

Development a quantitative segmentation model to assess the effect of comorbidity on patients with COVID-19

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Research

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Abstract

Background: The coronavirus disease 2019 (COVID-19) has brought a global disaster. Quantitative lesions may provide the radiological evidence of the severity of pneumonia and further to assess the effect of comorbidity on patients with COVID-19.

Methods: 294 patients with COVID-19 were enrolled from February, 24, 2020 to June, 1, 2020 from six centers. Multi-task Unet network was used to segment the whole lung and lesions from chest CT images. This deep learning method was pre-trained in 650 CT images (550 in primary dataset and 100 in test dataset) with COVID-19 or community acquired pneumonia and Dice coefficients in test dataset were calculated. 50 CT scans of 50 patients (15 with comorbidity and 35 without comorbidity) were random selected to mark lesions manually. The results will be compared with the automatic segmentation model. Eight quantitative parameters were calculated based on the segmentation results to evaluate the effect of comorbidity on patients with COVID-19.

Results: Quantitative segmentation model was proved to be effective and accurate with all Dice coefficients more than 0.85 and all accuracies more than 0.95. Of the 294 patients, 52 (17.7%) patients were reported having at least one comorbidity, 14 (4.8%) having more than one comorbidity. Patients with any comorbidity were older ($P<0.001$), had longer incubation period ($P<0.001$), were more likely to have abnormal laboratory findings ($P<0.05$) and be in severity status ($P<0.001$). More lesions (including larger volume of lesion, consolidation and ground-glass opacity) were shown in patients with any comorbidity than patients without comorbidity (all $P<0.001$). The more comorbidities patients have, the poorer CT manifestation is. The median volume of lesion, consolidation and ground-glass opacity in diabetes mellitus group was largest among the three prevalently single comorbidity groups.

Conclusions: Multi-task Unet network can make quantitative CT analysis of lesions to assess the effect of comorbidity on patients with COVID-19, further to provide the radiological evidence of the severity of pneumonia. More lesions were found in CT images of cases with comorbidity. The more comorbidities patients have, the poorer CT manifestation is.

Background

The coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) broke out in Wuhan, China in December 2019. It has widely spread all over the world, which has led to a major concern [1-3]. Up to July 28, 2020, 16,341,920 and 650,805 confirmed and death cases have been reported worldwide. The United States, as the largest epicenter, had the confirmed and death cases of 4,209,509 and 146,331, respectively [4].

A sharp increase in the number of cases due to human-to-human transmission has resulted in high rates of hospitalization and intensive care unit (ICU) admission which cause an extreme shortage of medical resources [5]. Therefore, it is especially important to use limited medical resources and social support for people with serious condition and prone to serious outcomes.

Previous reports found that comorbidities (including hypertension, diabetes mellitus, respiratory system disease and cardiovascular) may be a risk factor COVID-19 progression and also correlated with poorer clinical outcomes [6-10]. The relationship between COVID-19 and comorbidity was mostly explored from a clinical perspective so far few studies have studied it use radiology.

Chest CT has played a pivotal diagnostic role in the assessment of the disease severity according to the number, extent, density of patchy ground-glass opacities (GGOs) and consolidation [11]. Differential diagnosis, severity rating, and prognosis prediction about COVID-19 have been investigated using a quantitative CT combined with artificial intelligence (AI) technology [12-14]. However, quantitative CT study about the effect of comorbidity on patients with COVID-19 has not been reported, which may provide the radiological evidence of the severity of pneumonia.

In our study, the multi-task Unet, a deep learning method, was used to develop a segmentation model to quantify the pneumonia lesion including the volume of GGOs and consolidation to assess the effect of comorbidity on patients with COVID-19.

Materials And Methods

Patients

This was a retrospective study that collected data from six centers (TongDe Hospital of ZheJiang Province, The Second People's Hospital of Neijiang, The First Affiliated Hospital of Bengbu Medical College, Wenzhou People's Hospital, Anqing Municipal Hospital and The First Affiliated Hospital of Xi'an Jiaotong University). This study was approved by the Ethics Committee of the above hospitals, and written informed consents were waived because the anonymized study did not alter any diagnosis and treatment of the patients. 294 patients were enrolled from 24, February, 2020 to 1, June, 2020. We included patients who satisfied the following criteria: (a) positive for nextgeneration sequencing or realtime RTPCR of SARS-CoV-2 in throat or nose swabs; (b) complete clinical data; (c) patients underwent CT scans. The exclusion criteria were (a) poor images with heavy breathing artifacts or metal artifacts; (b) patients had history of pulmonary surgery. In this case, the patient has series of CT examinations, the most severe sets of images were included in our study.

Clinical information

The basic data including gender, age, incubation period, comorbidity (hypertension, diabetes mellitus, cardiovascular disease, cerebrovascular disease, COPD, hepatitis B infection, malignancy, chronic kidney disease and immunodeficiency, etc.), severity status (severe or non-severe), laboratory examinations including C reactive protein (CRP), white blood cell count (WBC) and lymphocyte percentage were extracted from medical computerized database for all patients. The severity of COVID-19 includes four types: mild, common, severe, and critical according to the guideline of 2019-nCoV (trial version⁷) issued by the China National Health Commission [15]. In this study, we divided all the patients into severe group (including severe and critical) with 38 cases and non-severe group (mild and common) with 256 cases. Of all the patients, there were 52 patients with comorbidity, the remaining 242 patients without comorbidity. Among these patients with comorbidity, 34 (65.4%) patients were with hypertension, 15 (28.9%) with diabetes mellitus, 8 (15.4%) with hepatitis B infection, 4 (7.7%) with cardiovascular disease, 3 (5.8%) with COPD and 2 (3.8%) with cerebrovascular disease. Patient with malignancy, chronic kidney disease and immunodeficiency was 1(1.9%).

