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A simulated single ventilator/dual patient ventilation strategy for acute respiratory distress syndrome during the COVID-19 pandemic

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Article citation details

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Review timeline

Original submission: 6 April 2020
Revised submission: 12 August 2020
Final acceptance: 12 August 2020

Note: Reports are unedited and appear as submitted by the referee. The review history

appears in chronological order.

Review History

RSOS-200585.R0 (Original submission)

Review form: Reviewer 1

Is the manuscript scientifically sound in its present form? Yes

Are the interpretations and conclusions justified by the results? Yes

Is the language acceptable?

Yes

Do you have any ethical concerns with this paper?

Have you any concerns about statistical analyses in this paper? No

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Recommendation?

Accept with minor revision (please list in comments)

Comments to the Author(s)

The authors describe a numerical model of ventilation of two patients using a single ventilator with modification to account for potential differences in lung compliance between the two patients. The model uses an electrical analogue approach to model the system. A standard splitter and modified splitter configuration are examined and compared. The modified splitter includes variable resistors and check valves to allow targeting of specific tidal volumes during mechanical ventilation of two patients with different lung compliance.

The manuscript is generally clearly written with appropriate figures provided. Some details are missing and some aspects of the work are not discussed in the current version. The following issues should be addressed:

The current approach does not allow either respiratory rate or I:E ratio to be set independently for each paper, the authors should comment on whether this presents any issues.

Poiseuille assumptions have been used to provide resistance values. What is Re value at the flow rates and for the dimensions of tubing in question? Is it reasonable to apply Poiseuille eqn. To determine these values, i.e. laminar conditions?

The patient model assumes no change in resistance in the presence of disease, the authors should comment on the appropriateness of this assumption.

There is no description in figure 1 or 2 or the methods of the role of the sensors in the Simullink model and how these are used, e.g. is this purely to provide output data for monitoring, or is this data also used to inform operation of the model at future time steps (e.g. feedback processes)?

Do the authors expect any variations to be observed if considering B-C, B-D, C-D configurations of patients which are not simulated in the paper currently?

No details of the solution timestep used for the numerical scheme are given? How does this relate to the time step used to post-process output data, which appears to be relatively coarse from the plots (e.g. Figure 6) where outputs have markers at intervals of ~ 0.3 seconds?

Several references refer to news articles and other documents and appear to be rather sparsely referenced, specific dates are not provided and it would be helpful to provide URLs to avoid ambiguity in locating these documents.

Review form: Reviewer 2 (Aaron B. Morton)

Is the manuscript scientifically sound in its present form? Yes

Are the interpretations and conclusions justified by the results? Yes

Is the language acceptable?

Do you have any ethical concerns with this paper?

No

Have you any concerns about statistical analyses in this paper? $N_{\rm O}$

Recommendation?

Accept as is

Comments to the Author(s)

Summary

The study entitled "A Simulated Single Ventilator / Dual Patient Ventilation Strategy for Acute Respiratory Distress Syndrome During the COVID-19 Pandemic" by Solis-Lemus and colleague addresses a critical shortcoming in former modeling investigations of multi patient ventilation. In particular, former modeling of multi patient ventilation assume equal lung physiology between all patients and, therefore, are likely irrelevant in a clinical setting. Indeed, mechanical ventilation (MV) use is broad, encompassing spinal cord injury, drug overdose, heart failure, and acute respiratory distress syndrome (ARDS). Patients from many of these conditions are likely to present with co-morbidities as well, making the matching of lung physiology between patients especially difficult. With this in mind, the authors a) present two electronic circuits (analogous in theory to the closed circuit of a single ventilator and two independent patients), b) created two mathematical models (standard splitter and modified splitter), c) demonstrate and quantify the pitfalls of previous multi patient MV models, d) provide a simple solution to the issues (variable flow restrictors added into the inspiration arms to control pressure for each patient and one-way valves on the expiration arms to prevent backflow), and e) made this software code freely available to the public for further testing and experimentation. Applying this model to a more clinically relevant model of multi patient ventilation, that is, pairing patients with different lung compliances on a single ventilator, the authors demonstrate that tidal volume could differ between patients as much as 38%. The authors then demonstrate through the simple addition of variable flow restrictors in the inspiratory tubes and one-way valves in the expiratory tubes, that it may be possible to ventilate two patients with mismatched lung physiology on a single ventilator.

