



# Polyphorm Nature-inspired reconstruction and visualization of transport networks

Oskar Elek



Joseph Burchett Angus Forbes Carlos Maltzahn UCSC/NMSU UCSC UCSC/CROSS





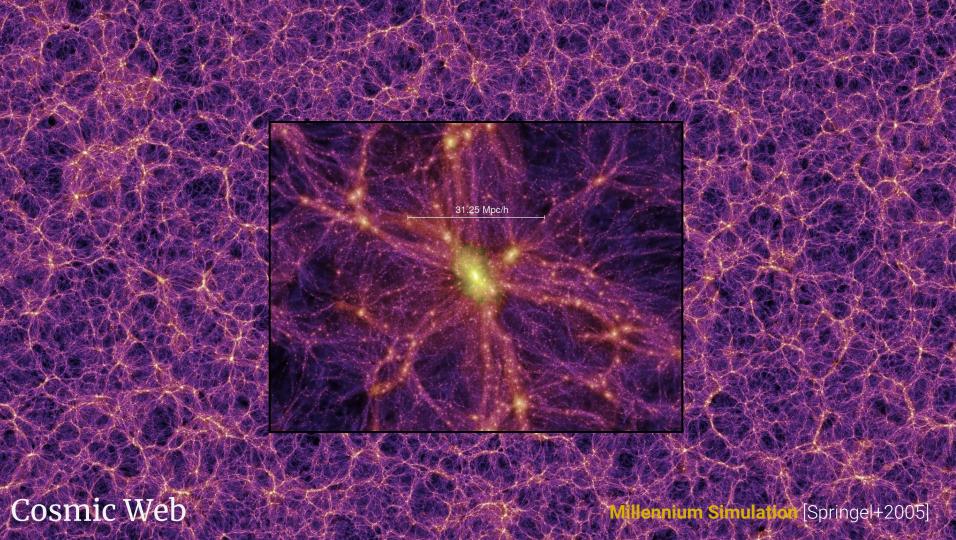


**Polyphorm** is an open source software that enables new types of analysis of complex scientific data using the MCPM algorithm.

In the past 2.5 years we used this project to make breakthroughs in analyzing data from astrophysics and cosmology.

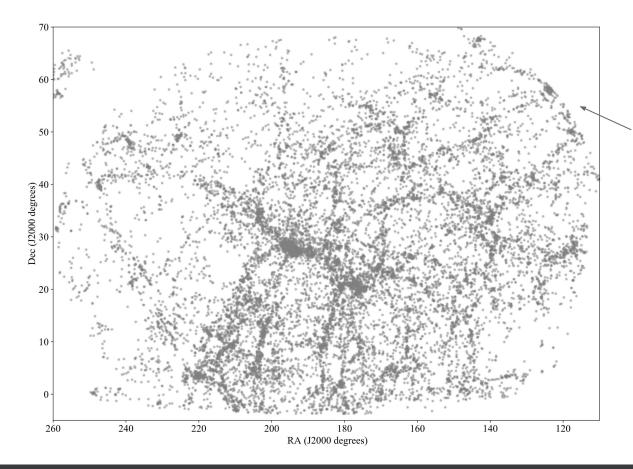
**PolyPhy** will make our methods available to a wider community of researchers and data scientists as a redesigned multi-platform Python port.

# Polyphorm and MCPM



, Cosmic Web: Where is it?

Hubble Deep Field



**Sloan Digital Sky Survey** 37.6k galaxies (3D points)

Redshifts .018 to .038

...density of data not sufficient to reconstruct the Cosmic Web without additional considerations

#### Cosmic Web: Where is it?



Unicellular, multi-nuclear protist

Searches it environment by climbing chemical marker gradients (emitted by food and also itself)