CT image acquisition

The non-contrast chest CT were performed using three multi-detector CT scanners with 64 or 128 channels (Somatom Definition AS+, Siemens Healthineers, Forchheim, Germany; GE Medical Systems, China Branch, Beijing, China or Philips Ingenuity Core128, Philips Medical Systems, Best, the Netherlands). The scanning range was from apex to the base of lungs. The detailed parameters for CT acquisition were as follows: tube voltage, 120 kVp; tube current, standard (reference mAs, 60–120) to low-dose (reference mAs, 30) with automatic exposure control; slice thickness, 1.0 or 1.25 mm; reconstruction interval, 1.0-3.0 mm; noise index (NI), 25 and matrix 512×512. A lung window was with a width of 2000 HU and a level of -600HU, and a mediastinal window with a width of 350 HU and a level of 40HU.

CT image segmentation and quantitative analysis

The deep learning method used in our study is Unet neural network [16] which has been reported to have a good performance on the segmentation of the biomedical images. Here, we used a multi-task Unet with a single encoder and two parallel decoders to learn to predict and segment the region of lung and lesions. The decoder containing attention block was used to learn to segment the lesions while the decoder containing stacked dilated convolutions was used to learn the lung segmentation, which provided a more efficient feature encoding and a regularizing effect. This neural network were implemented in Dr. pecker cloud platform (<http://www.jianpeicn.com/category/yuepianjqiren>) and our segmentation results were acquired from the platform. Our platform is open and free to all public research institutions in the world.

In order to make the neural network to learn to predict lesion and lung regions, labelled lesion samples lung samples were required. The lesion and lung regions were manually segmented using ITK-SNAP software (version 2.2.0; <http://www.itksnap.org>) in lung window with a width of 2000 HU and a level of -600 HU. Our segmentation system was pre-

trained by 650 annotated CT images (550 in primary dataset and 100 in test dataset) with COVID-19 or community acquired pneumonia. This neural network extracted CT image features, segmented lung and lesions, and classified whether the lesion was consolidation or GGO. The volumes of the lesions as the results in underlying disease group and non-underlying disease group were outputted finally. The general flow of this study was shown in Figure 1.

The specific image segmentation steps were as follows. First, a threshold value of -450 HU was used to distinguish GGO and consolidation (Figure 2&3. B). The margin of the lesion in each axial slice was delineated (Figure 2&3. C-D). Then, a 3D region of interest (ROI) was obtained based on delineated results including lung erosion diagram (Figure 2&3. E-F) and lesion diagram (Figure 2&3. G-H). The SimpleITK software tool (<http://www.simpleitk.org>) was used to quantify the mean HU of lung and lesions, volumes and numbers of lesions automatically.

Dice coefficient (between 0 and 1) in test dataset was used as an index to evaluate the quantitative segmentation effect of this model. Higher Dice coefficient presents better model.

To assess the accuracy of segmentation on our cohort, two cardiothoracic radiologists (C.Z. and J.W. with 5 and 12 years of experience, respectively) manually marked the infected area of 50 CT scans of 50 patients (15 with comorbidity and 35 without comorbidity) and compared the results with those of segmentation model.

In total, the following 8 quantitative parameters were acquired for further analysis: (a) the volume of the whole lung (LUV); (b) the volume of lesion (LEV); (c) the ratio of volume of lesion to whole lung (LEV/LUV); (d) the volume of consolidation (COV); (e) the ratio of volume of consolidation to whole lung (COV/LUV); (f) the volume of GGO (GGOV); (g) the ratio of volume of GGO to whole lung (GGOV/LUV) and (h) the number of lesions (LEN).

Statistical Analysis

Continuous variables were presented as median (IQR) and categorical variables were shown as n (%). Mann-Whitney U test, χ^2 test or Fisher's exact test were used to compare differences between patients with and without comorbidity. Statistical analyses were performed with SPSS (ver. 22.0; SPSS Inc., Chicago, IL, USA). Two-sided $P < 0.05$ was considered statistically significant.

Results

Segmentation effect of this model

The average Dice coefficient in test dataset was 0.973 for the right lung, 0.985 for left lung, and 0.864 for lesion segments (all > 0.85), suggesting the good performance of our neural network in lung and lesion segmentation task. The accuracy of segmentation was 0.987 for all lesions, 0.968 for consolidation and 0.953 for GGO (all > 0.95).

Characteristics of patients with or without any comorbidity

Of the 294 cases, 52 (17.7%) patients were reported having at least one comorbidity. Patients with any comorbidity were older (median: 55.50 vs. 44.00 years) ($P < 0.001$), had longer incubation period (median: 6.00 vs. 3.00 days) ($P < 0.001$). They were more likely to have abnormal CPR (75.0% vs. 53.7%) ($P < 0.01$), WBC (32.7% vs. 18.6%) ($P < 0.05$), and lymphocyte percentage (65.4% vs. 41.7%) ($P < 0.01$) and be in severity status (42.3% vs. 6.6%) ($P < 0.01$) (Table 1). As for the quantitative CT images analysis, larger LEV (median: 235.01 vs. 127.82 cm^3) ($P < 0.001$), LEV/LUV (median: 0.07 vs. 0.04) ($P < 0.001$), COV (median: 94.58 vs. 40.45 cm^3) ($P < 0.001$), COV/LUV (median: 0.03 vs. 0.01) ($P < 0.01$), GGOV (median: 150.26 vs. 74.71 cm^3) ($P < 0.001$), GGOV/LUV (median: 0.04 vs. 0.02) ($P < 0.001$) and LEN (median: 30.00 vs. 19.00) ($P < 0.01$) were shown in patients with at least one comorbidity than patients without comorbidity. The difference in gender distribution was not significant between two groups (Table 1).

