1	Title: Prognostic factors for COVID-19 pneumonia progression to severe
2	symptom based on the earlier clinical features: a retrospective analysis
3	Running title: Prognostic factors for severe COVID-19
4	Author: Huang Huang ¹ *, Shuijiang Cai ¹ *, Yueping Li ¹ , Youxia Li ¹ , Yinqiang
5	Fan ¹ , Linghua Li ¹ , Chunliang Lei ¹ , Xiaoping Tang ¹ , Fengyu Hu ¹ , Feng Li ^{1†} ,
6	Xilong Deng ^{1†} .
7	
8	Affiliation:
9	1. Guangzhou Eighth People's Hospital, Guangzhou Medical University,
10	Guangzhou, China.
11	
12	* These authors contributed equally to this work.
13	[†] These authors contributed equally to this work as senor authors.
14	Correspondence
15	Feng Li, E-mail: gz8h lifeng@126.com, Tel/Fax: 86-20-83844171
16	Xilong Deng, E-mail: gz8hdxl@126.com
17	Address: Guangzhou Eighth People's Hospital, Guangzhou Medical University
18	8 Huaying Rd, Guangzhou, Guangdong Province, 510440, China
19	
20	Words account: Abstract, 245 words; Text, 2334 words.
21	

- 22 Summary: With our successful experience of treating COVID-19 patients, we
- 23 retrospectively found that routine clinical features could reliably predict severe
- 24 pneumonia development, thus provide quick and affordable references for
- 25 physicians to save the otherwise fatal patients with the limited medical
- 26 resource.
- 27

28 Abstract

Approximately 15-20% of COVID-19 patients will develop severe 29 pneumonia, about 10 % of which will die if not properly managed. Earlier 30 discrimination of the potential severe patients basing on routine clinical and 31 laboratory changes and commencement of prophylactical management will not 32 only save their lives but also mitigate the otherwise overwhelmed health care 33 burden. In this retrospective investigation, the clinical and laboratory features 34 were collected from 125 COVID-19 patients, who were classified into mild (93 35 cases) or severe (32 cases) groups according to their clinical outcomes after 3 36 to 7-days post-admission. The subsequent analysis with single-factor and 37 multivariate logistic regression methods indicated that 17 factors on admission 38 39 differed significantly between mild and severe groups, but that only comorbid with underlying diseases, increased respiratory rate (>24/min), elevated C-40 reactive protein (CRP >10mg/liter), lactate dehydrogenase 41 and (LDH >250U/liter), were independently associated with the later disease 42 development. Finally, we evaluated their prognostic values with the receiver 43 operating characteristic curve (ROC) analysis and found that the above four 44 factors could not confidently predict the occurrence of severe pneumonia 45 individually, but that a combination of fast respiratory rate and elevated LDH 46 significantly increased the predictive confidence (AUC= 0.944, sensitivity= 47 0.941, and specificity= 0.902). A combination consisting of 3- or 4-factors could 48 further increase the prognostic value. Additionally, measurable serum viral RNA 49

50	post-admission independently predicted the severe illness occurrence. In
51	conclusion, a combination of general clinical characteristics and laboratory
52	tests could provide high confident prognostic value for identifying potential
53	severe COVID-19 pneumonia patients.
54	
55	Keywords: COVID-19, SARS-CoV-2, risk factor, clinical manifestation,
56	prognostic factor

