

# Could there be a link between oral hygiene and the severity of SARS-CoV-2 infections?

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## Key points

Describes what COVID-19 is and the associated risk factors for developing complications.

Investigates whether bacteria contribute to COVID-19 complications by causing bacterial superinfections.

Examines whether there is a link between the oral microbiome and COVID-19 complications.

## Abstract

On 30 January 2020, the World Health Organisation identified COVID-19, caused by the virus SARS-CoV-2, to be a global emergency. The risk factors already identified for developing complications from a COVID-19 infection are age, gender and comorbidities such as diabetes, hypertension, obesity and cardiovascular disease. These risk factors, however, do not account for the other 52% of deaths arising from COVID-19 in often seemingly healthy individuals. This paper investigates the potential link between SARS-CoV-2 and bacterial load, questioning whether bacteria may play a role in bacterial superinfections and complications such as pneumonia, acute respiratory distress syndrome and sepsis. The connection between COVID-19 complications and oral health and periodontal disease is also examined, as the comorbidities at highest risk of COVID-19 complications also cause imbalances in the oral microbiome and increase the risk of periodontal disease. We explore the connection between high bacterial load in the mouth and post-viral complications, and how improving oral health may reduce the risk of complications from COVID-19.

## Introduction

On 31 December 2019, the World Health Organisation (WHO) was informed of a cluster of 27 cases of pneumonia with no known cause linked to a wet animal wholesale market in Wuhan City, China.<sup>1</sup> By 7 January 2020, the causative virus was identified as SARS-CoV-2, which causes the disease COVID-19, and was declared a global emergency by the WHO on 30 January 2020.

Zhu *et al.* (2020) were able to identify and characterise SARS-CoV-2 using unbiased sequencing samples from infected patients with pneumonia.<sup>2</sup> The viral genome revealed SARS-CoV-2 to be 75–80% identical to severe acute respiratory syndrome coronavirus

(SARS-CoV) and several bat coronaviruses. Prior to this discovery, six coronaviruses were known to cause human disease. Four viruses (229E, OC43, NL63 and HKU1) are prevalent and cause common cold symptoms. The two other strains are SARS-CoV and Middle East respiratory syndrome Coronavirus (MERS-CoV), both of which are zoonotic in origin and fatal. SARS-CoV-2 has joined as the seventh member of the family of coronaviruses that infect humans.<sup>2</sup> No specific treatment is yet accessible; however, management involves restriction of travel, patient isolation and the support of medical supervision.<sup>3</sup>

COVID-19 affects people in different ways, with patients exhibiting a range of symptoms and severity. While risk factors such as age, gender and comorbidities have been highlighted as increasing the risk of complications and mortality, there is still a high proportion of patients with no identified risk factors who suffer from severe side effects and complications. As much as 10–15% of people under 60 years old with no risk factors are exhibiting moderate to severe reactions to COVID-19.<sup>1</sup> While COVID-19 has a viral

origin, it is suspected that in severe cases, bacterial superinfections may contribute to causing complications such as pneumonia and acute respiratory distress syndrome (ARDS). We explore the connection between high bacterial load in the mouth and post-viral complications, and how improving oral health may reduce the risk of complications from COVID-19.

## Symptoms

The clinical symptoms of COVID-19 appear after an incubation period of approximately 5.2 days<sup>4</sup> and range from fever (98.6%), fatigue (69.6%), dry cough (59.4%), myalgia (34.8%) and sore throat (17.4%).<sup>5</sup> Diarrhoea has also been shown to be a key distinguishing symptom of COVID-19 compared with SARS-CoV and MERS-CoV.<sup>6,7</sup>

COVID-19 infections can present with mild, moderate or severe illness. The severity of the illness, the rate of decline and the risk of mortality are significantly dependent on risk factors highlighted by Zhou *et al.* (2020) in the largest retrospective cohort study among