Characteristics of patients with 1 comorbidity or ≥ 2 comorbidities

Of the 52 patients with any comorbidity, 38 (73.1%) patients have one comorbidity, the remaining 14 (26.9%) had more than one comorbidity. Older age (median: 53.00 vs. 44.00 and 60.00 vs. 44.00 years), longer incubation period (median: 6.00 vs. 3.00 and 8.00 vs. 3.00 days), more likely having abnormal lymphocyte percentage (63.2% vs. 41.7% and 71.4 vs. 41.7%), more likely in severe state (36.8% vs. 6.6% and 57.1 vs. 6.6%) were exhibited both in cases with 1 comorbidity or ≥ 2 comorbidities compared to cases without any comorbidity. In terms of quantitative analysis of CT imaging, both patients with 1 comorbidity and more than one comorbidity were detected more lesions (LEV: 235.87 vs. 127.82 cm³ and 237.17 vs. 127.82 cm³; LEV/LUV: 0.07 vs. 0.04 and 0.09 vs. 0.04; COV: 97.86 vs. 40.45 cm³ and 94.89 vs. 40.45 cm³; COV/LUV: 0.03 vs. 0.01 and 0.04 vs. 0.01; GGOV: 135.80 vs. 74.71 cm³ and 197.58 vs. 74.71 cm³; GGOV/LUV: 0.03 vs. 0.02 and 0.05 vs. 0.02). More tendency to have abnormal CPR and more numbers of lesions were found in patients with one comorbidity, while, there was no difference between patients with ≥ 2 comorbidities and without comorbidity. Neither patients with one comorbidity nor with ≥ 2 comorbidities have significant difference in gender distribution and abnormal WBC (Table 2).

Characteristics of patients with different comorbidities

In order to assess the impact of a certain underlying disease with a higher incidence on COVID-19, we separately compared the clinical and imaging indicators of patients. Of the 52 patients with comorbidities, 22 (16.9%), 7 (3.7%) and 8 (1.9%) patients were reported only having hypertension, diabetes and hepatitis B infections, respectively. These three series all have more possibility in severity status compared with series without comorbidity (all $P < 0.001$). It is worth noting that almost all quantitative indicators except GGOV and LEN indicated that patients with hypertension and diabetes mellitus had more lesions than patients without any comorbidity. However, there were no differences between patients with hepatitis B infections and without comorbidity in almost all clinical and quantitative CT characteristics except appearance ratio of abnormal lymphocyte percentage and severity status (Table 3).

Discussion

In this study, we tried a multi-task Unet network, a segmentation tool to build a quantitative segmentation model to assess the impact of comorbidity on patients with COVID-19. Comorbidities may be a risk factor for severe status and pneumonia progression. What's more, they are also associated to poor clinical outcomes which consists of admission to ICU, or invasive ventilation, or death [6-10]. However, most previous studies were focused on the relationship between COVID-19 and comorbidity from a clinical perspective. As a radiologist with a very important position during the outbreak, we firstly assessed the impact of comorbidity with intuitive and quantitative data. It is of great necessity to provide the radiological evidence of the severity of pneumonia before treatment, which may greatly determine the clinical management and prognosis.

Our neural network model was trained in primary dataset, and tested on test dataset in order to confirm the robustness of it. The average Dice coefficients in test dataset were all more than 0.85 for lung and lesion segments, which suggested good performance in lung and lesion segmentation task. The lesion regions of all 294 cases are segmented by our multi-task Unet network firstly, then a part of segmentation results was checked manually. More than 95% lesion regions were segmented accurately, which indicates stability and accuracy of our Unet model segmentation.

From the quantitative CT images analysis, larger lesion volumes (including consolidation and GGOs) were found in patients with any comorbidity than without comorbidity. It's worth taking note of that the more comorbidity patients have, the poorer CT manifestation is. The more the lung parenchymal is involved, the more severe condition it would be. The appearance of GGO indicates that alveolar cavity is partially filled by fluid and cells, while the appearance of consolidation demonstrates that further accumulation of exudates in alveolar cavity and aggravation of interstitial edema, which always means disease progresses [17]. The increasing numbers of GGOs and densities of consolidation always indicates disease deterioration [18].

The pathogenesis of COVID-19 may be associated to underlying diseases due to their susceptibility conditions. Researchers reported that similar results were found in MERS [19]. Comorbidities are characterized by proinflammatory state, and the attenuation of the immune response [20-21]. For instance, a possible cause of diabetes mellitus is the accumulation of activated innate immune cells in metabolic tissues leading to the release of inflammatory mediators, thereby promoting insulin

resistance and β -cell damage [22]. What's more, metabolic disorders may lead to decreased immune function because of impaired macrophage and lymphocyte function [23]. Impaired immune function eventually may make patients more susceptible to other diseases [19]. What's more, High expression of angiotensin converting enzyme 2 (ACE2) in diabetes and hypertension may potentially facilitate viral uptake [24-25]. Our study found that patients with diabetes mellitus has the worst CT findings, however, the specific mechanism still remains in-depth study. In addition, patients with more than one comorbidity have greater possibility to have impaired immune function—which may be linked to more severe lung CT images.

Unexpectedly, we found that neither consolidation nor GGO volumes were not significantly different when comparing patients with hepatitis B infections and without comorbidity—although they were more likely in severity status. This may suggest that the severity of COVID-19 is not completely consistent with the CT findings. Meta-analyses from Wang et al [9] and Lippi G et al [26] did not provide sufficient evidence that liver disease was relevant to COVID-19 progress. However, the credibility of our results may be limited by the small number of cases with liver disease.

To our surprise, the results revealed that incubation period of individuals with comorbidity was longer than that of patients without comorbidity, which may have relationship with the insensitivity reaction to COVID-19 in comorbidity groups. The rapid progresses of pneumonia and poorer CT manifestation may be partly on account of delayed timely treatment.

This study had several limitations. First of all, small sample sizes of patients with a certain underlying disease alone may affect the credibility of some results and conclusions. Secondly, the most severe CT images included in our study cannot show the change of the COVID-19 in patients with or without comorbidities, which can prompt us to investigate and study in the next-step. Thirdly, the novelty of the algorithm of the quantitative assessment model may not be very outstanding compared to previously reported ones.

Conclusions

In conclusion, Multi-task Unet network can make quantitative CT analysis of lesions to assess the effect of comorbidity on patients with COVID-19. More lesions were found in CT images of cases with comorbidity. The more comorbidities patients have, the poorer CT manifestation is.