Amazing range of behaviors - look up "BBC slime mold" for a great time-lapse

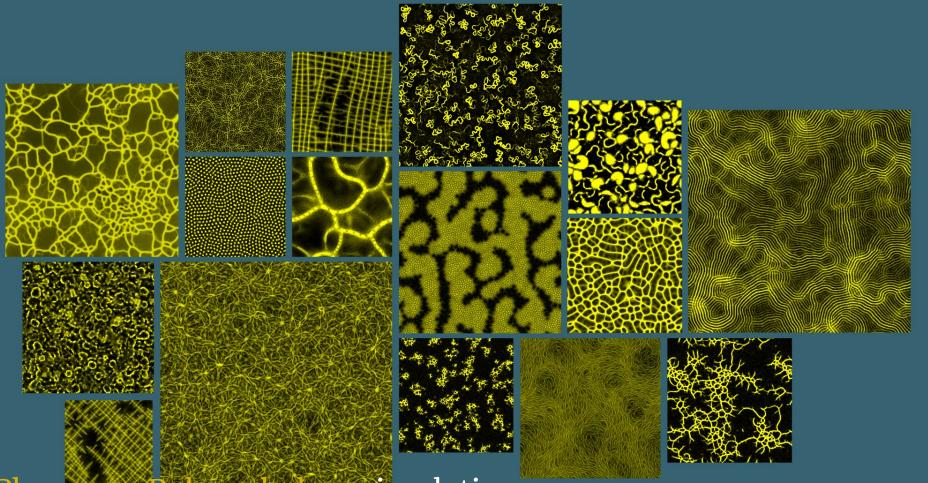
Forms approximate optimal transport networks over available food sources (NP-hard)



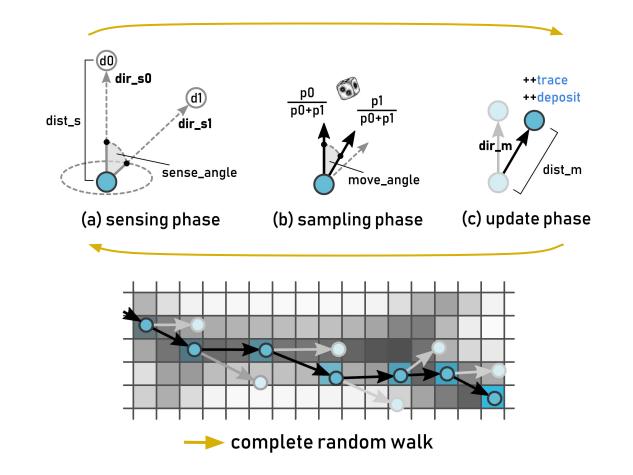
[Tero+2010]

#### Physarum Polycephalum – model organism



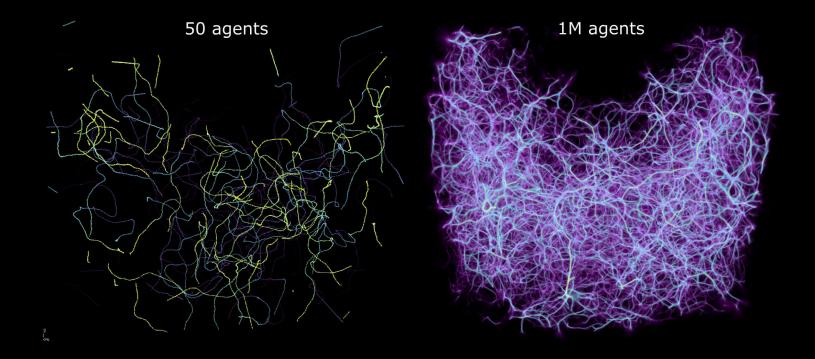


Physarum Polycephalum simulation



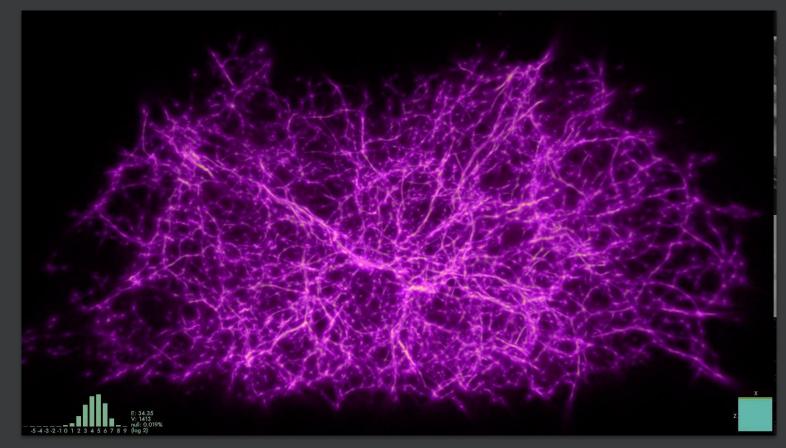
#### Monte Carlo Physarum Machine: Agent based simulation

