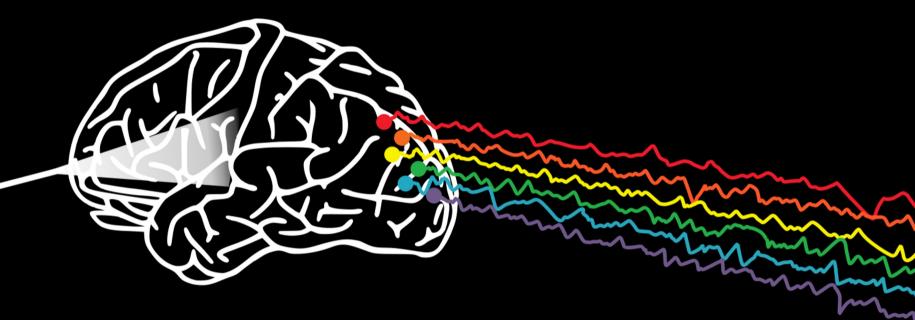
Studying the brain as a complex and nonlinear dynamical system with time-series analysis

CSYS5040 17 September 2024

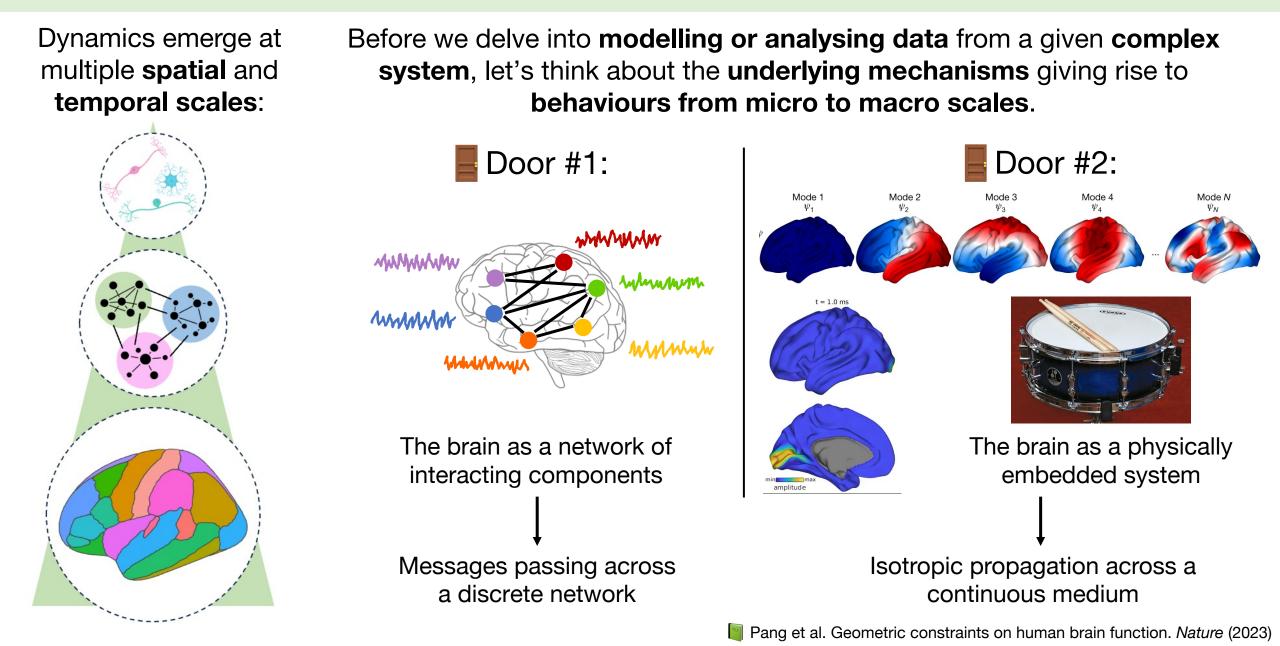


Annie G. Bryant PhD Candidate Dynamics and Neural Systems Lab

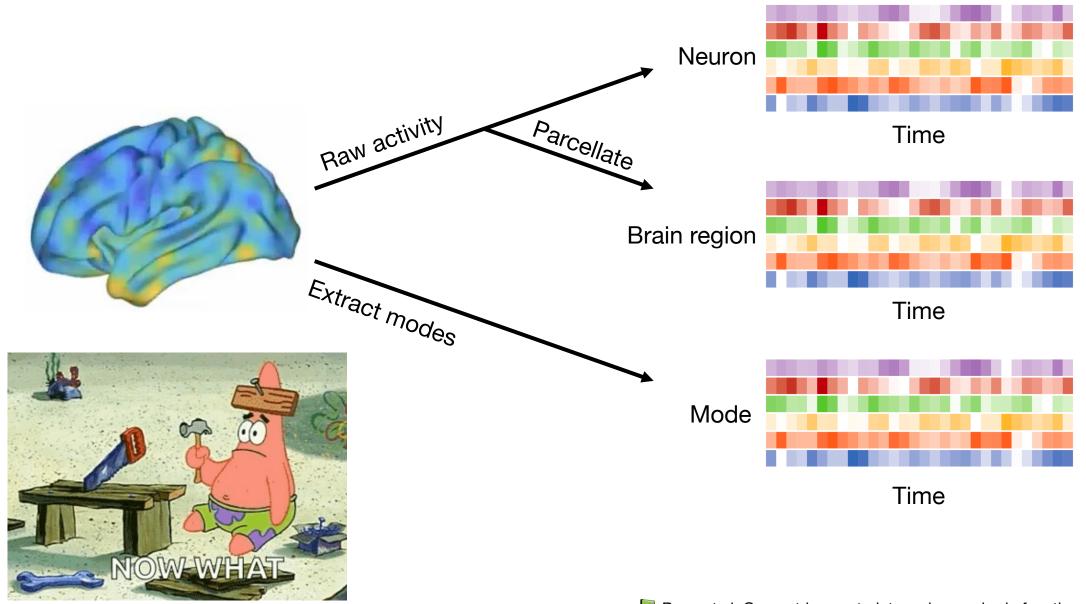




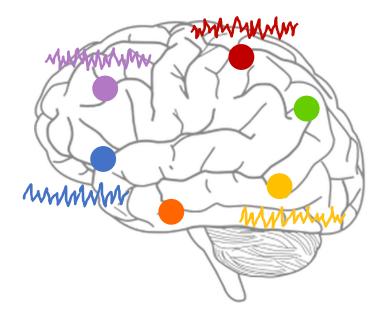
Mechanisms of brain activity across scales



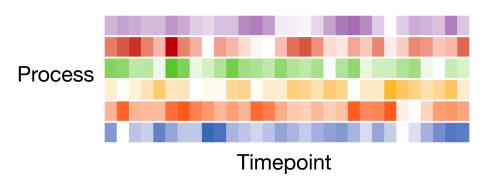
Different ways of representing brain dynamics



Not just the brain: time-varying complex systems are everywhere!



Multivariate time series (MTS)



City structure

City properties: density, traffic patterns, crime rates, culture

•

Economics

National economy: economic growth, recession

Physics



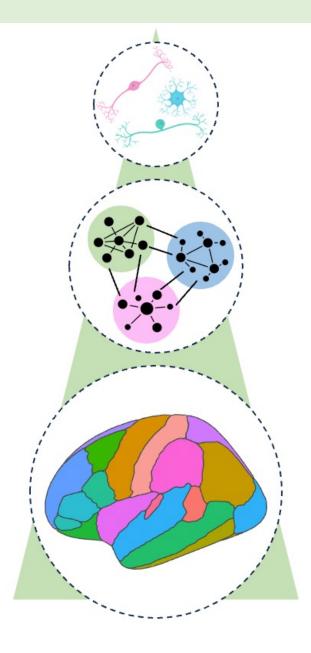
Fluid dynamics: vortices, turbulence

Social networks

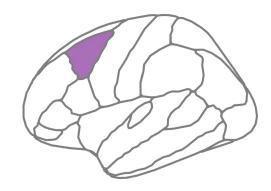


Facebook friends: community formation

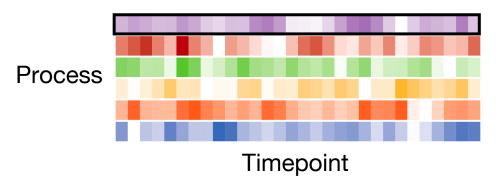
Treating the brain as a complex system of great biological interest



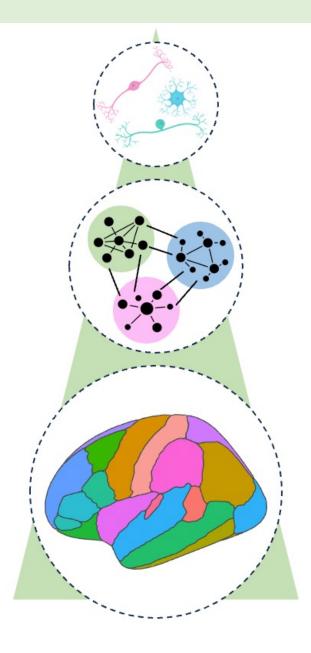


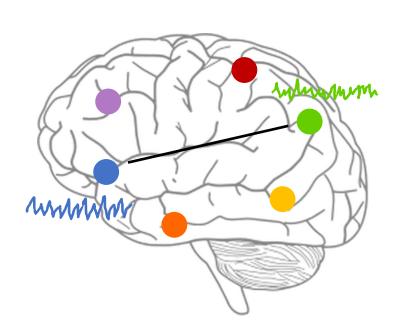


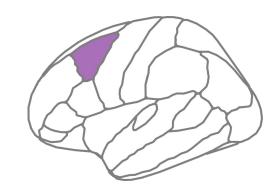
Multivariate time series (MTS)