The paper is well written, and the logic is sound. Due to the high potential need of this work and the absence of major or minor issues, we recommend this paper for publication in Royal Society Open Science.

General Comments

We want to thank the authors for performing such an important work at this time. Also, we appreciate the analogous effort of utilizing electronic circuit theory to represent respiratory physiology and the open access of these mathematical models.

Major Comments

none

Minor Comments

None

Decision letter (RSOS-200585.R0)

We hope you are keeping well at this difficult and unusual time. We continue to value your support of the journal in these challenging circumstances. If Royal Society Open Science can assist you at all, please don't hesitate to let us know at the email address below.

Dear Dr Niederer

On behalf of the Editors, I am pleased to inform you that your Manuscript RSOS-200585 entitled "A Simulated Single Ventilator / Dual Patient Ventilation Strategy for Acute Respiratory Distress

Syndrome During the COVID-19 Pandemic" has been accepted for publication in Royal Society Open Science subject to minor revision in accordance with the referee suggestions. Please find the referees' comments at the end of this email.

The reviewers and handling editors have recommended publication, but also suggest some minor revisions to your manuscript. Therefore, I invite you to respond to the comments and revise your manuscript.

• Ethics statement

If your study uses humans or animals please include details of the ethical approval received, including the name of the committee that granted approval. For human studies please also detail whether informed consent was obtained. For field studies on animals please include details of all permissions, licences and/or approvals granted to carry out the fieldwork.

Data accessibility

It is a condition of publication that all supporting data are made available either as supplementary information or preferably in a suitable permanent repository. The data accessibility section should state where the article's supporting data can be accessed. This section should also include details, where possible of where to access other relevant research materials such as statistical tools, protocols, software etc can be accessed. If the data has been deposited in an external repository this section should list the database, accession number and link to the DOI for all data from the article that has been made publicly available. Data sets that have been deposited in an external repository and have a DOI should also be appropriately cited in the manuscript and included in the reference list.

If you wish to submit your supporting data or code to Dryad (http://datadryad.org/), or modify your current submission to dryad, please use the following link: http://datadryad.org/submit?journalID=RSOS&manu=RSOS-200585

• Competing interests

Please declare any financial or non-financial competing interests, or state that you have no competing interests.

• Authors' contributions

All submissions, other than those with a single author, must include an Authors' Contributions section which individually lists the specific contribution of each author. The list of Authors should meet all of the following criteria; 1) substantial contributions to conception and design, or acquisition of data, or analysis and interpretation of data; 2) drafting the article or revising it critically for important intellectual content; and 3) final approval of the version to be published.

All contributors who do not meet all of these criteria should be included in the acknowledgements.

We suggest the following format:

AB carried out the molecular lab work, participated in data analysis, carried out sequence alignments, participated in the design of the study and drafted the manuscript; CD carried out the statistical analyses; EF collected field data; GH conceived of the study, designed the study, coordinated the study and helped draft the manuscript. All authors gave final approval for publication.

• Acknowledgements

Please acknowledge anyone who contributed to the study but did not meet the authorship criteria.

• Funding statement

Please list the source of funding for each author.

Please ensure you have prepared your revision in accordance with the guidance at https://royalsociety.org/journals/authors/author-guidelines/ -- please note that we cannot publish your manuscript without the end statements. We have included a screenshot example of the end statements for reference. If you feel that a given heading is not relevant to your paper, please nevertheless include the heading and explicitly state that it is not relevant to your work.

Because the schedule for publication is very tight, it is a condition of publication that you submit the revised version of your manuscript before 07-Aug-2020. Please note that the revision deadline will expire at 00.00am on this date. If you do not think you will be able to meet this date please let me know immediately.

To revise your manuscript, log into https://mc.manuscriptcentral.com/rsos and enter your Author Centre, where you will find your manuscript title listed under "Manuscripts with Decisions". Under "Actions," click on "Create a Revision." You will be unable to make your revisions on the originally submitted version of the manuscript. Instead, revise your manuscript and upload a new version through your Author Centre.

