

# μLED-based optical cochlear implants for spectrally selective activation of the auditory nerve

Alexander Dieter, Eric Klein, Daniel Keppeler, Lukasz Jablonski, Tamas Harczos, Gerhard Hoch, Vladan Rankovic, Oliver Paul, Marcus Jeschke, Patrick Ruther, and Tobias Moser  
**DOI: 10.15252/emmm.202012387**

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## Review Timeline:

|                     |             |
|---------------------|-------------|
| Submission Date:    | 24th Mar 20 |
| Editorial Decision: | 27th Apr 20 |
| Revision Received:  | 23rd May 20 |
| Accepted:           | 2nd Jun 20  |

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*Editor: Celine Carret*

## Transaction Report:

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27th Apr 2020

Dear Prof. Moser,

Thank you for the submission of your manuscript to EMBO Molecular Medicine. We have now heard back from the three referees whom we asked to evaluate your manuscript.

You will see from the set of comments pasted below, that the three referees are very positive about the study and only request minor revisions. Therefore, I would like to ask you to perform these minor revisions and, in order to gain time, to also consider the below editorial amendments before submitting your next version:

1) Please address the minor comments from the three referees.

Provide a point-by-point letter INCLUDING my comments as well as the reviewer's reports and your detailed responses to their comments (as Word file).

2) Please carefully check the authors guidelines for formatting your supplemental information:

Expanded view and Appendix (see:

<https://www.embopress.org/page/journal/17574684/authorguide#expandedview>).

An Appendix pdf file should be submitted separately, including all the supplementary information.

Please note the different nomenclature and content (i.e. supplementary methods should be moved into the main article), update call outs.

3) Figures: separate figure files should be provided

4) In the main manuscript file, please do the following:

- correct/answer the track changes suggested by our data editors and myself by working from the attached document
- add up to 5 keywords
- change "Methods" for "Materials and Methods"
- ethics: provide the gerbils gender, handling conditions and origin of purchase. Age at which they were used should be indicated in the article as well.
- in M&M, the statistical paragraph should reflect all information that you have filled in the Authors checklist, especially regarding randomisation, blinding, replication.
- indicate in legends exact n= and exact p= values, not a range, along with the statistical test used. Some people found that to keep the figures clear, providing an Appendix table Sx with all exact p-values was preferable. You are welcome to do this if you want to.

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(<http://embomolmed.embopress.org/authorguide#editorial3>) to be submitted with all revised manuscripts. Please use the checklist as guideline for the sort of information we need WITHIN the manuscript. This is particularly important for animal reporting, the use of human samples, antibody dilutions and exact p- and n-values that should be indicated instead of a range, along with the right justified statistical test.

This file will be published along the Review Process File.

6) For more information: There is space at the end of each article to list relevant web links for further consultation by our readers. Could you identify some relevant ones and provide such information as well? Some examples are patient associations, relevant databases, OMIM/proteins/genes links, author's websites, etc...

7) The Paper Explained: EMBO Molecular Medicine articles are accompanied by a summary of the articles to emphasize the major findings in the paper and their medical implications for the non-specialist reader. Please provide a draft summary of your article highlighting

- the medical issue you are addressing, = Problem
- the results obtained = Results
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This may be edited to ensure that readers understand the significance and context of the research. Please refer to any of our published articles for an example.

8) Every published paper now includes a 'Synopsis' to further enhance discoverability. Synopses are displayed on the journal webpage and are freely accessible to all readers. They include a short stand first (maximum of 300 characters, including space) as well as 2-5 one sentence bullet points that summarise the paper. Please write the bullet points to summarise the key NEW findings. They should be designed to be complementary to the abstract - i.e. not repeat the same text. We encourage inclusion of key acronyms and quantitative information (maximum of 30 words / bullet point). Please use the passive voice. Please attach these in a separate file or send them by email, we will incorporate them accordingly.

You are also encouraged to suggest a striking image or visual abstract to illustrate your article. If you do please provide a jpeg file 550 px-wide x (250-400)-px high.

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In the event of acceptance, this file will be published in conjunction with your paper and will include the anonymous referee reports, your point-by-point response and all pertinent correspondence relating to the manuscript. Let us know whether you agree with the publication of the RPF.

Please note that the Authors checklist will be published at the end of the RPF.

10) Data and software availability. Please add the following:

"Data and software availability" (within the M&M)

"The datasets and computer code produced in this study are available in the following databases:

- Modeling computer scripts: GitHub (<https://github.com/SysBioChalmers/GECKO/releases/tag/v1.0>)
- [data type]: [full name of the resource] [accession number/identifier] ([doi or URL or identifiers.org/DATABASE:ACCESSION])"

Please submit the MATLAB codes/scripts to repositories or provide them as source data within this submission, referenced in this section.

11) Dr. Ruther must obtain an ORCID number and associate it to this manuscript.

I look forward to seeing a revised form of your manuscript as soon as possible (however, given the exceptional current circumstances, we perfectly understand that delays are to be expected. If you don't mind, just update us after 3 months to let us know where you are standing with the revision).

Yours sincerely,

Celine Carret

Celine Carret, PhD  
Senior Editor  
EMBO Molecular Medicine

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**\*\* PLEASE NOTE \*\*** As part of the EMBO Publications transparent editorial process initiative (see our Editorial at <https://www.embopress.org/doi/pdf/10.1002/emmm.201000094>), EMBO Molecular Medicine will publish online a Review Process File to accompany accepted manuscripts.

In the event of acceptance, this file will be published in conjunction with your paper and will include the anonymous referee reports, your point-by-point response and all pertinent correspondence relating to the manuscript. If you do NOT want this file to be published, please inform the editorial office at [contact@embomolmed.org](mailto:contact@embomolmed.org).