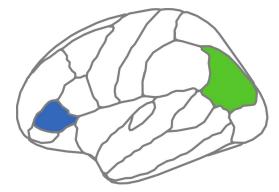


Treating the brain as a complex system of great biological interest

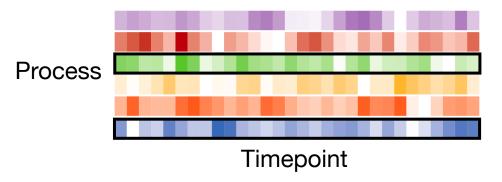




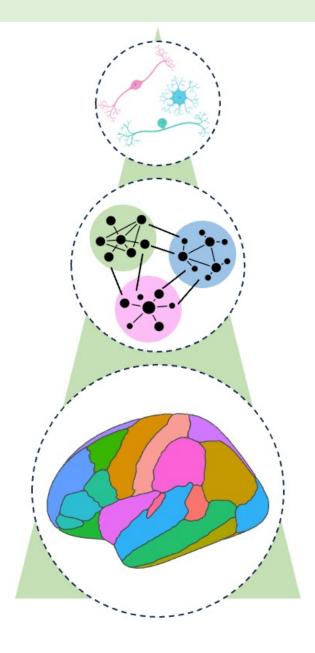


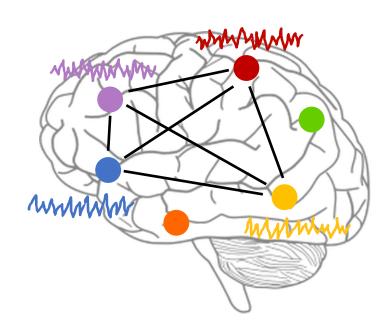


Multivariate time series (MTS)

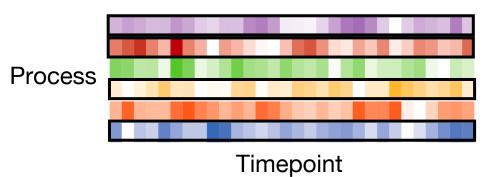


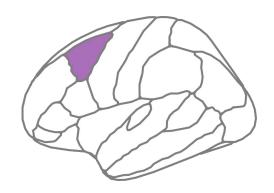
Treating the brain as a complex system of great biological interest

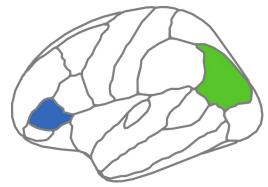


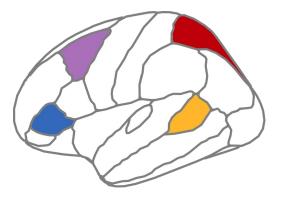


Multivariate time series (MTS)

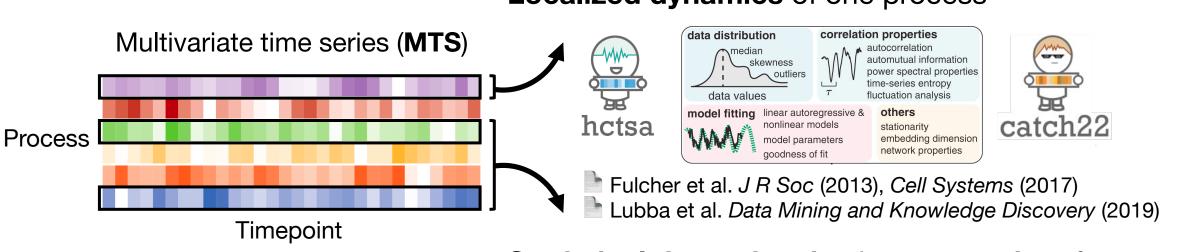






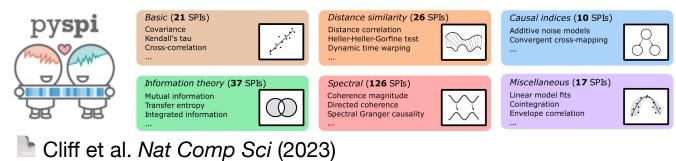


What does this generalized representation offer us?

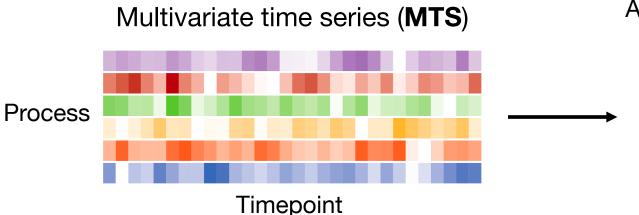


Localized dynamics of one process

Statistical dependencies between pairs of processes



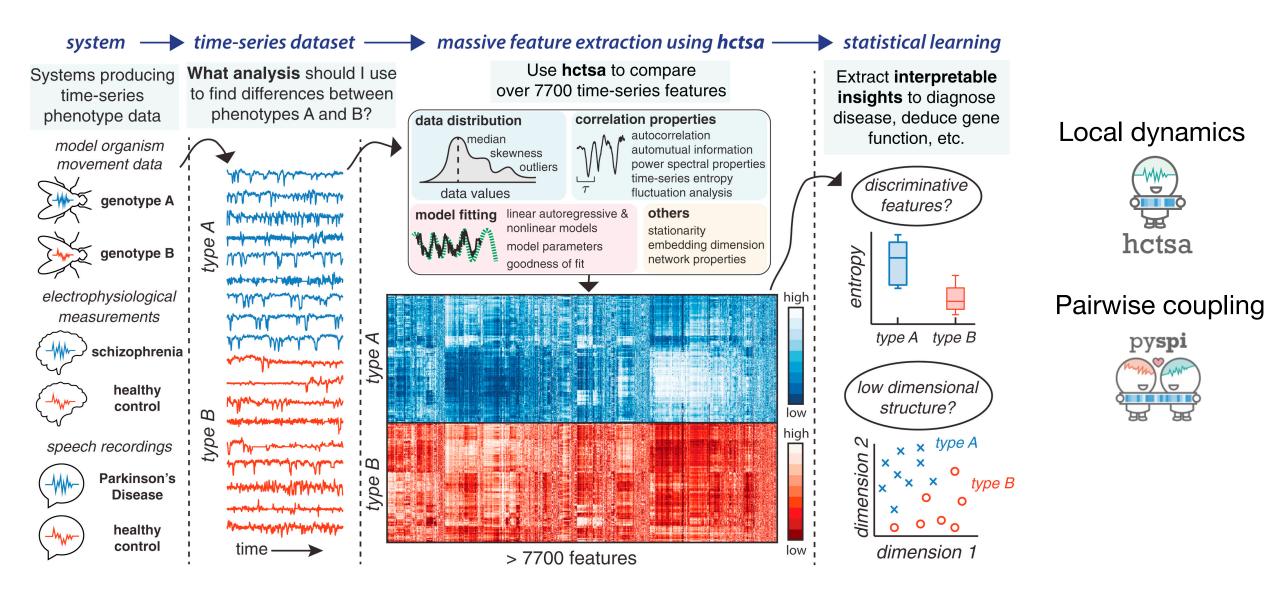
Quantifying the desired structure in a multivariate time series:



A set of statistical properties based on **clear scientific algorithms** and **interpretable theory**, which are **informative of interesting structure(s)** in our data

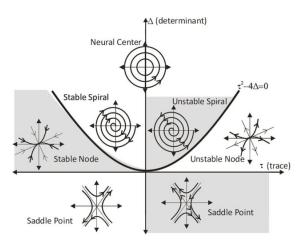
The highly comparative time-series feature analysis approach compares across a large library of scientific algorithms (spanning a large and interdisciplinary theoretical literature **[] [] [] [] []**)

This boils down to a common goal



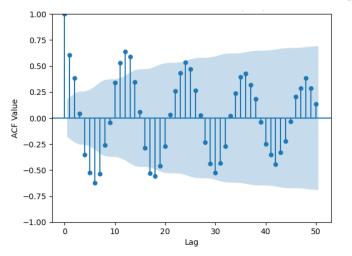
Fulcher & Jones. *hctsa*: A Computational Framework for Automated Time-Series Phenotyping Using Massive Feature Extraction. *Cell Systems* (2017)

hctsa and pyspi include both linear and nonlinear features



The arsenal of **linear time series analysis features** begin by **assuming a system with linear structure**, and we think about what such a linear system would do

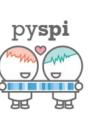
A linear system is **fully** captured by its **autocorrelation function (ACF)**





Local dynamics

ACF at different lags, local forecast based on rolling average, AR models, basic distributional properties (mean, variance)

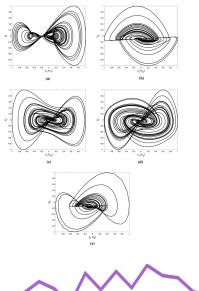


Pairwise coupling

(mm)

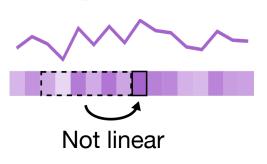
Pearson correlation, Granger causality, Euclidean distance, linear model fits, power envelope correlation

hctsa and pyspi include both linear and nonlinear features

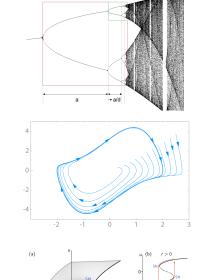


Nonlinear time series analysis

features do not make assumptions about the structure of the system, which can be additionally summarized by e.g., polynomial functions



Small changes to the input can give rise to:



Chaos

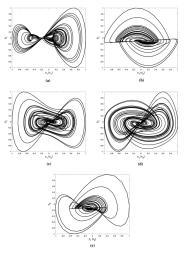
Oscillations



Bifurcations

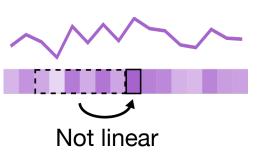
JLi et al. Nonlinear Dynamics (2021); Wikipedia; PhysicsOpenLab; Frey & Brauns arXiv (2021)

hctsa and pyspi include both linear and nonlinear features

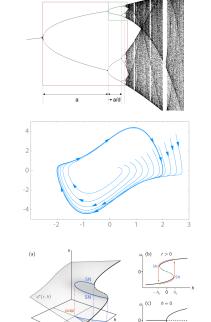


Nonlinear time series analysis

features do not make assumptions about the structure of the system, which can be additionally summarized by e.g., polynomial functions



Small changes to the input can give rise to:



Chaos

Oscillations



Local dynamics

Automutual information, Lyapunov exponent, fractional dimensionality, phasespace entropies, embedding distance



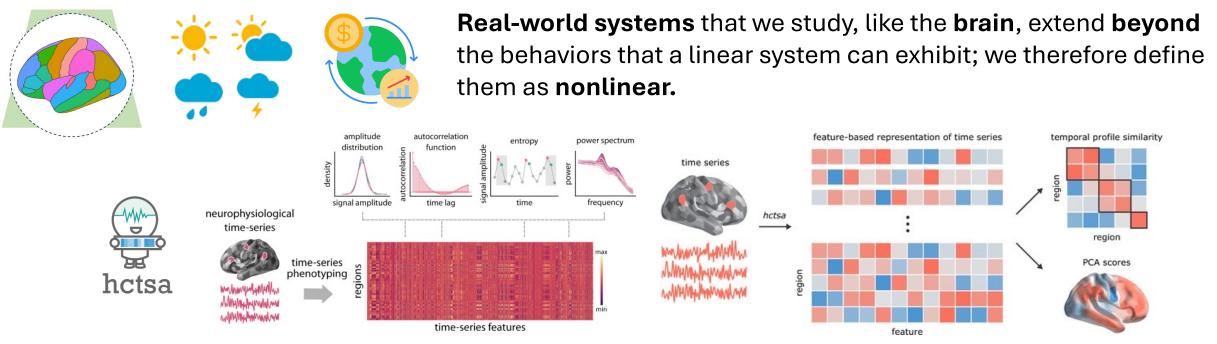
Pairwise coupling

Transfer entropy, dynamic time warping, additive noise modelling, phase lag index

Bifurcations

Ei et al. Nonlinear Dynamics (2021); Wikipedia; PhysicsOpenLab; Frey & Brauns arXiv (2021)

Nonlinearity in brain dynamics (?)

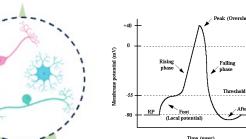


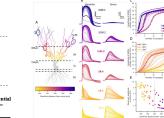
Applications of *hctsa* to **MEG (left)** and **fMRI (right)** data have shown that **linear properties** of local dynamics tend to dominate the **principal brain-wide axes of temporal variation** of the brain at rest.



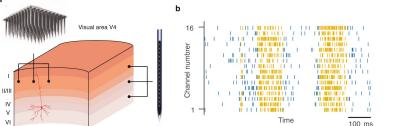
When your brain is at rest – as in not actively performing a cognitive task – it is sitting close to an equilibrium, where the **governing dynamics are approximately linear**. To detect nonlinear dynamics with **functional neuroimaging**, do we need to increase the **temporal precision** and/or **perturb** the brain away from resting equilibrium?

Do we still see **nonlinearity** at the **macroscale**?





Time (msec)



Article Open access Published: 11 December 2023

Macroscopic resting-state brain dynamics are best described by linear models

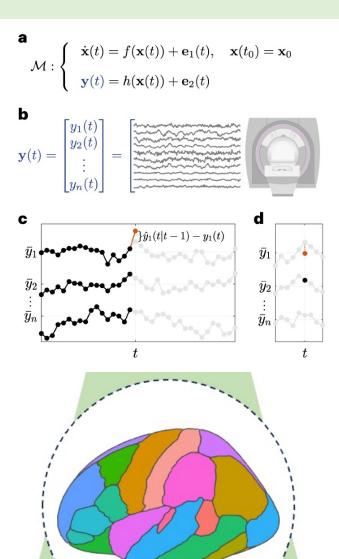
Erfan Nozari, Maxwell A. Bertolero, Jennifer Stiso, Lorenzo Caciagli, Eli J. Cornblath, Xiaosong He, Arun S. Mahadevan, George J. Pappas & Dani S. Bassett

Nature Biomedical Engineering 8, 68–84 (2024) Cite this article

8010 Accesses | 12 Citations | 18 Altmetric | Metrics

<u>https://www-archiv.fdm.uni-hamburg.de/b-online/library/crone/3028/membrane/mempot.html;</u> Humphries et al. *Neuroscience* (2023); Shi et al. *Nat Comms* (2022)

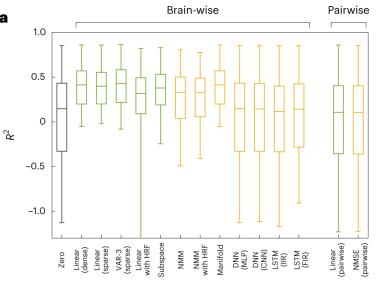
Do we still see **nonlinearity** at the **macroscale**?



а

With an AR(1) process: $x_{t+1} = f(x_t)$ with a linear function, fully specified by the autocorrelation function at lag 1

AC(1) is a "sufficient statistic" here – the summary of past observations that contains info needed to predict the future values of x



Article Open access Published: 11 December 2023

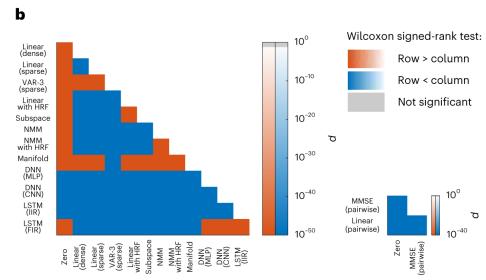
Macroscopic resting-state brain dynamics are best described by linear models

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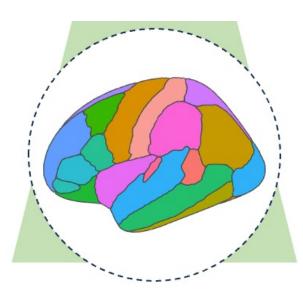
Linear autoregressive (AR) models

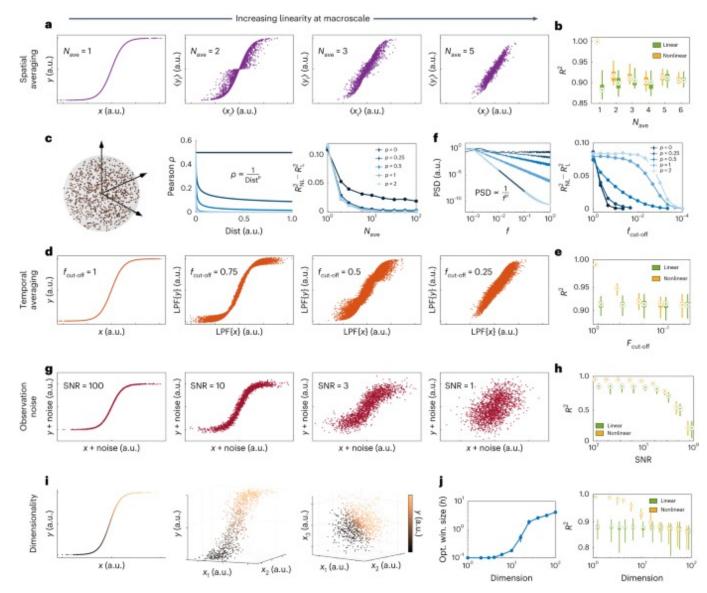


Do we still see **nonlinearity** at the **macroscale**?

4 "linearizing effects" that we can observe with functional neuroimaging at the whole-brain scale:

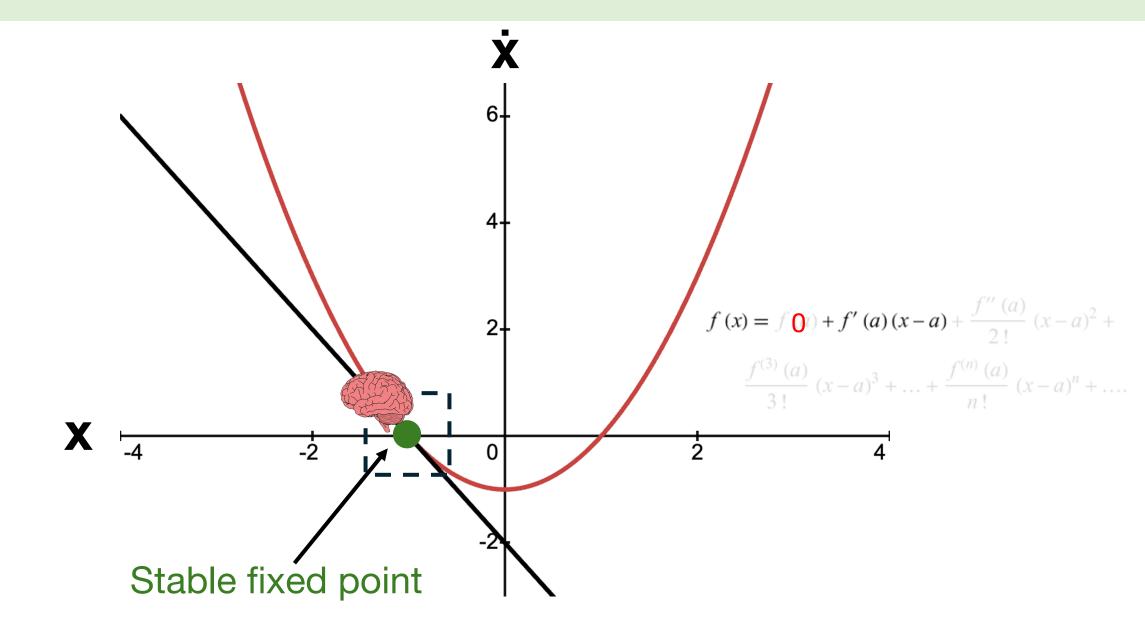
- 1. Spatial averaging
- 2. Temporal averaging
- 3. Observational noise
- 4. Dimensionality



