

POLYTECHNIQUE
MONTRÉAL

LE GÉNIE
EN PREMIÈRE CLASSE



INSTITUT DE
CARDIOLOGIE
DE MONTRÉAL

NeuroPoly



NeuroLibre: Taking papers to their logical conclusion

Nikola Stikov

École Polytechnique/Montreal Heart Institute
University of Montreal

Center for Advanced Interdisciplinary Research
University Ss Cyril and Methodius

My path

Stanford | Magnetic Resonance Systems Research Laboratory

OVERVIEW PUBLICATIONS PEOPLE ALUMNI PICTURES FINDING US PRIVATE



$$SNR \propto \sqrt{Time}$$

My publishing timeline



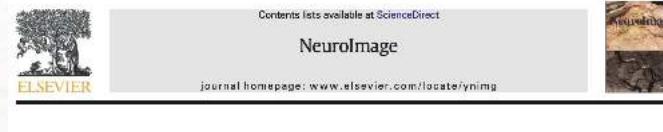
Bound pool fractions complement diffusion measures to describe white matter micro and macrostructure

Nikola Stikov^a, , Lee M. Perry^b, Aviv Mezer^b, Elena Rykhlevskaia^b, Brian A. Wandell^{a, b}, John M. Pauly^a, Robert F. Dougherty^b



Promise and pitfalls of g-ratio estimation with MRI

Jennifer S.W. Campbell^{a, b}, , Ilana R. Leppert^a, Sridar Narayanan^a, Mathieu Boudreau^a, Tanguy Duval^b, Julien Cohen-Adad^{b, e}, G. Bruce Pike^c, Nikola Stikov^{b, d}



In vivo histology of the myelin g-ratio with magnetic resonance imaging

Nikola Stikov^{a,d,*1}, Jennifer S.W. Campbell^{a,1}, Thomas Stroh^a, Mariette Lavelée^a, Stephen Frey^a, Jennifer Novek^a, Stephen Nuara^a, Ming-Kai Ho^a, Barry J. Bedell^a, Robert F. Dougherty^b, Ilana R. Leppert^a, Mathieu Boudreau^a, Sridar Narayanan^a, Tanguy Duval^b, Julien Cohen-Adad^d, Paul-Alexandre Picard^e, Alicja Gasecka^f, Daniel Côté^f, G. Bruce Pike^{a,c}



36086||
Data Article
Quantitative analysis of the myelin g-ratio from electron microscopy images of the macaque corpus callosum
Nikola Stikov^{a, d, *1}, Jennifer S.W. Campbell^{a, 1}, Thomas Stroh^a, Mariette Lavelée^a, Stephen Frey^a, Jennifer Novek^a, Stephen Nuara^a, Ming-Kai Ho^a, Barry J. Bedell^a, Robert F. Dougherty^b, Ilana R. Leppert^a, Mathieu Boudreau^a, Sridar Narayanan^a, Tanguy Duval^b, Julien Cohen-Adad^d, Paul-Alexandre Picard^e, Alicja Gasecka^f, Daniel Côté^f, G. Bruce Pike^{a, c}

My 10-year battle with 10-page PDFs

I got my PhD from Stanford University without having published a single journal article. I am now tenured and ready to share-ish my experiences with the publish-or-perish culture that is academia.

Nikola Stikov
Dec 28, 2021



<https://qantarot.substack.com/p/10-page-pdfs>

Science communication



www.humanbrainmapping.org/blog



www.ismrm.org/mrm



blog.ismrm.org



naukazadeca.mk



qantarot.substack.com

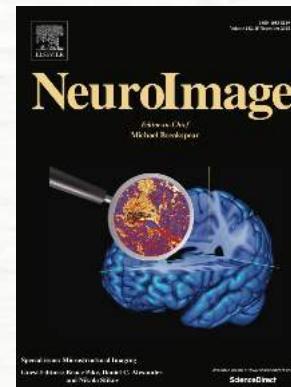
MAGNETIC RESONANCE IN MEDICINE



Bridging the macro-micro gap: biophysical MR modeling of the central nervous system

Christine Tardif, PhD
Douglas Mental Health Research Institute
McGill University

Nikola Stikov, PhD
École Polytechnique / Montreal Heart Institute
University of Montreal



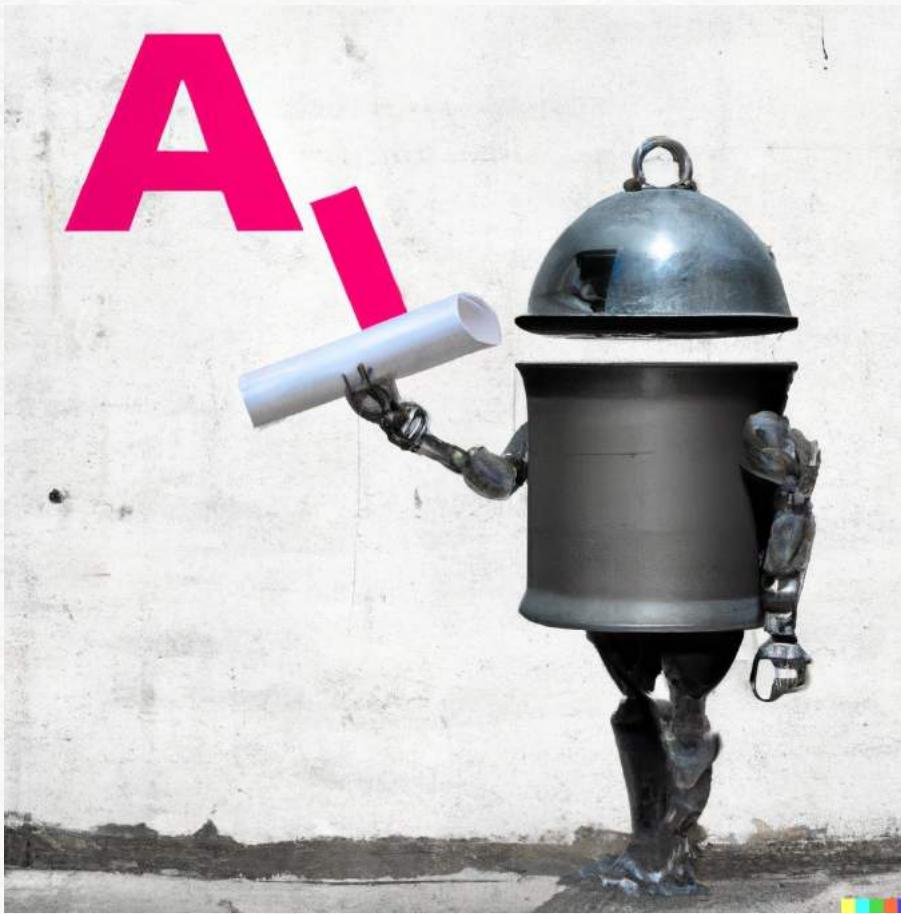
MR Imaging of Brain Microstructure

Guest Editors:

Prof. Bruce Pike [✉ bruce.pike@ucalgary.ca]

Prof. Daniel C. Alexander [✉ d.alexander@ucl.ac.uk]

Prof. Nikola Stikov [✉ nikola.stikov@polymtl.ca]



Prompt: Artificial intelligence swallowing an academic article, Banksy style. AI image generated by DALL-E

SHARE-ISH

ChatGPT and Galactica are taking scientific papers to their logical conclusion

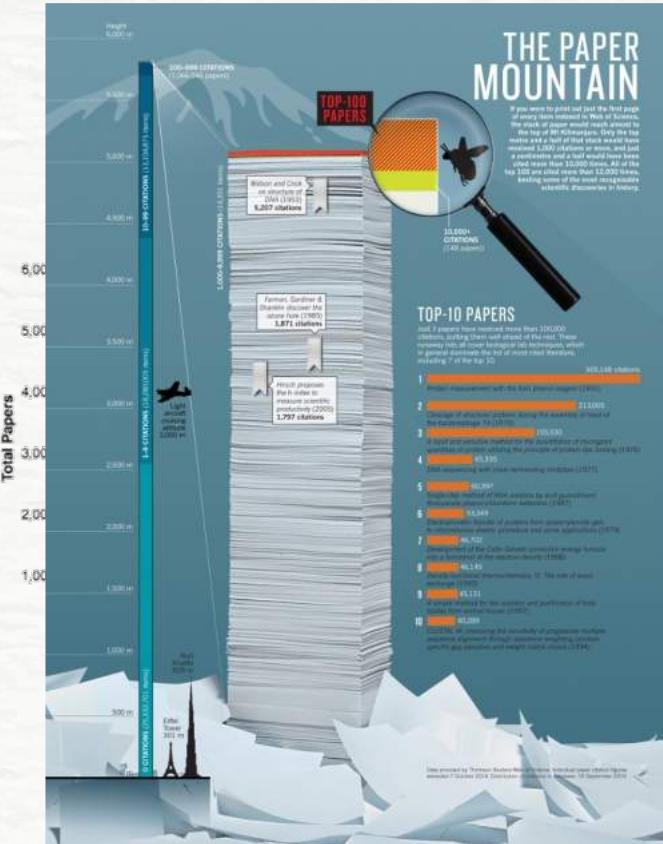
“... taken to its logical conclusion, every story is sad, because at the end everyone dies.”