- 57
- 58
- 59

60 Background

The novel coronavirus (SARS-CoV-2) seems to sweep across the globe 61 ever since its first successful jump from bat to the human being through a yet 62 unknown intermediate(s) approximately in late Nov 2019, still showing a 63 tendency of explosive number increase worldwide [1-3]. The SARS-CoV-2 virus 64 seems more contagious than its sibling virus, severe acute respiratory 65 syndrome (SARS) virus which outbroke in 2003, because over 120,000 66 individuals contract COVID-19 pneumonia within three months by March 11, 67 68 which was about 15 times of total SARS cases (8000 in 7 months) [4]. The surging increase of COVID-19 patients within a short time window severely will 69 absorb and occupy the limited medical resource, including physicians, nurses, 70 protective suits, masks, and goggles. Data from the China mainland showed 71 that the majority of total infected patients will recover under simple supervision 72 management, such as guarantined in compartment hospital isolated ward, but 73 74 that the overall case fatality rate was 2.3% [5]. For the clinical treatment of COVID-19 patients under shortage of enough medical supplies, the critical 75 76 issue and priority are to treat the severe COVID-19 patients (about 20% of the whole population [5]) and to save their lives with preventive and intensive 77 medical care. However, The clinical presentation of COVID-19 patients differed 78 substantially, including asymptomatic infection, mild upper respiratory tract 79 80 illness, and severe viral pneumonia[2, 6-8]. Therefore, the most crucial issue is to identify these patients and prioritize their treatment strategy by applying 81 Page 5

prophylactically medical treatment and management before they progress to
 the severe stage.

As known, the respiratory function worsens in the severe stage. In the 84 clinical practice, saturated oxygen (< 93% in rest state), reparatory rates (>30 85 times/min), and deteriorated chest radiology imaging (X-Ray and CT more high 86 resolution) provide references to confirm their severity [5, 9, 10]. Because of 87 the hypoxia stress, most patients will experience an over reactivated immune 88 89 storm, including elevated their expression level of some specific immunological 90 cytokines and changes of certain types of immune cell counts [6, 11]. Biopsy analysis also showed that the lung bilateral diffuse alveolar damage with 91 cellular fibromyxoid exudates [12]. However, the CT imaging and immunology 92 detection is not only expensive but also far unavailable as for the explosive 93 increase of suspected cases, in particular in those hospitals not well equipped. 94 Can some routine clinical characteristics or/and laboratory measurement, or 95 their combinations predict the occurrence of severe cases? 96

In this study, we retrospectively analyzed the clinical characteristics of those patients who progressed to severe pneumonia later and found that five simple clinical features and laboratory detection at an earlier time point could serve as prognostic factors facilitating discrimination of severe cases in advance.

103 Methods

104 **Patients**

105 COVID-19 diagnosis was according to the criteria in the new Coronavirus 106 pneumonia diagnosis and treatment plan (trial version 6) issued by the National 107 Health and Health Commission [13]. All 298 COVID-19 patients admitted to 108 Guangzhou Eighth People's Hospital from January 20 to February 29, 2020, 109 were included in this study. This study complied with the medical ethics of 110 Guangzhou Eighth People's Hospital. We obtained written consent from the 111 patients.

For this analysis, inclusion criteria were as the following: 1. diagnosed as 112 mild or ordinary on admission; 2. The length of hospitalization > 3 days, and the 113 overall duration of the disease > 7 days. Then, gualified patients were classified 114 into mild symptom group and severe symptom group based on the clinical 115 116 manifestation. The severe symptom diagnosis was according to criteria as following: 1) Respiratory distress, RR \geq 30 times/min in the resting state; 2) 117 Oxygen saturation \leq 93% in the resting state; 3) Arterial blood oxygen partial 118 pressure (PaO2) / oxygen concentration (FiO2) \leq 300mmHg). The rest of the 119 patients were in the mild group. 120

121 Data collection

Patient general information including gender, age, underlying diseases,
 epidemic history, etc. and their clinical data including symptoms, signs, clinical
 Page 7

classification (course duration> 7 days), laboratory test results and SARS-CoV2 viral test results were obtained with standardized data collection forms from
electronic medical records.