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When submitting your revised manuscript, please include:

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- 2) separate figure files\*
- 3) supplemental information as Expanded View and/or Appendix. Please carefully check the authors guidelines for formatting Expanded view and Appendix figures and tables at <https://www.embopress.org/page/journal/17574684/authorguide#expandedview>
- 4) a letter INCLUDING the reviewers' reports and your detailed responses to their comments (as Word file)

Also, and to save some time should your paper be accepted, please read below for additional information regarding some features of our research articles:

5) The paper explained: EMBO Molecular Medicine articles are accompanied by a summary of the articles to emphasize the major findings in the paper and their medical implications for the non-specialist reader. Please provide a draft summary of your article highlighting

- the medical issue you are addressing,
- the results obtained and
- their clinical impact.

This may be edited to ensure that readers understand the significance and context of the research. Please refer to any of our published articles for an example.

6) For more information: There is space at the end of each article to list relevant web links for further consultation by our readers. Could you identify some relevant ones and provide such information as well? Some examples are patient associations, relevant databases, OMIM/proteins/genes links, author's websites, etc...

7) Author contributions: the contribution of every author must be detailed in a separate section (before the acknowledgments).

8) EMBO Molecular Medicine now requires a complete author checklist (<https://www.embopress.org/page/journal/17574684/authorguide>) to be submitted with all revised manuscripts. Please use the checklist as a guideline for the sort of information we need WITHIN the manuscript as well as in the checklist. This is particularly important for animal reporting, antibody dilutions (missing) and exact p-values and n that should be indicated instead of a range.

9) Every published paper now includes a 'Synopsis' to further enhance discoverability. Synopses are displayed on the journal webpage and are freely accessible to all readers. They include a short stand first (maximum of 300 characters, including space) as well as 2-5 one sentence bullet points that summarise the paper. Please write the bullet points to summarise the key NEW findings. They should be designed to be complementary to the abstract - i.e. not repeat the same text. We encourage inclusion of key acronyms and quantitative information (maximum of 30 words / bullet point). Please use the passive voice. Please attach these in a separate file or send them by email, we will incorporate them accordingly.

You are also welcome to suggest a striking image or visual abstract to illustrate your article. If you do please provide a jpeg file 550 px-wide x 400-px high.

10) A Conflict of Interest statement should be provided in the main text

11) Please note that we now mandate that all corresponding authors list an ORCID digital identifier. This takes <90 seconds to complete. We encourage all authors to supply an ORCID identifier, which will be linked to their name for unambiguous name identification.

Currently, our records indicate that the ORCID for your account is 0000-0001-7145-0533.

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Each figure should be given in a separate file and should have the following resolution:

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Figures are not edited by the production team. All lettering should be the same size and style; figure panels should be indicated by capital letters (A, B, C etc). Gridlines are not allowed except for log plots. Figures should be numbered in the order of their appearance in the text with Arabic numerals. Each Figure must have a separate legend and a caption is needed for each panel.

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\*\*\*\*\* Reviewer's comments \*\*\*\*\*

Referee #1 (Remarks for Author):

This work by Dieter, Klein et al. describes the application of multi-channel  $\mu$ LED optical cochlea implants (oCIs) to evoke neuronal activity in optogenetically modified spiral ganglion neurons (SGNs). The overall translational goal is to overcome the limitations of electrical cochlear implants for hearing restoration, for example current spread and limited spectral selectivity of SGNs activation. This manuscript represents an important proof of concept showing optically evoked tonotopic activity of the auditory pathway and spectral selectivity of optogenetic SGN stimulation by 16  $\mu$ LED-based oCIs. Of note, the biological (efficient optogenetic transduction of SGN), engineering (biocompatible and stable  $\mu$ LED-based oCIs) and surgical (oCI transplantation) aspects are extremely challenging, require a precise interplay and adjustment and have been mastered by the authors. There are some minor points that need to be addressed and discussed in more detail. Minor comments:

- 1) Please define "oABR" the first time it is mentioned, not every interested reader knows that optically-evoked auditory brainstem responses are meant.
- 2) oCIs may offer advantages over eCIs, however so far, "high" spectral selectivity could be misleading due to the fact that single  $\mu$ LED stimulation was not successful and block-illumination was required (simultaneous shining with bunch of  $\mu$ LEDs). It would be great to discuss this aspect in more detail in regard to the performances of eCIs.
- 3) Does the evoked ICC activity by individual or block  $\mu$ LEDs follow particular patterns coupled to the optical stimulation? Are there any differences in activity depending on each individual  $\mu$ LED?
- 4) Is there a distribution/standard deviation of the light emission power among single LEDs of an array with respect to the same applied current?

- 5) The emission characteristics such as wavelength/bandwidth of  $\mu$ LEDs are not stated.
- 6) All signal profiles have been represented as an average of 1000 pulses. It would be great to show at least one example of a signal profile at baseline and during light stimulation.
- 7) How many SGN neurons are activated by one LED channel?
- 8) Why is the rate of opsin function after transfection in animals (12/32) so low? Please discuss/report the challenges for the general readers.