— Margaret Atwood, [The Blind Assassin](#)



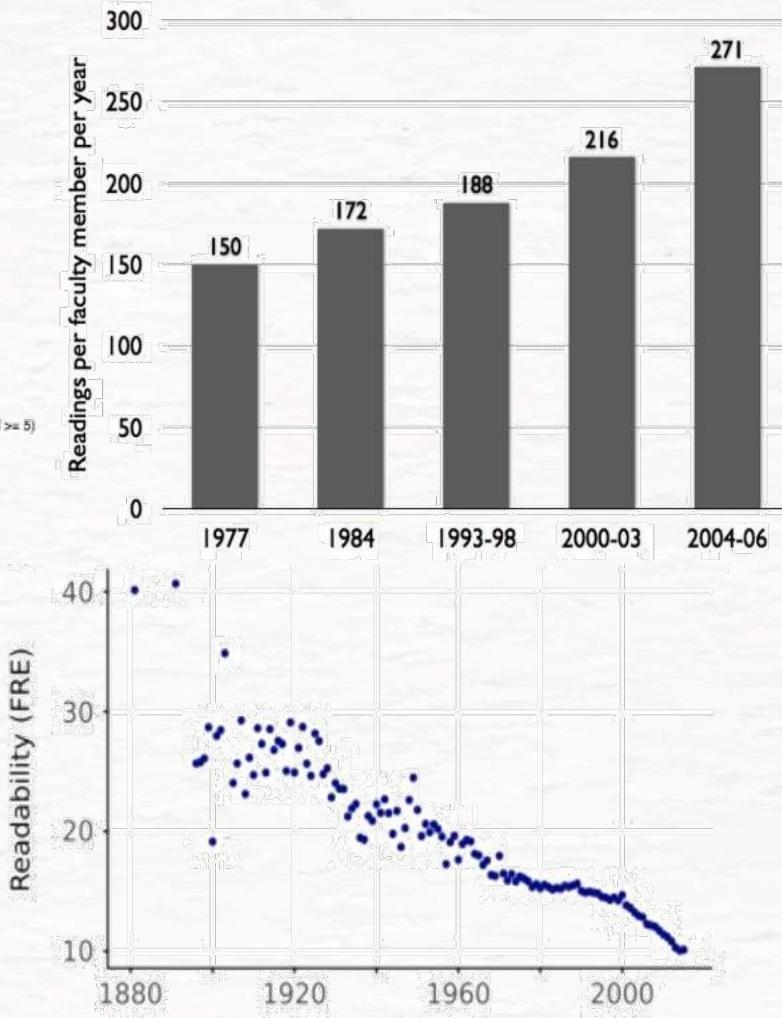
<https://qantarot.substack.com/p/chatgpt-and-galactica-are-taking>

The PDF crisis



<https://www.nature.com/news/the-top-100-papers-1.16224>

https://www.stm-assoc.org/2015_02_20_STM_Report_2015.pdf



<https://elifesciences.org/articles/27725>

<https://academic.oup.com/gigascience/article/8/6/giz053/5506490>



Research Debt

<https://distill.pub/2017/research-debt/>

The Scientific Paper Is Obsolete

Here's what's next.



**Genomic analysis of elongated skulls
and extensive female-biased immigration
in early Medieval Bavaria**

Krishna R. Veeramah^a, Andreas Rott^{b,1}, Melanie Groß^{c,1}, Lucy van Dorp^d, Saioa López^e, Karola Kirsanow^c, Christian Sell^c, Jens Blöcher^c, Daniel Wegmann^{f,g}, Vivian Link^{f,g}, Zuzana Hofmanová^{f,g}, Joris Peters^{b,h}, Bernd Trautmann^b, Anja Gairhosⁱ, Jochen Haberstroh^j, Bernd Päffgen^k, Garrett Hellenthal^d, Brigitte Haas-Gebhardⁱ, Michaela Harbeck^{b,2,3}, and Joachim Burger^{c,2,3}

^aDepartment of Ecology and Evolution, Stony Brook University, Stony Brook, NY 11794-5245; ^bState Collection for Anthropology and Palaeoanatomy, Bavarian Natural History Collections, 80333 Munich, Germany; ^cPalaeogenetics Group, Institute of Organismic and Molecular Evolution, Johannes Gutenberg University Mainz, 55099 Mainz, Germany; ^dUCL Genetics Institute, Department of Genetics, Evolution and Environment, University College London, WC1E 6BT London, United Kingdom; ^eCancer Institute, University College London, WC1E 6DD London, United Kingdom; ^fDepartment of Biology, University of Fribourg, 1700 Fribourg, Switzerland; ^gSwiss Institute of Bioinformatics, 1700 Fribourg, Switzerland; ^hArchaeoBioCenter and Institute for Palaeoanthropology, Domestication Research and the History of Veterinary Medicine, Ludwig-Maximilian University, 80539 Munich, Germany; ⁱBavarian State Collection for Anthropology and Palaeoanatomy, 80333 Munich, Germany; ^jDepartment of Archaeology, University of Bayreuth, 95444 Bayreuth, Germany; and ^kInstitute of Prehistory and Protohistory, University of Cologne, 50674 Cologne, Germany

Edited by Eske Willerslev, University of Copenhagen, Copenhagen, Denmark, and approved January 30, 2018 (received for review November 21, 2017)

Modern European genetic structure demonstrates strong correlations with geography, while genetic analysis of prehistoric humans has indicated at least two major waves of immigration to form in the 5th century AD, and that it emanated from a combination of the romanized local population of the border province of the former Roman Empire and immigrants from north of the Danube. Our whole-genome analysis of elongated skull individuals from early Medieval Bavaria shows that they were predominantly female, and that they had ancestry from both the local population and immigrants from the north. This pattern is consistent with the historical record, which indicates that the local population was heavily affected by female-biased immigration during the Migration Period. Our results also indicate that the genetic variation in the elongated skull individuals was similar to that found in modern Europeans, suggesting that they were part of the same genetic pool as the modern European population. This study provides new insights into the genetic history of Europe and the impact of immigration on the genetic variation of the local population.

<https://www.theatlantic.com/science/archive/2018/04/the-scientific-paper-is-obsolete/556676/>

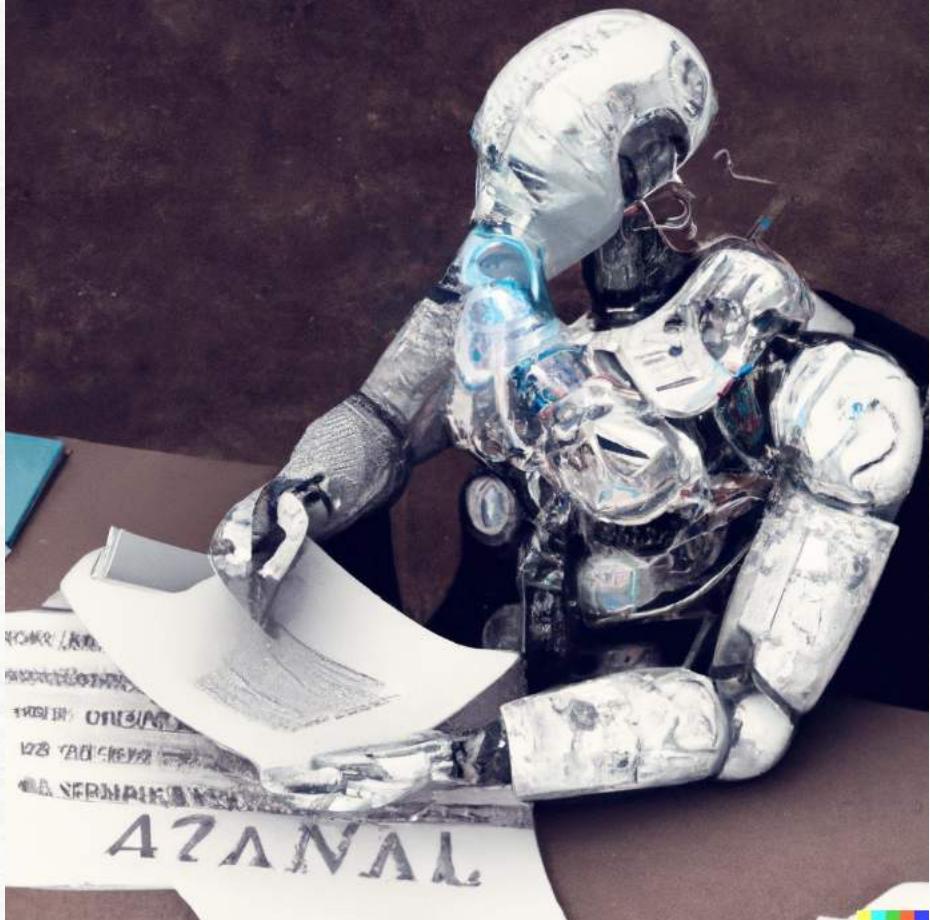
The Business of Extracting Knowledge from Academic Publications

December 7, 2021. Send your thoughts via [twitter](#) or [mail](#)

Close to nothing of what makes science actually work is published as text on the web

<https://markusstrasser.org/extracting-knowledge-from-literature/>

Taking papers to their logical conclusion



Prompt: Artificial intelligence eating a word salad and outputting an academic paper, digital art. AI image generated by DALL-E



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Can summarize academic literature, solve math problems, generate Wiki articles, write scientific code, annotate molecules and proteins, and more.