#### 3D "trace" density fields (Bolshoi-Planck dataset, 450k objects)



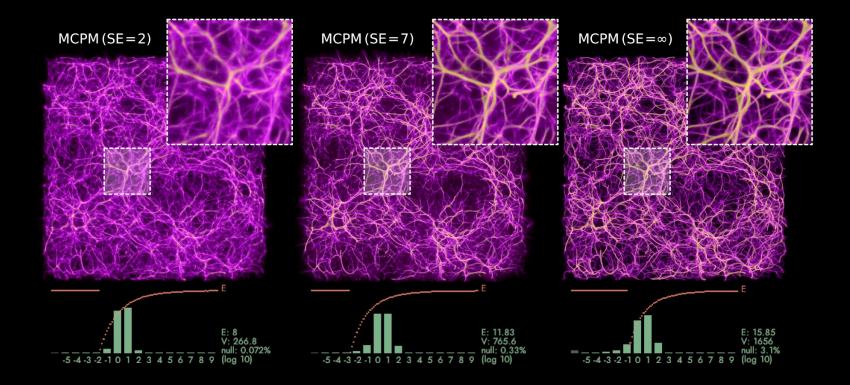
#### Monte Carlo Physarum Machine: Agent parallelism

#### Sloan Digital Sky Survey (SDSS) data, 300 Mpc wide, 37.6k galaxies



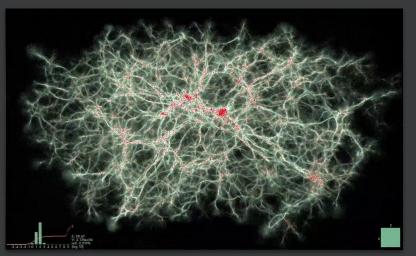
#### Monte Carlo Physarum Machine: Real-time fitting

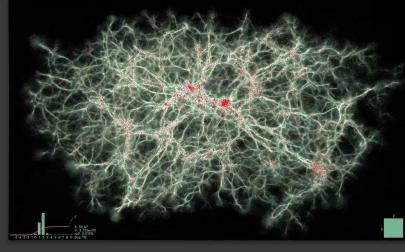
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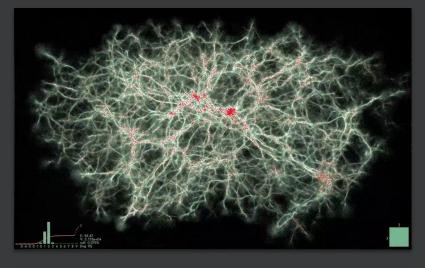


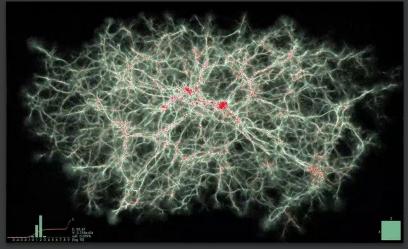
Monte Carlo Physarum Machine: Probabilistic sampling

# **Polyphorm:** Interactive Vis

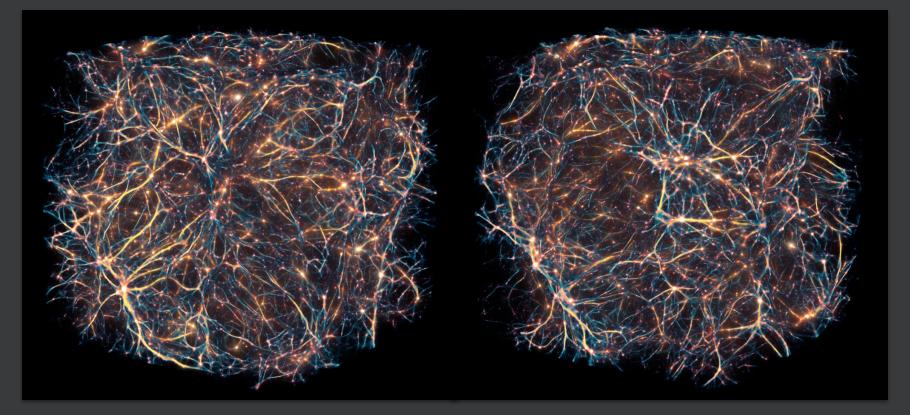








#### Physically plausible global illumination (using path-tracing) in Illustris TNG-100 data



### **Polyphorm:** Aesthetics

with David Abramov @ Vis Astro Data Challenge 2020

# What can Polyphorm/MCPM do?

#### Burchett J.N., Elek O., Tejos N., Prochaska J.X., Tripp T.M., Bordoloi R., Forbes A.G.

*Revealing the dark threads of the Cosmic Web* The Astrophysical Journal Letters, **2020**, Vol. 891(2)

