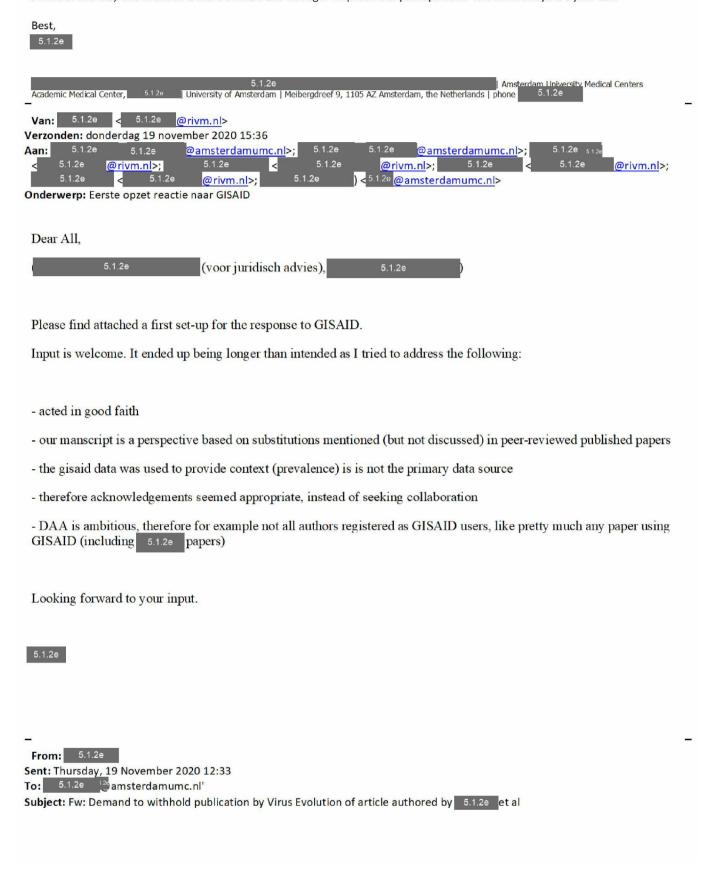


Hi 5.1.2e and all, I've made some edits to the latest version of the letter. Please see attached. As always, it's your call. I'm only trying to help. All the best, Department of Medical Microbiology | Amsterdam University Medical Center, University of Amsterdam From: 5.1.2e < 5.1.2e @rivm.nl> Date: Friday, 20 November 2020 at 11:37 To: " 5.1.2e 5.1.2e @amsterdamumc.nl>, " @amsterdamumc.nl>, 5.1.2e 5.1.2e 5 1 2e 5.1.2e 5.1.2e @rivm.nl>, 5.1.2e < @rivm.nl>, @rivm.nl>, @rivm.nl>, < 5.1.2e @amsterdamumc.nl> Subject: RE: Eerste opzet reactie naar GISAID 5.1.2e Wederom veel dank voor jouw zeer waardevol inzichten suggesties. En nieuwe opzet nav jouw suggesties (helft zo lang) Meer een open brief nu ipv een rebuttal * From: 5.1.2e < 5.1.2e @amsterdamumc.nl> Sent: donderdag 19 november 2020 19:04 5.1.2€ @rivm.nl>; @amsterdamumc.nl>; 5.1.2e @rivm.nl>; To: 5.1.2e @rivm.nl>; @rivm.nl>;) < 5.1.2e @amsterdamumc.nl> @rivm.nl>; Subject: Re: Eerste opzet reactie naar GISAID Hi 5.1.2e and all,

I believe the response is too defensive, too (scientifically) explanatory and (much) too long. Personally I would respond more or less along the same lines and length as the response to the journal, e.g.:

- start by acknowledging the obvious importance of GISAID as an open science platform and the importance of trust and rules to keep this afloat, and emphasize that your insitute contributes substantially to the success of GSAID, both directly by depositing human flu and Sars-cov-2 sequences as well as indirectly by facilitating sequence analyses by collaborating institutes (incl EMC) through sample collections within the Dutch public health network of GGDs.
- followed by part of the first paragraph of the previous response (We should emphasize that we have been acting in good faith, with no intented wrongdoing, in our research and the writing of our perspective. In this spirit we have openly communicated our findings in a national public health meeting attended by members of the Dutch 'mink consortium' in early October and more recently in a meeting attended by 5.1.2e the latter of which seems to have ignited the current predicament.) Add the bit about your failed attempts to preprint.
- then the ambiguity part of the previous response, ending with that you're looking forward to engage with GISAIDS to understand the exact rules of use and relieve the ambiguity for future reference.
- and then explain your solution for the current situation in order to not further delay sharing your important insights with the scientific community, particluarly in relation to the conclusions of Koopman's recent Science paper, i.e. that observed mutations might be associated with mink-adaptation and have possible effects on the binding of neutralizing antibodies.

I think in this way this could be a more concise and stronger response but perhaps other feel differently.. It's your call!



From:	5.1.5	<	5.1.5	@gisaid.org>

Sent: Monday, 16 November 2020 02:53

Cc: Legal at GISAID

Subject: Demand to withhold publication by Virus Evolution of article authored by 5.1.2e et al

Please see the attached letter from the GISAID Secretariat.

Kindly note that this matter is urgent. We ask that you respond as soon as practicable, and no later than by 20th November 2020.

a Public-Private-Partnership with the Federal Republic of Germany Lipowskystraße 10 | 81373 Munich | Germany District Court Munich Reg: VR204844 GISAID, from Vision to Reality

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