.27	3.57	<0.001*
Left Upper Lobe	3	36 (43.90%)	107 (68.15%)		1.01	2.73	0.001*
Left Lower Lobe	4	12 (52.44%)	123 (84.71%)		1.62	5.03	<0.001*
Main distribution:							
Anterior Part of Lungs	1	9 (23.17%)	25 (15.92%)		-0.47	0.63	0.172
Posterior Part of Lungs	4	15 (54.88%)	123 (78.34%)		1.06	2.88	<0.001*
Location of lesions: #							
Peripheral	4	19 (59.76%)	109 (69.43%)		0.43	1.53	0.135
Central	1	2 (14.63%)	4 (2.55%)		-1.88	0.15	0.002*
Both	2	21 (25.61%)	44 (28.02%)		0.12	1.13	0.690
Extent of lesions: #							
* <i>P</i> value < 0.05 indicates statistical signit	ficance,						

[#]Set as dummy variables in feature selection and Logistic model analysis

Abbreviations: WBC: White blood cell count; LDH: Lactate dehydrogenase; CRP: C-reactive protein; PCT: Procalcitonin; ALT: Alanine aminotransferase; AST: Aspartate aminotransferase

Features	Non-COVID-19	COVID-19	Coefficient	OR	Pvalue
	(n=82)	(n=157)			
Unifocal	41 (50.00%)	17 (10.83%)	-2.11	0.12	<0.001*
Multi-focal	28 (34.15%)	113 (71.97%)	1.60	4.95	<0.001*
Diffuse	13 (15.85%)	27 (17.20%)	0.10	1.10	0.792
Extent score:	4.41±5.32	5.48±3.59	0.07	1.07	0.072
Density of lesions: #					
GGO	35 (42.68%)	42 (26.75%)	-0.71	0.49	0.013*
Mixed (Mainly GGO)	26 (31.70%)	72 (45.86(%)	0.60	1.82	0.036*
Mixed (Mainly Consolidation)	18 (21.95%)	39 (24.84%)	0.16	1.18	0.619
Consolidation	3 (3.66%)	4 (2.54%)	-0.37	0.69	0.631
Shape of lesions: #					
Nodular	0 (0%)	1 (0.63%)	13.92	1113402.31	0.987
Linear	0 (0%)	5 (3.18%)	14.95	3106188.55	0.982
Patchy	56 (68.29%)	106 (66.88%)	-0.07	0.94	0.825
Large patchy	26 (31.71%)	46 (29.30%)	-0.11	0.89	0.700
Halo sign	22 (26.83%)	45 (28.66%)	0.09	1.10	0.765
Reverse halo sign	2 (2.44%)	9 (5.73%)	0.89	2.43	0.263
Reticulation	11 (13.41%)	50 (31.85%)	1.10	3.02	0.003*
Air bronchogram	22 (26.83%)	63 (31.85%)	0.60	1.83	0.043*
Bronchiectasis	8 (9.76%)	17 (10.83%)	0.12	1.12	0.797
Vascular enlargement	14 (17.07%)	68 (43.31%)	1.31	3.71	<0.001*
Pleural thickening	17 (20.73%)	84 (53.50%)	1.48	4.40	<0.001*
Pleural traction	16 (19.51%)	44 (28.03%)	0.47	1.61	0.152
Pleural effusion	9 (10.98%)	3 (1.91%)	-1.85	0.16	0.007*
Mediastinal Lymphadenopathy	20 (24.39%)	3 (1.91%)	-2.81	0.06	<0.001*
Liver-spleen CT value ratio	1.18(1.02-1.29)	1.17(1.06-1.35)	0.11	1.12	0.826
* <i>P</i> value < 0.05 indicates statistical signi	ficance,				
[#] Set as dummy variables in feature selection and Logistic model analysis					
Abbreviations: WBC: White blood cell cou Aspartate aminotransferase	ınt; LDH: Lactate dehydrogenase	e; CRP: C-reactive prot	ein; PCT: Procalcitonin; ALT	: Alanine aminotrar	nsferase; AST:

3.4. Feature selection

In LASSO model, the λ value of 0.0376 with log (λ) of -3.280 chosen (1-SE criteria), and a total of 38 features were reduced to 11 potential features with nonzero coefficients on the basis of 239 patients (21.7:1 ratio; Figure 2). These features were further incorporated in the multivariate logistic analysis (Table 4). Eight features were found to be statistically significant. COVID-19 group tended to have more fever (OR, 4.22; 95% CI [confidence interval], 1.09-18.63; *P* = 0.043), less probability of no contact history (meaning higher probability of indirect or direct contact history [OR, 284.73; 95% CI, 38.17-4214.18; *P* < 0.001]), lower WBC count (OR, 0.63; 95% CI, 0.47-0.77; *P* < 0.001), more involving left lower lobe (OR, 9.42; 95% CI, 1.95-62.80; *P* = 0.010), more exhibiting multifocal lesions (OR, 8.98; 95% CI, 1.58-61.36; *P* = 0.017), more pleural thickening (OR, 5.59; 95% CI, 1.32-28.85; *P* = 0.026), less located in central part (OR, 0.09; 95% CI, 0.01-0.75; *P* = 0.043), and less mediastinal lymphadenopathy (OR, 0.037; 95% CI, 0.00-0.29; *P* = 0.004).

Table 4 Multivariate logistic regression analysis of features for differentiating COVID-19 patients

Features	Coefficient	OR	95%CI	Pvalue		
Fever	1.44	4.22	(1.09,18.63)	0.043*		
Epidemiological history: None contact	-5.65	0.00	(0.00,0.03)	<0.001*		
WBC count	-0.47	0.63	(0.48,0.77)	<0.001*		
Lesion involvement: Unifocal	0.11	1.12	(0.12,10.58)	0.919		
Lesion involvement: Multi-focal	2.19	8.98	(1.59,61.36)	0.017*		
Involved lobes: Right Upper lobe	1.12	3.05	(0.75,13.21)	0.121		
Involved lobes: Left Upper Lobe	0.77	2.16	(0.51,9.52)	0.295		
Involved lobes: Left Lower Lobe	2.24	9.42	(1.95,62.80)	0.010*		
Pleural thickening	1.72	5.59	(1.32,28.85)	0.026*		
Mediastinal lymphadenopathy	-3.30	0.04	(0.00,0.29)	0.004*		
Distribution Central	-2.45	0.09	(0.01,0.75)	0.043*		
* <i>P</i> value < 0.05 indicates statistical significance,						
Abbreviations: WBC: White blood cell count.						

3.5. Nomogram

A nomogram was constructed based on the multivariate Logistic analysis model. The adjusted C-index of the nomogram was 0.97 (Figure 3A). The calibration curve was determined with bootstrap analysis to get bias-corrected estimation. It indicated great agreement between the prediction and the actual diagnosis in the probability (Figure 3B). The HL goodness-of-fit test showed good calibration as well (P = 0.4797). The CT images of two cases illustrated the application of the nomogram. (Figure 4)

3.6. External validation

The validation cohort included 59 cases with 43 COVID-19 and 16 non-COVID. The baseline data were collected in Table 1 and Table 2. 56 out of 59 cases were correctly predicted using the nomogram, reaching an accuracy of 94.91%. Calibration was good (P = 0.9956 for the HL goodness-of-fit test).

4. Discussion:

An ongoing outbreak of COVID-19 originated from Hubei province in China has been spreading worldwide. Experts in infectious and respiratory diseases, critical care, and radiology from all over the world have been making a joint effort to contain the epidemic situation.(14) Presently, RT-PCR is the standard confirmative method in spite of a few flaws including long turnaround time for the results in underdeveloped regions and low sensitivity especially in the early phase of the disease.(14)(15) On the contrary, chest CT scan is able to recognize the lesions at earlier stages with high sensitivity, thus is considered an important tool for screening.(8) Typical radiographic features on chest CT in COVID-19 patients were reported to predominantly include bilateral and peripheral GGOs and consolidative pulmonary opacities.(8, 16–18) Less typical signs included linear opacities, "crazy-paving" pattern and the reverse halo sign, etc.(8, 19–23) Before the RT-PCR results are attainable, the quarantine needed, but the isolation site is insufficient, and it possibly delays essential treatment. Thus, we investigated the differential values of clinical characteristics, laboratory results and CT features to better distinguish COVID-19 patients.