Declarations

Acknowledgements

Not applicable.

Authors' contributions

JW, GZY and CZ participated in the design of this study, and they all performed the statistical analysis. ZHX, YMG and ZYX carried out the study and collected important background information. CXC, HW and HFS collected raw data. GHC performed data processing on deep learning software. CZ drafted the manuscript. All authors read and approved the final manuscript.

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Availability of data and materials

All data generated or analyzed during this study are included in this published article.

Ethics approval and consent to participate

This study was approved by the ethical committee of Tongde Hospital of ZheJiang Province, The Second People's Hospital of Neijiang, The First Affiliated Hospital of Bengbu Medical College, Wenzhou People's Hospital, Anqing Municipal Hospital and The

First Affiliated Hospital of Xi'an Jiaotong University.

Consent for publication

Not applicable.

Competing interests

The authors declare no conflict of interest.

References

1. Velavan TP, Meyer CG. The COVID-19 epidemic. *Trop Med Int Health*. 2020; 25(3):278-280.
2. Holshue ML, DeBolt C, Lindquist S, Lofy KH, John Wiesman J, Bruce H, Spitters C, Ericson K, Wilkerson S, Tural A, et al. First case of 2019 novel coronavirus in the United States. *N Engl J Med*. 2020; 382(10):929-36.
3. Grasselli G, Zangrillo G, Zanella A, Zanella A, Antonelli M, Cabrini L, Castelli A, Cereda D, Coluccello A, Foti G, Fumagalli R, et al. Baseline Characteristics and Outcomes of 1591 Patients Infected With SARS-CoV-2 Admitted to ICUs of the Lombardy Region, Italy. 2020; *JAMA* 323:1574-81.
4. Coronavirus disease (COVID-19) Dashboard. <http://covid19.who.int/>
5. Grasselli G, Pesenti A, Cecconi M. Critical care utilization for the COVID-19 outbreak in Lombardy, Italy: early experience and forecast during an emergency response. *JAMA* 2020 Mar 13. doi: 10.1001/jama.2020.4031. [Epub ahead of print]
6. Yang J, Zheng Y, Gou X, Pu K, Chen ZF, Guo QH, Ji R, Wang HJ, Wang YP, Zhou YN. et al. Prevalence of Comorbidities and Its Effects in Coronavirus Disease 2019 Patients: A Systematic Review and Meta-Analysis. *Int J Infect Dis*. 2020; 94:91-5.
7. Emami A, Javanmardi F, Pirbonyeh N, Akbari A. Prevalence of underlying diseases in hospitalized patients with COVID-19: a systematic review and meta-analysis. *Arch Acad Emerg Med*. 2020; 8: e35.
8. Wang BL, Li RB, Lu Z, Hung Y. Does comorbidity increase the risk of patients with COVID-19: evidence from meta-analysis. *Aging (Albany NY)*. 2020; 12:6049-57.
9. Wu J, Li W, Shi X, Chen Z, Jiang B, Liu J, Wang D, Liu C, Meng Y, Cui L, et al. Early antiviral treatment contributes to alleviate the severity and improve the prognosis of patients with novel coronavirus disease (COVID-19). *J Intern Med*. 2020; 288(1):128-38.
10. Guan WJ, Liang WH, Zhao Y, Liang HR, Chen ZS, Li YM, Liu XQ, Chen RC, Tang CL, Wang T, et al. Comorbidity and its impact on 1590 patients with Covid-19 in China: a nationwide analysis. *Eur Respir J*. 2020; 55(5):2000547.
11. Kanne JP. Chest CT findings in 2019 novel coronavirus (2019-nCoV) infections from Wuhan, China: key points for the radiologist. *Radiology*. 2020; 295(1):16-7.
12. Shen C, Yu N, Cai S, Zhou J, Sheng J, Liu K, Zhou H, Guo Y, Niu G. Quantitative computed tomography analysis for stratifying the severity of Coronavirus Disease 2019. *J Pharm Anal*. 2020; 10(2):123-9.
13. Ni QQ, Sun ZY, Qi L, Chen W, Yang Y, Wang L, Zhang XY, Yang L, Fang Y, Xing ZJ, et al. A deep learning approach to characterize 2019 coronavirus disease (COVID-19) pneumonia in chest CT images. *Eur Radiol*. 2020; 2:1-11.
14. Wang Y, Chen Y, Wei Y, Li M, Zhang YZ, Zhang N, Zhao S, Zeng HJ, Deng W, Huang ZX, et al. Quantitative analysis of chest CT imaging findings with the risk of ARDS in COVID-19 patients: a preliminary study. *Ann Transl Med*. 2020; 8(9):594.
15. Diagnosis and treatment protocol for novel coronavirus pneumonia (trial version 7). National Health Commission of People's Republic of China Web site. <http://en.nhc.gov.cn/2020-03/29/c78469.htm>.
16. Ronneberger O, Fischer P, Brox T. U-Net: Convolutional networks for biomedical image segmentation. Paper presented at: International conference on medical image computing and computer-assisted intervention 2015.
17. Jin YH, Cai L, Cheng ZS. A rapid advice guideline for the diagnosis and treatment of 2019 novel coronavirus (2019-nCoV) infected pneumonia (standard version). *Mil Med Res Military*. 2020; 7(1):4.