Referee #2 (Remarks for Author):

The manuscript " $\mu$ LED-based optical cochlear implants for spectrally selective activation of the auditory nerve" by Dieter et al is the latest advance from this research group in developing optical cochlear implants, which could have a profound impact on individuals with hearing loss. The authors use optogenetic tools that they previously developed to infect both hearing and deafened gerbils with a fast channelrhodopsin variant targeted to cochlear spiral ganglion neurons (SGN), which comprise the ascending auditory nerve. In previous work the channelrhodopsins were activated using optical fibers to excite the auditory nerve. In this work, the authors demonstrate a significant advance in the method of light delivery to the cochlea, using micro-LEDs in a multi-channel array. The array could be positioned in different locations in the cochlea based upon type of implantation surgery. Further, the authors demonstrate that optical stimulation of the SGN has a reduced "spectral spread", indicating improved tonotopic activation of the auditory nerve, compared to traditional electrical cochlear implants. Overall, the work is solid and polished. Some analyses are complex and difficult to understand (see a comment below), but figures are well done, and analyses are appropriate.

1. The function of the optical cochlear implants compared to electrical cochlear implants is well done. Please also discuss how the function relates to the spectral selectivity of sound activation of the auditory system in general, or IC neurons in particular for this study. Based on the first sentence on page 4, you have this data.
2. First sentence page 4 - were the acoustic stimulation experiments performed in the same animals as the optical stimulation? This is not clear. When was this performed in relation to oCI insertion?
3. Is it known why oABRs failed in 20/32 animals? Was this an expression problem?
4. In figure 1D', it would be helpful to label the IHC location to distinguish from the eYFP in the type I dendrite
5. Figure 2 and 3 should refer to the "example" or "exemplar" data, as "exemplary" is used to mean the best or perfect example
6. Figure 3B is a particularly difficult to understand figure, but there is results text dedicated to explaining it. Further improvements could be made by indicating which data points in 3B refer to the panels in 3A, and in clearly detailing why some data points are negative on the y-axis.

Referee #3 (Remarks for Author):

This study is addressing the feasibility of a new generation of a cochlear implant using optogenetics instead of electrical stimulation. The team of T.Moser is a leader in the field. They already published two major papers in the field (Science trans med, 2018 and nature comm, 2019). In this paper, they go further using a fibre equipped with 16 micro-LEDs to allow a spectral selective

activation of the nerve fibre. To assess this selectivity, they illuminate SGN at different locations and then recorded corresponding inferior colliculus regions. This study uses cutting edge technologies from various fields of research (AAV, electrophysiological recordings, optogenetics, etc.). The study is well constructed and nicely described and constitutes a crucial preclinical breakthrough that allows going closer to the clinics. Also, they are aware that this study is not perfect and that they need to improve the power output of the LEDs if they want to apply optogenetic CI to human.

One concern is the low number of animals used and the discordance in the size of the different groups (i.e. 9 controls and 3 deafened AAV injected gerbils) even if finally, it does not change the conclusions of the findings.

Material and methods should be carefully reviewed and more detailed to allow reproducibility of the experiments in the scientific community.

The titer of AAV: the authors should explain and justify the titer used (mentioned precisely what is GC, how they obtained  $6.99 \times 10^{12}$  GC/ml and why do they injected only a final concentration of  $1.4 - 2.1 \times 10^{10}$  GC as in the previously cited studies they used much more (i.e. in Wrobel paper they injected  $3.2 \times 10^{12} - 2.7 \times 10^{13}$  genome copies/ $\mu$ l

ABRs are not detailed at all and do not refer to any other published paper

Age of the animal for deafening?

- Discordance in the number of animals used: 16 gerbils are mentioned in the "mat and meth" section. In results, the authors said: "After verifying opsin function (robust oABRs could be evoked in 12/32 animals)". Could they explain the different numbers?

Figure 1J: yellow is not visible

Figure 2 All the example of STCs should be presented at the same scale for radiant flux (mainly to compare panels C and G that are strikingly different and should be explained).

Statistical analysis should be better described (for example the authors wrote: "As for response strengths, also the number of recruited multi-units did not differ between hearing and deafened AAV-injected animals ( $p = 0.03/0.24/0.14/0.52$  for individual/block/all/fiber stimulation, Bonferroni-corrected, two-sample t-test,  $p = 0.05/4 = 0.0125$ "). Is it counterintuitive to say that there is no difference if  $p=0.03$  or  $0.05$ ? In addition, the authors should mention that they performed an ANOVA if this is the case and why did they perform t-test or any other test such as Wilcoxon ((figure 3 legend: "( $n = 9/3$  in  $N = 9/3$  hearing/deaf gerbils;  $p = 0.58$ , Wilcoxon rank sum test)"). To avoid a surcharge of the text, an additional table with all the statistics clearly explained should be performed.

Referee #1 (Remarks for Author):

This work by Dieter, Klein et al. describes the application of multi-channel  $\mu$ LED optical cochlea implants (oCIs) to evoke neuronal activity in optogenetically modified spiral ganglion neurons (SGNs). The overall translational goal is to overcome the limitations of electrical cochlear implants for hearing restoration, for example current spread and limited spectral selectivity of SGNs activation. This manuscript represents an important proof of concept showing optically evoked tonotopic activity of the auditory pathway and spectral selectivity of optogenetic SGN stimulation by 16  $\mu$ LED-based oCIs. Of note, the biological (efficient optogenetic transduction of SGN), engineering (biocompatible and stable  $\mu$ LED-based oCIs) and surgical (oCI transplantation) aspects are extremely challenging, require a precise interplay and adjustment and have been mastered by the authors. There are some minor points that need to be addressed and discussed in more detail.

We are grateful to Referee 1 for the appreciation of our work, as well as for the comments, which helped us to further improve our manuscript.

Minor comments:

1) Please define "oABR" the first time it is mentioned, not every interested reader knows that optically-evoked auditory brainstem responses are meant.

Thanks for pointing this out, we have added this information.

2) oCIs may offer advantages over eCIs, however so far, "high" spectral selectivity could be misleading due to the fact that single  $\mu$ LED stimulation was not successful and block-illumination was required (simultaneous shining with bunch of  $\mu$ LEDs). It would be great to discuss this aspect in more detail in regard to the performances of eCIs.