127 Statistical analysis

Quantitative data was firstly tested to be normality distribution with the 128 Kolmogorov-Smirnov method. Then, for normalized distributed data, t-test and 129 Tamhane T2 methods were used for variance even and uneven data, 130 respectively. For non-normal data, which was expressed as the median 131 (quartile) [M (P25, P75)], the Mann-Whitney U test was employed. The chi-132 square test (or Fisher exact probability method) was utilized for analyzing 133 qualitative data. Logistic regression analysis and the receiver operating 134 characteristic curve (ROC) analysis was employed to analyze the independent 135 risk factors. The difference was statistically significant at P <0.05. All analysis 136 was performed using SPSS software (version 20.0). 137

138

139 **Results**

140 **Patient general Information**

298 COVID-19 cases (about 85% of total cases in Guangzhou, China) were
admitted to Guangzhou Eighth People's Hospital for treatment from January 20
to February 29, 2020 (Fig. 1). According to the inclusion criteria, 173 cases were

excluded for reasons that 23 cases were already in severe symptom stage, 52 144 cases for a short hospitalization time of < 7 days, and 98 patients for other 145 defects, such as short of a complete set of detection. Finally, 125 cases, 146 including 63 males and 62 females, were qualified to be included for further 147 investigation, and all their disease courses were over seven days, with a 148 maximum of 32 days. Based on the severity of disease at 3-days post-149 admission, 93 patients fell in the mild group (general 38 cases and mild 55 150 cases) and 32 patients in severe group (severe 25 cases and critical 7 cases). 151

152 All included patients aged from 1.5 to 91 years (averaged 44.87 ± 18.55) years) (Table 1). Among them, 37 cases had at least one underlying disease, 153 including 20 cases with hypertension, 8 cases with diabetes, 5 cases with 154 coronary heart disease, 2 cases with chronic obstructive pulmonary disease, 2 155 cases with chronic kidney disease, 2 cases with chronic liver disease, 2 cases 156 with sleep apnea syndrome. Five individuals with two or more basic disorders 157 and 7 cases with obesity (BMI> 26). Epidemiologically, 88 cases had a history 158 of traveling to or living in the Hubei epidemic area before disease onset. 159 160 Interestingly, we observed that seven patients developed serum SARS-CoV-2 viral RNA positive after admission but ahead of diagnosis to be a severe 161 symptom. 162

163

Factors differed between the mild group and the severe group

164 The single-factor analysis was applied for each factor between the mild

group and the severe group (Table 1). More patients in the severe group were 165 old, with obese (BMI> 26), and with underlying diseases, especially with 166 hypertension and diabetes (P < 0.05) compared with the mild group. Among the 167 general factors, no significant difference showed in gender, history of traveling 168 to or living in the epidemic region, coughing, sneezing, muscle joint pain, 169 headache, fatigue, and gastrointestinal symptoms between these two groups 170 (P>0.05). However, more patients in the severe group were with high fever and 171 chest tightness and breath shortness (fast respiratory rate) (P < 0.05). The 172 173 serum concentration of C-reactive protein, procalcitonin, D-dimer, albumin, and lactate dehydrogenase (LDH) increased significantly in the severe group (P 174 <0.05). 175

Compared to the mild group, patients in the severe group had lower 176 absolute lymphocyte counts and higher eosinophil counts (P < 0.05), and similar 177 levels of other parameters, including white blood cells, neutrophils, platelets, 178 hemoglobin, prothrombin time, activated partial thromboplastin time, blood 179 lactic acid, blood creatinine, creatine kinase. Interestingly, the levels of 180 glutamate aminotransferase (ALT) and aspartate aminotransferase (AST) 181 significantly increased for severe patients (P<0.05). However, the median 182 values of ALT and AST were still within the normal range, indicating that most 183 of the severe COVID-19 patients had no significant liver damage. 184

185

Importantly, all seven patients with the presence of SARS-CoV-2 viral RNA

in blood during the hospitalization, but before being in the severe stage, finally
 progressed to severe stage, including two severe cases and five critical cases
 (P < 0.05).

