

# BMJ Open Clinical, laboratory and imaging predictors for critical illness and mortality in patients with COVID-19: protocol for a systematic review and meta-analysis

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## ABSTRACT

**Introduction** With the threat of a worldwide pandemic of COVID-19, it is important to identify the prognostic factors for critical conditions among patients with non-critical COVID-19. Prognostic factors and models may assist front-line clinicians in rapid identification of high-risk patients, early management of modifiable factors, appropriate triaging and optimising the use of limited healthcare resources. We aim to systematically assess the clinical, laboratory and imaging predictors as well as prediction models for severe or critical illness and mortality in patients with COVID-19.

**Methods and analysis** All peer-reviewed and preprint primary articles with a longitudinal design that focused on prognostic factors or models for critical illness and mortality related to COVID-19 will be eligible for inclusion. A systematic search of 11 databases including PubMed, EMBASE, Web of Science, Cochrane Library, CNKI, VIP, Wanfang Data, SinoMed, bioRxiv, Arxiv and MedRxiv will be conducted. Study selection will follow the Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines. Data extraction will be performed using the modified version of the Critical Appraisal and Data Extraction for Systematic Reviews of Prediction Modelling Studies checklist and quality will be evaluated using the Newcastle-Ottawa Scale and the Quality In Prognosis Studies tool. The association between prognostic factors and outcomes of interest will be synthesised and a meta-analysis will be conducted with three or more studies reporting a particular factor in a consistent manner. **Ethics and dissemination** Ethical approval was not required for this systematic review. We will disseminate our findings through publication in a peer-reviewed journal. **PROSPERO registration number** CRD 42020178798.

## INTRODUCTION

### Description of the condition

COVID-19, a newly emerged respiratory disease caused by SARS-CoV-2, was first reported in December 2019.<sup>1 2</sup> The infection has recently spread to at least 188 countries and regions, with more than 25 million

## Strengths and limitations of this study

- The evidence synthesis on prognostic factors and models of COVID-19-related critical conditions will play a pivotal role in assisting front-line clinical decision-making.
- The quality of included studies will be evaluated using a validated tool (Quality In Prognosis Studies) specifically developed to assess the risk of bias of prognosis studies.
- Given that primary studies can be conducted in different region, population or setting, prognostic factors or models can be assessed using different tools, heterogeneity in the pooled data may be a limitation of this review; however, subgroup analyses will help overcome this limitation.

confirmed cases and 850 000 deaths worldwide as of 1 September 2020.<sup>3</sup> The number of people infected is probably much higher due to the shortage of tests for COVID-19. Despite a variety of rapid public health responses aimed at containing the disease, many countries have been confronted with enormous challenges to the healthcare systems posed by the overwhelming number of patients requiring hospital admission, especially by those with progression to severe or critical illness according to the criteria in the WHO recommendations or the local guidelines.<sup>4-8</sup>

### Why is it important to do this review?

A report of 72 314 cases from the Chinese Center for Disease Control and Prevention showed that most of the patients with COVID-19 are asymptomatic or exhibit mild or moderate symptoms.<sup>9</sup> The vast majority of patients with mild and moderate symptoms are recommended to stay at home or are admitted in shelter/field hospitals.<sup>10-14</sup>

However, patients with mild symptoms may develop rapidly worsening respiratory failure that requires intubation.<sup>7</sup> Approximately 5%–29% of the patients progressed to a severe or critical condition such as acute respiratory distress syndrome or septic shock and/or multiple organ failure that required admission to the intensive care unit.<sup>9 15–18</sup> Patients who exhibited severe or critical symptoms or patients at high risk to develop severe conditions were the main reason behind the overwhelming number of patients who required admission or even intensive care. Hence, it is crucial to determine the prognostic factors associated with the risk of a subsequent critical outcome among patients with non-critical COVID-19. Prognostic factors and prediction models for severe or critical COVID-19 have many potential uses in various settings including informing individuals about the future course of their illness, aiding triage and referral, early management of modifiable factors, treatment and other factors related to clinical decision-making.