When submitting your revised manuscript, you will be able to respond to the comments made by the referees and upload a file "Response to Referees" in "Section 6 - File Upload". You can use this to document any changes you make to the original manuscript. In order to expedite the processing of the revised manuscript, please be as specific as possible in your response to the referees. We strongly recommend uploading two versions of your revised manuscript:

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- 2) A 'clean' version of the new manuscript that incorporates the changes made, but does not highlight them.

When uploading your revised files please make sure that you have:

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- 2) A separate electronic file of each figure (EPS or print-quality PDF preferred (either format should be produced directly from original creation package), or original software format);
- 3) Included a 100 word media summary of your paper when requested at submission. Please ensure you have entered correct contact details (email, institution and telephone) in your user account;
- 4) Included the raw data to support the claims made in your paper. You can either include your data as electronic supplementary material or upload to a repository and include the relevant doi within your manuscript. Make sure it is clear in your data accessibility statement how the data can be accessed;
- 5) All supplementary materials accompanying an accepted article will be treated as in their final form. Note that the Royal Society will neither edit nor typeset supplementary material and it will be hosted as provided. Please ensure that the supplementary material includes the paper details where possible (authors, article title, journal name).

Finally, please ensure that email addresses for all co-authors are up-to-date. At present, the following email address appears to be incorrect:

d.taylor@imperial.ac.uk

Supplementary files will be published alongside the paper on the journal website and posted on the online figshare repository (https://rs.figshare.com/). The heading and legend provided for each supplementary file during the submission process will be used to create the figshare page, so please ensure these are accurate and informative so that your files can be found in searches.

Files on figshare will be made available approximately one week before the accompanying article so that the supplementary material can be attributed a unique DOI.

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Once again, thank you for submitting your manuscript to Royal Society Open Science and I look forward to receiving your revision. If you have any questions at all, please do not hesitate to get in touch.

Kind regards, Lianne Parkhouse Editorial Coordinator Royal Society Open Science openscience@royalsociety.org

on behalf of the Associate Editor and Professor R. Kerry Rowe (Subject Editor) openscience@royalsociety.org

Associate Editor Comments to Author:

Thank you for submitting your research to Royal Society Open Science. Firstly, we apologise for the delays incurred during the initial peer-review process. We have now received the reviewer reports and we would be glad to accept your manuscript subject to the minor revisions requested by the referees. In your revised submission, please ensure to provide a point-by-point response to their comments, and also provide a tracked changes version of your paper to highlight the changes made. Lastly, we ask that you please archive your GitHub code within the Zenodo repository: https://guides.github.com/activities/citable-code/. Please then ensure that the Zenodo DOI is added to the data accessibility statement in addition to the GitHub URL

We look forward to receiving your revised submission.

Reviewer comments to Author:

Reviewer: 1 Comments to the Author(s)

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Do the authors expect any variations to be observed if considering B-C, B-D, C-D configurations of patients which are not simulated in the paper currently?

No details of the solution timestep used for the numerical scheme are given? How does this relate to the time step used to post-process output data, which appears to be relatively coarse from the plots (e.g. Figure 6) where outputs have markers at intervals of ~ 0.3 seconds?

Several references refer to news articles and other documents and appear to be rather sparsely referenced, specific dates are not provided and it would be helpful to provide URLs to avoid ambiguity in locating these documents.

Reviewer: 2 Comments to the Author(s)