Binary Logistic Regression Analysis of COVID-19 Severe Risk Factors

Next, all categorical variables were converted into covariates, including age, 190 presence of underlying diseases (Yes or No), hypertension (Yes or No), 191 diabetes (Yes or No), obesity (Yes or No), Temperature (<37.4 °C, 37.4-192 193 38.5°C, >38.5°C), fast respiratory rate (Yes or No), elevated C-reactive protein (>10 mg/liter), decreased lymphocyte count (<1.1*10E9/L) and eosinophil count 194 (<0.02*10E9/L), elevated procalcitonin (>0.05ng/L), elevated D-dimer (>=2.25 195 ug/L), decreased albumin (<35g/L), and elevated lactate dehydrogenase 196 (LDH, >250U/L), and subjected to single-factor logistic regression together with 197 multiple independent variables. Those variables with statistical significance 198 were chosen for subsequent binary logistic regression analysis testing the 199 200 model coefficients, goodness-of-fit, and multicollinearity. Four factors identified to be significantly relevant to the severity of COVID-19 were underlying 201 diseases (X1), fast respiratory rate (>24times/min) (X2), elevated C-reactive 202 protein level (CRP >10mg/L) (X3), and elevated lactate dehydrogenase level 203 (LDH >250U/L) (X4) (Table 2). Finally, the multifactor logistic regression 204 equation was obtained to be logit P = -6.488 + 2.752X1 + 4.056X2 + 2.424X3205 206 + 5.392X4. The β values and odds ratios (OR) for each factor were shown (table

207 2). The result indicated that elevated LDH ranks the highest correlated with 208 severe symptom development (OR=219.608), followed by the fast respiratory 209 rate (OR=57.726), with underlying diseases (OR=15.67) and elevated CRP 210 (OR=11.289).

211

212 The prognostic capacity for severe symptom development

To better evaluate the prediction capacity of each independent risk factors, 213 we plotted their receiver operating characteristic curve (ROC) for the 214 development of severe COVID-19 pneumonia, and calculated their area under 215 the ROC curve (AUC value), sensitivity, specificity, Cut-off value, Youden index 216 and p-value (Table 3). According to the general standard that AUC values 217 between 0.7 and 0.9 means a medium level of diagnostic values, and over 0.9 218 219 means a high level of diagnostic values, we observed that all the factors 220 (AUC<0.9) failed to provide high prognostic value when used alone. Then, we two-factor combination test showed that the combination of fast respiratory rate 221 and elevated LDH could provide a high confident prediction (AUC=0.944, 222 sensitivity=0.941, and specificity=0.902) (Table 3). The AUC values of elevated 223 LDH plus underlying diseases or plus elevated CRP were both over 0.9, but 224 their sensitivity or specificity was lower than 0.9. Then, triple factor combination 225 significantly increased the prognostic efficacy, and all combinations had 226 increased sensitivity and specificity (Table 3). Finally, we calculated the 227

228	prognostic value of the combination of all four factors and found that the AUC
229	value was significantly increased to 0.985 (95% CI 0.968 ~ 1.000), the
230	sensitivity to 0.912, and the specificity to 0.957 (Table 3).

231

232 Discussion

Our study showed that underlying disease, fast respiratory rate (>24 233 times/min), elevated serum C-reactive protein level (CRP, >10mg/L), and 234 235 elevated lactate dehydrogenase level (LDH, >250U/L) were four independent risk factors for predicting the progression of some COVID-19 patients from mild 236 to severe conditions. Firstly, elevated lactate dehydrogenase level ranked as 237 the number 1 (OR=219.332), and fast respiratory rate ranked as the number 2 238 (OR=57.726) among the four factors (Table 2). Interestingly, elevated lactate 239 dehydrogenase level was associated with severe SARS infection [14], which 240 241 outbroke in 2003, but was absent in the severe MERS infection[15] which is still circulating. When used individually, all four factors have a moderate prediction 242 value for their low specificity and sensitivity (AUC values <0.9) (Table 3). 243 Secondly, we found that the combination of two factors, fast respiratory rate 244 plus elevated LDH, could provide a high prognostic value for severe symptom 245 (AUC=0.944, sensitivity=0.941, development and specificity=0.902). 246 Combinations of triple factors could significantly increase the prognostic value 247 (AUC>0.9). Finally, a combination of all four factors, provide an excellent 248

prognostic efficacy, achieving AUC=0.985 (95% CI 0.968 ~ 1.000) with high
sensitivity (0.953), and specificity (0.968).