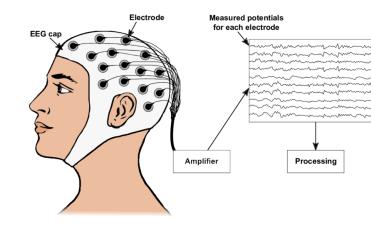


Nozari et al. Macroscopic resting-state brain dynamics are best described by linear models. *Nat Biomed Eng* (2024)

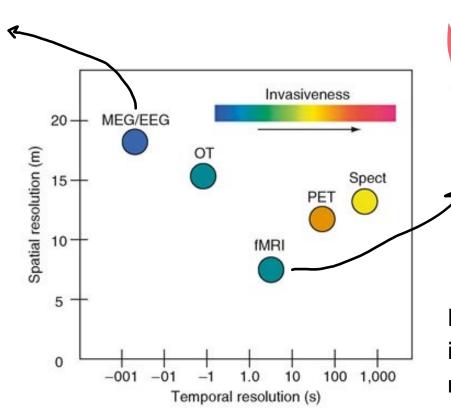
Nonlinear systems can be locally linear

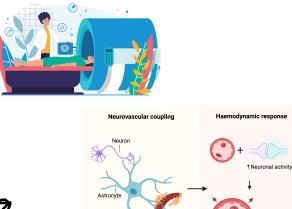


Brain imaging modalities exhibit a spatial/temporal resolution tradeoff



Electroencephalography (**EEG**) directly measures activity of neural masses (populations of neurons) at high temporal precision, but spatial resolution is limited to the cortical surface





Functional magnetic resonance imaging (**fMRI**) **indirectly** measures neural activity from the **cortex down to deep gray matter nuclei** via **hemodynamic coupling**

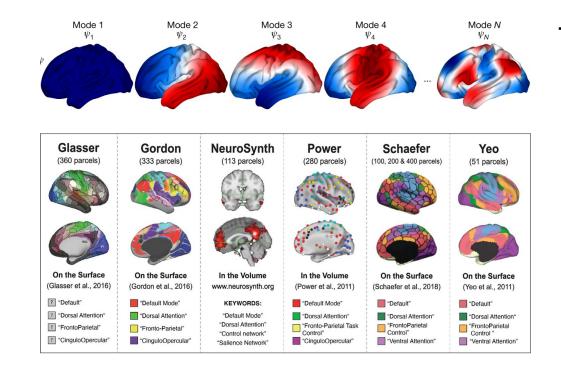


🕇 Temporal 🛛 🛉 Spatial

Time-series analysis with realistic brain activity datasets

>7,000 features





Tens to hundreds of modes or parcels





Sample sizes of a few hundred at the most in neuroimaging studies in general

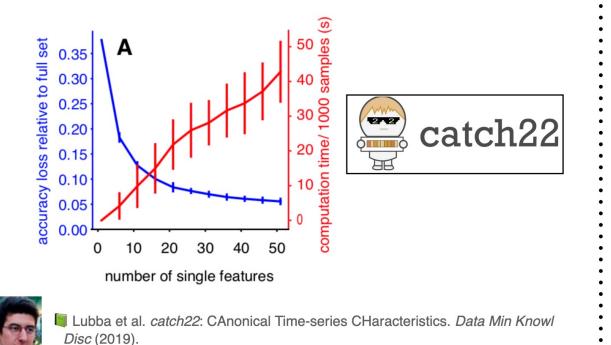
Bryce et al. Brain parcellation selection: An overlooked decision point with meaningful effects on individual differences in resting-state functional connectivity. Neuroimage (2021).

Time-series analysis with realistic brain activity datasets

catch22

Evaluate features across 93 diverse time-series classification tasks

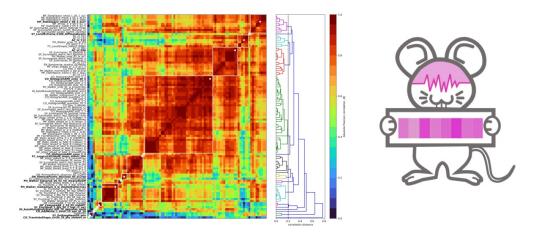
We can usually get away with using just 22 features.



catchaMouse16

Feature evaluation across **12 mouse fMRI classification tasks**

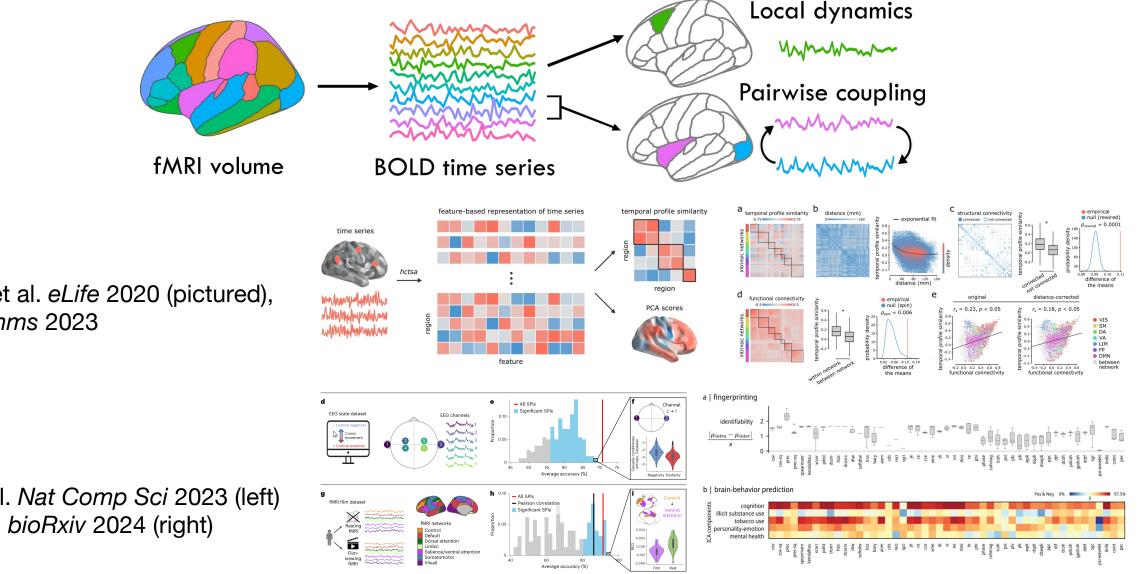
We can usually get away with using just 16 features.



Alam et al. Canonical time-series features for characterizing biologically informative dynamical patterns in fMRI. *bioRxiv* (2024).

Slide courtesy of Dr Ben Fulcher

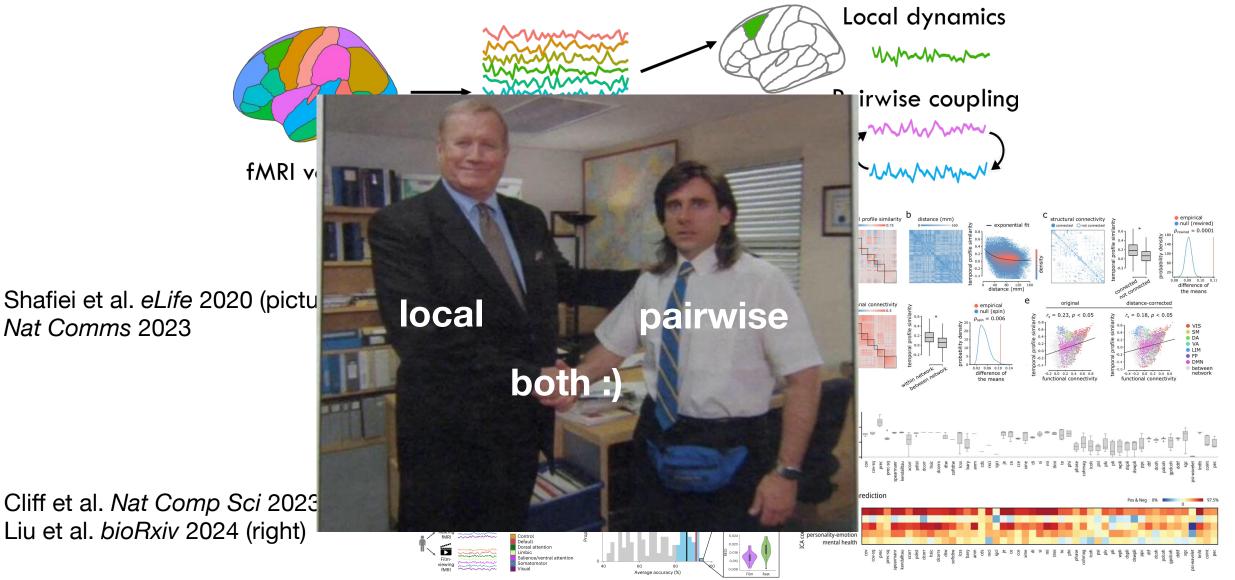
A shameless plug case study on time-series analysis for fMRI



Shafiei et al. eLife 2020 (pictured), Nat Comms 2023

Cliff et al. Nat Comp Sci 2023 (left) Liu et al. *bioRxiv* 2024 (right)