Summary

The study entitled "A Simulated Single Ventilator / Dual Patient Ventilation Strategy for Acute Respiratory Distress Syndrome During the COVID-19 Pandemic" by Solis-Lemus and colleague addresses a critical shortcoming in former modeling investigations of multi patient ventilation. In particular, former modeling of multi patient ventilation assume equal lung physiology between all patients and, therefore, are likely irrelevant in a clinical setting. Indeed, mechanical ventilation (MV) use is broad, encompassing spinal cord injury, drug overdose, heart failure, and acute respiratory distress syndrome (ARDS). Patients from many of these conditions are likely to present with co-morbidities as well, making the matching of lung physiology between patients especially difficult. With this in mind, the authors a) present two electronic circuits (analogous in theory to the closed circuit of a single ventilator and two independent patients), b) created two mathematical models (standard splitter and modified splitter), c) demonstrate and quantify the pitfalls of previous multi patient MV models, d) provide a simple solution to the issues (variable flow restrictors added into the inspiration arms to control pressure for each patient and one-way valves on the expiration arms to prevent backflow), and e) made this software code freely available to the public for further testing and experimentation. Applying this model to a more clinically relevant model of multi patient ventilation, that is, pairing patients with different lung compliances on a single ventilator, the authors demonstrate that tidal volume could differ between patients as much as 38%. The authors then demonstrate through the simple addition of variable flow restrictors in the inspiratory tubes and one-way valves in the expiratory tubes, that it may be possible to ventilate two patients with mismatched lung physiology on a single ventilator.

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General Comments

We want to thank the authors for performing such an important work at this time. Also, we appreciate the analogous effort of utilizing electronic circuit theory to represent respiratory physiology and the open access of these mathematical models.

Major Comments

none

Minor Comments

None

Author's Response to Decision Letter for (RSOS-200585.R0)

See Appendix A.

Decision letter (RSOS-200585.R1)

We hope you are keeping well at this difficult and unusual time. We continue to value your support of the journal in these challenging circumstances. If Royal Society Open Science can assist you at all, please don't hesitate to let us know at the email address below.

Dear Dr Niederer,

It is a pleasure to accept your manuscript entitled "A Simulated Single Ventilator / Dual Patient Ventilation Strategy for Acute Respiratory Distress Syndrome During the COVID-19 Pandemic" in its current form for publication in Royal Society Open Science.

Please ensure that an alternative email address is supplied for Dr Taylor, as d.taylor@imperial.ac.uk is not accepting messages from the Royal Society.

COVID-19 rapid publication process: We are taking steps to expedite the publication of research relevant to the pandemic. If you wish, you can opt to have your paper published as soon as it is ready, rather than waiting for it to be published the scheduled Wednesday.

This means your paper will not be included in the weekly media round-up which the Society sends to journalists ahead of publication. However, it will appear in the COVID-19 Publishing Collection which journalists will be directed to each week (https://royalsocietypublishing.org/topic/special-collections/novel-coronavirus-outbreak)

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Thank you for your fine contribution. On behalf of the Editors of Royal Society Open Science, we look forward to your continued contributions to the Journal.

Kind regards, Andrew Dunn Royal Society Open Science Editorial Office Royal Society Open Science openscience@royalsociety.org

on behalf of Prof R. Kerry Rowe (Subject Editor) openscience@royalsociety.org

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Appendix A

We would like to thank the reviewers and editors for their positive comments and constructive comments. We have addressed each of these in the document below. We have included the initial editor and reviewer comments in black. Responses are in blue and any changes to the text are in in italics and indented.

Associate Editor Comments to Author:

Thank you for submitting your research to Royal Society Open Science. Firstly, we apologise for the delays incurred during the initial peer-review process. We have now received the reviewer reports and we would be glad to accept your manuscript subject to the minor revisions requested by the referees. In your revised submission, please ensure to provide a point-by-point response to their comments, and also provide a tracked changes version of your paper to highlight the changes made. Lastly, we ask that you please archive your GitHub code within the Zenodo repository: https://guides.github.com/activities/citable-code/. Please then ensure that the Zenodo DOI is added to the data accessibility statement in addition to the GitHub URL

We have now updated our Github repository and created a version 0.2 of the code that we have uploaded onto Zenodo and we have updated the reference to the Zenodo repository in the text.

Reviewer: 1

The authors describe a numerical model of ventilation of two patients using a single ventilator with modification to account for potential differences in lung compliance between the two patients. The model uses an electrical analogue approach to model the system. A standard splitter and modified splitter configuration are examined and compared. The modified splitter includes variable resistors and check valves to allow targeting of specific tidal volumes during mechanical ventilation of two patients with different lung compliance.

The manuscript is generally clearly written with appropriate figures provided. Some details are missing and some aspects of the work are not discussed in the current version. The following issues should be addressed:

1. The current approach does not allow either respiratory rate or I:E ratio to be set independently for each paper, the authors should comment on whether this presents any issues.