In response to this comment, we have avoided the claim of "high spectral selectivity" throughout the MS. Specifically, in most cases, we have followed the advice of the reviewer to discuss it in comparison to the eCI. The information on the spread of excitation with mono- and bipolar electrical stimulation and acoustic stimulation (taken from our previous study that used the same set-up for comparison to fiber-based optogenetic stimulation) is provided in the discussion and we have added the spread of excitation of acoustic stimulation to Fig. 3C, which supports the notion of "near physiological" spectral selectivity.

3) Does the evoked ICC activity by individual or block  $\mu$ LEDs follow particular patterns coupled to the optical stimulation? Are there any differences in activity depending on each individual  $\mu$ LED?

At this point, we do not have an analysis regarding the particular pattern of individual  $\mu$ LEDs or  $\mu$ LED blocks. However, we do observe substantial variability among the responses, probably due to different patterns of opsin expression, oCI-design (pitch, pattern of functional  $\mu$ LEDs), and oCI position upon implantation (see also response to reviewer 3). We have commented on this variability now in the discussion.

4) Is there a distribution/standard deviation of the light emission power among single LEDs of an array with respect to the same applied current?

Characterizing a batch of 16 exemplary oCI samples with 16  $\mu$ LEDs each using optimal positioning toward the detector we extracted at a  $\mu$ LED current of 10 mA applied at a pulse duration of 1 ms an averaged optical output power of  $0.932 \pm 0.010$  mW (average  $\pm$  standard deviation (SD);  $n = 256$ ), respectively. The SD of the individual oCIs varied between 0.6 and 2% of the respective oCI-averaged output power.

5) The emission characteristics such as wavelength/bandwidth of  $\mu$ LEDs are not stated.

Done, 462 nm, we quote measurements in Klein et al., Front Neurosci 2018 (Figure 13 therein)

6) All signal profiles have been represented as an average of 1000 pulses. It would be great to show at least one example of a signal profile at baseline and during light stimulation.

The thousand repetitions are averaged for oABR data, which is correct. However, due to their small signal amplitude, no response can be observed in an individual trial. Multiunit activity-data – or the resulting  $d'$  values of them – originate from 30 repetitions only. We have now included an individual trace, as well as a corresponding scatter plot in response to 30 stimulus presentations, for a multi-unit recorded from the ICC in the supplementary figure S2.

7) How many SGN neurons are activated by one LED channel?

This is a great question, but difficult to answer. As indicated by the spread of excitation measurements shown for blocks of 4  $\mu$ LEDs in Fig. 3, this will heavily depend of the radiant flux of the  $\mu$ LED. Assuming i) 24.000 type I SGNs (early postnatal count, (Richter *et al*, 2011), ii) a uniform distribution along the tonotopic axis (8.8 octaves (Müller, 1996), and iii) a fraction of 30% of SGNs expressing CatCh across all tonotopic places (Wrobel *et al*, 2018), we approximate that maximally ~800 SGNs (3% of total) are recruited for the lowest spread of excitation obtained with blocks of 4  $\mu$ LEDs (~ 1 octave at  $d' = 1$ ).

8) Why is the rate of opsin function after transfection in animals (12/32) so low? Please discuss/report the challenges for the general readers.

Yes, the low fraction of opsin expressing SGNs upon intramodiolar AAV-injection into the adult gerbil cochlea is troublesome and we have extended the discussion of this point in response to the reviewer's comment. In brief, we suspect that AAV access to the SGN represents the major bottleneck. While we and others are actively working on improving AAV-mediated transduction of adult SGN e.g. by using more potent AAV vectors (e.g. AAV-PHP.B, this study) and application of larger volumes of AAV suspension to the scala tympani, we have not yet managed the reproducibility and transduction rates we find with early postnatal AAV injections into rodent cochleae (100% of surviving animals, 40-9X% of SGNs transduced). Within a previous study (Wrobel et al, Science Translational Medicine, 2018) we have extensively looked at the cochleae of oABR-negative animals, and could not find robust opsin expression. Unfortunately, we did not process the tissue of most of the negative animals in this study, but we have had a look at one of these cochleae, and have included this data to the MS now (Fig. EV2). While PV-staining revealed the presence of SGNs, the GFP-expression indicating *CatCh* was really sparse.

Referee #2 (Remarks for Author):

The manuscript "μLED-based optical cochlear implants for spectrally selective activation of the auditory nerve" by Dieter et al is the latest advance from this research group in developing optical cochlear implants, which could have a profound impact on individuals with hearing loss. The authors use optogenetic tools that they previously developed to infect both hearing and deafened gerbils with a fast channelrhodopsin variant targeted to cochlear spiral ganglion neurons (SGN), which comprise the ascending auditory nerve. In previous work the channelrhodopsins were activated using optical fibers to excite the auditory nerve. In this work, the authors demonstrate a significant advance in the method of light delivery to the cochlea, using micro-LEDs in a multi-channel array. The array could be positioned in different locations in the cochlea based upon type of implantation surgery. Further, the authors demonstrate that optical stimulation of the SGN has a reduced "spectral spread", indicating improved tonotopic activation of the auditory nerve, compared to traditional electrical cochlear implants. Overall, the work is solid and polished. Some analyses are complex and difficult to understand (see a comment below), but figures are well done, and analyses are appropriate.

We thank Referee 2 for the appreciation of our work, as well as for the comments, which helped us to further improve our manuscript.