Our hospital has treated over 80% of COVID-19 patients in Guangzhou city, 251 298 cases as of February 29, 2020, including 55 severe cases, but only one 252 death case. All the patients except two patients recovered as of March 15. A 253 retrospective analysis of all the cases revealed that the extremely low fatality 254 rate in our hospital (1 of 298 cases, 0.0336%, which was pretty lower than the 255 overall fatality rate (2.3%) in China [5]) was largely attributed to the effect of an 256 257 expert panel, consisting of physicians from multiple disciplines, including infectious diseases, respiratory diseases, and intensive care unit (ICU), and 258 radiology. Patients newly admitted were reviewed by the panel, and patients 259 who meet several of the following criteria were transferred immediately to the 260 ICU isolation ward for close supervision, including, (1) the illness onset has 261 entered 7-10 days; (2) over 50 years old; (3) obesity, pregnant women, children; 262 263 (4) with underlying diseases, especially hypertension, diabetes, COPD; (5) fast respiratory rate : (6) obvious decline in spirit and appetite: (7) Significant 264 265 reduction and/or progressive decline of peripheral blood lymphocytes; (8) Decreased in albumin; (9) elevated C-reactive protein (10) elevated lactate 266 dehydrogenase; and (11) quickly deteriorated, or with two or more lesions in 267 lungs revealed by chest imaging. Once they progressed to the severe stage, 268 they received treatment immediately. The above four prognostic factors, as 269 routine and affordable clinical characteristics, were included in these criteria 270 Page 14

and facilitated their immediate and preventive therapy from a retrospective aspect.

All the seven patients who were detected to be serum viral RNA positive 273 developed severe symptoms very soon, which further confirmed our previous 274 observation that detectable 2019-nCoV viral RNA in Blood is a reliable indicator 275 for the further clinical severity [16]. However, as the viral RNA positive rate were 276 low high (7cases of 32 cases, 21.8%) in this study and from other reports [17] 277 and viral RNA detection is expensive, we do not recommend to continuously 278 279 detecting viral RNA. In this regard, we suggest reserving the precious reagent for confirming virus infection. 280

In conclusion, our study indicated that underlying disease, fast respiratory 281 rate, elevated serum C-reactive protein level, and elevated lactate 282 dehydrogenase level significantly correlated to the development of severe 283 COVID-19 pneumonia, and that elevated lactate dehydrogenase and fast 284 respiratory rate or plus one or two more other factors can serve as prognostic 285 factors for the discriminating potential severe cases among the mild COVID-19 286 patients. Our study provided convenient, reliable, and affordable references for 287 both patients and physicians to make a high confident decision to commence 288 289 management and treatment safely.

290 **NOTES**

291 Contributors

Huang Huang, Feng Li, and Xilong Deng conceived the study, and wrote the manuscript. Huang Huang, Yongjiang Cai and collected data and performed data analysis. Huang Huang, Yongjiang Cai, Yueping Li, Youxia Li, Yinqiang Fan, and Xilong Deng participated in the clinical treatment. Linghua Li, Chunliang Lei, and Xiaoping Tang supervised the clinical treatment. Fengyu Hu analyzed the results. All authors read the manuscript and approved the final version.

299 Acknowledgments

300 The authors would like to thank all nurses in the treatment team for taking care

301 of the patients and doctors in the expert panel for their treatment guidance.

302 Disclosure

303 The contents of this article are solely the responsibility of the authors and do 304 not necessarily represent the views of any organization.

305 Funding

- 306 This work was supported by National Natural Science Foundation of China
- 307 (No. 81670536 and 81770593) and by the Chinese National Grand Program on
- 308 Key Infectious Disease Control (2017ZX10202203-004-002 and 309 2018ZX10301404-003-002).