The most common symptom in the patients we enrolled is fever, followed by cough and chest distress. As a differential feature, fever is significant in both univariate and multivariate analysis. This echoes previous studies, and fever is the leading symptom listed in the case definition for surveillance of COVID-19 by the Chinese Health Commission.(12, 24, 25) Therefore, it is necessary to monitor body temperature and at-home temperature measurement is a useful and easy way for the public to early notice. Additionally, we noticed a small portion of the patients with digestive disorders like diarrhea and anorexia, and it occurred more in the COVID-19 group. Increasing evidence shows the manifestation of COVID-19 is not always confined to respiratory symptoms, but may also involve other systems, e.g., the central nervous system.(26)(27)

The contact history is another valuable factor for COVID-19, including direct contact with COVID-19 patients, direct exposure in Hubei Province or other districts with confirmed cases, and indirect contact with those who were exposed.(28) According to the National Health Commission of China, a patient with one exposure or contact history and two clinical conditions can be regarded as a suspected case. (12) However, with the swift spread of the disease, some contact history is unrevealed, making it harder to contain the epidemic.(29) More active precaution and isolation is needed.

Among the laboratory parameters, WBC count is significantly lower in COVID-19 group in both univariate and multivariate analysis, and lymphocyte count is lower in univariate analysis. This is consistent with previous findings and the criteria by the IChinese Health Commission.(1, 10, 12) We also found lower levels of CRP and PCT in the COVID-19 group. They are useful indicators of infection or inflammation, and CRP was previously reported to increase in COVID-

19 patients by some researchers.(8)(30) Our finding may result from higher extent of increased levels of these indices in non-COVID-19 patients since they had other inflammatory conditions including bacterial infection, while other studies used healthy controls.

The location of the lesions varied among studies, yet the peripheral site is most frequently reported. In this study, COVID-19 lesions were less seen in the central lung compared with non-COVID-19 group in both univariate and multivariate analyses, and the parameter of the left lower lobe involvement is a significant distinct finally included in nomogram construction. Besides, compared with non-COVID-19 cases, COVID-19 is more likely to exhibit multifocal distribution rather than unifocal changes, and more likely to have reticulated changes, vascular enlargement, and pleural thickening. COVID-19 patients are also less likely to have pleural effusion and mediastinal lymphadenopathy, which is consistent with prior researches.(19)

Fever, contact history, decreased WBC count, left lower lobe location, pleural thickening, multifocal lesions, peripheral distribution, and absence of mediastinal lymphadenopathy were found to be features independently associated to COVID-19 patients. On the basis of these parameters, a nomogram was built to better interpret our findings, which is popular in cancer research these years.(31) According to our nomogram, the point of each feature adds up to a total score with a corresponding probability of COVID-19.

A nomogram can be validated by both internal and external validation.(32) In this study, internal validation used the data of the same cohort for the generation of the nomogram, and external validation used the data from another institution. Both internal and external validation indicated good agreement between the prediction and the actual diagnosis in the probability.

In summary, this study is the first to investigate the features to distinguish confirmed COVID-19 patients from CT-suspected cases, which is a critical challenge in clinical practice before RT-PCR results are available. The nomogram can be used as an instant tool able to provide practical reference for individualized management for every suspected patient and is likely to offer effective and scientific basis for empirical treatment.

Our study had several limitations. Firstly, in this multi-center study, the normal range and results of the laboratory data might be different due to the differences in the kits, equipment, and environmental conditions. However, three institutions are all China's Grade-A Tertiary Hospitals, with laboratories of the highest qualifications, and similar protocols are adhered, thus the results are relatively stable. Secondly, the sample size is relatively small since no data was obtained from the epicenter of the outbreak. Besides, despite being the standard confirmative test, RT-PCR has false-negative probabilities, therefore our results might be biased since non-COVID-19 group might include infected patients. Further prospective investigation with larger sample size and evolved diagnostic techniques is expected.

5. Conclusion:

In conclusion, fever, contact history, decreased WBC count, left lower lobe involvement, pleural thickening, multifocal lesions, peripheral distribution, and absence of mediastinal lymphadenopathy are able to distinguish COVID-19 patients from other suspected patients. The nomogram based on these features is a useful tool in the clinical practice.

