

ETWORK NEURO SCIENCE

an open access 🔓 journal





Citation: Expert, P., Lord, L. D., Morten L. Kringelbach, M. L., & Petri, G. (2019). Editorial: Topological Neuroscience. Network Neuroscience, 3(3), 653-655 https://doi.org/10.1162/netn_e_00096

https://doi.org/10.1162/netn_e_00096

Received: 9 May 2019

Competing Interests: The authors have declared that no competing interests

Corresponding Author: Paul Expert paul.expert08@imperial.ac.uk

Handling Editor: **Olaf Sporns**

Copyright: © 2019 Massachusetts Institute of Technology **Published under a Creative Commons** Attribution 4.0 International (CC BY 4.0) license



FOCUS FEATURE: Topological Neuroscience

Editorial: Topological Neuroscience

Paul Expert^{1,2,3,4}, Louis-David Lord⁵, Morten L. Kringelbach^{5,6}, and Giovanni Petri^{7,8}

¹Department of Mathematics, Imperial College London, London, UK ²EPSRC Centre for Mathematics of Precision Healthcare, Imperial College London, London, UK ³Department of Neuroimaging, Institute of Psychiatry, Psychology and Neuroscience, Kings College London, London, UK ⁴Global Digital Health Unit, School of Public Health, Faculty of Medicine, Imperial College London, London, UK ⁵Department of Psychiatry, University of Oxford, Oxford, UK ⁶Center for Music in the Brain, Aarhus University, Aarhus, Denmark ⁷ISI Foundation, Turin, Italy ⁸ISI Global Science Foundation, New York, New York, USA

Keywords: Topological data analysis, Neuroscience, Multiple scales, Higher order interactions

ABSTRACT

Topology, in its many forms, describes relations. It has thus long been a central concept in neuroscience, capturing structural and functional aspects of the organization of the nervous system and their links to cognition. Recent advances in computational topology have extended the breadth and depth of topological descriptions. This Focus Feature offers a unified overview of the emerging field of topological neuroscience and of its applications across the many scales of the nervous system from macro-, over meso-, to microscales.

From the early drawings of Ramon y Cajal to today, topological descriptions have played a central role in neuroscience. In recent years, thanks to advancements in both mathematical tools and data availability, the range and diversity of such descriptions are expanding rapidly, spanning theoretical, computational, and experimental approaches to brain connectivity. This Focus Feature on "Topological Neuroscience" aims at presenting the breadth of applicability of topological data analysis (TDA) methods in neuroscience across scales and modalities.

Computational topology offers new frameworks for both the analytical description and the understanding of brain function. A common denominator to these new tools is their ability to find meaningful simplifications of high-dimensional data. As such, TDA aims to capture mesoscale patterns of disconnectivity and explicitly encode higher order interactions, that is, interactions between more than two regions or components (Giusti, Ghrist, & Bassett, 2016). In addition to the description of the shape of spaces derived from neuroimaging data, topology might play an even more fundamental role in brain organization, as indicated by mounting evidence for how the brain encodes space and memories (Dabaghian, Mémoli, Frank, & Carlsson, 2012). Finally, the intrinsic robustness of TDA methods and the features they identify make them powerful candidates not only to characterize healthy brain function but also potentially as biomarkers for disease (Romano et al., 2014).

Recent seminal research has shown the potential and impact of topological approaches. Topological differences have been found at the population and individual levels in functional connectivity (Lee, Chung, Kang, Kim, & Lee, 2011; Lee, Kang, Chung, Kim, & Lee, 2012) in both healthy and pathological subjects. Higher dimensional topological features have been employed to detect differences in brain functional configurations in neuropsychiatric disorders and altered states of consciousness relative to controls (Chung et al., 2017; Petri et al., 2014), and to characterize intrinsic geometric structures in neural correlations (Giusti, Pastalkova, Curto, & Itskov, 2015; Rybakken, Baas, & Dunn, 2017). Structurally, persistent homology techniques have been used to detect nontrivial topological cavities in white-matter networks (Sizemore et al., 2018), discriminate healthy and pathological states in developmental (Lee et al., 2017) and neurodegenerative diseases (Lee, Chung, Kang, & Lee, 2014), and also to describe the brain arteries' morphological properties across the lifespan (Bendich, Marron, Miller, Pieloch, & Skwerer, 2016). Finally, the properties of topologically simplified activity have identified backbones associated with behavioral performance in a series of cognitive tasks (Saggar et al., 2018).

