Pathways for Bradykinin Formation and Interrelationship with Complement as a Cause of Edematous Lung in COVID-19 Patients

Allen P. Kaplan, M.D., Berhane Ghebrehiwet, D.V.M., D.Sc.

PII: S0091-6749(20)31502-5

DOI: https://doi.org/10.1016/j.jaci.2020.10.025

Reference: YMAI 14815

To appear in: Journal of Allergy and Clinical Immunology

Received Date: 22 September 2020

Revised Date: 15 October 2020

Accepted Date: 16 October 2020

Please cite this article as: Kaplan AP, Ghebrehiwet B, Pathways for Bradykinin Formation and Interrelationship with Complement as a Cause of Edematous Lung in COVID-19 Patients, *Journal of Allergy and Clinical Immunology* (2020), doi: https://doi.org/10.1016/j.jaci.2020.10.025.

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1	Pathways for Bradykinin Formation and Interrelationship with Complement as a Cause of
2	Edematous Lung in COVID-19 Patients
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4	Allen P. Kaplan, M.D. and Berhane Ghebrehiwet, D.V.M., D.Sc.
5	
6	From the Pulmonary and Critical Care Division, The Medical University of South Carolina,
7	Charleston, SC and The Division of Rheumatology, Allergy, and Clinical Immunology, SUNY
8	Stony Brook
9	
10	Corresponding Author:
11 12 13 14 15 16 17 18 19	Dr. Allen P. Kaplan, MD Medical University of South Carolina Dept of Med., Pulmonary 96 Jonathan Lucas St. Suite 812-CSB` Charleston, SC. 29425-2220 Primary Phone Number: 843-722-1253 E-mail Address: kaplana@musc.edu
20	Conflict of Interest: I receive no compensation for writing the editorial, Dr. Ghebrehiwet and I
21	are the sole authors, Dr. Kaplan has not ties to pharmaceutical companies who have products
22	related to this subject, and Dr. Ghebrehiwet receives royalties from the sale of detection kit for
23	gC1qR and monoclonal antibodies 60.11 and 74.5.2.
24	
25	Key Words: Bradykinin
26	Kallikrein
27	Edema

COVID 28

29	Abbreviations:	gC1qR – Receptor for the globular heads of C1q
30		u-PAR – Urokinase plasminogen activator receptor

- CP-N Carboxypeptidase N 31
- 32

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33 The beta coronavirus SARS-CoV-2 infects lung alveolar type II epithelial calls by attaching to 34 angiotensin converting enzyme 2 (ACE-2) expressed at the cells surface via its viral spike 35 protein(1). A transmembrane serum protease (TMPRSS2) activates the viral spike protein and 36 enables cell entry. The more severe manifestations of the resultant inflammatory response 37 include dry cough, dyspnea, tachypnea, a feeling of drowning, pulmonary edema, unilateral or bilateral pneumonia, mottling and ground glass opacifies on CT scan, and progression to the 38 39 Acute Respiratory Distress Syndrome (ARDS) requiring ventilatory support(2). Hypoxemia is 40 particularly prominent throughout and a hyaline membrane of dead cells can be observed at 41 autopsy. Once infection takes hold, a cascade of inflammatory events is initiated including the 42 release of cytokines such as I-1, IL-6, IP-10, MCP-1, TNFa(3) (and many more) which has been referred to as a "cytokine storm". In addition, the prominent edema seen throughout the lung and 43 44 the association of ACE inhibition with severe angioedema has focused attention on another 45 innate inflammatory cascade; namely, the overproduction of bradykinin(3) which is the focus of 46 this editorial.

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There are two general pathways for the production of bradykinin, the first being the release of cellular tissue kallikrein which cleaves low molecular weight kininogen (LK) to release lysbradykinin (Fig. 1). Tissue kallikrein is secreted as an active enzyme (i.e. processed intracellularly) and is a particularly prominent product of the lung, pancreas, kidney, salivary glands, and the prostate. There are 15 homologous gene products, three of which can produce bradykinin (KLK 1, 2, and 12), KLK1 being the most prominent.