18. Li Yan, Xia LM. Coronavirus Disease 2019 (COVID-19): Role of Chest CT in Diagnosis and Management. *AJR Am J Roentgenol.* 2020; 214(6):1280-6.
19. Badawi A, Ryoo SG. Prevalence of comorbidities in the Middle East respiratory syndrome coronavirus (MERS-CoV): a systematic review and meta-analysis. *Int J Infect Dis.* 2016; 49:129-33.
20. Zheng YY, Ma YT, Zhang JY, Xie X. COVID-19 and the cardiovascular system. *Nat Rev Cardiol.* 2020; 17(5):259-60.
21. Chen NS, Zhou M, Dong X, Qu JM, Gong FY, Han Y, Qiu Y, Wang JL, Liu Y, Wei Y, et al. Epidemiological and clinical characteristics of 99 cases of 2019 novel coronavirus pneumonia in Wuhan, China: a descriptive study. *Lancet.* 2020; 395(10223):507-13.
22. Odegaard JI, Chawla A. Connecting type 1 and type 2 diabetes through innate immunity. *Cold Spring Harb Perspect Med.* 2012; 2(3): a007724.
23. Dooley KE, Chaisson RE. Tuberculosis and diabetes mellitus: convergence of two epidemics. *Lancet Infect Dis.* 2009; 9(12):737-46.
24. Von der Thüsen J, Van der Eerden M. Histopathology and genetic susceptibility in COVID-19 pneumonia. *Eur J Clin Invest.* 2020 Apr 30; e13259. doi: 10.1111/eci.13259. [Epub ahead of print].
25. Zhang J, Miao Yu M, Song Tong S, Liu LY, TangLV. Predictive factors for disease progression in hospitalized patients with coronavirus disease 2019 in Wuhan, China. *J Clin Virol.* 2020 Jun; 127:104392. doi: 10.1016/j.jcv. [Epub ahead of print].
26. Lippi G, Maria Santos de Oliveira H, Michael Henry B. Chronic liver disease is not associated with severity or mortality in Coronavirus disease 2019 (COVID-19): a pooled analysis. *Eur J Gastroenterol Hepatol.* 2020 Apr 10. doi: 10.1097/MEG.0000000000001742. [Epub ahead of print].

Tables

Table 1 Characteristics of patients with or without any comorbidity

Variables	Total (n=294)	Without comorbidity (n = 242)	With any comorbidity (n=52)	P value
Gender				0.149
Male	134(45.6%)	127(52.5%)	19(36.5%)	
Female	160(54.4%)	115(47.5%)	33(63.5%)	
Age(years)	46.00(34.75-54.00)	44.00(32.00-53.00)	55.50(47.00-65.25)	0.000
Incubation period (days)	5.00(3.00-7.00)	3.00(2.00-7.00)	6.00(4.00-8.00)	0.000
CRP abnormal				0.005
Yes	169(57.5%)	130(53.7%)	39(75.0%)	
No	125(42.5%)	112(46.3%)	54(38.85%)	
WBC abnormal				0.024
Yes	62(21.1%)	45(18.6%)	17(32.7%)	
No	232(78.9%)	197(81.4%)	35(67.3%)	
LY % abnormal				0.002
Yes	135(45.9%)	101(41.7%)	34(65.38%)	
No	159(54.1%)	141(58.3%)	18(34.62%)	
severity status				0.000
Yes	38(12.9%)	16(6.6%)	22(42.3%)	
No	256(87.1%)	226(93.4%)	30(57.7%)	
Quantitative CT analysis				
LUV(cm ³)	3728.28(2944.22-4642.55)	3709.28(2939.74-4538.04)	4180.53(3027.12-4868.62)	0.218
LEV(cm ³)	139.25(58.28-297.34)	127.82(50.50-265.91)	235.01(89.65-658.18)	0.000
LEV/LUV	0.04(0.02-0.09)	0.04(0.01-0.08)	0.07(0.02-0.16)	0.000
COV(cm ³)	45.00(12.97-119.82)	40.45(11.82-96.27)	94.58(23.93-208.55)	0.000
COV/LUV	0.01(0.00-0.04)	0.01(0.00-0.03)	0.03(0.02-0.06)	0.001
GGOV (cm ³)	80.02(36.95-169.48)	74.71(31.08-150.72)	150.26(50.36-305.02)	0.000
GGOV /LUV	0.02(0.00-0.05)	0.02(0.01-0.04)	0.04(0.02-0.09)	0.000
LEN	22.00(11.00-39.00)	19.00(10.00-37.25)	30.00(21.25-44.75)	0.001

Note—CRP: C reactive protein; WBC: white blood cell count; LY %: lymphocyte percentage; LUV: the volume of the whole lung; LEV: the volume of lesion; COV: the volume of consolidation; GGOV: the volume of GGO; LEN: the number of lesions. P values written in bold indicate a significant difference.

Table 2 Characteristics of patients with 1 comorbidity or ≥ 2 comorbidities

Variables	Without comorbidity (n = 242)	1 comorbidity (n=38)	≥ 2 comorbidities (n=14)	P1 value	P2 value
Gender				0.126	0.734
Male	127(52.5%)	13(34.2%)	6(42.9%)		
Female	115(47.5%)	25(65.8%)	8(57.1%)		
Age(years)	44.00(32.00-53.00)	53.00(47.00-61.00)	60.00(52.00-67.00)	0.000	0.000
Incubation period (days)	3.00(2.00-7.00)	6.00(4.00-7.00)	8.00(3.00-13.00)	0.029	0.041
CRP abnormal				0.021	0.072
Yes	130(53.7%)	28(73.7%)	11(78.6%)		
No	112(46.3%)	10(26.3%)	3(21.4%)		
WBC abnormal				0.138	0.064
Yes	45(18.6%)	11(28.9%)	6(42.9%)		
No	197(81.4%)	27(71.1%)	8(57.1%)		
LY % abnormal				0.014	0.030
Yes	101[41.7%]	24(63.2%)	10(71.4%)		
No	141[58.3%]	14(36.8%)	4(28.6%)		
severity status				0.000	0.000
Yes	16(6.6%)	14(36.8%)	8(57.1%)		
No	226(93.4%)	22(42.3%)	6(42.9%)		
Quantitative CT analysis					
LUV(cm ³)	3709.28(2939.74-4538.04)	3885.28(2846.22-4916.78)	4170.63(3129.45-4697.75)	0.606	0.252
LEV(cm ³)	127.82(50.50-265.91)	235.87(85.14-629.24)	237.17(161.81-837.53)	0.001	0.009
LEV/LUV	0.04(0.01-0.08)	0.07(0.02-0.15)	0.09(0.04-0.27)	0.001	0.017
COV(cm ³)	40.45(11.82-96.27)	97.89(23.65-200.86)	94.86(24.80-289.90)	0.001	0.022
COV/LUV	0.01(0.00-0.03)	0.03(0.01-0.06)	0.04(0.01-0.11)	0.001	0.046
GGOV (cm ³)	74.71(31.08-150.72)	135.80(49.43-281.20)	197.58(66.95-377.93)	0.003	0.006
GGOV /LUV	0.02(0.01-0.04)	0.03(0.02-0.10)	0.05(0.02-0.15)	0.003	0.013
LEN	19.00(10.00-37.25)	31.00(20.00-46.00)	30.00(23.00-32.00)	0.001	0.202