This Focus Feature offers a unified overview of this emerging field of topological neuroscience and of its applications across many scales of the nervous system from macro-, over meso-, to microscales. First, Sizemore, Phillips-Cremins, Ghrist, and Bassett (2019) provide an accessible introduction to the language of topological data analysis and investigate its potential in structural and genetic connectivity datasets. Chung, Lee, DiChristofano, Ombao, and Solo (2019) focus instead on differences in whole-brain functional topology in a cohort of twins and propose a novel topological metric that captures the heritability of topological features. In the context of event-related fMRI, Ellis, Lesnick, Henselman-Petrusek, Keller, and Cohen (2019) investigate the feasibility of topological techniques for recovering signal representations under different conditions. At the mesoscopic scale, Babichev, Morozov, and Dabaghian (2019) propose a computational model to assess the effect of memory replays in parahippocampal networks on the development and stabilization of hippocampal topological maps of space. At an even smaller scale, Bardin, Spreemann, and Hess (2019) show that topological features of spiketrain data can be used to understand how individual neurons give rise to network dynamics, and hence to classify topologically such emergent behaviors. From a methodological point of view, Patania, Selvaggi, Veronese, Dipasquale, Expert, and Petri (2019) build topological gene expression networks that robustly capture the relationships between genetic pathways and brain function. Finally, Geniesse, Sporns, Petri, and Saggar (2019) present open-source tools designed to explore graphical representations of high-dimensional neuroimaging data extracted using topological data analysis at the individual level and without spatial nor temporal averaging.

It is now high time to put topological neuroscience center stage and to bring together the growing but often separate communities involved in applied topological analysis. Still, numerous challenges and questions remain before TDA methods become widely accepted and can come to realize their full potential. Notably, more research is needed both in terms of contextualization and functional interpretation of topological features (Lord et al., 2016; Verovsek, Kurlin, & Lesnik, 2017), and of scalability and computability of some of these features (Otter, Porter, Tillmann, Grindrod, & Harrington, 2017). However, there are already encouraging signs coming from academic conferences and schools in related fields (e.g., Netsci, Conference on Complex Systems, Applied Machine Learning Days), where tracks or satellites dedicated to TDA methods are already being organized. In this context, and considering that network-based methods sit in the larger realm of TDA, the journal Network Neuroscience is a natural venue to nurture and grow topological neuroscience in the coming years.

REFERENCES

- Babichev, A., Morozov, D., & Dabaghian, Y. (2019). Replays of spatial memories suppress topological fluctuations in cognitive map. *Network Neuroscience*, *3*(3), 707–724.
- Bardin, J. B., Spreemann, G., & Hess, K. (2019). Topological exploration of artificial neuronal network dynamics. *Network Neuroscience*, *3*(3), 725–743.
- Bendich, P., Marron, J. S., Miller, E., Pieloch, A., & Skwerer, S. (2016). Persistent homology analysis of brain artery trees. *Annals of Applied Statistics*, *10*(1), 198.
- Chung, M. K., Villalta-Gil, V., Lee, H., Rathouz, P. J., Lahey, B. B., & Zald, D. H. (2017). Exact topological inference for paired brain networks via persistent homology. In *International*

