

HYPOTHESES

A hypothesized role for dysregulated bradykinin signaling in COVID-19 respiratory complications

Joseph A. Roche¹ | Renuka Roche²

¹Eugene Applebaum College of Pharmacy and Health Sciences, Wayne State University, Detroit, MI, USA

²College of Health and Human Services, Eastern Michigan University, Ypsilanti, MI, USA

Correspondence

Joseph A. Roche, Eugene Applebaum College of Pharmacy and Health Sciences, Wayne State University, 259 Mack Ave., Detroit, MI 48201, USA.
 Email: joseph.roche@wayne.edu

Funding information

Eugene Applebaum College of Pharmacy and Health Sciences, Wayne State University (EACPHS, WSU)

Abstract

As of April 20, 2020, over time, the COVID-19 pandemic has resulted in 157 970 deaths out of 2 319 066 confirmed cases, at a Case Fatality Rate of ~6.8%. With the pandemic rapidly spreading, and health delivery systems being overwhelmed, it is imperative that safe and effective pharmacotherapeutic strategies are rapidly explored to improve survival. In this paper, we use established and emerging evidence to propose a testable hypothesis that, a vicious positive feedback loop of des-Arg(9)-bradykinin- and bradykinin-mediated inflammation → injury → inflammation, likely precipitates life threatening respiratory complications in COVID-19. Through our hypothesis, we make the prediction that the FDA-approved molecule, icatibant, might be able to interrupt this feedback loop and, thereby, improve the clinical outcomes. This hypothesis could lead to basic, translational, and clinical studies aimed at reducing COVID-19 morbidity and mortality.

KEYWORDS

bradykinin, bradykinin receptor, coronavirus, icatibant, inflammation, injury

According to data reported by the World Health Organization through its COVID-19 homepage, as of April 20, 2020, 2:00 AM CEST, out of 2 319 066 confirmed cases over time, there have been 157 970 deaths, putting the Case Fatality Rate at ~6.8%.¹ As the COVID-19 pandemic is rapidly spreading, and health delivery systems are being overwhelmed by the large numbers of patients needing acute care for breathing difficulty, it is imperative that safe and effective pharmacotherapeutic strategies are rapidly explored to improve survival.^{2,3} Since time is of the essence to reduce mortality in

patients with COVID-19 respiratory complications, repurposing FDA-approved drugs that have a good safety profile for off-label and/or compassionate use should be a strategic priority.⁴

It is in this context that we propose a testable hypothesis for dysregulated bradykinin (BK) signaling in COVID-19 respiratory complications. Through our hypothesis, we hope that researchers and clinicians would be able to identify candidate drugs for off-label and/or compassionate use in patients with unremitting respiratory distress from COVID-19.

Abbreviations: ACE, angiotensin converting enzyme; APP, aminopeptidase-P; B1R, bradykinin-B1-receptor; B2R, bradykinin-B2-receptor; BK, bradykinin; CoV, coronavirus; COVID-19, coronavirus disease 19; DABK, des-Arg(9)-bradykinin; DPP4, dipeptidyl peptidase-4; ER, endoplasmic reticulum; FDA, United States Food and Drug Administration; HAE, hereditary angioedema; IL, interleukin; SARS, severe acute respiratory syndrome.

We wish to acknowledge that prior to submission of this article, we had posted an earlier version of this hypothesis in the form of an Open Letter (https://figshare.com/articles/AN_OPEN_LETTER_TO_THE_SCIENTIFIC_COMMUNITY_ON_THE_POSSIBLE_ROLE_OF_DYSREGULATED_BRADYKININ_SIGNALING_IN_COVID-19_RESPIRATORY_COMPLICATIONS/12093696). We apologize for not having disclosed this, as is required by the FASEB Journal's submission policy.

This article was fast-tracked under a recently instituted interim policy in which the editors may, at their discretion, accept coronavirus-related manuscripts submitted for the Review, Perspectives and Hypotheses categories without additional review.