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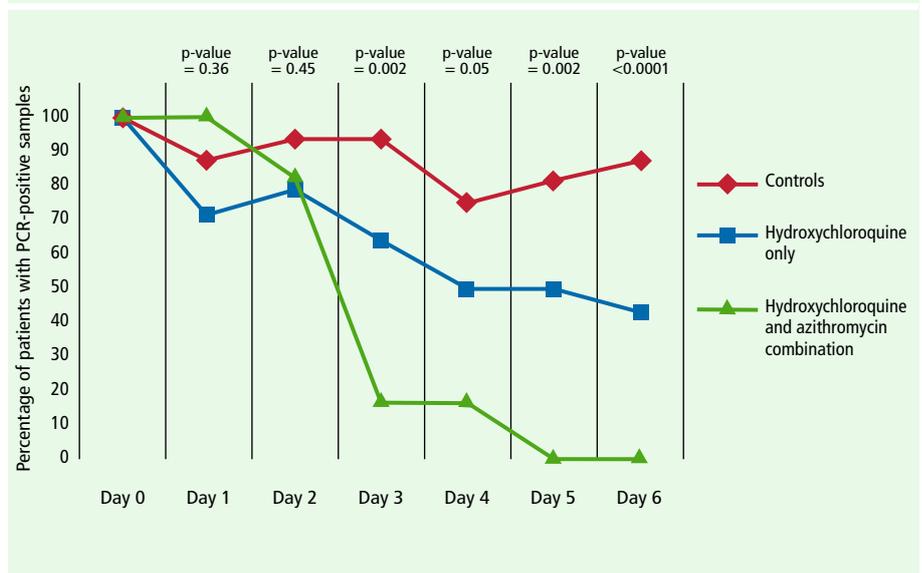
patients with COVID-19.<sup>8</sup> These risk factors include age (the mean age was 69 years old), gender (men represented 70% of deaths) and an underlying comorbidity in 48% of cases (hypertension 30%, diabetes 19%, or heart disease 8%). In April, research from Europe added obesity as a risk factor for developing complications from COVID-19, with 47.6% of patients in intensive care units (ICUs) having a BMI over 30 kg/m<sup>2</sup>.<sup>9</sup> The main complications of severe illness are blood clots, pneumonia, sepsis, septic shock and ARDS.<sup>1</sup> It has also been discovered that, in some cases, a dysregulated immune reaction triggers overproduction of early response pro-inflammatory cytokines (tumour necrosis factor [TNF], IL-6, and IL-1β), resulting in a 'cytokine storm'.<sup>10</sup> This exposes patients to increased risk of vascular hyperpermeability and multi-organ failure, particularly to the heart and kidneys.<sup>11,12,13</sup> IL-6 levels are shown to be comparatively higher in non-survivors than those who survive.<sup>10</sup>

When pneumonia spreads to the lungs and blood oxygen levels fall, patients require assisted ventilation, shifting their diagnosis from pneumonia to ARDS. Of 201 severely affected inpatients in China, 41.8% developed ARDS. Of these, 52.4% died due to respiratory failure, making ARDS the primary cause of death in infected patients.<sup>14</sup> This is supported by a country report covering Italy, showing that 96.5% of complications arising from COVID-19 infection were ARDS, followed by acute renal failure (29.2%).<sup>15</sup> This suggests that patients are more likely to die from post-viral infection complications than from COVID-19. ARDS was also the biggest cause of death for SARS-CoV<sup>16</sup> and MERS-CoV infections,<sup>17</sup> further confirming the connection between the three coronaviruses. As of 3 March 2020, COVID-19 was predicted to have a 3.4% mortality rate.<sup>5</sup> The period from the onset of symptoms to death averages 14 days.

## Discussion

A significant unanswered question is why some patients suffer from COVID-19 more severely than others. Though patients with risk factors such as age, gender and comorbidities have an increased rate of complications and mortality, there is still a high proportion of seemingly young and healthy infected patients with no identified risk factors who suffer from severe side effects and complications, while some infected patients never go on to develop anything more than anosmia.<sup>18</sup> It is common

**Fig. 1 The different outcomes of different treatment options for COVID-19 patients. Reproduced from Gautret P, Lagier J-C, Parola P *et al.*, Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial, *Int J Antimicrob Agents*, 2020, Elsevier<sup>23</sup>**



for respiratory viral infections to predispose patients to bacterial superinfections, leading to increased disease severity and mortality; for example, during the influenza pandemic in 1918, where the primary cause of death was not from the virus itself but from bacterial superinfections. The same was apparent in the 2009 H1N1 influenza pandemic, where again bacterial superinfections were the primary cause of death as opposed to the virus itself.<sup>19</sup> Despite the proven importance of superinfections in the severity of respiratory disease, they are often understudied during respiratory infection outbreaks, mainly due to the fact that diagnosing a superinfection is complex.<sup>19</sup>