Network Neuroscience 654

- Conference on Information Processing in Medical Imaging (pp. 299–310).
- Chung, M. K., Lee, H., DiChristofano, A., Ombao, H., & Solo, V. (2019). Exact topological inference of the resting-state brain networks in twins. *Network Neuroscience*, *3*(3), 674–694.
- Dabaghian, Y., Mémoli, F., Frank, L., & Carlsson, G. (2012). A topological paradigm for hippocampal spatial map formation using persistent homology. *PLoS Computational Biology*, 8(8), e1002581.
- Ellis, C. T., Lesnick, M., Henselman-Petrusek, G., Keller, B., & Cohen, J. D. (2019). Feasibility of topological data analysis for event-related fMRI. *Network Neuroscience*, *3*(3), 695–706.
- Geniesse, C., Sporns, O., Petri, G., & Saggar, M. (2019). Generating dynamical neuroimaging spatiotemporal representations (DyNeuSR) using topological data analysis. *Network Neuroscience*, *3*(3), 763–778.
- Giusti, C., Ghrist, R., & Bassett, D. S. (2016). Two's company, three (or more) is a simplex. *Journal of Computational Neuroscience*, 41(1), 1–14.
- Giusti, C., Pastalkova, E., Curto, C., & Itskov, V. (2015). Clique topology reveals intrinsic geometric structure in neural correlations. *Proceedings of the National Academy of Sciences*, 112(44), 13455–13460.
- Lee, H., Chung, M. K., Kang, H., Kim, B.-N., & Lee, D. S. (2011). Discriminative persistent homology of brain networks. In 2011 IEEE International Symposium on Biomedical Imaging: From Nano to Macro (pp. 841–844).
- Lee, H., Chung, M. K., Kang, H., & Lee, D. S. (2014). Hole detection in metabolic connectivity of Alzheimer's disease using *k*-Laplacian. In *International Conference on Medical Image Computing and Computer-Assisted Intervention* (pp. 297–304).
- Lee, H., Kang, H., Chung, M. K., Kim, B.-N., & Lee, D. S. (2012).
 Persistent brain network homology from the perspective of dendrogram. *IEEE Transactions on Medical Imaging*, 31(12), 2267–2277.
- Lee, H., Kang, H., Chung, M. K., Lim, S., Kim, B.-N., & Lee, D. S. (2017). Integrated multimodal network approach to pet and MRI

- based on multidimensional persistent homology. *Human Brain Mapping*, 38(3), 1387–1402.
- Lord, L.-D., Expert, P., Fernandes, H. M., Petri, G., Van Hartevelt, T. J., Vaccarino, F., . . . Kringelbach, M. L. (2016). Insights into brain architectures from the homological scaffolds of functional connectivity networks. *Frontiers in Systems Neuroscience*, 10, 85.
- Otter, N., Porter, M. A., Tillmann, U., Grindrod, P., & Harrington, H. A. (2017). A roadmap for the computation of persistent homology. *EPJ Data Science*, 6(1), 17.
- Patania, A., Selvaggi, P. L., Veronese, M., Dipasquale, O., Expert, P., & Petri, G. (2019). Topological gene expression networks recapitulate brain anatomy and function. *Network Neuroscience*, 3(3), 744–762.
- Petri, G., Expert, P., Turkheimer, F., Carhart-Harris, R., Nutt, D., Hellyer, P. J., & Vaccarino, F. (2014). Homological scaffolds of brain functional networks. *Journal of The Royal Society Interface*, *11*(101), 20140873.
- Romano, D., Nicolau, M., Quintin, E.-M., Mazaika, P. K., Light-body, A. A., Cody Hazlett, H., . . . Reiss, A. L. (2014). Topological methods reveal high and low functioning neuro-phenotypes within fragile X syndrome. *Human Brain Mapping*, *35*(9), 4904–4915.
- Rybakken, E., Baas, N., & Dunn, B. (2017). Decoding of neural data using cohomological feature extraction. *arXiv preprint* arXiv:1711.07205.
- Saggar, M., Sporns, O., Gonzalez-Castillo, J., Bandettini, P. A., Carlsson, G., Glover, G., & Reiss, A. L. (2018). Towards a new approach to reveal dynamical organization of the brain using topological data analysis. *Nature Communications*, *9*(1), 1399.
- Sizemore, A. E., Giusti, C., Kahn, A., Vettel, J. M., Betzel, R. F., & Bassett, D. S. (2018). Cliques and cavities in the human connectome. *Journal of Computational Neuroscience*, *44*(1), 115–145.
- Sizemore, A. E., Phillips-Cremins, J., Ghrist, R., & Bassett, D. S. (2019). The importance of the whole: Topological data analysis for the network neuroscientist. *Network Neuroscience*, *3*(3), 656–673.
- Verovsek, S. K., Kurlin, V., & Lesnik, D. (2017). Higher-dimensional skeletonization problem. *arXiv:1701.08395*.

Network Neuroscience 655