310 Conflicts of interests

311 We declare that we have no conflicts of interest.

312 Conflicts of interests

- 313 This study complied with the medical ethics of Guangzhou Eighth People's
- Hospital. We obtained written consent from the patients.
- 315

316 **References**

- 317 1. Organization WH. Coronavirus disease (COVID-2019) situation reports Available at:
- 318 <u>https://www.who.int/emergencies/diseases/novel-coronavirus-2019/situation-reports/</u>.
- 2. Chen N, Zhou M, Dong X, et al. Epidemiological and clinical characteristics of 99 cases of 2019
- novel coronavirus pneumonia in Wuhan, China: a descriptive study. Lancet **2020**; 395:507-13.
- 321 3. Li Q, Guan X, Wu P, et al. Early Transmission Dynamics in Wuhan, China, of Novel Coronavirus-
- 322 Infected Pneumonia. N Engl J Med **2020**.
- 4. Aiping Wu, Peihua Niu, Lulan Wang, et al. Mutations, Recombination and Insertion in the
- 324 Evolution of 2019-nCoV. bioRxiv **2020**.
- 325 5. Team TNCPERE. Vital Surveillances: The Epidemiological Characteristics of an Outbreak of 2019
- Novel Coronavirus Diseases (COVID-19) China, 2020. China CDC Weekly **2020**; 2:113-22.
- 327 6. Huang C, Wang Y, Li X, et al. Clinical features of patients infected with 2019 novel coronavirus
- 328 in Wuhan, China. Lancet **2020**; 395:497-506.
- 329 7. Wang D, Hu B, Hu C, et al. Clinical Characteristics of 138 Hospitalized Patients With 2019 Novel
- 330 Coronavirus-Infected Pneumonia in Wuhan, China. JAMA **2020**.
- 8. Guan WJ, Ni ZY, Hu Y, et al. Clinical Characteristics of Coronavirus Disease 2019 in China. N Engl
- 332 J Med **2020**.

- 9. Xu X, Yu C, Qu J, et al. Imaging and clinical features of patients with 2019 novel coronavirus
- 334 SARS-CoV-2. Eur J Nucl Med Mol Imaging 2020.
- 10. Pan Y, Guan H, Zhou S, et al. Initial CT findings and temporal changes in patients with the novel
- 336 coronavirus pneumonia (2019-nCoV): a study of 63 patients in Wuhan, China. Eur Radiol **2020**.
- 337 11. Zhou, Y, Binqing Fu, Xiaohu Zheng, et al. Aberrant pathogenic GM-CSF+ T cells and
- 338 inflammatory CD14+CD16+ monocytes in severe pulmonary syndrome patients of a new
- 339 coronavirus. bioRxiv **2020**.
- 12. Xu Z, Shi L, Wang Y, et al. Pathological findings of COVID-19 associated with acute respiratory
- distress syndrome. Lancet Respir Med **2020**.
- 342 13. nhc.gov.cn. Notice on Issuing a New Coronary Virus Pneumonia Diagnosis and Treatment Plan
- 343 (Trial Version 6, in Chinese). Available at:
- 344 http://www.nhc.gov.cn/yzygj/s7653p/202002/8334a8326dd94d329df351d7da8aefc2.shtml.
- 345 14. Liu CL, Lu YT, Peng MJ, et al. Clinical and laboratory features of severe acute respiratory
- 346 syndrome vis-a-vis onset of fever. Chest **2004**; 126:509-17.
- 347 15. Ko JH, Park GE, Lee JY, et al. Predictive factors for pneumonia development and progression
- to respiratory failure in MERS-CoV infected patients. J Infect **2016**; 73:468-75.
- 16. Chen W, Lan Y, Yuan X, et al. Detectable 2019-nCoV viral RNA in blood is a strong indicator
- 350 for the further clinical severity. Emerg Microbes Infect **2020**; 9:469-73.
- 17. Wang W, Xu Y, Gao R, et al. Detection of SARS-CoV-2 in Different Types of Clinical Specimens.
- 352 JAMA **2020**.