55 The second pathway is present in plasma and consists of Factor XII, plasma prekallikrein (PK), 56 and high-molecular weight kininogen (HK)(4). Prekallikrein circulates primarily as a 57 bimolecular complex with HK (about 75-80% bound) as does coagulation factor XI (95% is 58 bound). They compete for a single overlapping binding site but there is sufficient HK present to 59 bind both. Both factor XII and prekallikrein possess minute levels of proteolytic activity relative to their respective active enzymes which may be the initial spark needed for activation to 60 61 proceed. All three proteins are also bound to bimolecular sites on the surface of endothelial cells 62 (Figure 1). Factor XII binds primarily to u-PAR-cytokeratin 1 (CK-1) while HK binds to 63 gC1qR-cytokeratin 1 with PK attached to the HK (Fig. 1). Once activation proceeds, Factor XII 64 is converted to two forms of the activated enzyme, factor XIIa (80 Kd) and factor XIIf (28.5-30 Kd; β FXIIa). Both can convert prekallikrein to kallikrein and kallikrein digests HK to release 65 66 bradykinin (Arg-pro-pro-gly-phe-ser-pro-phe-arg). Factor XII activation proceeds by a relatively 67 slow autoactivation process to produce a small amount of factor XIIa and a very rapid positive feedback in which the initial kallikrein formed activates all remaining factor XII in seconds to 68 69 yield factor XIIa and then factor XIIf. Tissue kallikrein does not activate factor XII. The larger 70 80 Kd factor XIIa is the clotting factor that converts factor XI to factor XIa to continue the 71 intrinsic coagulation pathway (Fig. 1). Factor XIIf, lacks a surface binding site, loses 96-98% of 72 the clotting activity, but gains a new function i.e. activation of C1r to initiate the classical 73 complement cascade (Fig. 1). This is not surprising since cross-activation by enzymes of the 74 complement system and the coagulation pathway proteins is known to occur when either 75 pathway is activated. For example, plasmin, Factor Xa and Factor XIa are able to cleave C3 and 76 C5 to generate the potent chemoattractants C3a and C5a which, in turn are able to recruit and 77 activate leukocytes to produce pro-inflammatory cytokines. This contributes to the cytokine

storm that is the hallmark of many inflammatory processes including those induced by SARSCoV-2.

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81 Bradykinin causes vasodilation and increases vascular permeability by interacting with 82 constitutively expressed B-2 receptors on small venules. The same is true of lys-bradykinin 83 produced by tissue kallikrein (Fig. 1) although the lys is rapidly removed by aminopeptidase P. 84 Bradykinin is degraded primarily by ACE, a dipeptidase which removes the C-terminal phe-arg, 85 which inactivates it, followed by removal of ser-pro. An alternative process requires carboxypeptidase activity (carboxypeptidase N in plasma and carboxypeptidase M on pulmonary 86 vascular endothelial cells) to first remove the C-terminal arg from either bradykinin (plasma 87 cascade) or lys-bradykinin (tissue kallikrein product) (Fig. 1). This leaves des-arg⁹ bradykinin 88 (Arg-pro-pro-gly-phe-ser-pro-phe) which is minimally reactive with B-2. However this peptide 89 90 binds to the B-1 receptor which also mediates vasodilation and vascular permeability. The B-1 receptor is not normally present but is induced by IL-1 or TNFa (produced by febrile viral 91 illnesses such as COVID-19) as well as gC1qR. It's ligands are des-arg⁹ bradykinin(3) as well as 92 des-arg⁹ lys-bradykinin (Fig. 1). 93

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There are many observations and theories regarding a prominent role for bradykinin and perhaps des-arg⁹ bradykinin in the pathogenesis of the pulmonary dysfunction of COVID-19 which is linked in part to changes in the renin-angiotensin system (RAS). Studies of gene expression in bronchoalveolar lavage specimens of COVID-19 patients(5), when compared to normal control specimens, reveal upregulation of multiple components that lead to bradykinin production and downregulation of factors that control the process. All "kallikreins and kininogens" are