Other strands of the coronavirus have shown to enhance streptococcal adherence to epithelial cells along the respiratory tract, causing complications such as pneumonia<sup>20</sup> and inflammatory damage in the lungs that inhibit the clearance of bacteria. Zheng *et al.*<sup>21</sup> compared the levels of granulocytes in patients suffering from mild COVID-19 infections to those suffering from severe infections. Patients suffering from severe infections had a remarkably higher neutrophil count and a significantly lower lymphocyte count than in mild patients. A high neutrophil count is abnormal for a viral infection, but very common for a bacterial infection, resulting in Zheng *et al.* suspecting that, in severe cases of COVID-19, bacterial superinfection is common. Zheng *et al.* argued that abnormally

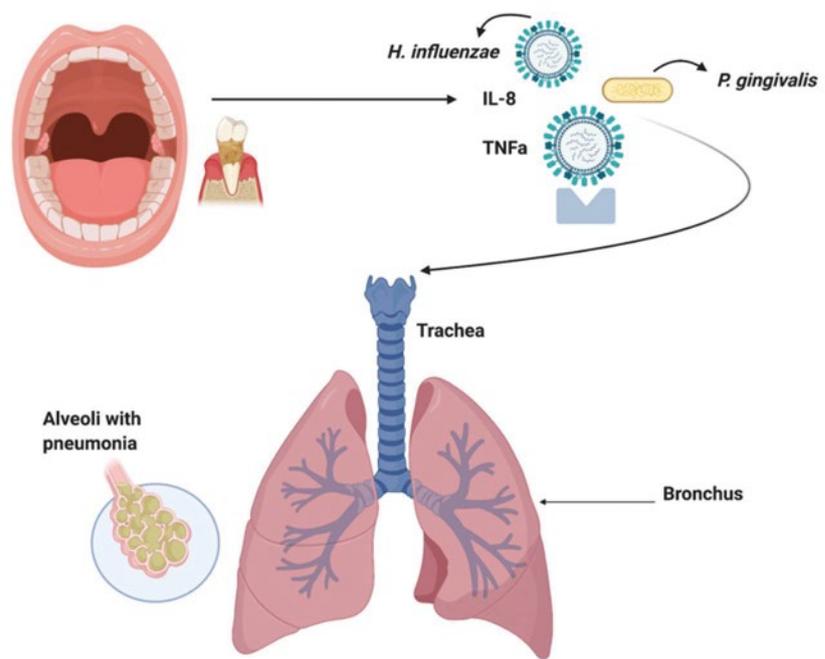
low levels of lymphocytes, which are usually the main line of defence against viral infections, indicates either functional exhaustion of cells<sup>21</sup> or potentially that, in severe cases of the disease, a bacterial superinfection supersedes the original viral infection. This is supported by the fact that Zhou *et al.* noted 50% of patients with severe COVID-19 died with the presence of a secondary bacterial infection.

When lymphocyte levels reduce, severity increases. This was also seen by Liu *et al.*,<sup>22</sup> who deduced that the neutrophil-to-lymphocyte ratio predicts the severity of the patient's response to COVID-19. They found that in severe cases, over 80% of patients had an exceptionally high bacterial load secondary to a bacterial superinfection and required antibiotics. This is supported by the promising research performed in France,<sup>23</sup> where a combination of hydroxychloroquine (antiviral) and azithromycin (antibiotic) was shown to cure 100% of patients virologically after six days, compared to patients who had hydroxychloroquine alone (57.1%) and those who had no treatment (12.5%). While the use of hydroxychloroquine has been under much dispute, it is important to understand that antibiotics clearly play an influential role in treatment outcome, improving the success rate by 42.9% in this case. A retrospective report of 1,060 patients in May 2020 confirms the efficacy of using a combination of an antibiotic with an antiviral, with 91.7% virologically cured after ten days.<sup>24</sup> Furthermore, Chen

*et al.* reported that 71% of COVID-19 patients in China needed antibiotics upon admission to hospital,<sup>25</sup> while in Italy, 84% of patients needed antibiotics,<sup>26</sup> illustrating the potential importance of antibiotics and therefore bacteria in the course of a COVID-19 complication.<sup>19</sup> See Figure 1 for the outcomes of different treatment options for COVID-19.