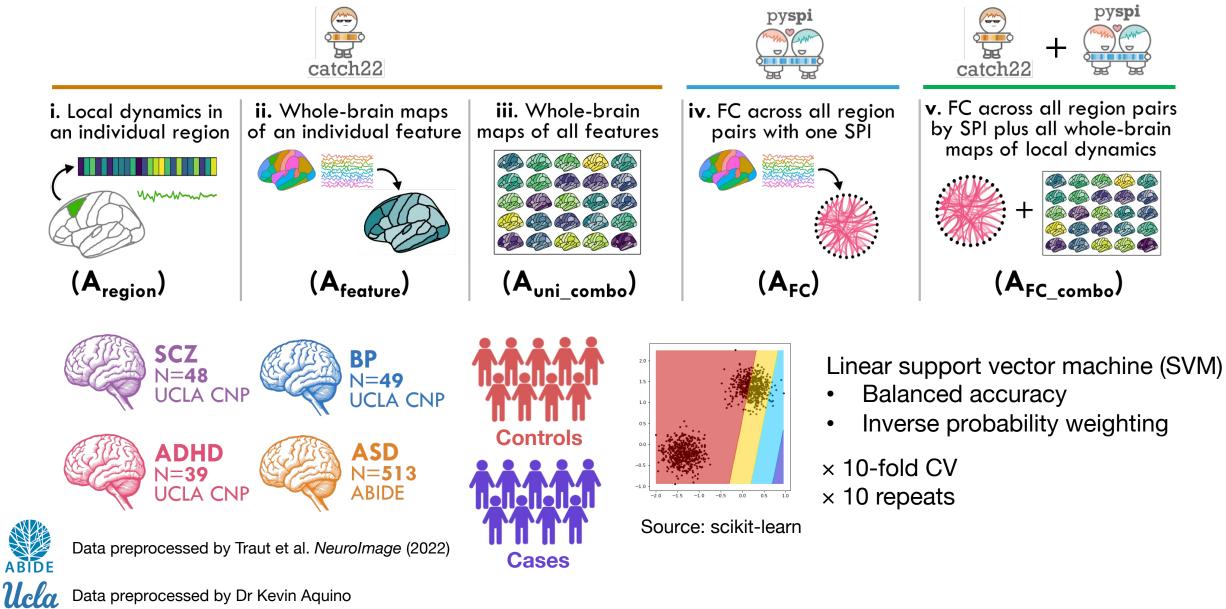
A shameless plug case study on time-series analysis for fMRI



Nat Comms 2023

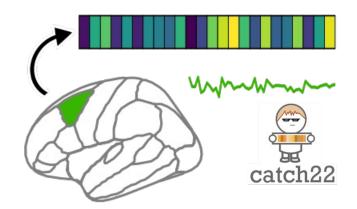
Cliff et al. Nat Comp Sci 2023 Liu et al. *bioRxiv* 2024 (right)

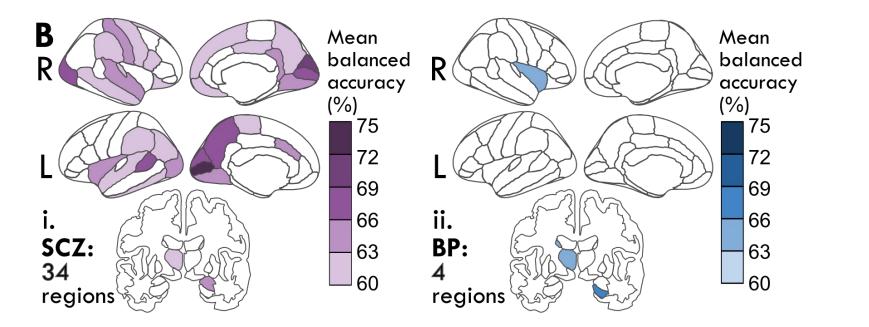
Classifying neuropsychiatric disorder cases vs. healthy controls



Interpretable spatial maps of region-specific dysfunction

Dynamical signatures of resting-state activity in **individual brain regions** can distinguish patients from controls in schizophrenia and bipolar disorder



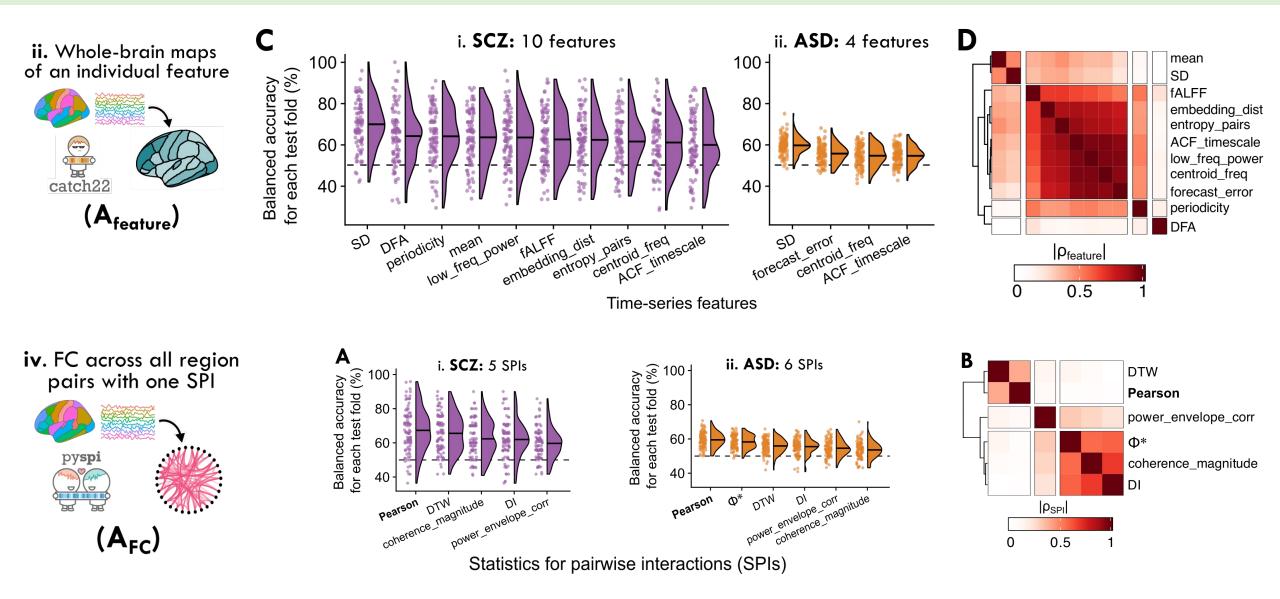


Gene expression 🧬

Anatomical changes 🥰

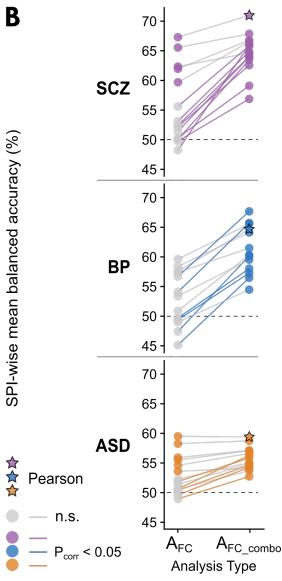
Stimulation analysis

Highlighting linear features for resting-state fMRI analysis



The benefit of **integrating** local dynamics **and** pairwise coupling

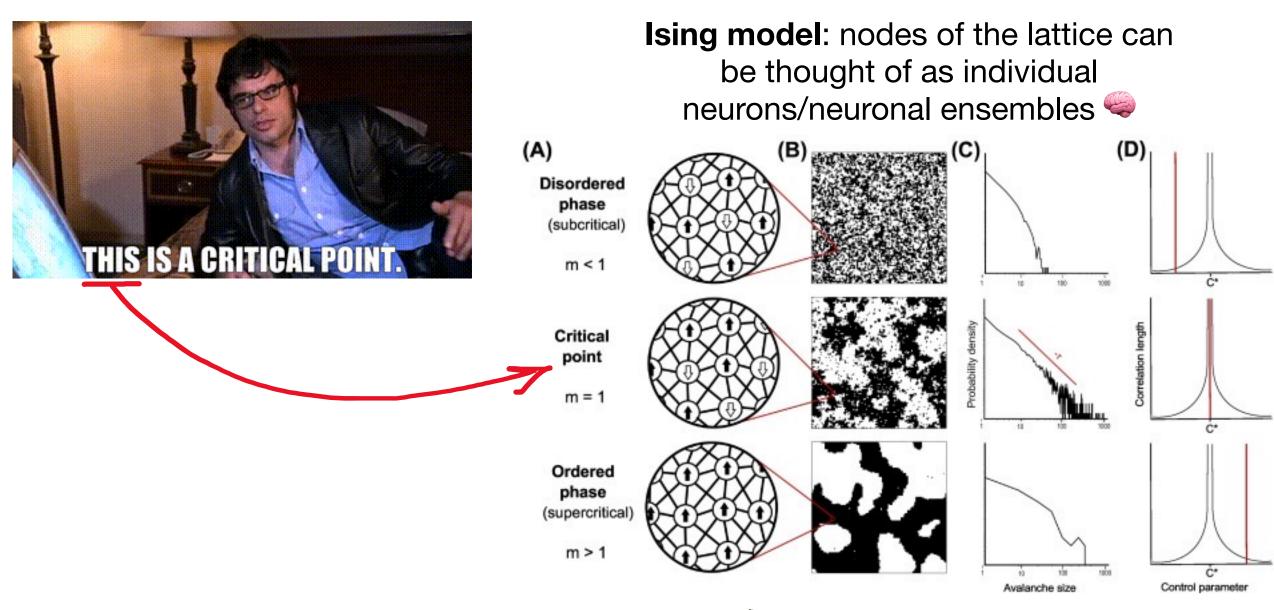
SPI-wise mean balanced accuracy (%)