1. The function of the optical cochlear implants compared to electrical cochlear implants is well done. Please also discuss how the function relates to the spectral selectivity of sound activation of the auditory system in general, or IC neurons in particular for this study. Based on the first sentence on page 4, you have this data.

It is true that we have acoustic data from these animals for confirming the recording position of the electrode array in the ICC. However, we do not think that comparison to this data is fair, as the mapping has been performed after cochlear surgery, and thus hearing in these animals is already compromised and not as good as in wildtype animals (as shown in Dieter et al, Nature Communications, 2019, thresholds across the hearing range of the gerbil are elevated by  $\geq 20$  dB in animals that underwent cochlear surgery). However, we have recorded data in response to pure tone acoustic stimulation in naïve gerbils in this previous study, and have now added the average spread of excitation obtained from this data to Fig. 3C of the current manuscript for ease of orientation.

2. First sentence page 4 - were the acoustic stimulation experiments performed in the same animals as the optical stimulation? This is not clear. When was this performed in relation to oCI insertion?

We have added a sentence to outline the experimental flow more clearly. Acoustic stimulation has been done in all non-deafened animals. It was, however, done after performing cochlear surgeries which most likely have compromised hearing (by a threshold shift of  $\sim 20$  dB across the whole hearing range, as shown in Dieter et al, Nat. Comm. 2019) and is thus not comparable to natural hearing in a naïve animal.

3. Is it known why oABRs failed in 20/32 animals? Was this an expression problem?

Yes, the low fraction of opsin expressing SGNs upon intramodiolar AAV-injection into the adult gerbil cochlea is troublesome and we have extended the discussion of this point in response to the reviewer's comment. In brief, we suspect that AAV access to the SGN represents the major bottleneck. While we and others are actively working on improving AAV-mediated transduction of adult SGN e.g. by using more potent AAV vectors (e.g. AAV-PHP.B, this study) and application of larger volumes of AAV suspension to the scala tympani, we have not yet managed the reproducibility

and transduction rates we find with early postnatal AAV injections into rodent cochleae (100% of surviving animals, 40-9X% of SGNs transduced). Within a previous study (Wrobel et al, Science Translational Medicine, 2018) we have extensively looked at the cochleae of oABR-negative animals, and could not find robust opsin expression. Unfortunately, we did not process the tissue of most of the negative animals in this study, but we have had a look at one of these cochleae, and have included this data to the MS now (Fig. EV2). While PV-staining revealed the presence of SGNs, the GFP-expression indicating *CatCh* was really sparse.

4. In figure 1D', it would be helpful to label the IHC location to distinguish from the eYFP in the type I dendrite.

Done.

5. Figure 2 and 3 should refer to the "example" or "exemplar" data, as "exemplary" is used to mean the best or perfect example.

Thanks for pointing this out, we have changed the phrase as suggested.

6. Figure 3B is a particularly difficult to understand figure, but there is results text dedicated to explaining it. Further improvements could be made by indicating which data points in 3B refer to the panels in 3A, and in clearly detailing why some data points are negative on the y-axis.

Thanks for helping us to make these results more accessible. We have now indicated the data points for best electrodes in figure 3A in figure 3B. As before, there were examples in figure 3A which were taken from cochleostomy and round window implantation (i.e. two different experiments), we have now chosen an example of blockwise stimulation with one oCI. While this might not shift the BE as much as different oCI insertions, we can now refer to four data points from the same implant in figure 3B, and thereby hopefully contribute to a more intuitive understanding of this figure. Also, we have added a short sentence about the origin of negative data points.

Referee #3 (Remarks for Author):

This study is addressing the feasibility of a new generation of a cochlear implant using optogenetics instead of electrical stimulation. The team of T. Moser is a leader in the field. They already published two major papers in the field (Science trans med, 2018 and nature comm, 2019). In this paper, they go further using a fibre equipped with 16 micro-LEDs to allow a spectral selective activation of the nerve fibre. To assess this selectivity, they illuminate SGN at different locations and then recorded corresponding inferior colliculus regions. This study uses cutting edge technologies from various fields of research (AAV, electrophysiological recordings, optogenetics, etc..). The study is well constructed and nicely described and constitutes a crucial preclinical breakthrough that allows going closer to the clinics. Also, they are aware that this study is not perfect and that they need to improve the power output of the LEDs if they want to apply optogenetic CI to human.

One concern is the low number of animals used and the discordance in the size of the different groups (i.e. 9 controls and 3 deafened AAV injected gerbils) even if finally, it does not change the conclusions of the findings.

We thank Referee 3 for the appreciation of our work and the remarks, which helped us to further improve our manuscript. We want to note that the two different groups (9 hearing and 3 deafened animals) are not meant to be compared against each other. Rather, we have quantified results of the pooled data ( $n = 12$ ), but wanted to demonstrate in a subset ( $n = 3$ ) of animals, that our technology works in deafened animals as well. Unfortunately, we have used up all oCIs with this design for the current study, and oCIs of the next generation will have a different design (as mentioned in the discussion), which prevents us from doing additional experiments with the current settings.

Material and methods should be carefully reviewed and more detailed to allow reproducibility of the experiments in the scientific community.

Fair point, done and we have added the requested information.

The titer of AAV: the authors should explain and justify the titer used (mentioned precisely what is GC, how they obtained  $6.99 \times 10^{12}$  GC/ml and why do they injected only a final concentration of  $1.4 - 2.1 \times 10^{10}$  GC as in the previously cited studies they used much more (i.e. in Wrobel paper they injected  $3.2 \times 10^{12} - 2.7 \times 10^{13}$  genome copies/ $\mu$ l).

Thanks for catching this one!!

GC = genome copies, as this abbreviation is only used twice, we now spare this abbreviation.