Based on our examination of basic and clinical studies, we hypothesize that dysregulated BK signaling is involved in COVID-19 respiratory complications for the following reasons (also see Figure 1):

- The severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2), which causes COVID-19, is known to enter host cells in the respiratory system via the transmembrane protein, angiotensin converting enzyme 2 (ACE2)^{5,6}
- SARS-CoV infection depletes ACE2⁷

- ACE2 depletion increases levels of des-Arg(9)-bradykinin (DABK), which is a bioactive metabolite of BK that is associated with lung injury and inflammation⁸⁻¹⁰
- A possible role for BK in COVID-19 respiratory distress is consistent with established evidence that, BK, histamine, and serotonin, have for long been known as key mediators of acute lung inflammation and respiratory distress¹¹

Experimental evidence suggests that, most downstream effects of DABK are mediated through its binding to the BK-B1-receptor (B1R). However, DABK not only binds strongly

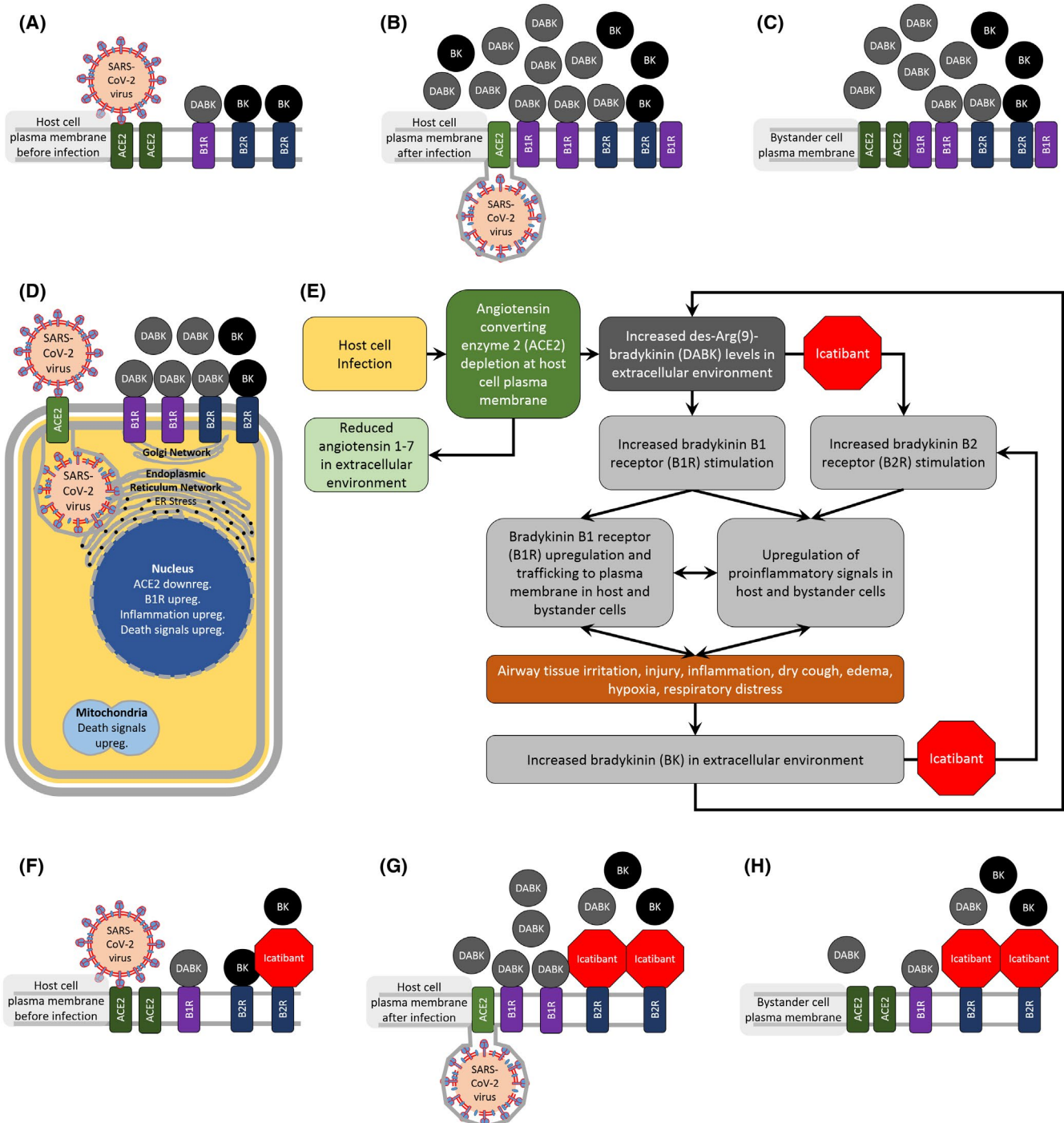


FIGURE 1 Hypothesized role for dysregulated bradykinin signaling in COVID-19 respiratory complications and the potential benefit of bradykinin receptor blockers. SARS coronavirus-2 (SARS-CoV-2), the virus that causes coronavirus disease 19 (COVID-19), is known to enter host cells in the respiratory system via the transmembrane protein, angiotensin converting enzyme 2 (ACE2)^{5,6} (Panel A). SARS-CoV infection depletes ACE2 at the plasma membrane of infected cells⁷ (Panel B). In the extracellular environment of both infected cells as well as neighboring bystander cells, ACE2 depletion increases the levels of des-Arg(9)-bradykinin (DABK), which is a bioactive metabolite of bradykinin (BK) that is associated with airway inflammation⁸ (Panels B, C). SARS-CoV infection severely affects host cell homeostasis,⁵³ by triggering endoplasmic reticulum stress,⁵⁴ mitochondrial death signaling,⁵⁵ downregulation of ACE2,⁷ upregulation of pro-inflammatory genes,⁵⁶ and nuclear death signals,⁵⁷ which ultimately lead to cell death⁵³ (Panels D, E). Cellular injury and inflammation induces BK-B1-receptor (B1R) upregulation and trafficking to the plasma membrane, which amplifies DABK-mediated inflammation and injury^{58,59} (Panel D). Tissue injury and inflammation also increases BK