354 Tables

Gender Gal 47 16 Male 63 47 16 Female 62 46 16 Age (years)§ 44.87±18.55 40.49±17.66 59.43±13.47 Underlying disease 37 17 20 Hypertension 20 7 13	0.96 <0.05 <0.05
Male 63 47 16 Female 62 46 16 Age (years)§ 44.87±18.55 40.49±17.66 59.43±13.47 Underlying disease 37 17 20 Hypertension 20 7 13	<0.05
Female 62 46 16 Age (years)§ 44.87±18.55 40.49±17.66 59.43±13.47 Underlying disease 37 17 20 Hypertension 20 7 13	<0.05
Age (years)§ 44.87±18.55 40.49±17.66 59.43±13.47 Underlying disease 37 17 20 Hypertension 20 7 13	<0.05
Underlying disease371720Hypertension20713	<0.05
Hypertension 20 7 13	10.00
	<0.05*
Diabetes 8 2 6	<0.05*
Obesity (BMI>26) 7 2 5	<0.05*
Travel to epidemic area886523	0.388
Temperature	<0.05
<37.4°C 57 48 9	
37.4°C-38.5°C 48 34 14	
>38.5°C 20 11 9	
Coughing 76 56 20	0.819
Running nose 21 14 7	0.533
Muscle joint pain27234	0.147
Headache 24 16 8	0.334
Fatigue 48 32 16	0.061
Digestive Symptoms 19 15 4	

355 Table 1. Characteristics of COVID-19 patients

Page 19

Fast respiratory rate	20	4	16	<0.05*	
Serum viral RNA positive	7	0	7	<0.05*	
White cells (10E9/L)	5.57±1.76	5.65±1.73	5.33±1.86	0.411	
Neutrophils $(10E9/L)^{\$}$	3.43±1.43	3.26±1.28	34.97±1.77	0.053	
	1.32(1.05-	1.43(1.23-	0.82(0.57-	-0.05**	
Leukocytes (TOE9/L)	2.18)	2.21)	1.05)	<0.05***	
Eosinophils (10E9/L) ^{&}	0.02 (0-0.09)	0.04 (0.1-0.12)	0(0-0)	<0.05**	
Platelets (10E9/L) [§]	200.56±56.24	206.01±55.61	182.46±55.47	0.052	
Hemoglobin (g/L) $^{\$}$	134.39±18.02	135.31±17.92	131.32±18.32	0.306	
Prothrombin time (sec)§	13.69±1.13	13.66±0.89	13.80±1.70	0.668	
Activated partial	39 30+4 74	38 93+4 49	40 49+5 37	0.13	
prothrombin time (sec) $^{\$}$	<i>39.3</i> 0 ⊥न ./ न	50.95± 1 . 1 9	+0.+9±3.37	0.15	
C-reactive protein (CRP)	6.32(1.63-	4.00(1.06-	46.345 (28.97-	<0.05**	
(mg/L) ^{&}	23.50)	12.41)	60.50)	<0.05	
D-dimer (ug/L) ^{&}	910(700-	780(560-	1760(1297.5-	<0.05**	
	1400)	1050)	3265)	~0.0 <i>3</i> · ·	
Procalcitonin (ng/ml) ^{&}	0.047(0.03-	0.037(0.027-	0.070(0.051-	<0.05**	
r rocaicitonin (ng/m)	0.076)	0.063)	0.145)	<0.05	
Lactic acid (mmol/L) §	1.78±0.71	1.72±0.76	1.93±0.55	0.207	
Alanine aminotransferase	18.90(13.40-	16.70(12.40-	27.5(19.70-	<0.05**	
(ALT, U/L) ^{&}	25.20)	22.15)	41.25)	\0.03	

Aspartate					
aminatronaforada (ACT	18.40(14.20-	17.20(13.75-	31.50(23.25-	<0.05**	
aminotransierase(AST,	27.15)	21.00)	37.75)	<0.03	
U/L) ^{&}					
Albumin (a/l)	28 20+5 20	20 82+1 40	22 40+4 70	<0.05	
Albumin (g/L)*	38.30±3.30	39.03±4.49	55.49 1 4.79	<0.03	
Creatinine (umol/L)§	67.15±28.21	64.02±26.95	77.56±42.19	0.271	
Creatine kinase (CK					
Cleatine Killase (CK,	82.89±48.39	77.93±46.05	100.27±59.73	0.08	
U/L)***					
Lactate dehydrogenase	175(150-		377(779 75-		
	175(150-	161 (145-192)	522(27):15-	<0.05**	
(LDH, U/L) ^{&}	241.5)		400)		