$6.99 \times 10^{12}$  GC/ml were measured by the qPCR AAV titration kit by TaKaRa, as we now have added, and  $1.4 - 2.1 \times 10^{10}$  does not refer to a concentration, but rather approximate the number of genome copies that have been injected, as calculated by the volume of the injected virus suspension (2-3  $\mu$ l). It is true that the viral titer differs from Wrobel et al. (2018), where we now realized that we have made a mistake. The reported titer there should be  $3.2 \times 10^{12} - 2.7 \times 10^{13}$  genome copies/ml, which compares to the titer used in the current study, we are about to submit a correction for Wrobel et al. (2018).

ABRs are not detailed at all and do not refer to any other published paper.

We have included additional information and a reference to a previous paper with detailed description.

Age of the animal for deafening? Has been added.

Discordance in the number of animals used: 16 gerbils are mentioned in the "mat and meth" section. In results, the authors said: "After verifying opsin function (robust oABRs could be evoked in 12/32 animals)". Could they explain the different numbers?

We apologize for not stating this clearly. We could evoke oABRs in 12 out of 32 opsin-injected animals and performed IC recordings in these we animals (9 hearing and 3 deafened). In addition, we have added 4 non-injected wildtype animals (two hearing and two deafened), which then adds up to the 16 gerbils mentioned in the materials and methods section. Actually, to complicate it even more, we have had 3 more animals showing oABRs (which increase the fraction of positive animals in our study) for which, however, we could not obtain IC recordings, either due to death of the animal, or to misplacement of the electrode array. We have now added these animals to the total number, and explained the number of animals more clearly in general, hopefully avoiding confusion.

Figure 1J: yellow is not visible.

As this is not essential, we have removed the sentence, as the visibility of the encapsulated cables and  $\mu$ LEDs might be compromised when using a more prominent color for the encapsulation.

Figure 2 All the example of STCs should be presented at the same scale for radiant flux (mainly to compare panels C and G that are strikingly different and should be explained).

We now present all the panels at the same radiant flux, which makes these figures more comparable. It is correct, that STCs in response to different oCIs and recorded from different animals are quite variable, which might result from different opsin expression patterns, different oCI-designs (in terms of pitch, and in terms of functional/unfunctional  $\mu$ LEDs on individual devices), and different positions of implants. We now chose more comparable STCs to avoid confusion, but we added these points to the discussion.

Statistical analysis should be better described (for example the authors wrote: "As for response strengths, also the number of recruited multi-units did not differ between hearing and deafened AAV-injected animals ( $p = 0.03/0.24/0.14/0.52$  for individual/block/all/fiber stimulation, Bonferroni-corrected, two-sample t-test,  $p = 0.05/4 = 0.0125$ "). Is it counterintuitive to say that there is no difference if  $p=0.03$  or  $0.05$ ? In addition, the authors should mention that they performed an ANOVA if this is the case and why did they perform t-test or any other test such as Wilcoxon ((figure 3 legend: "( $n = 9/3$  in  $N = 9/3$  hearing/deaf gerbils;  $p = 0.58$ , Wilcoxon rank sum test)"). To avoid a surcharge of the text, an additional table with all the statistics clearly explained should be performed. Thanks for pointing this out. We have discussed this, and adopted statistical tests to be more adequate (mainly Wilcoxon Rank Sum and ANOVA). We have stated the exact test in each figure legend, and have summarized the statistics in a supplementary table.

2nd Jun 2020

Dear Tobias,

Thank you very much for your reply and for providing the last set of items. We are pleased to inform you that your manuscript is accepted for publication and is now being sent to our publisher to be included in the next available issue of EMBO Molecular Medicine.

We would like to remind you that as part of the EMBO Publications transparent editorial process initiative, EMBO Molecular Medicine will publish a Review Process File online to accompany accepted manuscripts. If you do NOT want the file to be published or would like to exclude figures, please immediately inform the editorial office via e-mail.

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Celine Carret, PhD  
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|                            |                                            |
|----------------------------|--------------------------------------------|
| Corresponding Author Name: | Prof. Dr. Tobias Moser, Dr. Patrick Ruther |
| Journal Submitted to:      | EMBO Molecular Medicine                    |
| Manuscript Number:         | EMM-2020-12387-V2                          |

#### Reporting Checklist For Life Sciences Articles (Rev. June 2017)

This checklist is used to ensure good reporting standards and to improve the reproducibility of published results. These guidelines are consistent with the Principles and Guidelines for Reporting Preclinical Research issued by the NIH in 2014. Please follow the journal's authorship guidelines in preparing your manuscript.

#### A- Figures

##### 1. Data

**The data shown in figures should satisfy the following conditions:**

- the data were obtained and processed according to the field's best practice and are presented to reflect the results of the experiments in an accurate and unbiased manner.
- figure panels include only data points, measurements or observations that can be compared to each other in a scientifically meaningful way.
- graphs include clearly labeled error bars for independent experiments and sample sizes. Unless justified, error bars should not be shown for technical replicates.
- if  $n \leq 5$ , the individual data points from each experiment should be plotted and any statistical test employed should be justified
- Source Data should be included to report the data underlying graphs. Please follow the guidelines set out in the author ship guidelines on Data Presentation.