#### Elek O., Burchett J.N., Prochaska J.X., Forbes A.G.

*Polyphorm: Structural analysis of cosmological datasets via interactive Physarum polycephalum visualization* Transactions of Visualization and Computer Graphics, **2021**, Vol. 27(2), Presented at VIS 2020

#### Simha S., Burchett J.N., Prochaska J.X., Chittidi J.S., Elek O., Tejos N. et al.

*Disentangling the Cosmic Web toward FRB 190608* The Astrophysical Journal, **2020**, Vol. 901(2)

Zhou H., Elek O., Anand P., Forbes A.G.
Bio-inspired structure identification in language embeddings
Visualization for Digital Humanities, 2020

# Burchett J.N., Abramov D., Elek O., Forbes A.G. Volumetric Reconstruction for Interactive Analysis of the Cosmic Web IEEE Vis Astro, 2020, Winner of the DataVis Challenge

#### Elek O., Burchett J.N., Prochaska J.X., Forbes A.G.

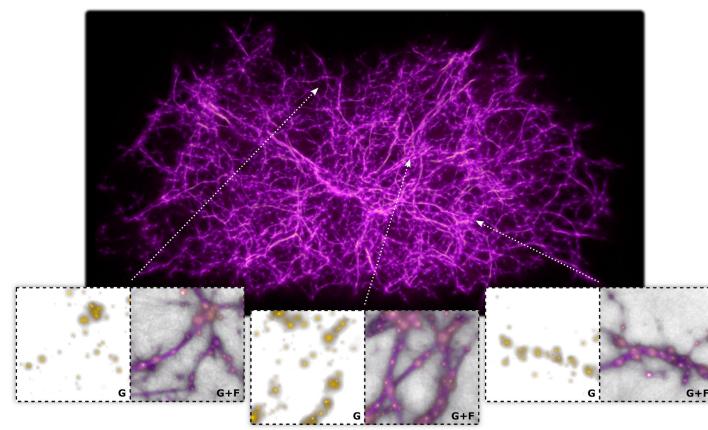
*Monte Carlo Physarum Machine: Characteristics of Pattern Formation in Continuous Stochastic Transport Networks* Artificial Life Journal, **2021**, Vol. 27(4), Will be presented at ALIFE 2022

#### Scientific record so far

Astronomy & cosmology

- Data visualization
- Procedural modeling
- Computational linguistics

Sloan Digital Sky Survey galaxy data, ~300 Mpc wide, 37k galaxies



Covered by:

- Hubble ST
- NASA
- ESA
- Ars Technica
- Planetary Society
- Popular Science
- SciShow
- Seeker
- Forbes

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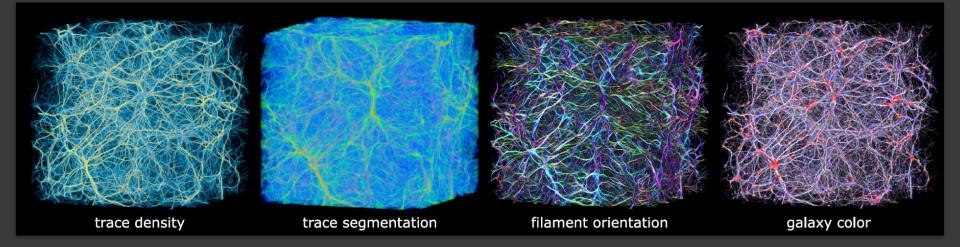
- Inverse
- Astronomy
- Universe Today
- Discover
- Reddit
- Space
- Entrepreneur

## First density map of the Cosmic Web!

Burchett, Elek et al.: Revealing the Dark Threads of the Cosmic Web (Astrophysical Journal Letters, March 2020)