levels and BK-B2-receptor (B2R) stimulation^{59,60} (Panels D, E). Our testable hypothesis for dysregulated BK signaling in COVID-19 respiratory complications is that, ACE2 depletion in SARS-CoV-2-infected cells causes DABK accumulation in the extracellular environment of infected and neighboring bystander cells, which triggers a vicious positive feedback loop of inflammation and injury leading to even greater levels of DABK- and BK-mediated inflammation and injury (Panel E). DABK not only binds strongly to B1Rs, through which it exerts downstream effects, but also binds weakly to B2Rs in certain tissues, and exerts effects that are blocked by the B2R blocker, icatibant^{12,13} (Panel E). Since there are currently no FDA-approved drugs that selectively block DABK signaling through B1Rs, we provide a testable prediction that, off-label use of FDA-approved icatibant, will at least partially interrupt the positive feedback loop of DABK- and BK-mediated inflammation → injury → inflammation, and improve clinical outcomes in patients with COVID-19 respiratory complications (Panels E-H). Bidirectional arrows suggest that, these processes are likely to aggravate each other and be part of smaller positive feedback loops

to B1Rs, but also binds weakly to the BK-B2-receptor (B2R) in certain tissues, and exerts downstream effects that are blocked by the B2R blocker, icatibant.^{12,13}

Therefore, the off-label use of the B2R blocker, icatibant, seems promising for patients with unremitting respiratory distress caused by COVID-19. Icatibant (Trade Name: FIRAZYR; Takeda, Tokyo, Japan) is a drug that has been approved by the United States Food and Drug Administration (FDA) and other regulatory bodies, for the treatment of angioedema episodes in patients (18 years and older) with hereditary angioedema (HAE).¹⁴ Icatibant is thought to work by binding to B2Rs and blocking the downstream activity of BK in a variety of cells, including those present in blood vessels and the airway.¹⁵ Icatibant is effective in treating breathing difficulty in patients presenting with angioedema, including angioedema caused by angiotensin converting enzyme (ACE) inhibitors taken for hypertension.¹⁶ It might be purely coincidental that COVID-19 causes a “dry cough”¹⁷—a rare but characteristic side effect of ACE inhibitors, which is linked to BK.¹⁸