Abbreviations:

COVID-19: Coronavirus Disease 2019 SARS-CoV-2: Severe Acute Respiratory Syndrome Coronavirus 2 WHO: World Health Organization PHEIC: Public Health Emergency of International Concern NAAT: Nucleic Acid Amplification Test RT-PCR: Reverse Transcriptase-polymerase Chain Reaction CT: Computed Tomography IRB: Institutional Review Board WBT: White Blood Count CPR: C-reactive protein LDH: Lactate Dehydrogenase PCT: Procalcitonin GGO: Ground-glass Opacity RHS: Reversed Halo Sign ROI: Region of Interest LASSO: Least Absolute Shrinkage and Selection Operator

OR: Odds Ratio

LR ratio: Liver/Spleen ratio

95% CI: 95% confidence interval

Declarations:

Ethics approval and consent to participate

This multi-center retrospective study was approved by the institutional review board (IRB) and the requirement of written informed consent was waived.

Consent for publication

All co-authors have read and approved of its submission to this journal.

Availability of data and materials

The datasets generated and/or analysed during the current study are not publicly available due to ethical restrictions but are available from the corresponding author on reasonable request.

Authors' Contributions

XL: drafted the work

YZ (Yajing Zhao): drafted the work

YL: substantively revised the work

- YZ (Yingyan Zheng): analysis
- NM: interpretation of data
- QH: interpretation of data
- ZR: analysis
- AX: acquisition
- XQ: acquisition
- DW: design of the work

BY: conception

All authors have read and approved the manuscript

Competing interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Ethics Accordance

This study was approved by the institutional review board (IRB), and the methods were carried out in accordance with the Declaration of Helsinki.

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Figures



Figure 1

Workflow of the whole study.



Figure 2

Feature selection using the least absolute shrinkage and selection operator (LASSO) binary logistic regression model. (A) The parameter (λ) in the LASSO model used 10-fold cross-validation based on minimum criteria. The mean squared error was plotted versus log(λ). Dotted vertical lines were drawn at the optimal values by using the minimum criteria and the 1 standard error of the minimum criteria (the 1-SE criteria). (B) The plot of LASSO coefficient profiles was produced against the log (λ) sequence. The dotted vertical line was drawn at the optimal values by using the minimum criteria, and the 1 standard error of the minimum criteria (the 1-SE criteria). (B) The plot of LASSO coefficient profiles the minimum criteria (the 1-SE criteria), and the latter was chosen with the λ value of 0.0376 and log (λ) of -3.280 according to the 10-fold cross-validation that resulted in 11 nonzero coefficients.



Figure 3

The nomogram and calibration curves based on significant features in multivariate analysis. (A) A nomogram was built on the basis of eight significant features in multivariate Logistic model. If a patient is suspected to be COVID-19 by radiological diagnosis, the data needed includes whether he has fever, contact history, decreased WBC count, left lower lobe involvement, pleural thickening, multifocal lesions, peripheral distribution or absence of mediastinal lymphadenopathy. The point of each feature adds up to a total score with a corresponding probability of COVID-19. (B) The calibration curve was determined with bootstrap analysis to get bias-corrected estimation. It indicated great agreement between the prediction and the actual grouping in the probability.



Figure 4

Two representative cases to illustrate the application of the nomogram. (A) A 40-year-old male patient complained of fever for 4 days (score \approx 80). He had travelled to Huangshi, a city in Wuhan Province, China a week before the onset (score \approx 100). His laboratory tests indicated leukocytopenia (1.99*10^9/L, score \approx 92). His chest CT showed patchy ground glass opacities with vascular enlargement and reticular changes on bilateral lower lobes (left lower lobe involvement: score \approx 83; multifocal: score \approx 85). Lesions were located both central and peripheral (score \approx 80). No mediastinal lymphadenopathy was observed (score \approx 80). Slight pleural thickening was observed (score \approx 85). Total estimated score reached around 687, indicating >99.8% probability to be a COVID-19 case. He was later confirmed by RT-PCR. (B) A 60-year-old female patient complained of fever for 3 days (score \approx 80). She claimed no contact or exposure history (score \approx 60). Her WBC count is slightly elevated (10.52*10^9/L, score \approx 60). Her chest CT showed unifocal (score \approx 80). Mediastinal lymphadenopathy was observed in mediastinal window (score \approx 60). No pleural thickening(score \approx 70) . Total estimated score reached around 541, indicating <0.2% probability to be a COVID-19 case. She was radiologically suspected as COVID-19, but the diagnosis of COVID-19 was ruled out by a negative RT-PCR. She was finally diagnosed with Respiratory syncytial virus infection.