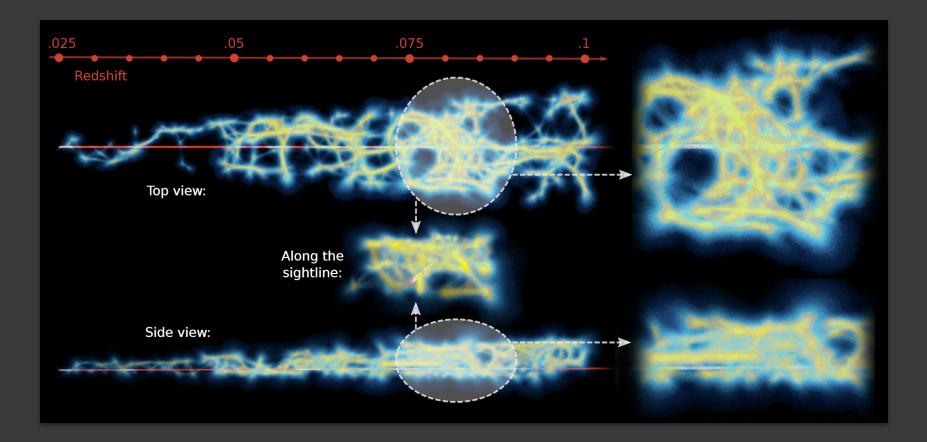
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#### Illustris TNG simulation, 100<sup>3</sup> Mpc volume, 4M halos



#### **Polyphorm:** Structural insights

with David Abramov @ Vis Astro Data Challenge 2020



#### Cosmic Web towards a Fast Radio Burst

Simha et al.: Disentangling the Cosmic Web towards FRB 190608 (Astrophysical Journal Letters, 2020)





## Natural language processing

Bio-inspired Structure Identification in Language Embeddings with Henry Zhou @ Vis 4 Digital Humanities

radical

well-regulated

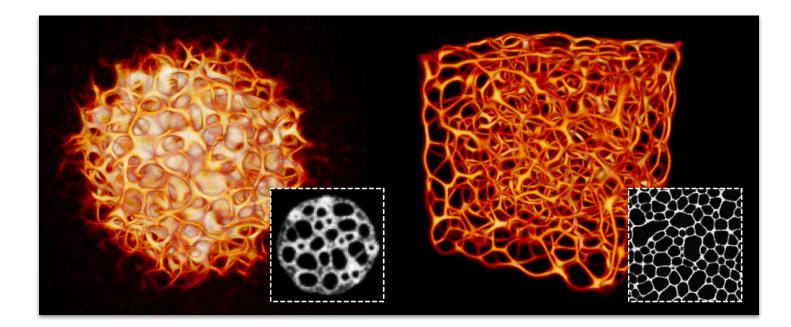
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credential

tool °

evolutionary · o ° neuroethic

non-expert

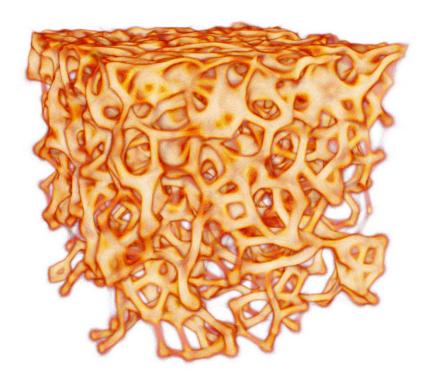


#### Elek O., Burchett J.N., Prochaska J.X., Forbes A.G.

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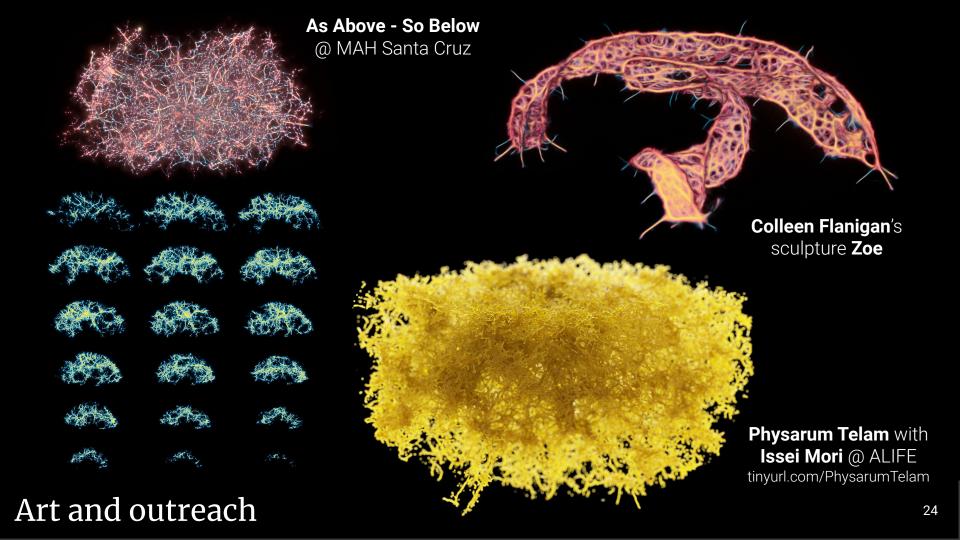
#### Artificial life approach to data modeling