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<https://twitter.com/paperswithcode/status/1592546933679476736>



Yann LeCun @ylecun · Nov 15

This tool is to paper writing as driving assistance is to driving. It won't write papers automatically for you, but it will greatly reduce your cognitive load while you write them.



Yann LeCun @ylecun · Nov 15

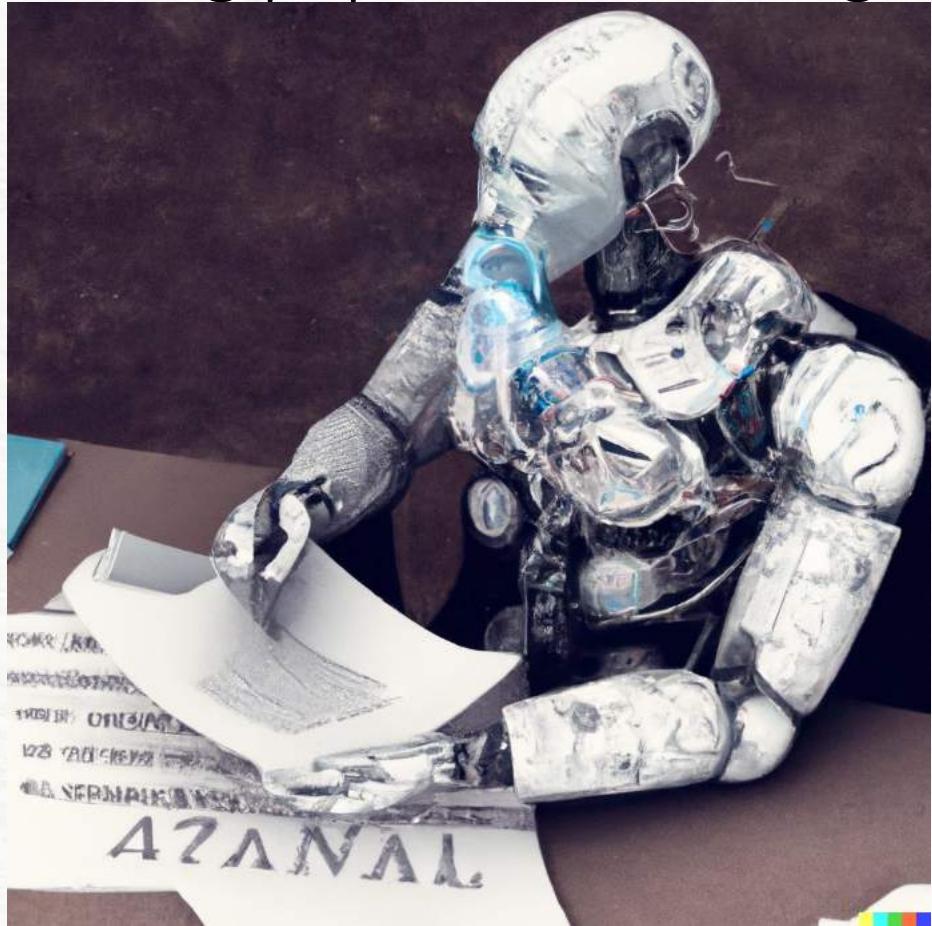
A Large Language Model trained on scientific papers. Type a text and galactica.ai will generate a paper with relevant references, formulas, and everything.

Amazing work by @MetaAI / @paperswithcode

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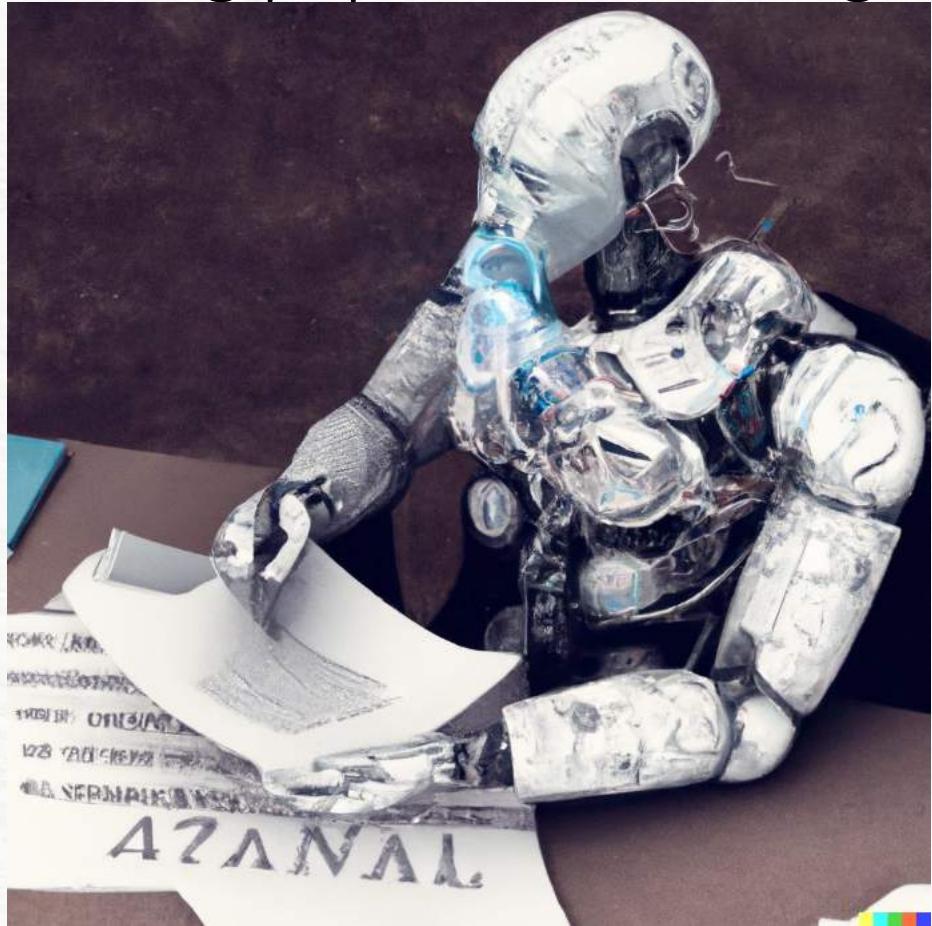
<https://twitter.com/paperswithcode/status/1592546933679476736>

 Michael Black
@Michael_J_Black

I asked **#Galactica** about some things I know about and I'm troubled. In all cases, it was wrong or biased but sounded right and authoritative. I think it's dangerous. Here are a few of my experiments and my analysis of my concerns. (1/9)

https://twitter.com/Michael_J_Black/status/1593133722316189696

Taking papers to their logical conclusion



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Vendor neutral MRI pulse sequences for qMRI

The vendor neutral MRI pulse sequences were developed by researchers at the National Institute of Standards and Technology (NIST) and the National Institutes of Health (NIH) as part of the Quantitative Imaging Network (QIN).

The sequences are designed to be compatible with both 1.5 T and 3 T scanners from multiple vendors and to facilitate the measurement of relaxation times and proton density (T1, T2, and PD) in the brain.

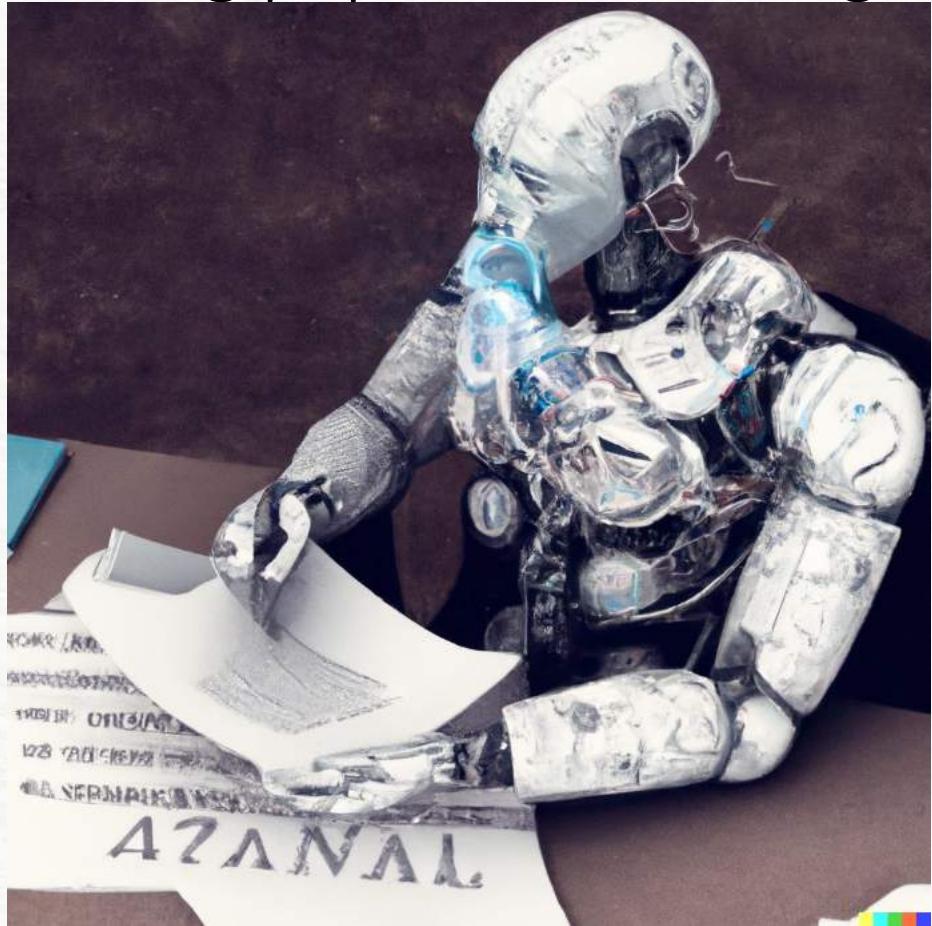
The pulse sequences and processing software are available for free download and use.