### Status of the current literature

Evidence is rapidly accumulating about prognostic factors and models for critical conditions or mortality related to COVID-19. Recently, two systematic reviews focusing on specific perspectives of COVID-19 have been published.<sup>19 20</sup> Henry and colleagues published a systematic review that included only the laboratory biomarkers and excluded the clinical and imaging predictors associated with severe illness and mortality in COVID-19.<sup>19</sup> Another review by Wynants and colleagues focused on the prediction models for diagnosis and prognosis of COVID-19 infection.<sup>20</sup> Eight studies regarding prognostic models for severe state or mortality were included. However, only the studies aimed at developing or validating a model or a scoring system were included, while those aimed at predictor findings were excluded from this systematic review.<sup>20</sup> In addition, since China was the first epicentre of COVID-19, many studies on the prediction of COVID-19 may have been published in Chinese journals. According to our preliminary results, more than 15 studies regarding prognostic factors or models have been published in four Chinese databases (China National Knowledge Infrastructure (CNKI), Wanfang, CBM and VIP) that were not included in the aforementioned systematic review. Limited data are available on the overview of evidence that focuses on clinical, laboratory and imaging prognostic factors for critical illness or mortality associated with COVID-19. Moreover, a huge number of recent articles have emerged with the worldwide pandemic. Many valuable articles on prognostic factors or models of COVID-19 have not been included in these published reviews. Among these, some high-quality papers have been published in leading journals,<sup>21 22</sup> which provided us with more evidence and insights into this topic. Therefore, there is a need for a systematic review to evaluate and synthesise the data from the current studies from a comprehensive perspective on clinical, laboratory

and imaging prognostic factors and prediction models for critical illness and mortality associated with COVID-19.

### Research aims

We aim to systematically assess the clinical, laboratory and imaging predictors as well as models for severe or critical illness and mortality in patients with COVID-19. Predictors and models for critical illness may be different from that of mortality, so it will be assessed according to different outcomes.

### METHODS

This systematic review protocol followed the Preferred Reporting Items for Systematic Review and Meta-Analysis Protocols (PRISMA-P) recommendations<sup>23</sup> and the Cochrane Handbook. The PRISMA-P checklist is presented in online supplemental appendix 1. This review protocol was started in early April and has been registered with the International Prospective Register of Systematic Reviews.<sup>24</sup>

### Search strategy

A systematic search of 11 public domain databases including PubMed, EMBASE, Web of Science, Cochrane Library, CNKI, Chinese Science and Technology Periodical Database (VIP), Wanfang database (Wanfang Data), China Biology Medicine disc (SinoMed), bioRxiv, Arxiv and MedRxiv will be performed. We will use exploded Medical Subject Headings and the appropriate corresponding keywords related to the population, combined with exposure and outcomes such as: 'COVID-19' OR 'SARS-CoV-2' OR '2019-nCoV' OR 'novel coronavirus' AND 'critically' OR 'severe' OR 'mortality' OR 'deterioration' AND 'predictor' OR 'prediction' OR 'prognostic' OR 'factor'. Additionally, a publication list of the COVID-19 Living Systematic Review<sup>25</sup> and other resources<sup>26</sup> will be screened for additional relevant references. There will be no restrictions on language or publication status (preprint or peer-reviewed articles). The research will be restricted to articles concerning humans from December 2019 to the present. We will include additional papers from other sources including the references of review articles or studies identified during screening. A sample search strategy for PubMed is shown in online supplemental appendix 2.

### Eligibility criteria

#### Participants

All patients with confirmed diagnosis of COVID-19, explicitly classified as mild, moderate, severe or critically ill according to accepted diagnostic criteria such as the WHO recommendations or the local guidelines, will be included. The criteria in the guidelines may be modified over time. Thus, the criteria in different periods or regions will be acceptable.

#### Exposures

Any data related to demographics, symptoms and signs, pulmonary functions, laboratory tests, radiological

findings, comorbidities and interventions will be considered potential predictors for critical illness or mortality in patients with COVID-19. This information may include factors such as the age, fever, shortness of breath, underlying diseases, mechanical ventilation, and dexamethasone or other interventions.

### Comparators

Based on the published studies, many factors including older age; underlying diseases such as hypertension, diabetes and cardiovascular diseases; and chest radiographic abnormalities were independent predictive factors for critical illness in hospitalised patients with COVID-19.<sup>21 22</sup> These potential variables will be considered the comparators. Participants with and without specific clinical, laboratory and imaging information will be compared to clarify the significance of this information in predicting critical illness and mortality associated with COVID-19.

### Outcomes

The outcomes will include deterioration, progression, severe critical illness or death related to COVID-19 according to accepted criteria.

### Timing and setting

There will be no restriction on the time point when the prognostic factors were under review as well as on the period when the outcomes were predicted. No restriction will be imposed on the setting.