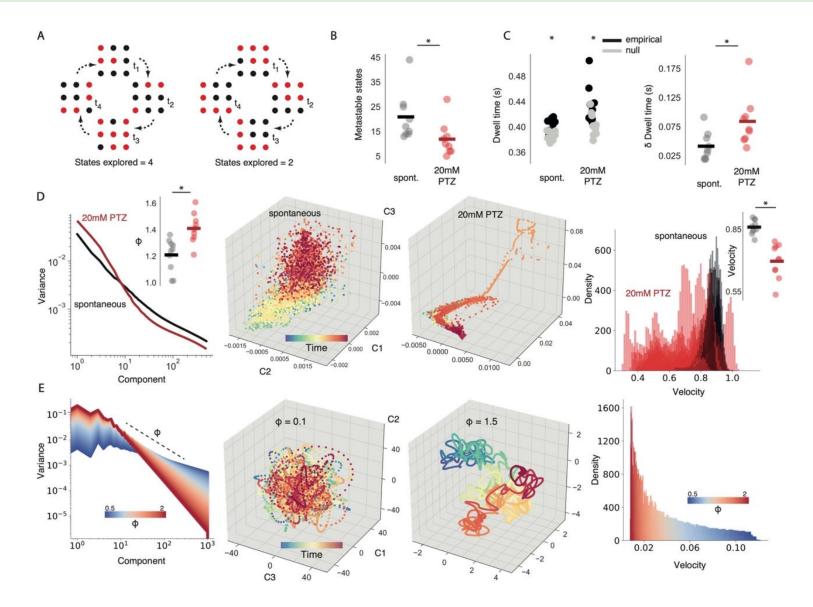
iv. FC across all region pairs with one SPI py**spi** iii. Whole-brain maps of all features catch22

Most functional **connectivity** metrics are more informative with the inclusion of brain-wide maps of local regional **dynamics**

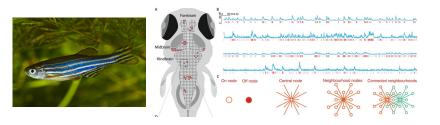
Segue to: Tracking the distance to criticality



We see patterns of **criticality** in **brain dynamics**



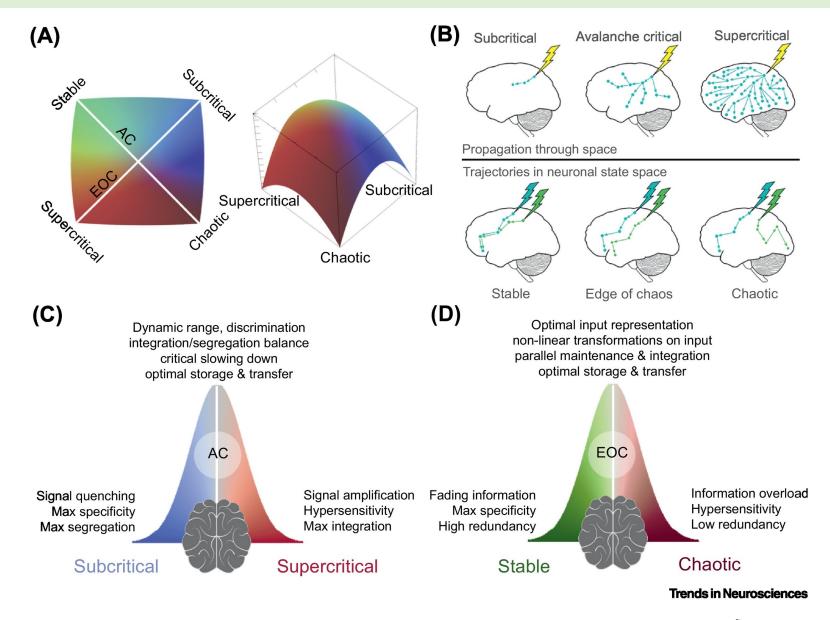
"(A) A critical system can explore a greater subset of its possible brain states ... while a non-critical system will explore a more limited subset."



Zebrafish models are used to study how **epilepsy** induces changes to **excitation/inhibition balances** that push the brain **away from criticality**.

Burrows et al. Single-cell Networks Reorganise to Facilitate Whole-brain Supercritical Dynamics During Epileptic Seizures. *bioRxiv* (2021)

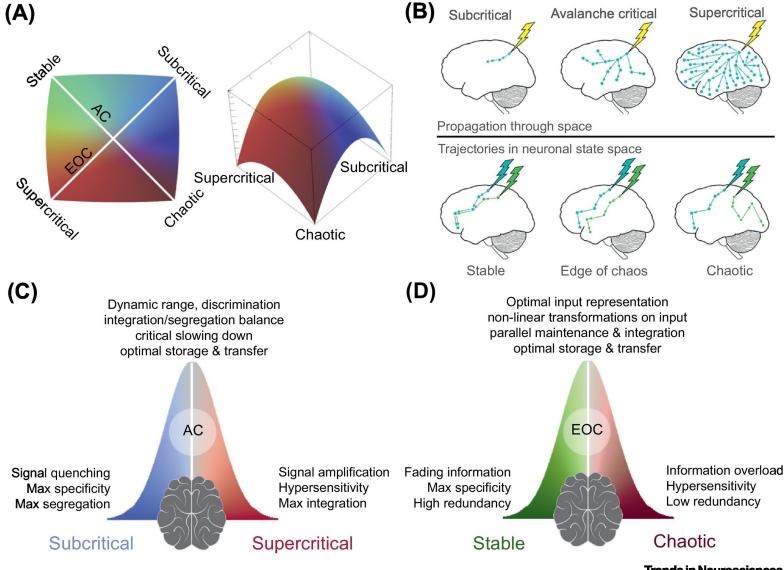
We see patterns of criticality in brain dynamics



"The distance to criticality presents a promising and underexploited biological parameter for characterizing cognitive differences and mental illness."

"...these ideas underscore the relevance of the **distance to criticality** for **cognition**: this distance may be **dynamically varied** at a moment-to-moment timescale in order to **flexibly adapt** to task requirements."

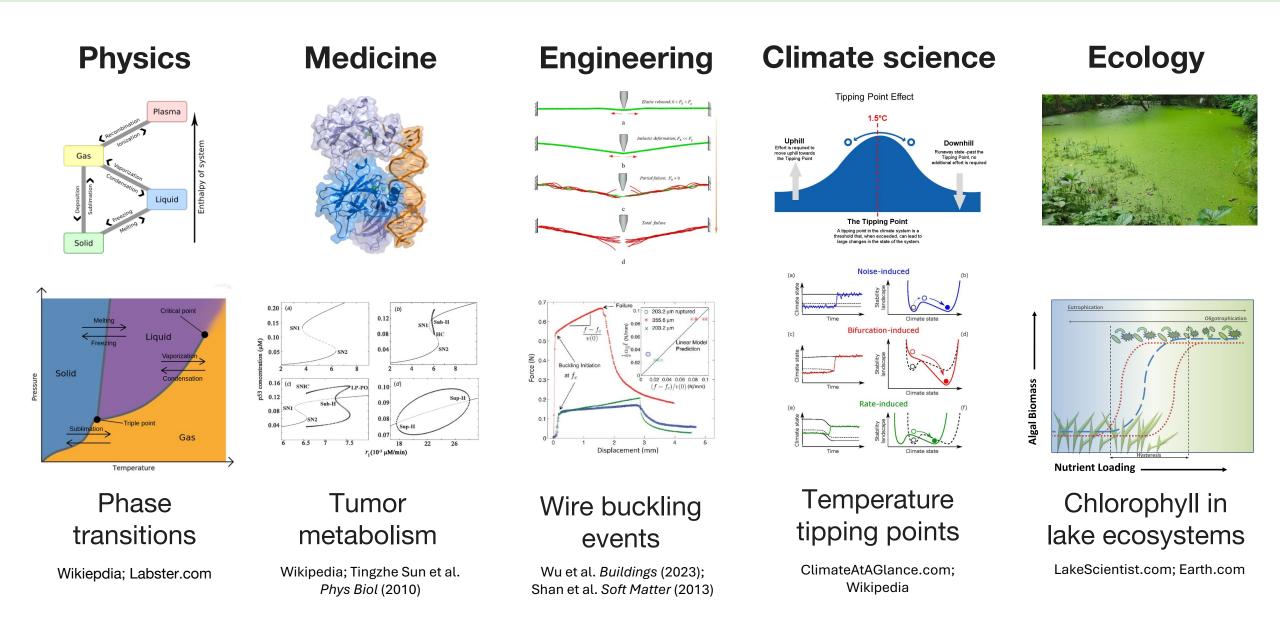
We see patterns of criticality in brain dynamics



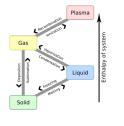
"[T]he distance to criticality may constitute a well-placed biological variable for understanding both healthy and pathological changes in cognition and behavior, one that forms a direct mechanistic link between neurons and neuronal ensembles at the micro scale and computation at the macro scale."

Trends in Neurosciences

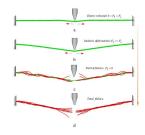
Distance to criticality is relevant beyond the brain

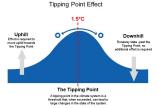


How is the distance to criticality typically measured?