We thank the Reviewer for highlighting this important point. Indeed, if two patients are to be ventilated from a single ventilator, then by necessity the I:E ratio and the respiratory rate must be identical for both patients. In the context of Acute Respiratory Distress Syndrome, I:E ratios may often be reduced from normal and variations in respiratory rate may need to be considered. However, when designing this study we felt that one reasonable approach could involve cohorting patients into groups based on body weight, disease severity and required I:E ratio/respiratory rate. Owing to this requirement for cohorting, the factor by which split ventilation would increase ventilatory capacity is likely to be less than 2. Ongoing research in our group is seeking to retrospectively analyse ventilatory requirements amongst Covid patients in order to estimate this factor; but the results will not be available until later in the year. In the meantime we have added the following paragraph to the manuscript explaining this limitation:

Given that the I:E ratio and respiratory rate would need to be identical for both patients, it will likely be necessary to cohort patients based on body weight, lung disease severity and required ventilatory parameters. Such cohorting is likely to reduce the factor by which split

ventilation could increase ventilatory capacity and would need to be considered in the design of clinical protocols seeking to implement this strategy.

2. Poiseuille assumptions have been used to provide resistance values. What is Re value at the flow rates and for the dimensions of tubing in question? Is it reasonable to apply Poiseuille eqn. To determine these values, i.e. laminar conditions?

Poiseulle flow is indeed an approximation, and at higher flow rates it is possible that flow is actually turbulent, especially given the fact tubing is often ribbed. Future work should consider this, and/or the inclusion of non-linear e.g. quadratic resistances. However, even with these modifications, tubing resistances are still likely to be orders of magnitude smaller than the resistances of the other components in the circuit, and hence should not change the over nature of the results. We have added a section to the new Limitations paragraph outlining the above considerations:

Finally, the assumption of Poiseulle flow within the ventilator tubing is an approximation, and at higher flow rates it is possible that flow becomes turbulent, especially given that the tubing is often ribbed. Future work should consider this, and/or the inclusion of non-linear e.g. quadratic resistances.

3. The patient model assumes no change in resistance in the presence of disease, the authors should comment on the appropriateness of this assumption.

We again thank the Reviewer for raising this important point. Increase in respiratory resistance may occur owing to patient factors or ventilatory circuit factors. In ARDS, inspiratory resistance is only slightly higher than in normal subjects and a significant increase in respiratory resistance may suggest ventilatory circuit factors (e.g. a blocked endotracheal tube). Whilst we have not simulated such a scenario in the present manuscript; the model as provided does allow for analysis of this situation which would be simulated by increasing the values of Rettl and Rettl. The model as provided also allows for analysis of the effects of increased small airways resistance which would be simulated by increasing the value R_{LX} in the patient submodules. As the model is currently implemented it does not, however, allow for the scenario where inspiratory and expiratory resistance are different, which may occur in ARDS. However, future development of the proposed model could include the addition of two resistances and two diodes within the patient module which would allow this more complicated scenario to be studied. Rather than conduct extensive simulations of every such scenario we preferred in the present manuscript to describe the model and publish the source code to make the model available to the research community. We have added a section to the new Limitations paragraph outlining the above considerations:

... we have assumed in this model that small airways resistance remains constant under disease conditions. This is a simplifying assumption. Whilst inspiratory resistance is only minimally increased in ARDS compared to normal subjects, increased expiration resistance has been observed. Further model development using flow directional resistances in the patient model would be required to capture this complexity.

4. There is no description in figure 1 or 2 or the methods of the role of the sensors in the Simullink model and how these are used, e.g. is this purely to provide output data for monitoring, or is this data also used to inform operation of the model at future time steps (e.g. feedback processes)?

In the manuscript, "sensors" are Simscape blocks used to denote which signals should be logged for analysis, plotting, etc. These do not represent actual sensors or model any physical aspects of real sensors. We have omitted these blocks in the figures to make for a less cluttered figure. In a clinical setting we would, of course, expect appropriate pressure and flow sensors to be in place for monitoring of the patient, which should then be used by the clinician to decide whether any settings should be adjusted as by the clinician. We have added the following text to the caption of Figure 1 to clarify what is meant with "sensor":

The text "sensor" on the patient blocks indicate that the respective signals are connected to appropriate Simscape sensor blocks, namely current and voltage sensors, to allow for these signals to be logged during simulation.