Note—CRP: C reactive protein; WBC: white blood cell count; LY %: lymphocyte percentage; LUV: the volume of the whole lung; LEV: the volume of lesion; COV: the volume of consolidation; GGOV: the volume of GGO; LEN: the number of lesions; P1 value: 1

comorbidity vs. without comorbidity; P2 value: ≥ 2 comorbidities vs. without comorbidity; P values written in bold indicate a significant difference.

Table 3 Characteristics of patients with different comorbidities

Variables	Without comorbidity (n = 242)	Hypertension (n=22)	Diabetes mellitus (n=7)	Hepatitis B infection(n=8)	P1 value	P2 value	P3 value
Gender					0.552	0.544	0.111
Male	127(52.5%)	13(59.1%)	5(71.4%)	7(87.5%)			
Female	115(47.5%)	9(40.9%)	2(28.6%)	1(12.5%)			
Age(years)	44.00(32.00-53.00)	56.00(48.50-62.25)	51.00(43.00-69.00)	46.00(43.00-52.00)	0.000	0.055	0.315
Incubation period (days)	3.00(2.00-7.00)	6.50(4.00-7.25)	6.00(5.00-7.00)	4.00(2.25-6.50)	0.076	0.208	0.835
CRP abnormal					0.003	1.000	0.586
Yes	130(53.7%)	19(86.4%)	4(57.1%)	3(37.5%)			
No	112(46.3%)	3(13.6%)	3(42.9%)	5(62.5%)			
WBC abnormal					0.773	0.041	0.379
Yes	45(18.6%)	3(13.6%)	4(57.1%)	3(37.5%)			
No	197(81.4%)	19(86.4%)	3(42.9%)	5(62.5%)			
LY % abnormal					0.047	1.000	0.027
Yes	101(41.7%)	14(63.6%)	3(42.9%)	7(87.5%)			
No	141(58.3%)	8(36.4%)	4(57.1%)	1(12.5%)			
severity status					0.000	0.000	0.010
Yes	16(6.6%)	7(31.8%)	3(42.9%)	3(37.5%)			
No	226(93.4%)	15(68.2%)	4(57.1%)	5(62.5%)			
Pleural effusion					1.000	0.859	0.935
Yes	13(5.4%)	1(4.5%)	1(14.3%)	1(12.5%)			
No	229(94.6%)	21(95.5%)	6(85.7%)	7(87.5%)			
Quantitative CT analysis							
LUV(cm ³)	3709.28(2939.74-4538.04)	3279.78(2647.81-4775.08)	4289.78(3043.78-5644.50)	4457.41(2895.61-5104.92)	0.545	0.400	0.393
LEV(cm ³)	127.82(50.50-265.91)	192.55(68.09-581.92)	570.00(206.43-1365.34)	187.85(76.33-361.09)	0.036	0.001	0.315
LEV/LUV	0.04(0.01-0.08)	0.05(0.02-0.15)	0.17(0.05-0.26)	0.04(0.02-0.09)	0.033	0.001	0.429
COV(cm ³)	40.45(11.82-96.27)	74.24(21.58-212.75)	178.87(119.98-605.27)	67.43(21.00-168.81)	0.039	0.001	0.345
COV/LUV	0.01(0.00-0.03)	0.02(0.01-0.06)	0.04(0.03-0.11)	0.02(0.00-0.05)	0.045	0.001	0.390

GGOV (cm ³)	74.71(31.08-150.72)	104.19(46.51-212.08)	369.13(66.95-765.53)	82.82(48.20-169.62)	0.075	0.004	0.471
GGOV/LUV	0.02(0.01-0.04)	0.03(0.02-0.08)	0.10(0.02-0.15)	0.02(0.01-0.04)	0.047	0.003	0.590
LEN	19.00(10.00-37.25)	37.00(20.00-46.25)	37.00(22.00-46.00)	24.50(16.75-45.25)	0.004	0.068	0.190

Note—CRP: C reactive protein; WBC: white blood cell count; LY %: lymphocyte percentage; LUV: the volume of the whole lung; LEV: the volume of lesion; COV: the volume of consolidation; GGOV: the volume of GGO; LEN: the number of lesions. P1 value: Hypertension vs. without comorbidity; P2 value: Diabetes mellitus vs. without comorbidity; P3 value: Hepatitis B infection vs. without comorbidity; P values written in bold indicate a significant difference.