### Types of study to be included

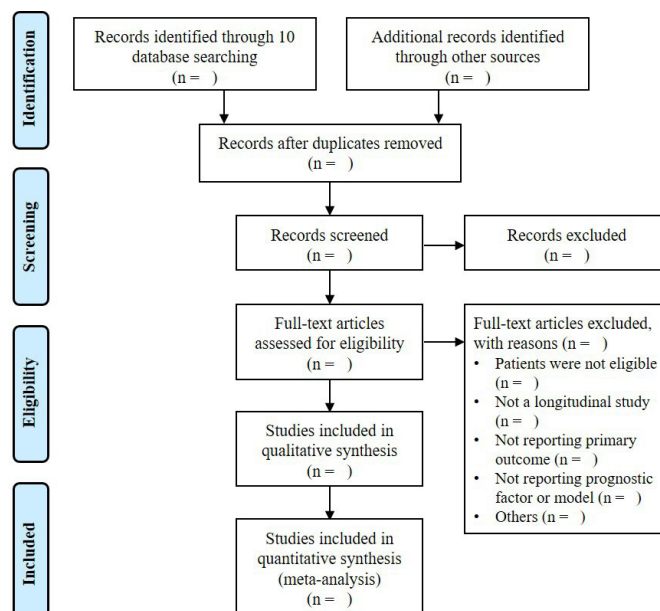
Both experimental and longitudinal observational studies including randomised controlled trials, cohort studies, case-control studies and registry studies will be included. Review articles, editorials, letters, comments, case reports, cross-sectional studies and studies that failed to investigate the prognostic factors or models for critical illness or mortality will be excluded.

### Study selection

Two reviewers (JL, LF) will independently perform the initial search and examine the titles, abstracts and full texts (if necessary) to identify eligible studies according to the inclusion and exclusion criteria. Disagreements between the reviewers will be resolved by consensus and by adjudication of a third reviewer (QL) in case of persistent disagreement. The selection process is illustrated in a Preferred Reporting Items for Systematic Reviews and Meta-Analyses diagram (figure 1).<sup>27</sup>

### Data extraction

Data extraction will be independently conducted by two reviewers (TZ, PJ) using a standard data extraction form developed according to the Critical Appraisal and Data Extraction for Systematic Reviews of Prediction Modelling Studies (CHARMS) checklist for prediction modelling studies and its modified version (CHARMS-PF)<sup>28 29</sup> as well as according to the Prediction Model Risk of Bias



**Figure 1** Flow diagram of the study selection process.

Assessment Tool.<sup>30</sup> For each included trial, the following key information will be extracted based on availability: name of the first author, year of publication, study location, study design, study setting, participants, sample size, follow-up period, outcomes of interest, risk and prognostic factors, missing data, summary statistics, results, interpretation and discussion. The authors of the studies will be contacted through email or telephone in case of missing relevant data.

### Assessment of the risk of bias

We will evaluate the risk of bias using the Newcastle-Ottawa Scale<sup>31</sup> and the Quality In Prognosis Studies checklist, which has been recommended by the Cochrane Group to assess the risk of bias in studies related to prognostic factors.<sup>32</sup> Quality assessment will be performed independently by two reviewers (TZ, LK) and discrepancies will be resolved through consensus.

### Data synthesis

Essential data will be summarised in tables for evaluation. Estimates of risk difference in terms of critical illness and mortality will be calculated. For categorical variables, ORs, relative risk or HRs will be analysed to compare these variables between mild/moderate and severe/critical COVID-19 cases. Studies reporting adjusted or unadjusted results will be analysed separately. Only the unadjusted effect estimates for prognostic factors will be combined, while effect estimates from multivariate models will be described qualitatively. With three or more studies reporting a particular factor in a consistent manner, a meta-analysis will be conducted using the Review Manager software (RevMan V.5.3, the Cochrane Collaboration, London, UK) to synthesise the association of prognostic factors and critical illness or mortality in patients with COVID-19. For severe or critical illness and mortality, the data will be synthesised according to



different outcomes. Heterogeneity among the included studies will be tested using the  $I^2$  statistic.<sup>33</sup> Forest plots will be presented as significant predictors. In case of substantial heterogeneity, subgroup analyses will be conducted to examine or to explore the causes of heterogeneity. Subgroup analysis will be based on the categories defined by the following characteristics: study location/region, risk of bias and particular population such as children and elderly people.

### Ethics and dissemination

Ethical approval was not required for this systematic review. We will disseminate our findings through publication in a peer-reviewed journal.

### Patient and public involvement

There is no patient or public involvement in the whole process of conducting this systematic review.

## DISCUSSION

With an unprecedented threat of a worldwide COVID-19 pandemic, there has been an increasing need for early identification of patients at higher risk of progression to critical illness or even death. This systematic review will comprehensively summarise the existing evidence on clinical, laboratory and imaging factors and models for predicting critical conditions and mortality in patients with COVID-19. The findings of this review will provide front-line clinicians an early surrogate for disease severity before the onset of critical illness, which may play a key role in assisting the clinicians in early management of modifiable factors, appropriate triaging of patients and optimising the use of limited healthcare resources.

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**Contributors** XL and YG conceived the research question. TZ, LF, PJ and LK developed the search strategy and performed the preliminary search, screening and data extraction. QL and YG contributed to the methodological development of the protocol. XL and JL drafted the manuscript. QL and YG revised the manuscript and all authors developed and approved the final manuscript before submission.

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