- 356 ***Fisher's Exact Test**
- 357 **Mann-Whitney U Test
- 358 § average±standard deviration(STD)
- 359 & average (95% confidence interval)

361 Table 2. Independent factors associated with severe symptom development in

362 COVID-19 patients

Va	riables	β	S.E.	chi-square	P value	OR (95% CI)
V	Underlying	2.752	1.066	6.666	0.01	15.67 (1.94-
▲1	diseases					126.55)
	Fast					
V	respiratory	4.056	1.183	11.76	0.001	57.726 (5.685-
X 2	rate (>24					586.191)
	times/min)					
V	CPR (>10	2.424	1.004	5.823	0.016	11.289 (1.577-
X 3	mg/liter)					80.838)
						219.608
X_4	LDH (>250	5.392	1.24	18.911	< 0.001	(19.332-
	U/liter)					2494.742)
Intercept		-6.488	1.499	18.738	0.001	0.002

363 S.E., standard error;

364 OR (95% CI), Odd Ratio (95% confidence interval).

365

366

		Factor	AUC (95% CI)	Sensitiv ity	Specif icity	Cut- off value	Youde n Index	P value
_		Underlying diseases (1)	0.722 (0.614~0.829)	0.618	0.826	0.367	0.444	< 0.001
	Single	Fast respiratory rate (2)	0.758 (0.648~0.867)	0.559	0.957	0.492	0.516	<0.001
	factor	Elevated CRP (3)	0.774 (0.685~0.864)	0.853	0.696	0.298	0.549	< 0.001
		Elevated LDH (4)	0.855 (0.766~0.944)	0.765	0.946	0.461	0.711	< 0.001
-		(1) + (2)	0.853 (0.767~0.939)	0.824	0.793	0.223	0.617	<0.001
		(1) + (3)	0.854 (0.779~0.928)	0.853	0.696	0.274	0.549	< 0.001
	Two	(1) + (4)	0.940 (0.894~0.987)	0.971	0.783	0.156	0.754	< 0.001
	factors	(2) + (3)	0.870 (0.795~0.944)	0.912	0.663	0.18	0.575	< 0.001
		(2) + (4)	0.944 (0.892~0.996)	0.941	0.902	0.315	0.843	<0.001
_		(3) + (4)	0.918 (0.856~0.981)	0.765	0.946	0.365	0.711	< 0.001
		(1) + (2) + (3)	0.910 (0.850~0.969)	0.765	0.902	0.253	0.667	< 0.001
	Three	(1) + (2) + (4)	0.976 (0.955~0.998)	0.912	0.935	0.411	0.846	<0.001
	factors	(1) + (3) + (4)	0.963 (0.933~0.993)	0.912	0.891	0.227	0.803	< 0.001
		(2) + (3) + (4)	0.964 (0.919~1.000)	0.912	0.934	0.355	0.847	<0.001
-	Four	(1) + (2) + (3) + (4)	0.985	0.912	0.957	0.374	0.869	<0.001
	Tactors	(3) + (4)	10.200~1.000)					

368 Table 3. Prognostic values for severe COVID-19 pneumonia development.

369

AUC (95% CI): Area under receiver operating characteristic curve (95% confidence interval).

370

372 Figure legends

373	Figure 1. Enrollment chart of COVID-19 patients. The clinical information of a
374	total of 298 patients admitted to the hospital was reviewed. Patients were
375	excluded according to the criteria 1) already in severe stages, 2) with a disease
376	course <7 days, and 3) other reasons such as incomplete detection panel.
377	Finally, 125 patients were further divided into two groups. The severe group
378	included the patients who developed severe COVID-19 pneumonia later (>3
379	days post-admission). The patients remaining were kept in the mild group.
380	
JOT	

382 Figures

383

Figure1 384