Figures

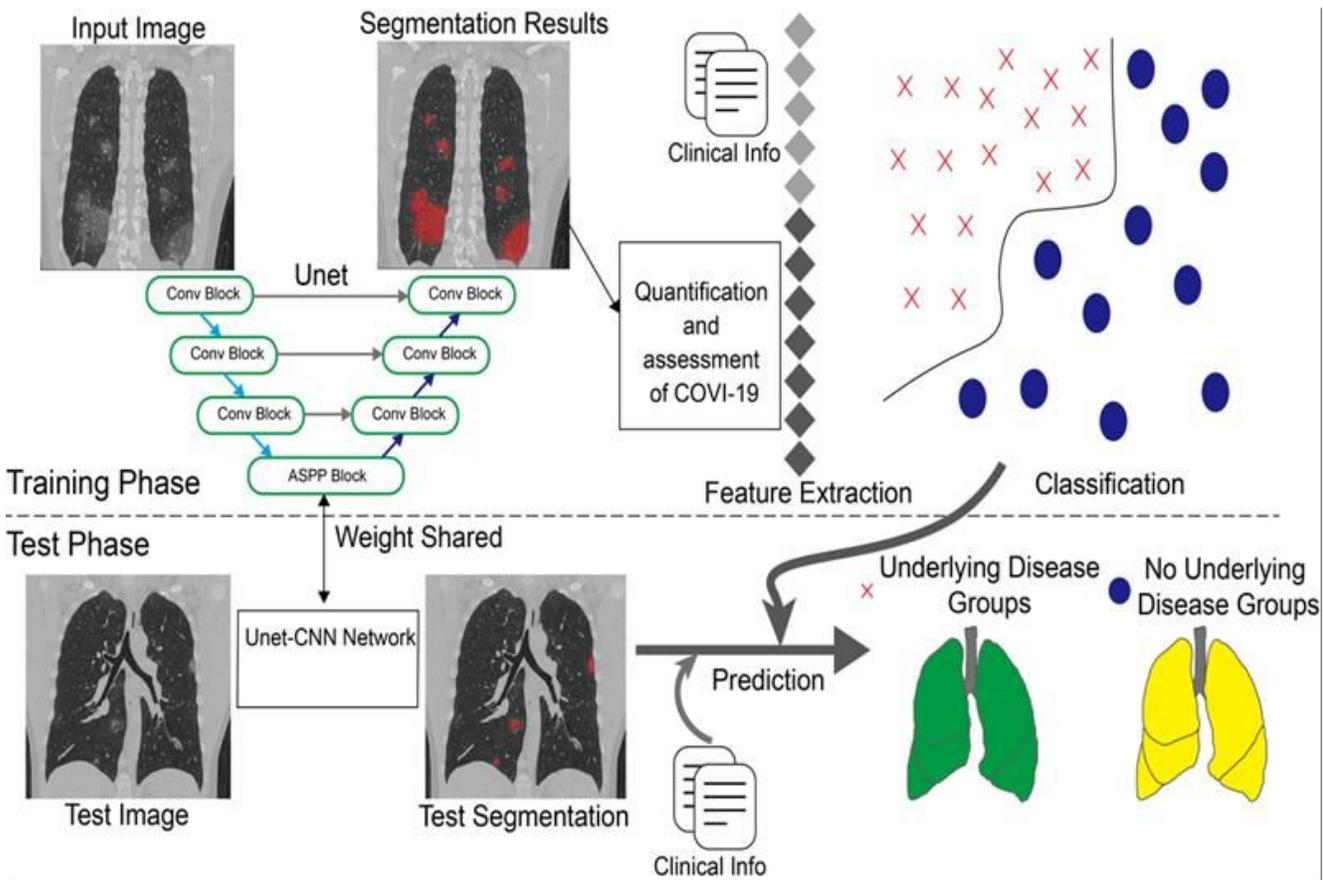


Figure 1

The general flow of Unet neural network to segment lung and lesions. Our neural network model was trained in the training dataset, and tested on test dataset. 550 CT images were split into primary dataset and 100 were primary dataset, respectively. First, CT images were inputted into this neural network to extract image features, segment lung and lesion and further classify whether the lesion was consolidation or GGO. The outputted results were the volumes of the lesions in underlying disease group and no underlying disease group.