#### Building bio-inspired scaffoldings for 3D printing (with Drew Ehrlich and Milad Hakimshafaei)





#### Computational fabrication and architecture



- First fully 3D simulation of Physarum networks
- Probabilistic 'Monte Carlo' sampling and continuous density reconstruction
- Natively parrelizable method, real-time on a single GPU, converges in 1-2 minutes
- Interactive visualization and expert supervision of the reconstruction process
- Easy to interpret and explain to non-experts

#### Monte Carlo Physarum Machine: Novel contributions

Future of MCPM: PolyPhy

# CRB55 CENTER FOR RESEARCH IN OPEN SOURCE SOFTWARE



# We have partnered with UCSC's **CROSS** to develop **PolyPhy** as an open source data-scientific software

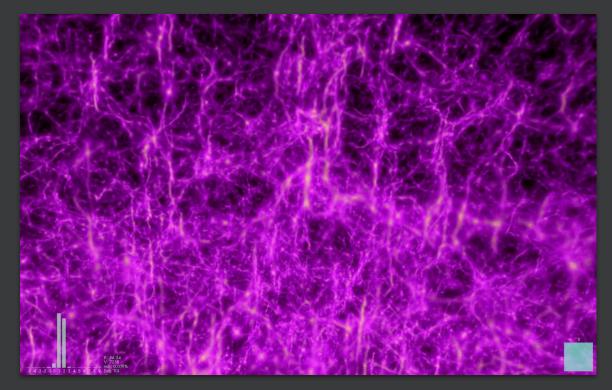
# **Project objectives:**

- 1. Develop PolyPhy for 3D and high–D data
- 2. Explore new scientific use cases
- 3. Build a community of domain experts and data scientists centered around this methodology

Expanding our search range to higher redshifts + cross-verification of the MCPM model with other modalities

Building on wide-sky surveys: SDSS and LRG

320k+ SDSS galaxies, redshift < 0.1



### Astronomy and cosmology

#### Using Polyphorm for reconstructing 3D maps of neuronal networks

Extension of the MCPM algorithm to operate on density fields rather than point clouds as input data

Friedmann et al.: Mapping Mesoscale Axonal Projections in the Mouse Brain Using A 3D Convolutional Network @ PNAS 2019

# TrailMa Ilastik TrailMap TrailMap Segmentation (0,1) Raw Data (16-bit) Skeletons (0-36) Probability Map (0-1)

#### Computational neuroscience

# Mapping genomic data of large human populations

Examining dispersion and clustering of genetic markers corresponding to distinct geographic or ethnic groups

IND/PK/AMB/AAB/WAA OEG WBC OEG WE WB WI

Diaz-Papkovich et al.: UMAP reveals cryptic population structure and phenotype heterogeneity in large genomic cohorts @ PLOS Genetics 2019

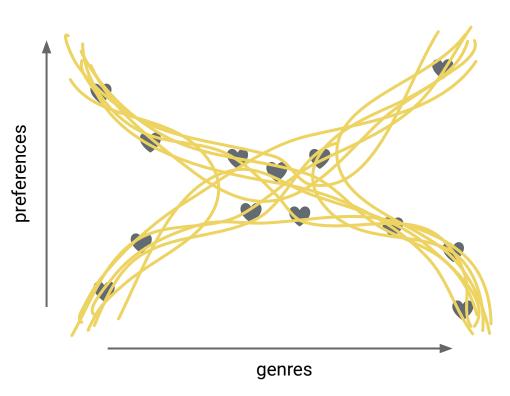
### Computational genomics

# The recommender system problem:

*"If I listen to some specific songs on Spotify, what others might I be interested in?"* 

Required tasks/queries:

- similarity
- interpolation
- extrapolation



# Navigating sparse data

- 1. How would you design a toolkit with such tight coupling between simulation and visualization?
- 2. Would you use **PolyPhy** for your science? How?
- 3. Any insights into the existing use cases?



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Polyphorm

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