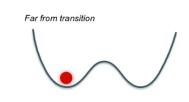


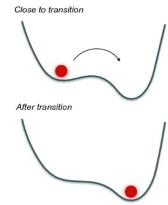




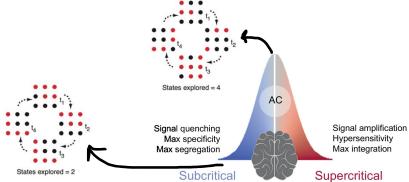
Critical slowing down

Dynamics are: More variable Evolving on a slower timescale





As you get closer to the critical point:



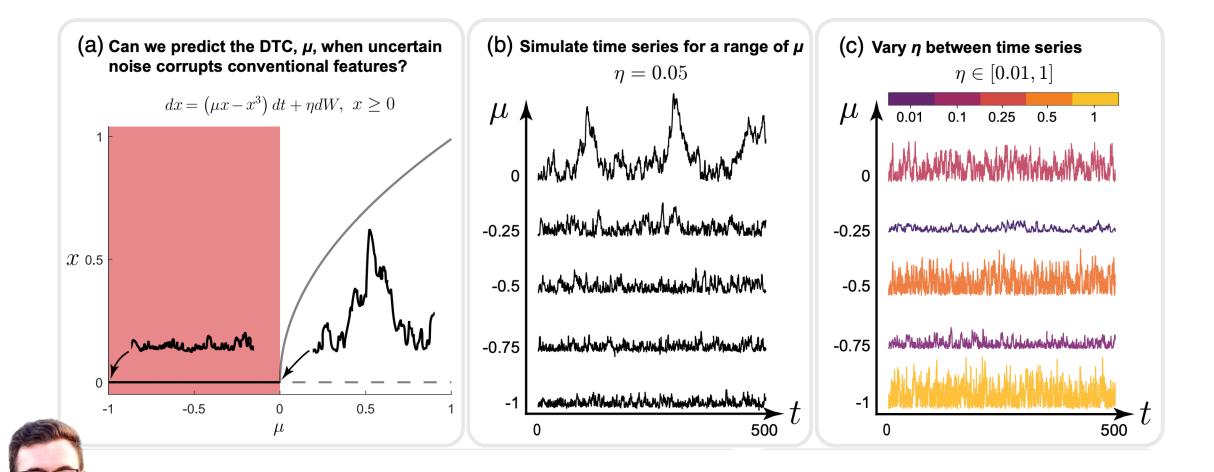
The system will explore a greater state domain closer to the critical point, which generally manifests as increase in the standard deviation (SD) of the time series.

Properties of the potential function naturally motivate timeseries features like the **SD or lag-1 autocorrelation** to track distance to the critical point, but both these measurements are **biased by noise**.

Physical systems like the brain exhibit noise that is **not trivial** relative to the scale of their **deterministic dynamics**.

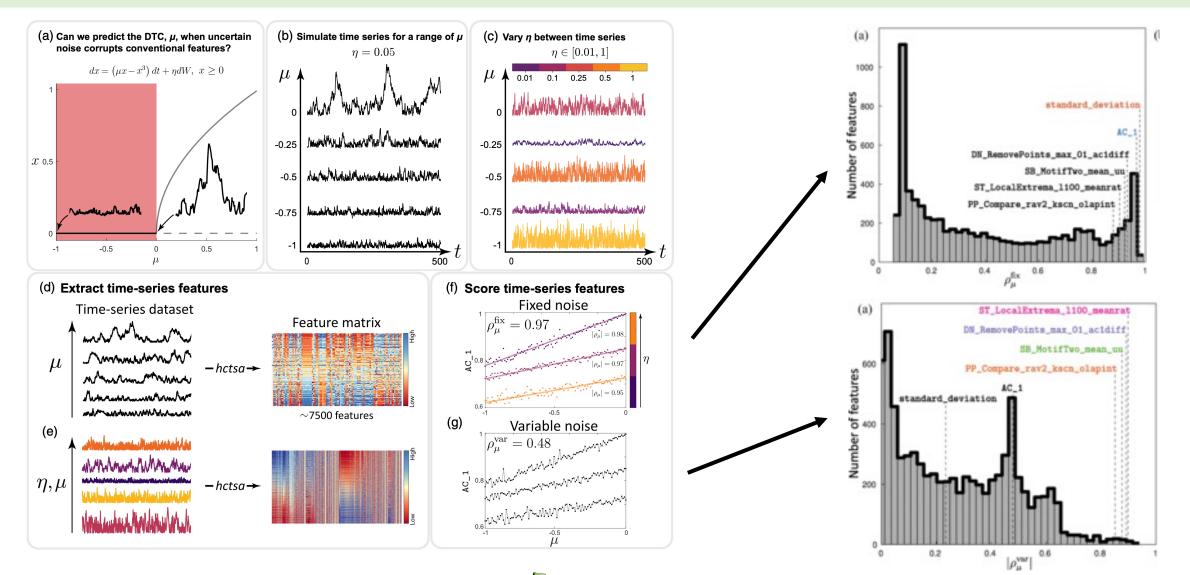
Meisel. From Neurons to Networks: Critical Slowing Down Governs Information Processing Across Vigilance States. (2022)

The search for a **DTC measure** that is **robust** across levels of **dynamical noise**



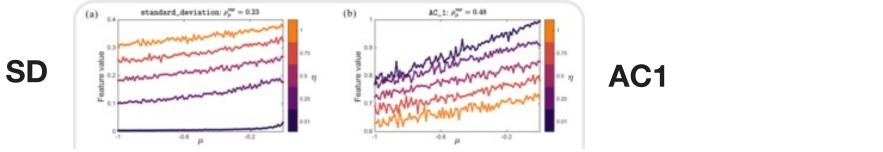
Harris, Gollo & Fulcher. Tracking the Distance to Criticality in Systems with Unknown Noise. Phys Rev X (2024)

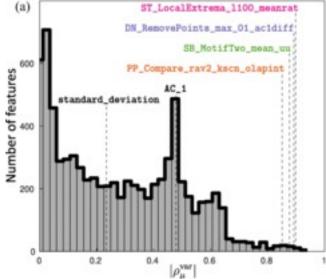
The search for a **DTC measure** that is **robust** across levels of **dynamical noise**



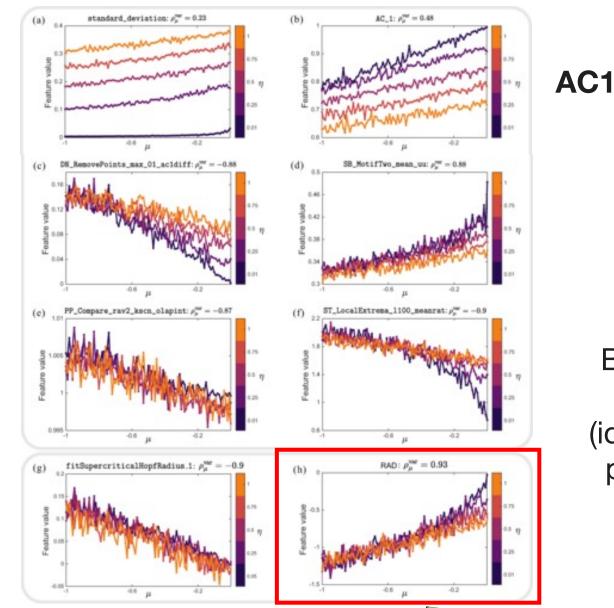
Harris, Gollo & Fulcher. Tracking the Distance to Criticality in Systems with Unknown Noise. Phys Rev X (2024)

Conventional DTC measures are sensitive to noise

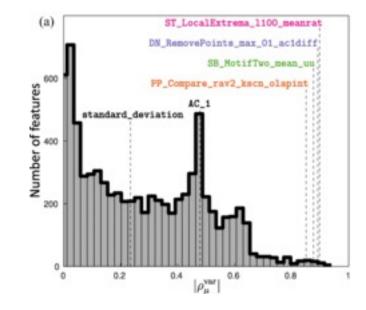




...but new noise-robust time-series features enter the villa



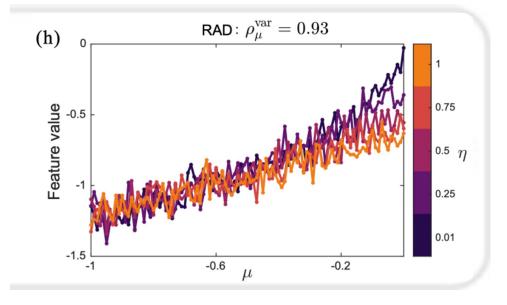
SD



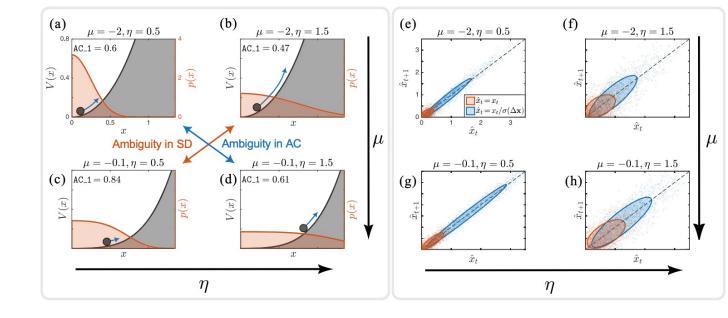
Brendan & co inspected properties of these noise-robust algorithms (identified via **data-driven analysis**!) to propose a new noise-robust index of the distance to criticality: **Rescaled autodensity**

Harris, Gollo & Fulcher. Tracking the Distance to Criticality in Systems with Unknown Noise. *Phys Rev X* (2024)