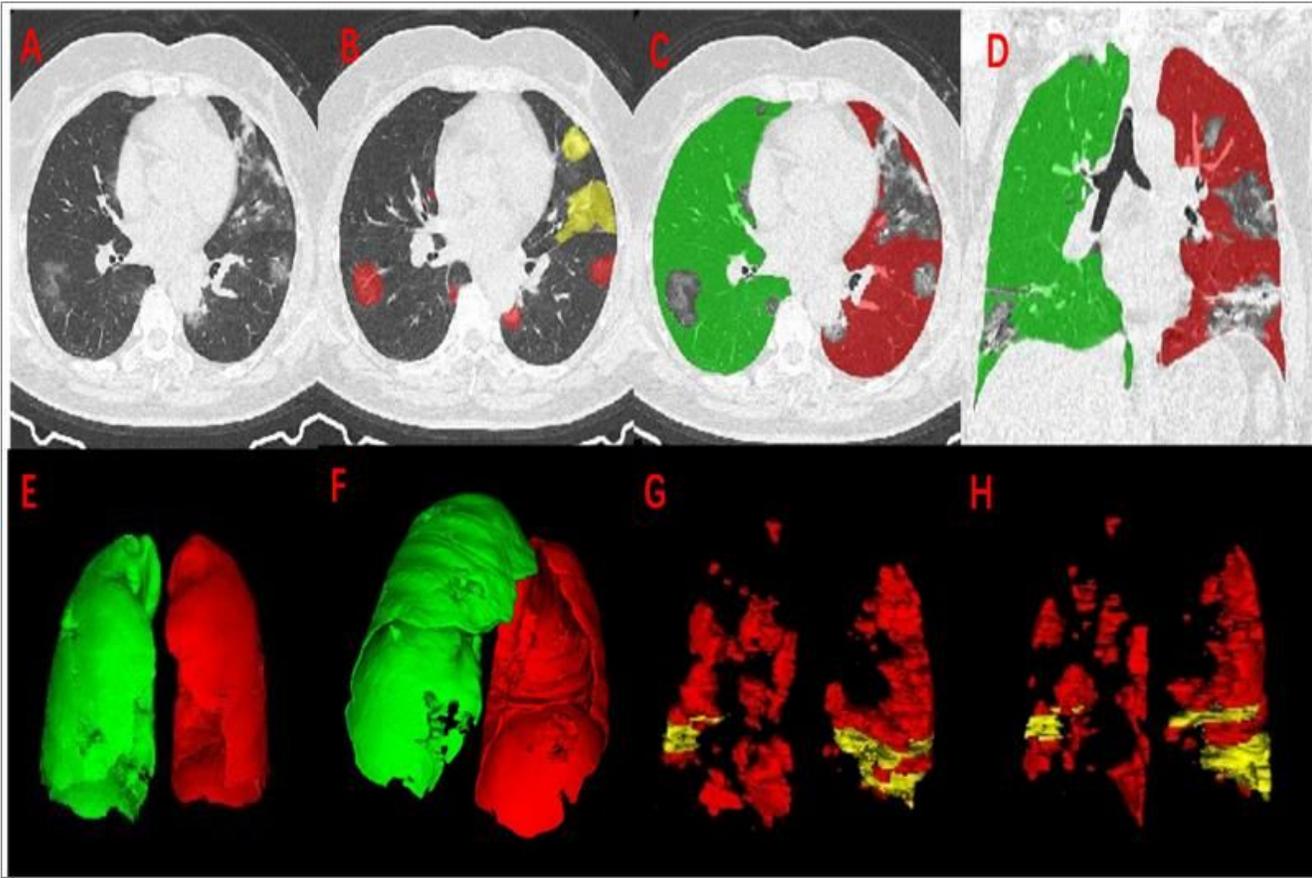


Figure 2

Novel coronavirus pneumonia in a 50-year-old woman in diabetes mellitus group with fever and cough for eight days before admission. A. Axial plane of CT scan in the lung window showed multiple irregular pieces of GGO (black arrow) and consolidation (white arrow). B. GGO (red area) and consolidation (yellow area) were identified and marked according to value of -450 HU. C-D. Axial and coronal plane of CT image showed the margin of the lesion in each slice was delineated. E-F. A 3D lung erosion diagram exhibited the regions of lesion erosion (white arrows). G-H. A 3D lesion diagram demonstrated the volume of GGO (red area) and consolidation (yellow area).

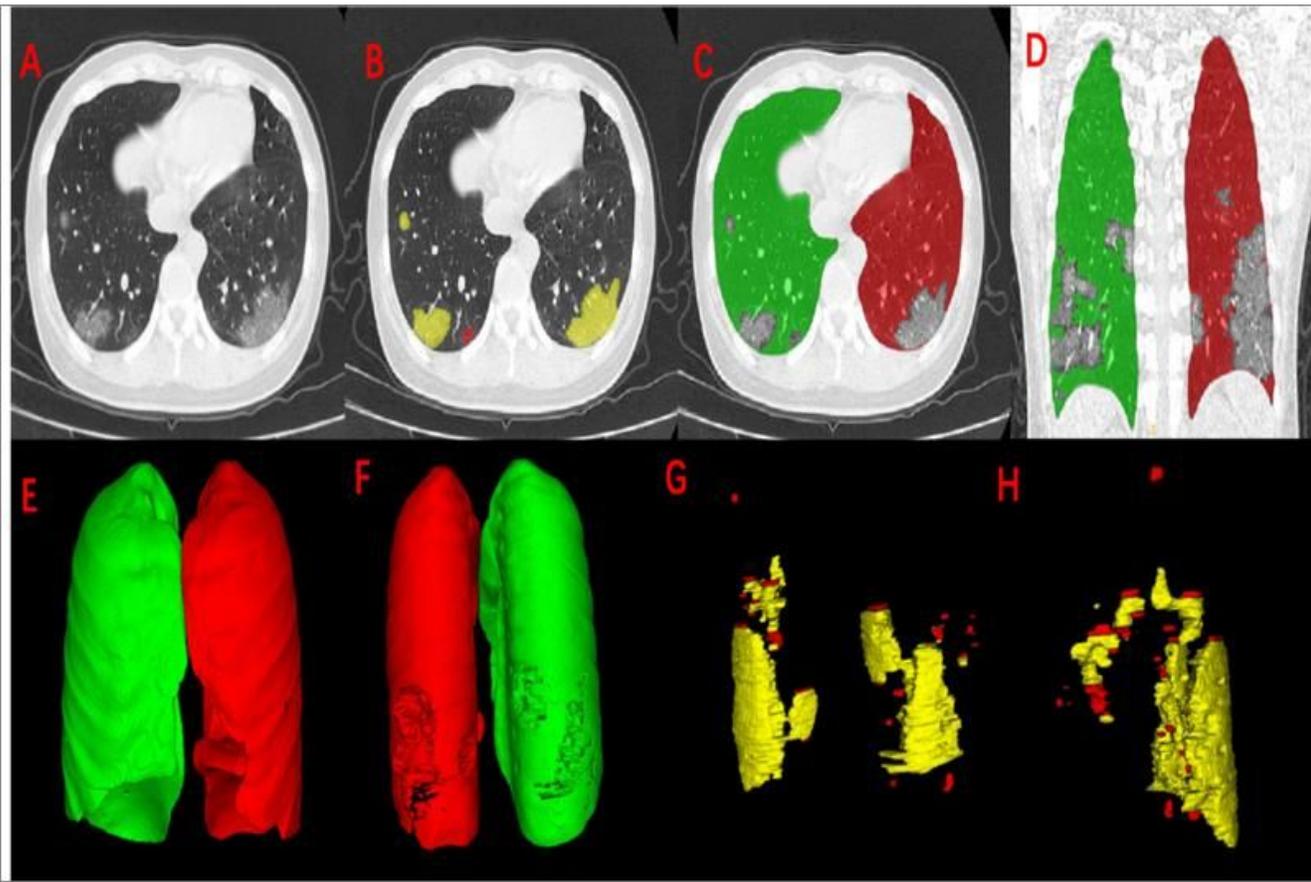


Figure 3

Novel coronavirus pneumonia in a 25-year-old woman without underlying disease. A-B. Axial plane of CT scan in the lung window showed most lesions were GGO (red area) with a little consolidation (yellow area) under the pleura. C-D. The margin of the lesion was delineated showed in axial and coronal plane of CT image. E-H. A 3D lung erosion diagram exhibited the regions of lesion erosion (white arrows). G-H. A 3D region of lesion (white arrows in E-F) was obtained based on delineated results including lung erosion diagram (E-F) and lesion diagram (G-H).