##### 2. Captions

**Each figure caption should contain the following information, for each panel where they are relevant:**

- a specification of the experimental system investigated (eg cell line, species name).
- the assay(s) and method(s) used to carry out the reported observations and measurements
- an explicit mention of the biological and chemical entity(ies) that are being measured.
- an explicit mention of the biological and chemical entity(ies) that are altered/varied/perturbed in a controlled manner.
- the exact sample size (n) for each experimental group/condition, given as a number, not a range;
- a description of the sample collection allowing the reader to understand whether the samples represent technical or biological replicates (including how many animals, litters, cultures, etc.).
- a statement of how many times the experiment shown was independently replicated in the laboratory.
- definitions of statistical methods and measures:
  - common tests, such as t-test (please specify whether paired vs. unpaired), simple  $\chi^2$  tests, Wilcoxon and Mann-Whitney tests, can be unambiguously identified by name only, but more complex techniques should be described in the methods section;
  - are tests one-sided or two-sided?
  - are there adjustments for multiple comparisons?
  - exact statistical test results, e.g., P values = x but not P values < x;
  - definition of 'center values' as median or average;
  - definition of error bars as s.d. or s.e.m.

Any descriptions too long for the figure legend should be included in the methods section and/or with the source data.

**In the pink boxes below, please ensure that the answers to the following questions are reported in the manuscript itself. Every question should be answered. If the question is not relevant to your research, please write NA (non applicable). We encourage you to include a specific subsection in the methods section for statistics, reagents, animal models and human subjects.**

#### B- Statistics and general methods

Please fill out these boxes ▼ (Do not worry if you cannot see all your text once you press return)

|                                                                                                                                                                                         |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 1.a. How was the sample size chosen to ensure adequate power to detect a pre-specified effect size?                                                                                     | The sample size of CatCh-injected animals was chosen according to estimations based on our previous studies (Dieter et al, Nature Communications 2019). As the number of experiments was limited by the amount of available optical cochlear implants, we have decided to add a reduced number of non-injected animals to evaluate non-specific auditory nerve stimulation and a reduced number of deafened animals (as a proof of concept) to support the main findings of high frequency resolution, as these experiments had to be performed in CatCh-injected (to enable optogenetic) and hearing (to enable frequency assessment of the inferior colliculus) animals. |
| 1.b. For animal studies, include a statement about sample size estimate even if no statistical methods were used.                                                                       | The sample size of CatCh-injected animals was chosen according to estimations based on our previous studies (Dieter et al, Nature Communications 2019). As the number of experiments was limited by the amount of available optical cochlear implants, we have decided to add a reduced number of non-injected animals to evaluate non-specific auditory nerve stimulation and a reduced number of deafened animals (as a proof of concept) to support the main findings of high frequency resolution, as these experiments had to be performed in CatCh-injected (to enable optogenetic) and hearing (to enable frequency assessment of the inferior colliculus) animals. |
| 2. Describe inclusion/exclusion criteria if samples or animals were excluded from the analysis. Were the criteria pre-established?                                                      | All animals that received viral injections to optogenetically transduce the auditory nerve have been tested for functional opsin expression by recording optically evoked auditory brainstem responses (oABRs). Only animals in which an obvious oABR could be evoked have been used for IC recordings. Out of these, only animals in which the electrode array could be placed adequately, as demonstrated by the tonotopy, have been used for analysis. These criteria have been pre-established as analysis of IC responses to optogenetic stimulation requires successful optogenetic manipulations and appropriate recording positions.                               |
| 3. Were any steps taken to minimize the effects of subjective bias when allocating animals/samples to treatment (e.g. randomization procedure)? If yes, please describe.                | No randomization or blinding was taken. However, all experiments performed in this study have been performed in a standardized manner.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
| For animal studies, include a statement about randomization even if no randomization was used.                                                                                          | As all animals underwent multiple surgeries, and animals undergoing surgeries must be tracked, no randomization was used. As no different genotypes, reagents, or similar are compared against each other, randomization was also not needed in this study.                                                                                                                                                                                                                                                                                                                                                                                                                |
| 4.a. Were any steps taken to minimize the effects of subjective bias during group allocation or/and when assessing results (e.g. blinding of the investigator)? If yes please describe. | No steps were taken to minimize the effect of subjective bias during group allocation. However, data has been obtained for all animals in an identical way and analysis has been done automatically by MATLAB-code, where objective measures were taken.                                                                                                                                                                                                                                                                                                                                                                                                                   |
| 4.b. For animal studies, include a statement about blinding even if no blinding was done                                                                                                | No blinding was performed.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| 5. For every figure, are statistical tests justified as appropriate?                                                                                                                    | statistical tests are justified, as all data meets the criteria for the used Wilcoxon Rank Sum test (comparison of two groups) or analysis of variance (comparison of more than two groups).                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |

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|                                                                                                                    |                                                                                                                                                                                                                            |
|--------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Do the data meet the assumptions of the tests (e.g., normal distribution)? Describe any methods used to assess it. | As the Wilcoxon Rank Sum test, as a non-parametric statistical test, does not require normal distribution, it is justified to be used in this study. Compared groups are independent.                                      |
| Is there an estimate of variation within each group of data?                                                       | Each group of data contains an estimate of variation, which is either reported as standard deviation, standard error of the mean, or in quartiles. Also, we have plotted all individual data points underlying the graphs. |
| Is the variance similar between the groups that are being statistically compared?                                  | The variance of compared groups is in a similar range.                                                                                                                                                                     |

### C- Reagents

|                                                                                                                                                                                                                                                                                                                                                                  |                                                                                                                                                                                                                                                                            |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 6. To show that antibodies were profiled for use in the system under study (assay and species), provide a citation, catalog number and/or clone number, supplementary information or reference to an antibody validation profile. e.g., Antibodypedia ( <a href="#">see link list at top right</a> ), 1DegreeBio ( <a href="#">see link list at top right</a> ). | parvalbumin (cat. #195004, Synaptic Systems, Göttingen, Germany); GFP (cat. #ab13970, Abcam, Cambridge, United Kingdom); Goat anti-guinea-pig-568 (cat. #A11075, Thermo Fisher Scientific, Waltham, US); goat anti-chicken-488 (cat. #A11039, MoBiTec, Göttingen, Germany) |
| 7. Identify the source of cell lines and report if they were recently authenticated (e.g., by STR profiling) and tested for mycoplasma contamination.                                                                                                                                                                                                            | NA                                                                                                                                                                                                                                                                         |