The sequences and processing software were designed to be used with phantoms, but can also be used for *in vivo* studies.

This sequence repository can be used for both research and educational purposes.

For more information on the sequences, please refer to the pulse sequence references listed below.

Taking papers to their logical conclusion



Prompt: Artificial intelligence eating a word salad and outputting an academic paper, digital art. AI image generated by DALL-E

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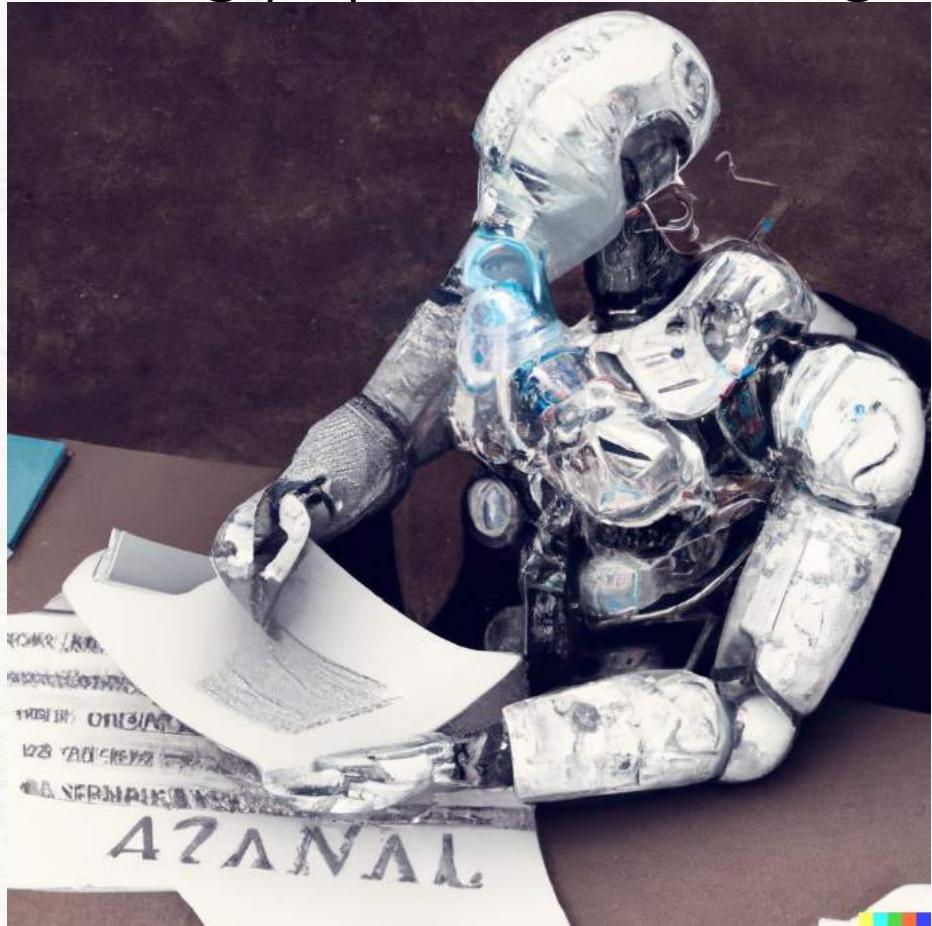
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<https://twitter.com/paperswithcode/status/1592546933679476736>

 Bojan Tunguz ✅
@tunguz

I believe that scientific papers are vestiges of the bygone era, especially in the fields of DS/ML/AI. IMHO, ML researchers should skip paper/arXiv altogether. Just publish the code and your data. Explain what code does. Publish a few benchmarking scripts. That's it.

Taking papers to their logical conclusion



Prompt: Artificial intelligence eating a word salad and outputting an academic paper, digital art. AI image generated by DALL-E

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<https://twitter.com/paperswithcode/status/1592546933679476736>

 vitalik.eth ✅
@VitalikButerin

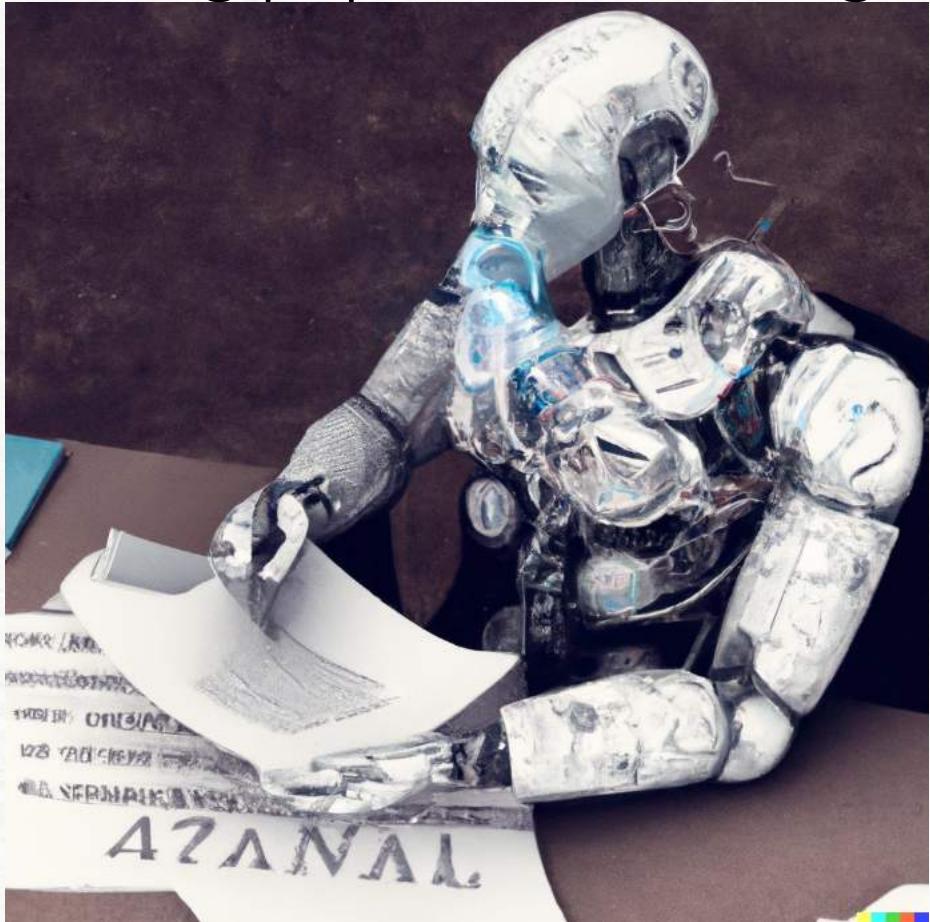
Replies to @tylervcowen

My theory is different: the AI-driven commoditization of articulateness in elite registers of the English language is going to degrade its value, and by extension further degrade the social status of those who depend on it as a status marker.

9:48 AM · Nov 23, 2022

<https://twitter.com/VitalikButerin/status/1595429084116520962>

Taking papers to their logical conclusion



Prompt: Artificial intelligence eating a word salad and outputting an academic paper, digital art. AI image generated by DALL-E

This is not necessarily a bad thing. As science is getting more computational, papers will either evolve or die. Either way, the academic article of the future will be nothing like the 10-page word salads that AI now so successfully emulates. The question is what will become of the academic paper once you take away the prose?

<https://qantarot.substack.com/p/chatgpt-and-galactica-are-taking>

WHAT REMAINS WHEN YOU TAKE AWAY THE PROSE?

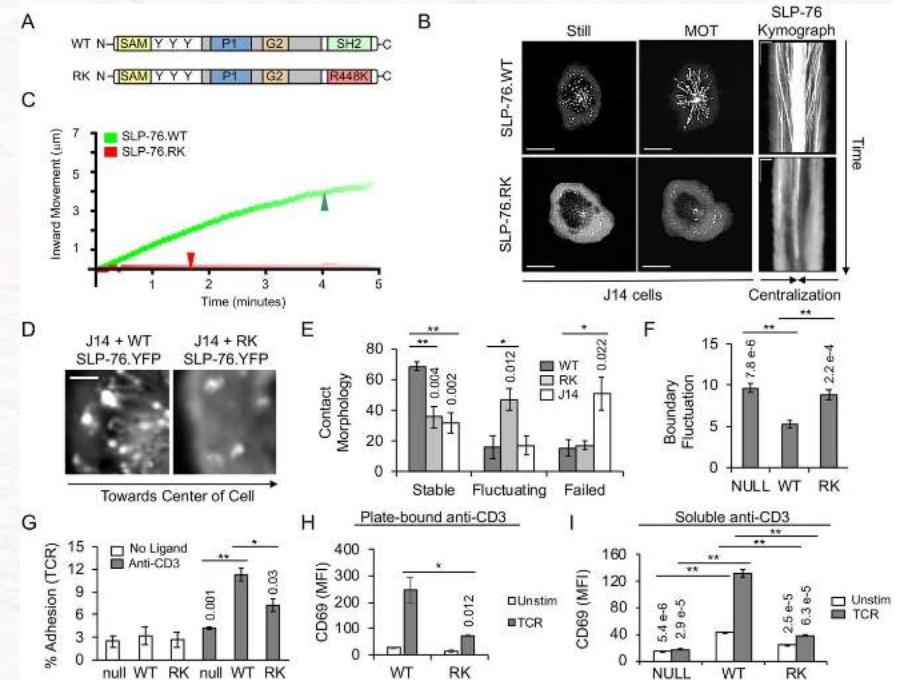


FIGURES



Nicolas Andry
L'orthopédie (1741)

Slide courtesy of Agâh Karakuzu



A typical figure in Nature



THE DATA

THE CODE

**THE
PROVENANCE**

Research Papers Used to Have Style. What Happened?