5. Do the authors expect any variations to be observed if considering B-C, B-D, C-D configurations of patients which are not simulated in the paper currently?

We ran the model across a wide range of combinations, including those suggested above. In all cases trends were similar to those presented for the combinations analysed in the initial manuscript. We have now included these results in Table 4 for completeness and noted that they give the same results as the healthy – ARDS pairing simulations.

6. No details of the solution timestep used for the numerical scheme are given? How does this relate to the time step used to post-process output data, which appears to be relatively coarse from the plots (e.g. Figure 6) where outputs have markers at intervals of ~ 0.3 seconds?

After extensive analysis, we were satisfied with the default Simulink and Simscape solver settings. We therefore decided to use the default settings for the presentation of the results and release of the code on GitHub. The key default solver selection parameters are: "Variable-step" and "Automatic solver selection". The following relevant settings are set to "auto" by default: max step size, min step size, initial step size, and absolute tolerance. The relative tolerance is set by default to 10^{-3} . The algorithm within Simulink ended up choosing the solver "ode45" with max step size 0.3. The time steps used to post-process data are the same as those generated by the solver. Since we have provided all files via GitHub, the readers will be free to change the max step size or other settings, should they wish to plot the data at a finer scale. We have added the following text to the Implementation section:

The default Simulink and Simscape settings were used to generate the data, namely automatic selection of solver and step sizes with a relative error tolerance of 10^{-3} . Simulink/Simscape was able to reduce the model to an ODE and chose to solve the ODE with the Dormand-Prince explicit, variable-step size Runge-Kutta (4,5) formula (the Matlab ode45 solver), with maximum step size of 0.3. The plots were generated using the resulting variable-step size data, but joined up with lines.

7. Several references refer to news articles and other documents and appear to be rather sparsely referenced, specific dates are not provided and it would be helpful to provide URLs to avoid ambiguity in locating these documents.

We thank the Review for highlighting these ambiguities. URLs have been added where relevant.

Reviewer: 2

Comments to the Author(s)

Summary

The study entitled "A Simulated Single Ventilator / Dual Patient Ventilation Strategy for Acute Respiratory Distress Syndrome During the COVID-19 Pandemic" by Solis-Lemus and colleague addresses a critical shortcoming in former modeling investigations of multi patient ventilation. In particular, former modeling of multi patient ventilation assume equal lung physiology between all patients and, therefore, are likely irrelevant in a clinical setting. Indeed, mechanical ventilation (MV) use is broad, encompassing spinal cord injury, drug overdose, heart failure, and acute respiratory distress syndrome (ARDS). Patients from many of these conditions are likely to present with comorbidities as well, making the matching of lung physiology between patients especially difficult. With this in mind, the authors a) present two electronic circuits (analogous in theory to the closed circuit of a single ventilator and two independent patients), b) created two mathematical models (standard splitter and modified splitter), c) demonstrate and quantify the pitfalls of previous multi patient MV models, d) provide a simple solution to the issues (variable flow restrictors added into the inspiration arms to control pressure for each patient and one-way valves on the expiration arms to prevent backflow), and e) made this software code freely available to the public for further testing and experimentation. Applying this model to a more clinically relevant model of multi patient ventilation, that is, pairing patients with different lung compliances on a single ventilator, the authors demonstrate that tidal volume could differ between patients as much as 38%. The authors then demonstrate through the simple addition of variable flow restrictors in the inspiratory tubes and one-way valves in the expiratory tubes, that it may be possible to ventilate two patients with mismatched lung physiology on a single ventilator.

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General Comments

We want to thank the authors for performing such an important work at this time. Also, we appreciate the analogous effort of utilizing electronic circuit theory to represent respiratory physiology and the open access of these mathematical models.

Major Comments none Minor Comments None

We thank the Reviewer for his or her supportive comments about our manuscript.