The lungs, similar to the oral cavity, are often referred to as an 'ecological community of commensal, symbiotic and pathogenic organisms'.<sup>27</sup> Microbial immigration and elimination are constant between the oral cavity and the lungs, allowing for a healthy microbiotic distribution.<sup>28</sup> Similar to the gut or the oral cavity, bacterial communities that colonise the lungs are recognised for their function in conserving tissue, immune and organ homeostasis. The lungs are unique, however, as they are oxygen-rich and therefore contain a fine equilibrium of lipid-rich surfactants that influence bacteriostatic activity. This equilibrium can alter drastically during illness, allowing microbial overgrowth and injury to the lungs. Lower respiratory infection is initiated by the contamination of the lower airway epithelium by inhalation of microorganisms encompassed in aerosolised droplets, or by aspiration of oral secretions associated with oral disease (containing microorganisms such as *P. gingivalis*, *F. nucleatum*, *P. intermedia*).<sup>29</sup> Periodontitis and decay are the two most common oral diseases associated with an imbalance of pathological bacteria in the mouth. Cytokines (such as IL-1 and TNF) from periodontally diseased tissues can infiltrate the saliva through the gingival crevicular fluid and be aspirated to cause inflammation or infection within the lungs (Fig. 2).<sup>30</sup> Therefore, inadequate oral hygiene can increase the risk of inter-bacterial exchanges between the lungs and the mouth, increasing the risk of respiratory infections and potentially post-viral bacterial complications.<sup>31</sup> In a nationwide population-based cohort study, 49,400 chronic periodontitis patients were treated with periodontal therapy over 11 years. The Kaplan-Meier plot demonstrated the total incidences of pneumonia significantly decreased over a 12-year follow-up in the group receiving periodontal therapy ( $p < 0.001$ ). The reduction or eradication of periodontal disease significantly reduces one's risk of pneumonia.

Good oral hygiene has been recognised as a means to prevent airway infections in patients, especially in those over the age of 70.<sup>32</sup> The oral cavity houses more than 700 bacteria, viruses



**Fig. 2** Bacteria that colonise the mouth are shed into the saliva. The pathogenic bacteria within the saliva can then be aspirated into the lower respiratory tract and cause or aggravate an infection. Reproduced with permission from Scannapieco F A, Role of Oral Bacteria in Respiratory Infection, *J Periodontol*, 1999, John Wiley and Sons.<sup>30</sup>

and fungi that can colonise the mouth.<sup>33</sup> Various microbiological habitats exist within the mouth; however, the primary bacterial inhabitants are *P. intermedia*, *S. mutans*, *F. nucleatum* and *P. gingivalis*.<sup>34</sup>

Several mechanisms have been proposed to explain the potential role of oral bacteria in the pathogenesis of a respiratory infection:<sup>30</sup>

1. Aspiration of oral pathogens into the lungs
2. Periodontal disease-associated enzymes may modify the mucosal surfaces to allow for adhesion and colonisation of respiratory pathogens
3. Periodontal disease-associated enzymes may destroy the salivary pellicles on bacteria to hinder clearance from mucosal surfaces
4. Respiratory epithelium may be altered by periodontal-associated cytokines to promote infection by respiratory pathogens.<sup>35</sup>

The bacteria in the oral biofilm are in prime position to be aspirated into the respiratory tract and help initiate or progress conditions such as pneumonia or sepsis. A high bacterial and viral load in the mouth can further complicate systemic diseases such as cardiovascular disease, cancer, neurodegenerative disease and autoimmune diseases,<sup>36</sup> further supporting the link between mouth and body. The established risk factors for COVID-19 (age, gender and

comorbidities)<sup>8</sup> are also heavily implicated in imbalances in the oral microbiome. Diabetes, hypertension and heart disease are associated with higher numbers of *F. nucleatum*, *P. intermedia* and *P. gingivalis*,<sup>37</sup> and associated with the progression of periodontal disease. Patients with periodontal disease are at a 25% increased risk of cardiovascular disease,<sup>38,39</sup> triple the risk of diabetes mellitus<sup>40</sup> and a 20% increased risk of hypertension.<sup>41</sup> Epithelial sensitisation and haematogenous spread of proinflammatory mediators such as cytokines, produced in the diseased periodontal tissue, can increase systemic inflammation and diminish airflow. This can be exacerbated by the stimulation of the liver to produce acute phase proteins, such as interleukin-6, which potentiate the inflammatory response of the lungs and the rest of the body.<sup>42</sup> Similarly, patients with severe COVID-19 infections also express systemic inflammation and significantly higher levels of interleukin-6, interleukin-2, interleukin-10, TNF and C-reactive protein.<sup>43</sup>