A Brief History of Scientific Writing



Roger's Bacon

Writes Secretorum · [Subscribe](#)

Aug 11

26

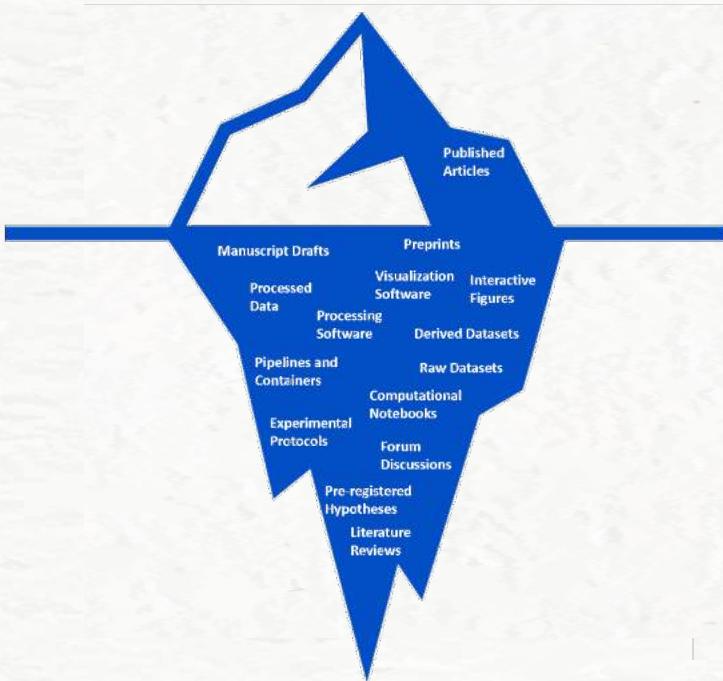
9



All this to say: Scientists in the pre-modern era wrote freely, despite calls to do away with that freedom. At some point, narrative and literary styles vanished and were replaced with rigid formats and impoverished prose. The question now is: **Have we gone too far in removing artistry from scientific writing?**

<https://newscience.substack.com/p/scientific-styles>

An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. The actual scholarship is the **complete software development environment** and the **complete set of instructions** which generated the figures.



-- Buckheit and Donoho
WaveLab and Reproducible Research, 1995

An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. The actual scholarship is the **complete software development environment** and the **complete set of instructions** which generated the figures.

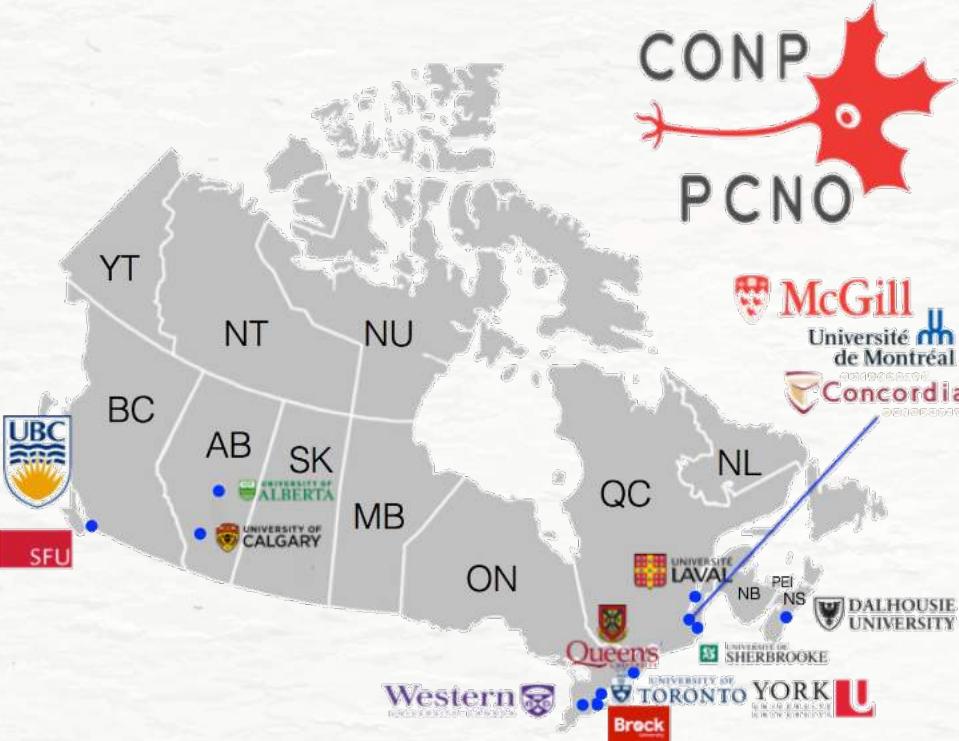


-- Buckheit and Donoho
WaveLab and Reproducible Research, 1995

The Canadian Open Neuroscience Platform: Catching Up to Plan S and Going Further

Posted August 15, 2019 by PLOS in Advocacy, Guest blog, In the News, Open Access, Open Science, Publishing

Guest Authors: Dylan Roskams-Edris (Open Science Consultant and Invited Scholar with the Center for Genomics and Policy,), JB Poline (Associate Professor, Neurology and Neurosurgery, McGill University), and Nikola Stikov (Associate Professor, Department of Electrical Engineering, Polytechnique Montreal).



Twitter: @NeuroLibre



Scientific director: Alan Evans

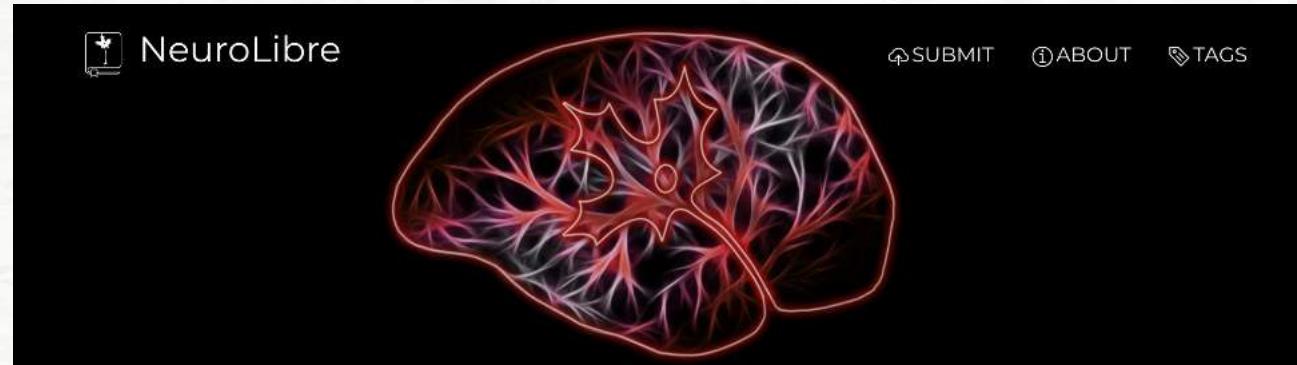
Steering Committee



Canadian Open
Neuroscience Platform



Reproducible research



<https://neurolibre.org>

PLOS COMPUTATIONAL BIOLOGY

OPEN ACCESS

PERSPECTIVE

Beyond advertising: New infrastructures for publishing integrated research objects

Elizabeth DuPre , Chris Holdgraf, Agah Karakuzu, Loïc Tetrel, Pierre Bellec, Nikola Stikov, Jean-Baptiste Poline 



NeuroLibre

NeuroLibre : A preprint server for full-fledged
reproducible neuroscience

AUTHORS

Agah Karakuzu, Elizabeth DuPre, Loic Tetrel, Patrick Bermudez, Mathieu Boudreau, Mary Chin, Jean-Baptiste Poline, Samir Das, Pierre Bellec, Nikola Stikov