Icatibant has been shown to be safe and effective, with side effects and adverse reactions being rare when used in the context of angioedema.¹⁹ A human study on the off-label use of icatibant to treat allergic rhinitis showed that, the drug significantly reduced grass pollen antigen-induced hyperresponsiveness to histamine, which was linked to icatibant inhibiting interleukin-8 (IL-8) release.²⁰ The fact that IL-8 is implicated in acute lung injury and respiratory distress, further supports the empirical use of icatibant in the treatment of unremitting respiratory distress in COVID-19.²¹

From a scientific standpoint, analyzing plasma levels of BK and DABK in patients with respiratory complications from COVID-19, might help support our hypothesis.^{22,23} It might also be useful to retroactively obtain data on patients who have been treated recently with icatibant for angioedema, while having COVID-19 as a comorbidity, to ascertain whether or not COVID-19 respiratory symptoms decreased after icatibant

administration. In addition, it would be worth closely monitoring outcomes in patients with COVID-19 who take ACE inhibitors (for hypertension), dipeptidyl peptidase-4 (DPP4) inhibitors (for diabetes mellitus), or neprilysin inhibitors (for heart failure), since these drugs are known to interfere with BK breakdown and thus increase the BK bioavailability.²⁴

It is possible that molecules other than icatibant, which act on BK signaling pathways, might also be able to reduce the respiratory distress in COVID-19. For example, blocking DABK's main target, B1R, might produce better outcomes.²⁵ However, at this time, B1R blockers (eg, orally-active BI-113823) have only been tested in animals and in limited human trials.²⁶⁻²⁸ Nonetheless, B2R blockade has been shown to be effective in the context of airway hyperresponsiveness and respiratory distress in animal models; and, DABK has been shown to act on the B2R in some tissues.^{12,13,25} Inhibiting BK production with the FDA-approved drug, ecallantide, also seems promising, although it carries a risk of anaphylaxis in some patients.²⁹ Increasing plasma levels of aminopeptidase-P (APP), an enzyme that degrades BK and DABK, could also be tested as a benign intervention aimed at accelerating BK and DABK degradation.³⁰

We speculate that dysregulated BK signaling might even explain some of the perplexing observations on COVID-19. Emerging data suggest that in the United States of America, morbidity and mortality among African Americans has been disproportionately higher compared to other ethnic groups.³¹⁻³³ It is possible that African Americans are more affected due to their increased sensitivity to BK, a greater susceptibility to ACE-inhibitor-induced angioedema, and a polymorphism (XPNPEP2 C-2399A) linked to ACE-inhibitor-induced angioedema in African American males.³⁴⁻³⁷ In addition, data suggest that respiratory complications are more often seen in males than females.³⁸⁻⁴¹ Interestingly, APP activity has been reported to be higher in females irrespective of the XPNPEP2 C-2399A polymorphism.³⁷ Furthermore, vasopressors are

required to stabilize some patients with COVID-19 critical illness^{17,38}, and, it is well-known that BK elevation reduces blood pressure.⁴²⁻⁴⁴ A loss of smell and/or taste has been reported by some patients with COVID-19⁴⁵⁻⁴⁷; and, a loss of smell has been reported in patients with HAE and in persons who take ACE inhibitors.^{48,49} There are anecdotes of individuals experiencing extreme thirst while having COVID-19^{50,51}; and, elevated BK and ACE inhibitors are associated with increased thirst.⁵² However, confounding factors might exist, and, therefore, these observations will need to be evaluated more objectively.

In summary, established and emerging evidence on SARS-CoV, ACE2, BK and DABK signaling, angioedema, and respiratory distress, has helped us develop a testable hypothesis, which may link dysregulated BK signaling to COVID-19 respiratory complications. There is a critical need to develop basic and clinical studies to test this hypothesis, since there are approved drugs that might be effective in interrupting the vicious feedback loop that might exist between dysregulated BK signaling and tissue injury.

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CONFLICT OF INTEREST

The authors have no conflicts to declare.

AUTHOR CONTRIBUTIONS

J.A. Roche and R. Roche were involved in reviewing the literature and writing this manuscript.

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