\* for all hyperlinks, please see the table at the top right of the document

### D- Animal Models

|                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                                                                                        |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------|
| 8. Report species, strain, gender, age of animals and genetic modification status where applicable. Please detail housing and husbandry conditions and the source of animals.                                                                                                                                                                                                                                                            | The requested information is added in the materials/methods.                                           |
| 9. For experiments involving live vertebrates, include a statement of compliance with ethical regulations and identify the committee(s) approving the experiments.                                                                                                                                                                                                                                                                       | A statement of compliance with ethical regulations as requested is found within the materials/methods. |
| 10. We recommend consulting the ARRIVE guidelines ( <a href="#">see link list at top right</a> ) (PLoS Biol. 8(6), e1000412, 2010) to ensure that other relevant aspects of animal studies are adequately reported. See author guidelines, under 'Reporting Guidelines'. See also: NIH ( <a href="#">see link list at top right</a> ) and MRC ( <a href="#">see link list at top right</a> ) recommendations. Please confirm compliance. | compliance is confirmed                                                                                |

### E- Human Subjects

|                                                                                                                                                                                                                                                                                                                                                        |    |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----|
| 11. Identify the committee(s) approving the study protocol.                                                                                                                                                                                                                                                                                            | NA |
| 12. Include a statement confirming that informed consent was obtained from all subjects and that the experiments conformed to the principles set out in the WMA Declaration of Helsinki and the Department of Health and Human Services Belmont Report.                                                                                                | NA |
| 13. For publication of patient photos, include a statement confirming that consent to publish was obtained.                                                                                                                                                                                                                                            | NA |
| 14. Report any restrictions on the availability (and/or on the use) of human data or samples.                                                                                                                                                                                                                                                          | NA |
| 15. Report the clinical trial registration number (at ClinicalTrials.gov or equivalent), where applicable.                                                                                                                                                                                                                                             | NA |
| 16. For phase II and III randomized controlled trials, please refer to the CONSORT flow diagram ( <a href="#">see link list at top right</a> ) and submit the CONSORT checklist ( <a href="#">see link list at top right</a> ) with your submission. See author guidelines, under 'Reporting Guidelines'. Please confirm you have submitted this list. | NA |
| 17. For tumor marker prognostic studies, we recommend that you follow the REMARK reporting guidelines ( <a href="#">see link list at top right</a> ). See author guidelines, under 'Reporting Guidelines'. Please confirm you have followed these guidelines.                                                                                          | NA |

### F- Data Accessibility

|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |                                                                                                                                                                                        |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 18. Provide a "Data Availability" section at the end of the Materials & Methods, listing the accession codes for data generated in this study and deposited in a public database (e.g. RNA-Seq data: Gene Expression Omnibus GSE39462, Proteomics data: PRIDE PXD000208 etc.) Please refer to our author guidelines for 'Data Deposition'.<br><br>Data deposition in a public repository is mandatory for:<br>a. Protein, DNA and RNA sequences<br>b. Macromolecular structures<br>c. Crystallographic data for small molecules<br>d. Functional genomics data<br>e. Proteomics and molecular interactions                                                                                                                                              | We have added a data availability statement. As data deposition is not mandatory for our type of data, we are storing it on our institute's server and make it available upon request. |
| 19. Deposition is strongly recommended for any datasets that are central and integral to the study; please consider the journal's data policy. If no structured public repository exists for a given data type, we encourage the provision of datasets in the manuscript as a Supplementary Document (see author guidelines under 'Expanded View' or in unstructured repositories such as Dryad ( <a href="#">see link list at top right</a> ) or Figshare ( <a href="#">see link list at top right</a> ).                                                                                                                                                                                                                                              | As data deposition is not mandatory for our type of data, we are storing it on our institute's server and make it available upon request.                                              |
| 20. Access to human clinical and genomic datasets should be provided with as few restrictions as possible while respecting ethical obligations to the patients and relevant medical and legal issues. If practically possible and compatible with the individual consent agreement used in the study, such data should be deposited in one of the major public access-controlled repositories such as dbGAP ( <a href="#">see link list at top right</a> ) or EGA ( <a href="#">see link list at top right</a> ).                                                                                                                                                                                                                                       | This study does not include human clinical and genomic datasets.                                                                                                                       |
| 21. Computational models that are central and integral to a study should be shared without restrictions and provided in a machine-readable form. The relevant accession numbers or links should be provided. When possible, standardized format (SBML, CellML) should be used instead of scripts (e.g. MATLAB). Authors are strongly encouraged to follow the MIRIAM guidelines ( <a href="#">see link list at top right</a> ) and deposit their model in a public database such as Biocompare ( <a href="#">see link list at top right</a> ) or JWS Online ( <a href="#">see link list at top right</a> ). If computer source code is provided with the paper, it should be deposited in a public repository or included in supplementary information. | This study does not include computational models.                                                                                                                                      |

### G- Dual use research of concern

|                                                                                                                                                                                                                                                                                                                                   |    |
|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----|
| 22. Could your study fall under dual use research restrictions? Please check biosecurity documents ( <a href="#">see link list at top right</a> ) and list of select agents and toxins (APHIS/CDC ( <a href="#">see link list at top right</a> )). According to our biosecurity guidelines, provide a statement only if it could. | No |
|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----|