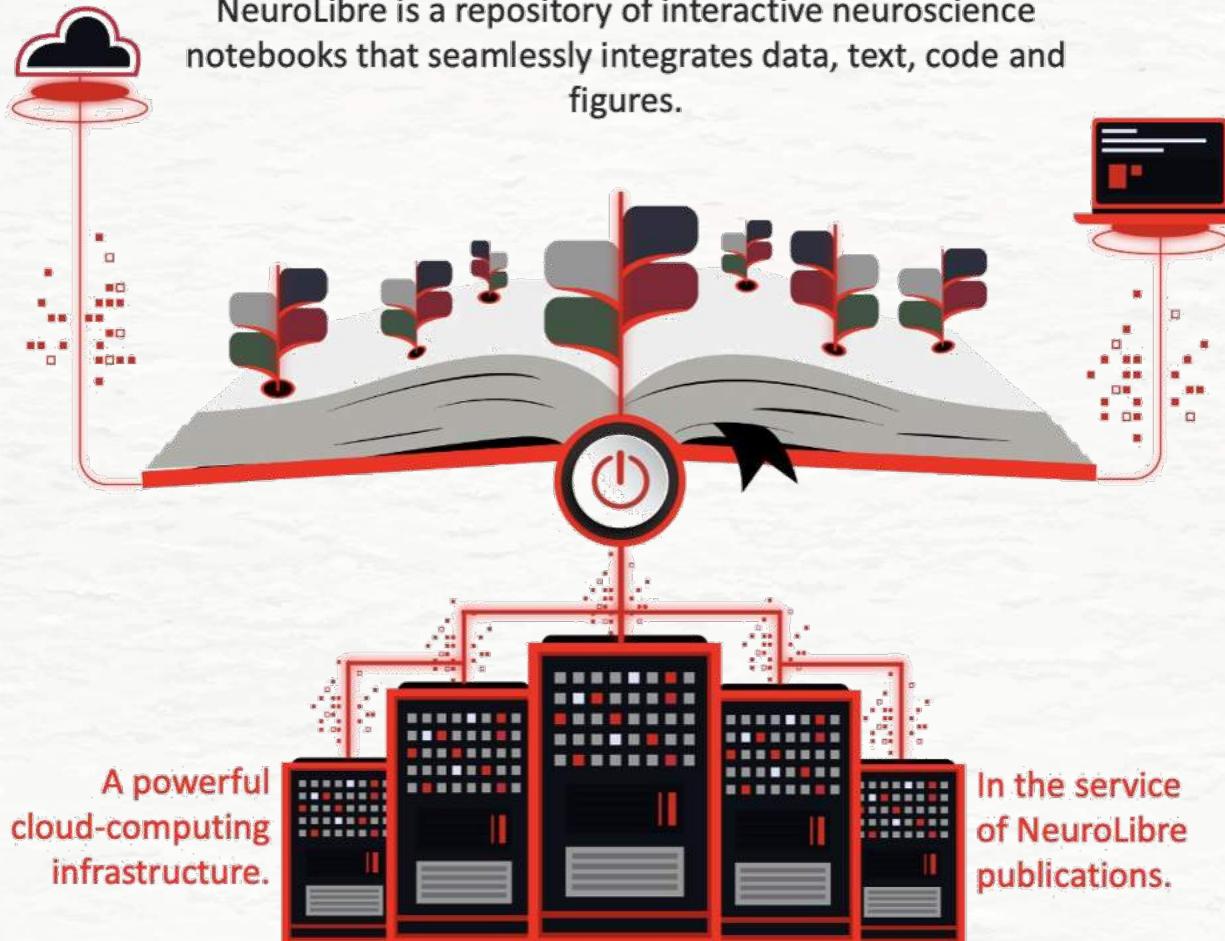
Agah Karakuzu





Agah Karakuzu

NeuroLibre is a repository of interactive neuroscience notebooks that seamlessly integrates data, text, code and figures.





Layer 1: A PDF compatible document

<https://neurolibre.org/papers/10.55458/neurolibre.00019>

The steady-state longitudinal magnetization of an inversion recovery experiment can be derived from the Bloch equations for the pulse sequence $\{\theta_{180} - TI - \theta_{90} - (TR-TI)\}$, and is given by:

$$M_z(TI) = M_0 \frac{1 - \cos(\theta_{180}) e^{-\frac{TR}{T_1}} - [1 - \cos(\theta_{180})] e^{-\frac{TI}{T_1}}}{1 - \cos(\theta_{180}) \cos(\theta_{90}) e^{-\frac{TR+TI}{T_1}}} \quad (1)$$

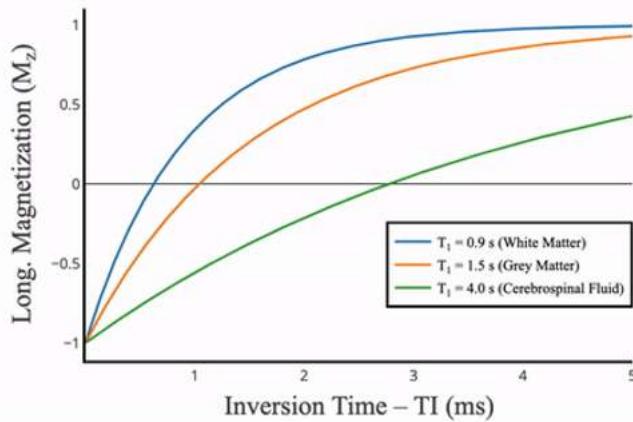
where M_z is the longitudinal magnetization prior to the θ_{90} pulse. If the in-phase real signal is desired, it can be calculated by multiplying Eq. 1 by $k \sin(\theta_{90}) e^{-TE/T_2}$, where k is a constant. This general equation can be simplified by grouping together the constants for each measurements regardless of their values (i.e. at each TI , same TE and θ_{90} are used) and assuming an ideal inversion pulse:

$$M_z(TI) = C(1 - 2e^{-\frac{TI}{T_1}} + e^{-\frac{TR}{T_1}}) \quad (2)$$

where the first three terms and the denominator of Eq. 1 have been grouped together into the constant C . If the experiment is designed such that TR is long enough to allow for full relaxation of the magnetization ($TR > 5T_1$), we can do an additional approximation by dropping the last term in Eq. 2:



Layer 2: Dynamic figures



Practically, Eq. 1 is the better choice for simulating the signal of an inversion recovery experiment, as the TRs are often chosen to be greater than $5T_1$ of the tissue-of-interest, which rarely coincides with the longest T_1 present (e.g. TR may be sufficiently long for white matter, but not for CSF which could also be present in the volume). Equation 3 also assumes ideal inversion pulses, which is rarely the case due to slice profile effects. Figure 3 displays the inversion recovery signal magnitude (complete relaxation normalized to 1) of an experiment with TR = 5 s and T_1 values ranging between 250 ms to 5 s, calculated using both equations.

Figure 3. Signal recovery curves simulated using Eq. 3 (solid) and Eq. 1 (dotted) with a TR = 5 s for T_1 values ranging between 0.25 to 5 s.

(View simulation code)

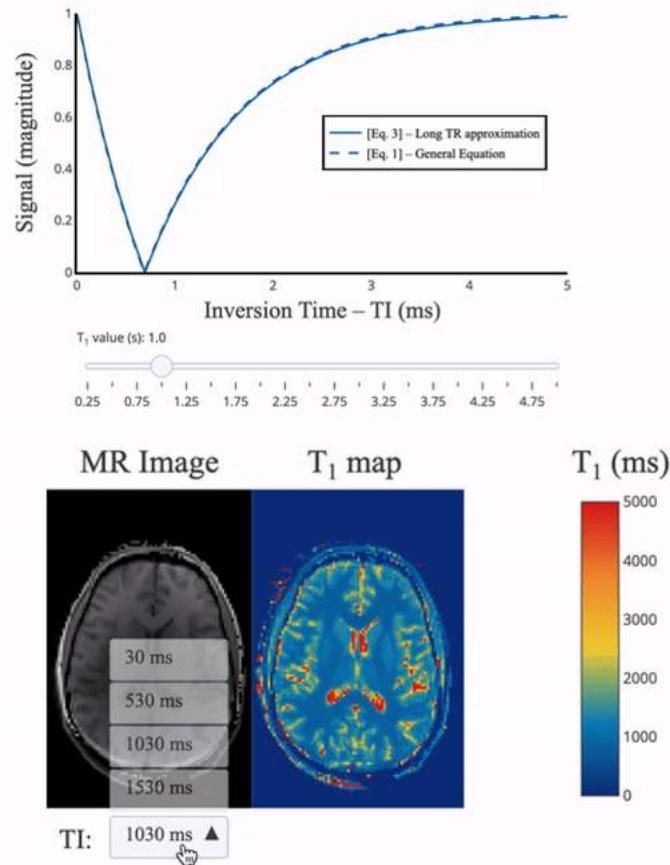
<https://neurolibre.org/papers/10.55458/neurolibre.00019>

You can observe the actual data points



NeuroLibre

Layer 3: Interactivity



<https://neurolibre.org/papers/10.55458/neurolibre.00019>

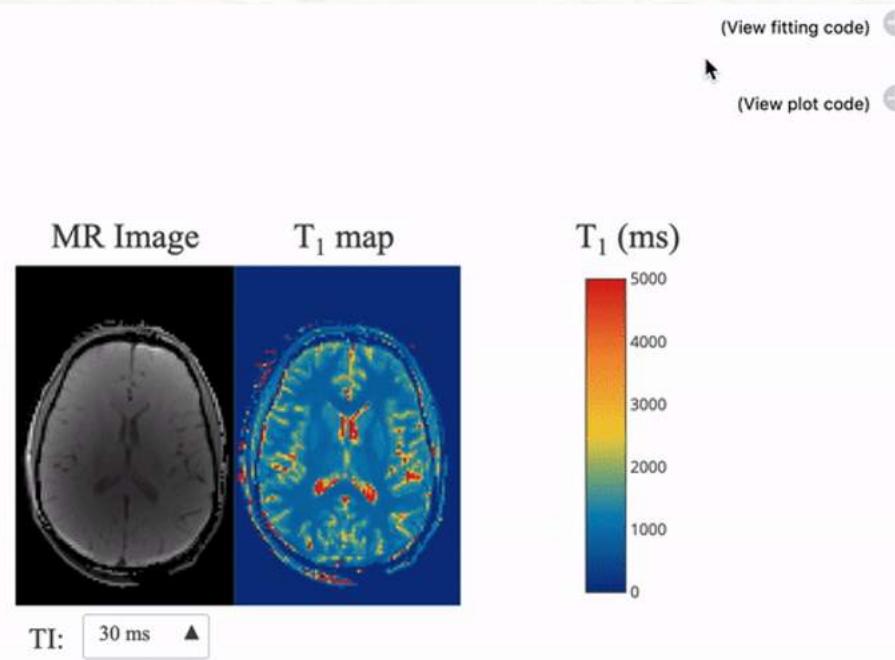
You can
explore the
phenomenon.