101	upregulated, the B-2 receptor was increased 207-fold and the B-1 receptor, 2945-fold. The gene
102	expression for C1-INH was decreased 33-fold which would render the plasma bradykinin
103	cascade labile and overreactive as we see in C1-INH deficiency (types I and II HAE) in which
104	enzymes not adequately inhibited by C1-INH include both forms of activated factor XII, plasma
105	kallikrein, and C1r. By contrast, gene expression for ACE was decreased 8-fold so that
106	bradykinin would not be inactivated normally. While viral binding to ACE-2 limits its
107	enzymatic activity(3) so that des-arg ⁹ bradykinin is not degraded (ACE-2 removes C-terminal
108	phe) and lowered ACE levels also limit des-arg ⁹ bradykinin degradation (it removes C-terminal
109	ser-pro-phe acting then as a tripeptidase rather than a dipeptidase). With the markedly
110	augmented bradykinin receptor production, a "bradykinin storm" can result.
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112	Our own preliminary observations (unpublished) reveal upregulation and secretion of gC1qR by
113	infected cells which creates the cell surface platform for activation of the bradykinin cascade and
114	secreted gC1qR also upregulates the B-1 receptor(6). The Renin-Angiotensin system(7) can also
115	be contributory in that decreased ACE limits formation of the vasoconstrictor angiotensin II from
116	angiotensin I. As angiotensin I accumulates, ACE-2 removes C-terminal phe to produce
117	angiotensin 1-9. This moiety stimulates angiotensin-2 receptors to cause vasodilation and can do
118	so synergistically with bradykinin(5). If significant amounts of angiotensin II were produced,
119	ACE-2 can then convert it to another vasodilator, angiotensin 1-7 active through the MAS
120	receptor(7). Here, the balance of decreased ACE-2 via viral binding and internalization(3) and
121	increased ACE-2, as seen when COVID-2 BAL fluids are examined(5), needs to be quantified at
122	the protein level (cell surface and interstitial fluid) rather than the DNA level to determine the net
123	enzymatic effect.

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125	There are numerous reports of a possible therapeutic role for antagonists of cytokines such as IL-
126	1 (Anakinra) or IL-6 (Tocilizumab) to treat COVID-19(2, 7, 8). We suggest use of lanadelumab
127	to block plasma kallikrein(9) and Icatibant to inhibit B-2 receptors as possible therapy for the
128	severe pulmonary manifestations of COVID-19. Preliminary observations employing Icatibant
129	(uncontrolled) indicate improved oxygenation. Antagonists of tissue kallikrein and the B-1
130	receptors have been employed for research purposes but are not approved for clinical use, but
131	there is a need for such agents. Simultaneous inhibition of B-1 and B-2 receptors is also
132	possible. Finally, a monoclonal antibody to gC1qR to disrupt bradykinin formation along
133	endothelial cell surfaces would be a novel, additional approach(10).
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164	<u>Fig. 1</u>
165	A diagram of important interrelationships involving the bradykinin-forming pathways in patients
166	with COVID-19. Pulmonary epithelial cells release tissue kallikrein which cleaves LK to release
167	lys-bradykinin that is rapidly converted to bradykinin. Both are ligands of the B-2 receptor. The
168	plasma cascade is recruited into the surrounding lung tissue and all the requisite proteins for
169	bradykinin formation exist bound to endothelial cells where activation can proceed along the

170 surface as well as in the fluid phase. In addition, activation of endothelial cells by IL-1 or $TNF\alpha$

171 can release HSP-90 and prolylcarboxypeptidase. Both convert PK to plasma kallikrein <u>if</u> PK is

- bound to HK. Formation of bradykinin and its des-arg⁹ degradation products stimulate B2 and
- 173 B1 receptors respectively, leading to lung edema.
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