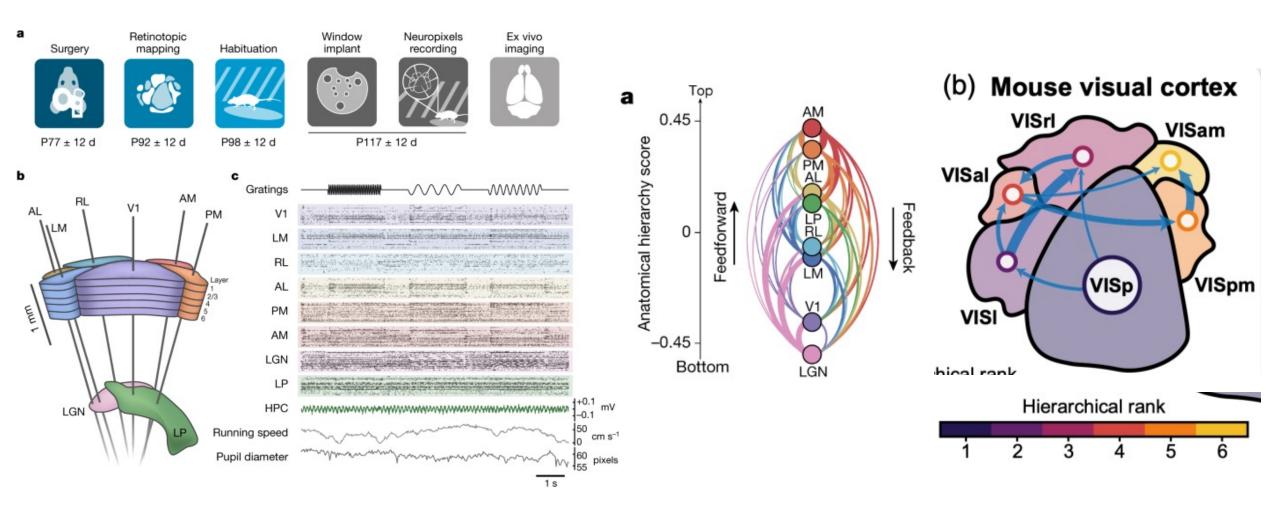
...but new noise-robust time-series features enter the villa



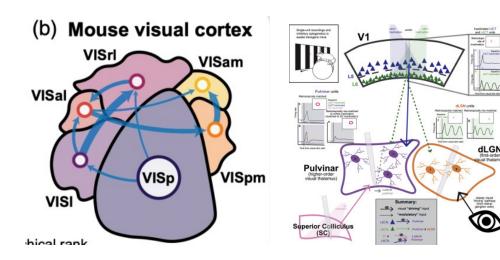




Rescaling time-series values by the spread of differenced values corrects for the confounding effect of a **variable-noise amplitude**, by capturing the **shape of the invariant density** (which depends on both the DTC and the noise amplitude) relative to the **spread of fast fluctuations** (which depends only on the noise amplitude)



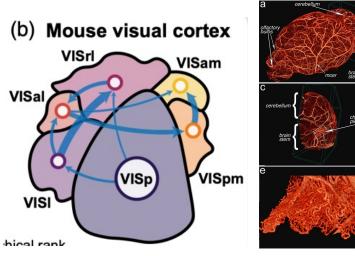
Hypothesis: **higher-order** regions exhibit **longer timescales** of neuronal activity because they are **closer to the critical point**

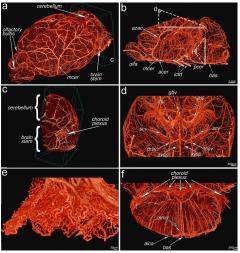


This is a good setting to test the noise robustness of RAD because we see variable noise levels across brain regions, attributable to:

• Differences in thalamic drives

Hypothesis: **higher-order** regions exhibit **longer timescales** of neuronal activity because they are **closer to the critical point**

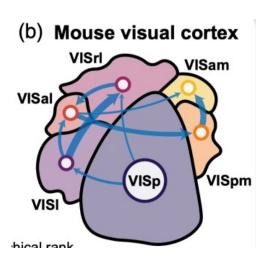


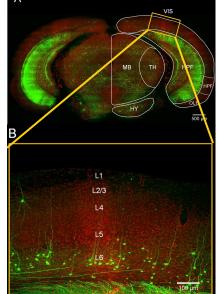


This is a good setting to test the noise robustness of RAD because we see variable noise levels across brain regions, attributable to:

- Differences in thalamic drives
- Differences in adjacent vasculature

Hypothesis: **higher-order** regions exhibit **longer timescales** of neuronal activity because they are **closer to the critical point**

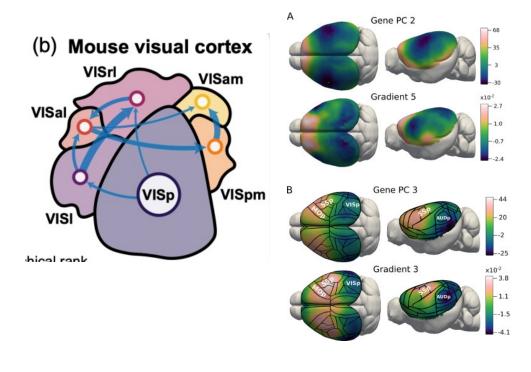




This is a good setting to test the noise robustness of RAD because we see variable noise levels across brain regions, attributable to:

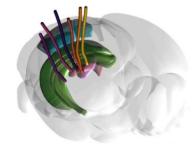
- Differences in thalamic drives
- Differences in adjacent vasculature
- Differences in cytoarchitecture

Hypothesis: **higher-order** regions exhibit **longer timescales** of neuronal activity because they are **closer to the critical point**



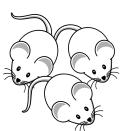
This is a good setting to test the noise robustness of RAD because we see variable noise levels across brain regions, attributable to:

- Differences in thalamic drives
- Differences in adjacent vasculature
- Differences in cytoarchitecture
- Differences in transcriptomic and neuromodulatory gradients

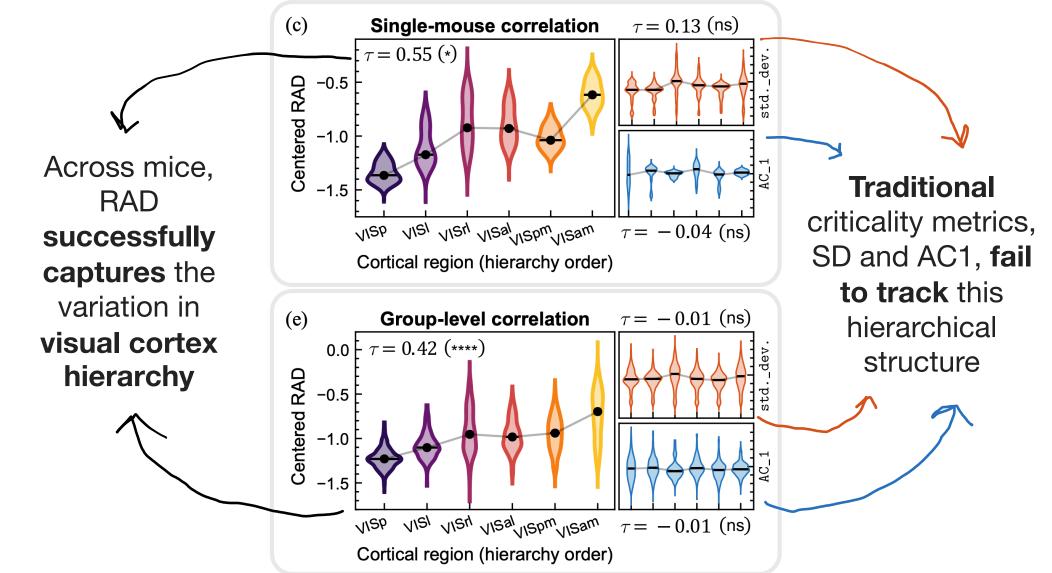








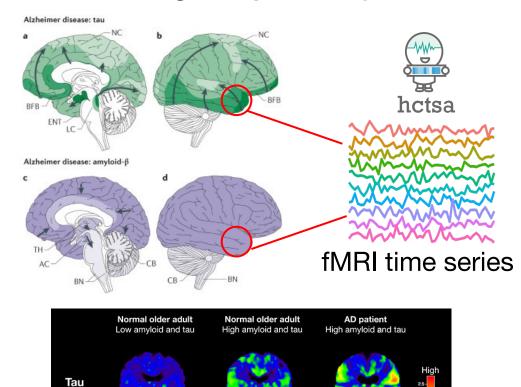
39 mice



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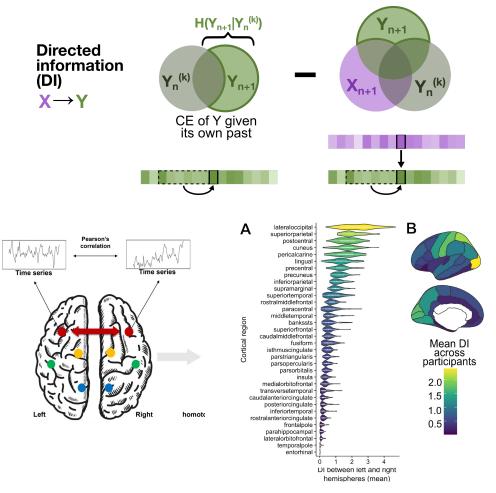
My final PhD projects with hctsa + pyspi

Project #1: Characterizing the effects of Alzheimer's disease pathology on localized region-specific dynamics



Amyloid

Project #2: **Directed information** flow between the left and right hemispheres in health and disease



Thanks to my research groups as always 👳



Dynamics & Neural Systems Group

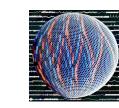
Dr Ben FulcherRishi MaranTrent HendersonBrendan HarrisKieran OwensJoshua MooreAria NguyenTeresa Dalle Nogare

Shine Lab

A/Prof Mac Shine Brandon Munn Eli Mueller Natasha Taylor Gabriel Wainstein Christopher Whyte Bella Orlando Joshua Tan



github.com/anniegbryant/ CSYS5040_Demo







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