You can
interact with
the real-world data.



NeuroLibre

Layer 4: Transparency



You can
SEE THE CODE
that
generates the
outputs.

<https://neurolibre.org/papers/10.55458/neurolibre.00019>



Layer 5:

Reproducibility

Practically, Eq. 1 is the better choice for simulating the signal of an inversion recovery experiment, as the TRs are often chosen to be greater than $5T_1$ of the tissue-of-interest, which rarely coincides with the longest T_1 present (e.g. TR may be sufficiently long for white matter, but not for CSF which could also be present in the volume). Equation 3 also assumes ideal inversion pulses, which is rarely the case due to slice profile effects. Figure 3 displays the inversion recovery signal magnitude (complete relaxation normalized to 1) of an experiment with TR = 5 s and T_1 values ranging between 250 ms to 5 s, calculated using both equations.

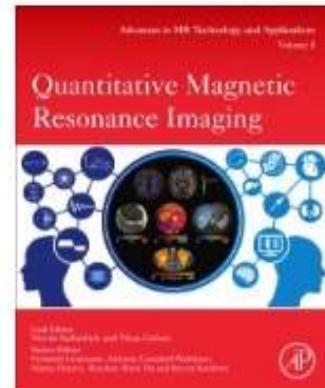
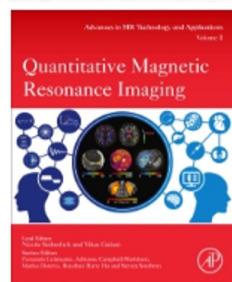
Figure 3. Signal recovery curves simulated using Eq. 3 (solid) and Eq. 1 (dotted) with a TR = 5 s for T_1 values ranging between 0.25 to 5 s.

```
%use octave  
  
% Verbosity level 0 overrides the disp function and supresses warnings.  
% Once executed, they cannot be restored in this session  
% (kernel needs to be restarted or a new notebook opened.)  
VERBOSITY_LEVEL = 0;  
  
if VERBOSITY_LEVEL==0
```

(View plot code)

You can
RUN the code that
generates the
outputs.

<https://neurolibre.org/papers/10.55458/neurolibre.00019>



Quantitative Magnetic Resonance Imaging, Volume 1

1st Edition

Q

 Write a review

Editors: Nicole Seiberlich, Vikas Gulani, Adrienne Campbell-Washburn, Steven Sourbron, Mariya Ivanova Doneva, Fernando Calamante, Houchun Harry Hu

Paperback ISBN: 9780128170571

Imprint: Academic Press

Published Date: 27th November 2020

Page Count: 1092

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^aSections 2.2–2.5 in this book chapter have been previously published under a creative commons license on the qMRLab blog. Visit the original blog posts for interactive versions of the figures: <https://qmrlab.org/jekyll/2018/10/23/T1-mapping-inversion-recovery.html>.
<https://qmrlab.org/jekyll/2018/12/11/T1-mapping-variable-flip-angle.html>.
<https://qmrlab.org/2019/04/08/T1-mapping-mp2rage.html>.
<https://qmrlab.org/2019/04/09/T1-mapping-t1rho.html>.



eLife

<https://elifesciences.org/61523>

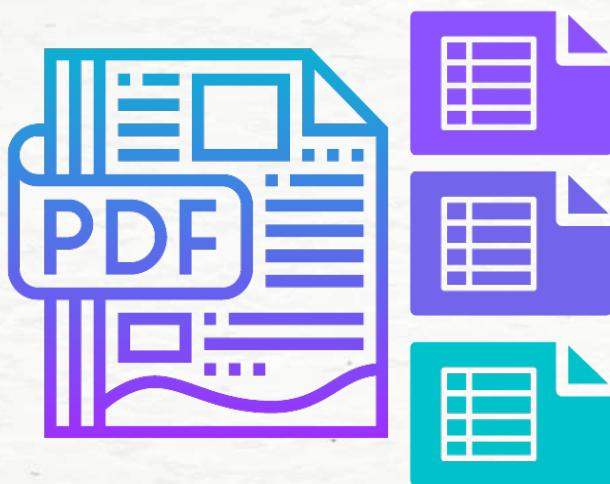
Online-executable interactive figures



<https://neurolibre.github.io/myelin-meta-analysis>

Mancini and Karakuzu et al. (2020) eLife 10.7554/eLife.61523

Myelin meta-analysis



Spreadsheets in a PDF?

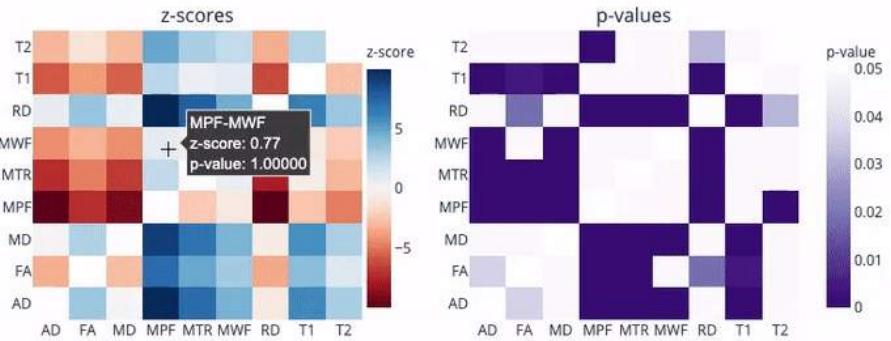
Figure 6

To visually represent these results, we used two heatmaps, one for the z-scores and one for the p-values: each element refers to the comparison between the measure on the x axis and the one on the y axis.

Figure 6

Click to show +

Figure 6: Statistical pairwise comparisons between R^2 estimates



Mancini, Karakuzu et al. 2020, [10.7554/eLife.61523](https://doi.org/10.7554/eLife.61523)

NeuroLibre
Interactive meta-analysis



Magnetic Resonance in Medicine



Mathieu Boudreau



EDITORIAL | Free Access

Reproducibility and the future of MRI research

Nikola Stikov , Joshua D. Trzasko, Matt A. Bernstein

First published: 10 August 2019 | <https://doi.org/10.1002/mrm.27939>

complex. For a truly international journal like MRM, it will be difficult to find a one-size-fits-all answer.

However, few would disagree that reproducibility accelerates scientific progress, as well as its translation to clinical practice. As tectonic technological changes shape research in the 21st century, we need to be careful not to damage its foundations, and instead focus on aspects of open science where we can reach consensus. The open-science movement needs to pick its battles. The battle for reproducible research is one that it cannot afford to lose.



EDITORIAL | Free Access

On the open-source landscape of *Magnetic Resonance in Medicine*

Mathieu Boudreau , Nikola Stikov, Peter Jezzard

First published: 28 July 2022 | <https://doi.org/10.1002/mrm.29366>

% of total papers that shared...	2019	2020	2021
code OR data	11	14	31
code	9	12	24
data	5	6	18

MAGNETIC RESONANCE IN MEDICINE



ISMRM

ONE
COMMUNITY
FOR CLINICIANS
AND SCIENTISTS

Whole-brain B₁-mapping using three-dimensional DREAM

Content

- Introduction
- Abstract
- Theory
- Code Example

Plots

- 2D slices
- 3D figure
- Multiple Views

GitHub repository

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Axial view slices

Phase unwrapping based on integrated Fourier-domain and image-domain analyses

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Phase Unwrapping

Time-optimized 4D phase contrast MRI with real-time convex optimization of gradient waveforms and fast excitation methods