Nagaoka *et al.*<sup>44</sup> investigated the effects of periodontopathic bacteria such as *F. nucleatum*, *P. gingivalis* and *P. intermedia* on pneumonia, and found that *P. intermedia* in particular induced severe pneumonia in subjects with higher levels of periodontopathic bacteria. Pneumonia and acute viral

respiratory infections are two of the most common airway infections in older patients<sup>45</sup> and the greatest causes of death in patients over 70 years. A randomised controlled trial in Japan investigated whether improved oral care reduced the incidence of pneumonia and pneumonia-related deaths. Four hundred and seventeen patients were provided with oral care after every meal and compared with the control group. Of the control group, 19% contracted pneumonia compared with only 11% who received oral care. Moreover, the post-pneumonia mortality rate in the control group was almost double that of the group prescribed oral care.<sup>45</sup> The link between good oral care and a reduced risk of acute viral respiratory infections has been established in a number of other studies,<sup>30,46,47,48</sup> including a systematic review that concluded one in ten pneumonia-related deaths in the elderly could be prevented by improving oral hygiene.<sup>32</sup> Furthermore, improved oral care can significantly reduce the incidence of ventilator-associated pneumonia in ICU patients.<sup>49</sup>

## Conclusion

The more severe the form of COVID-19, the higher the chance of complications such as pneumonia, ARDS, sepsis, septic shock and death. The development, severity and risk of complications following a COVID-19 infection depend on a number of host and viral factors that will affect a patient's immune response. While 80% of patients with COVID-19 infections have mild symptoms, 20% progress to have a severe form of infection associated with higher levels of inflammatory markers (IL-2, IL-6, IL-10), bacteria and neutrophil-to-lymphocyte count.

We suggest that the connection between the oral microbiome and COVID-19 complications should be investigated in the process of better understanding the outcomes of COVID-19 disease. The four main comorbidities associated with an increased risk of complications and death from COVID-19 are also associated with altered oral biofilms and periodontal disease, hence why the link between poor oral health and COVID-19 complications is suggested. Periodontopathic bacteria are implicated in systemic inflammation, bacteraemia, pneumonia and even death. Periodontopathic bacteria is also present in the metagenome of patients severely infected with SARS-CoV-2, where high reads for *Prevotella* (493 reads), *Staphylococcus* (1,659 reads) and

*Fusobacterium* (463 reads) were discovered.<sup>50</sup> It is clear that bacterial superinfections are common in patients suffering from a severe case of COVID-19, with more than 50% of deaths exhibiting bacterial superinfections.<sup>19</sup> Furthermore, it is common for respiratory viruses to predispose patients to bacterial superinfections, as seen in the influenza outbreaks in 1918 and 2009.<sup>19</sup> Over 80% of patients in ICU exhibited an exceptionally high bacterial load,<sup>22</sup> and treatment has been successful with a dual regime of an antiviral and an antibiotic.<sup>23</sup>

Despite the proven importance of superinfections in the severity of respiratory viral diseases, they are often understudied due to the complexity of diagnosis and culture-based microbiological testing being less sensitive once antibiotics are administered.<sup>19</sup>

More research should be performed on bacterial superinfections, and the connection, if any, between the oral microbiome and COVID-19 complications are urgently required to establish the importance of oral hygiene and pre-existing oral disease in the severity and mortality risk of COVID-19.

Meanwhile, we recommend that oral hygiene be maintained, if not improved, during a SARS-CoV-2 infection in order to reduce the bacterial load in the mouth and the potential risk of a bacterial superinfection. We recommend that poor oral hygiene be considered a risk to post-viral complications, particularly in patients already predisposed to altered biofilms due to diabetes, hypertension or cardiovascular disease. Bacteria present in patients with severe COVID-19 are associated with the oral cavity and improved oral hygiene may play a part in reducing the risk of complications.

### Conflict of interest

The authors declare no conflicts of interest related to the study.

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