Content

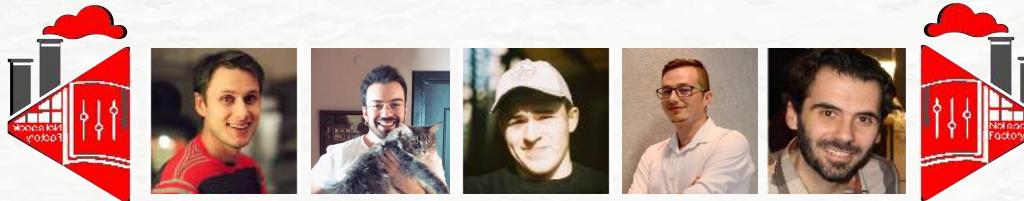
- Introduction
- Abstract
- Example

Original code

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```
# Repeat with no pms
trax = 100.0
N = 29 # Shorter
I = 0
d_M1 = E[1] + r_init_m1 + m1_shift
G_resid = gopt.optimize(d_N0, d_M1, dt=dt, n_it = n_it, cushion = cushion)
```

0.000536932276673977



NeuroLibre
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MRI Online Course

Educational Content

Interactive Jupyter Book GitHub Code

Quantitative T1 mapping

Educational Content

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Image processing with Spinal Cord Toolbox (SCT)

Educational Content

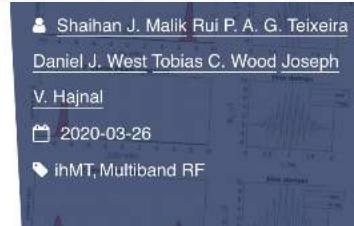
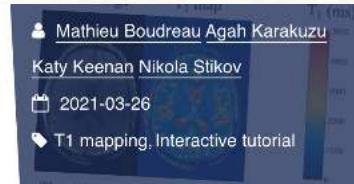
Interactive Jupyter Book GitHub Code

Steady-state imaging with inhomogeneous magnetization transfer contrast using multiband radiofrequency pulses

Abstract

Interactive Jupyter Book GitHub Code

MRM 10.1002/mrm.27984



A gradient optimization toolbox for general purpose time-optimal MRI gradient waveform design

Abstract

Interactive Jupyter Book GitHub Code

MRM 10.1002/mrm.28384

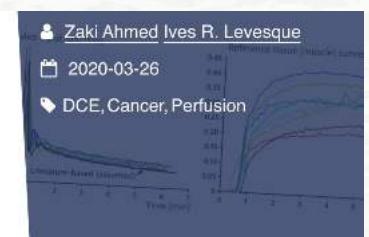


Pharmacokinetic modeling of dynamic contrast-enhanced MRI using a reference region and input function tail

Abstract

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MRM 10.1002/mrm.27913



Time-optimized 4D phase contrast MRI with real-time convex optimization of gradient waveforms and fast excitation methods

Abstract

Interactive Jupyter Book GitHub Code

MRM 10.1002/mrm.27716



The use of Fourier-domain analyses for unwrapping phase images of low SNR

Abstract

Interactive Jupyter Book GitHub Code

MRM 10.1002/mrm.27719



<https://ismrm.github.io/mrpub/>

PLOS COMPUTATIONAL BIOLOGY

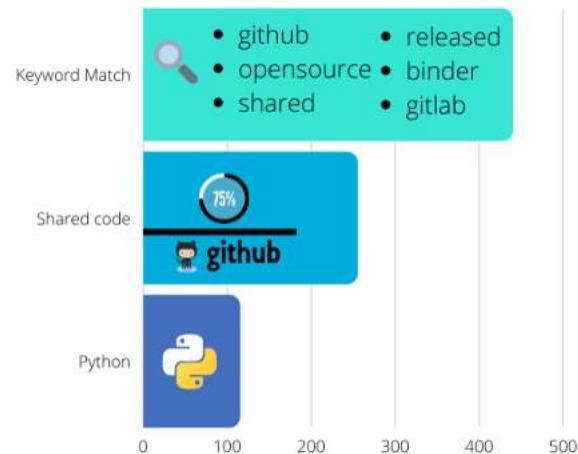
OPEN ACCESS

EDITORIAL

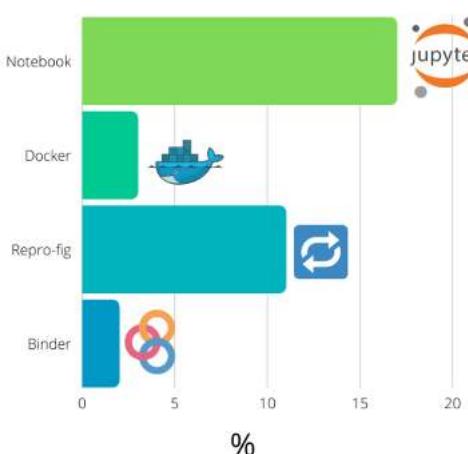
On the open-source landscape of PLOS Computational Biology

Mathieu Boudreau , Jean-Baptiste Poline, Pierre Bellec, Nikola Stikov

Published: February 11, 2021 • <https://doi.org/10.1371/journal.pcbi.1008725>



622 articles published in 2019





Bayes-optimal estimation of overlap between populations of fixed size

Content

Introduction

Abstract

Figures

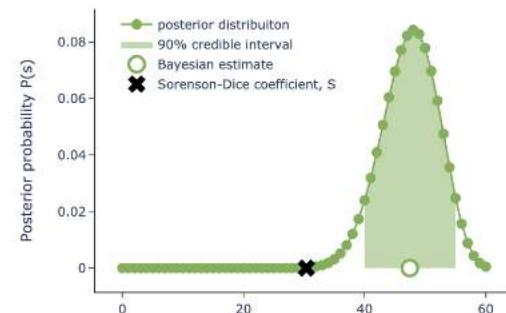
[Fig 2. Inference and uncertainty using the posterior](#)

[Fig 3. Bayesian repertoire overlap consistently estimates true overlap](#)

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true overlap open circle; Eq (7), and the interval accounting for at least 90% of the area under the posterior curve provides an equal-tailed 90% credible interval [shading; Eq (8)]. The \hat{S} estimate is shown for comparison [black cross; Eq (1)], and is typically less than or equal to \hat{s} .

Link to the published figure: <https://doi.org/10.1371/journal.pcbi.1006898.g002>



Efficient neural decoding of self-location with a deep recurrent network

Content

Introduction

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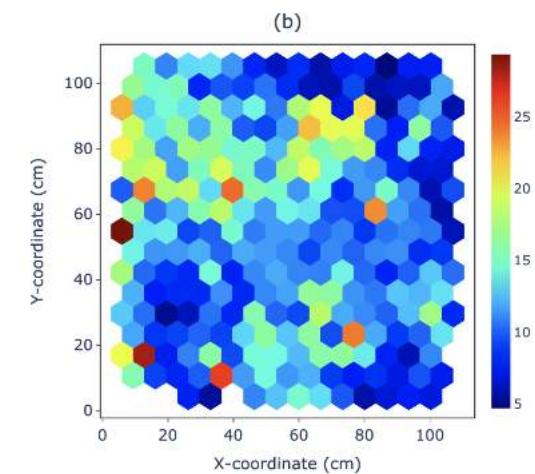
[Fig 1. Accurate decoding of position with a RNN](#)

[Fig 2. Comparison of RNN and Bayesian decoders](#)

[Fig 3. Spatial decoding across animals in 2D and 1D environments](#)

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(b) Mean prediction error for different regions for R2192



Crowdsourcing the research article of the future

I will only publish one research article this year. This is the one!



NIKOLA STIKOV
OCT 20, 2023



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NOTE: Mathieu Boudreau, Agâh Karakuzu and Pierre Bellec contributed to this post

<https://preprint.neurolibre.org/10.55458/neurolibre.00014/>



“Had he been a lunatic or an intellectually honourable man who'd thought things through to their logical conclusion? And was there any difference?”

— Margaret Atwood, *Oryx and Crake*



<https://qantarot.substack.com/p/chatgpt-and-galactica-are-taking>



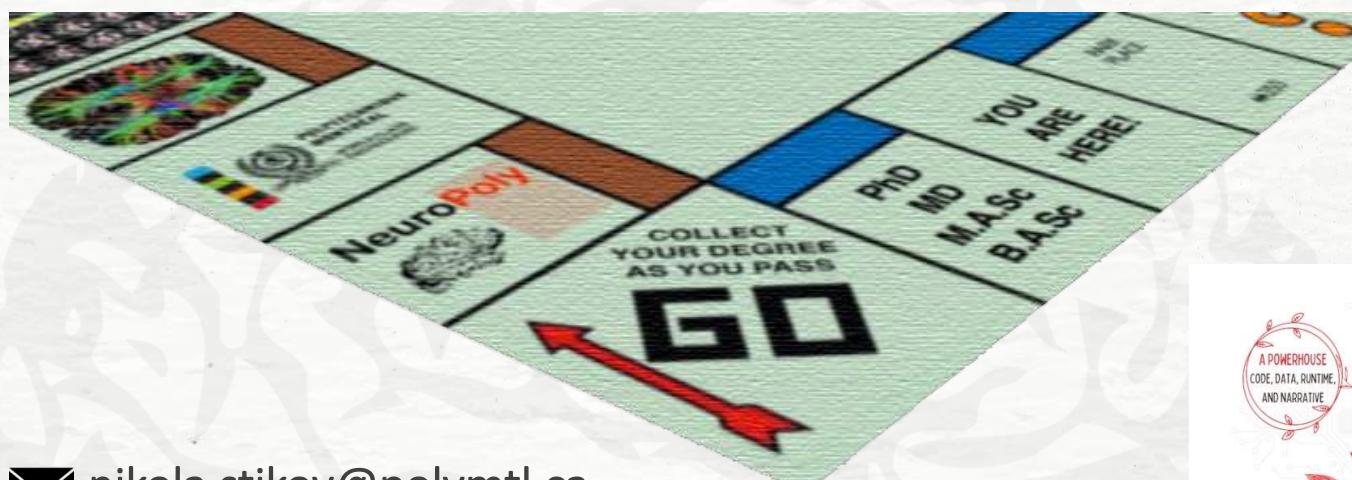
Thank you!

Agah Karakuzu
Mathieu Boudreau
Nadia Blostein
Julien Cohen-Adad
Eva Alonso Ortiz
Benjamin De Leener
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Kiril Zelenkovski
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Tanguy Duval
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Bruce Pike
Jennifer Campbell
Sridar Narayanan
Christine Tardif
Robert Brown
David Rudko
Robert Dougherty
Brian Wandell



Thank you!

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