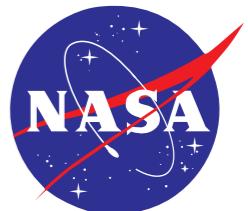


If you would like to participate in live polls today...
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Your Expertise (Select all that apply)



- Computer Science
- Biology
- Statistics
- Physics
- Data Visualization
- Software Development
- Instrumentation
- None of the above

Your interests (Select all that apply)



- Computer Science
- Biology
- Statistics
- Physics
- Data Visualization
- Software Development
- Instrumentation
- None of the above

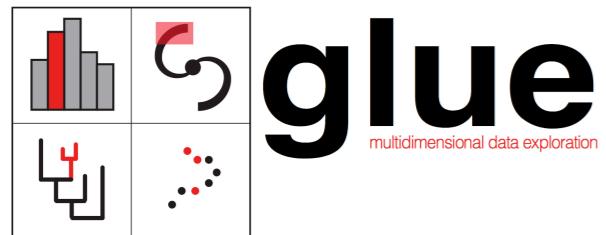


DISCOVERY THROUGH DATA DIVERSITY: EXPLORATORY DATA VISUALIZATION WITH GLUE

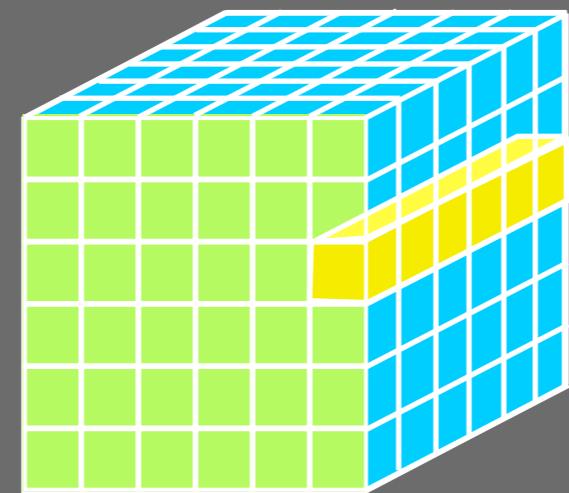


Alyssa A. Goodman, Harvard-Smithsonian Center for Astrophysics,
Radcliffe Institute for Advanced Study, and glue solutions, inc.

PRACTICALITY



PRINCIPLES



PHILOSOPHY

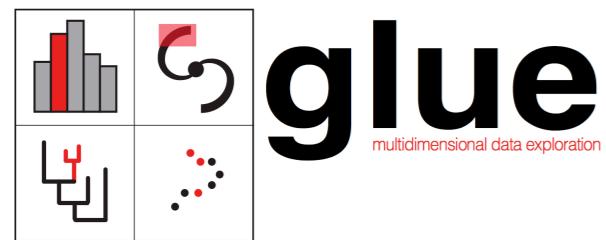


DIVERSE TOOLS

DIVERSE DATA

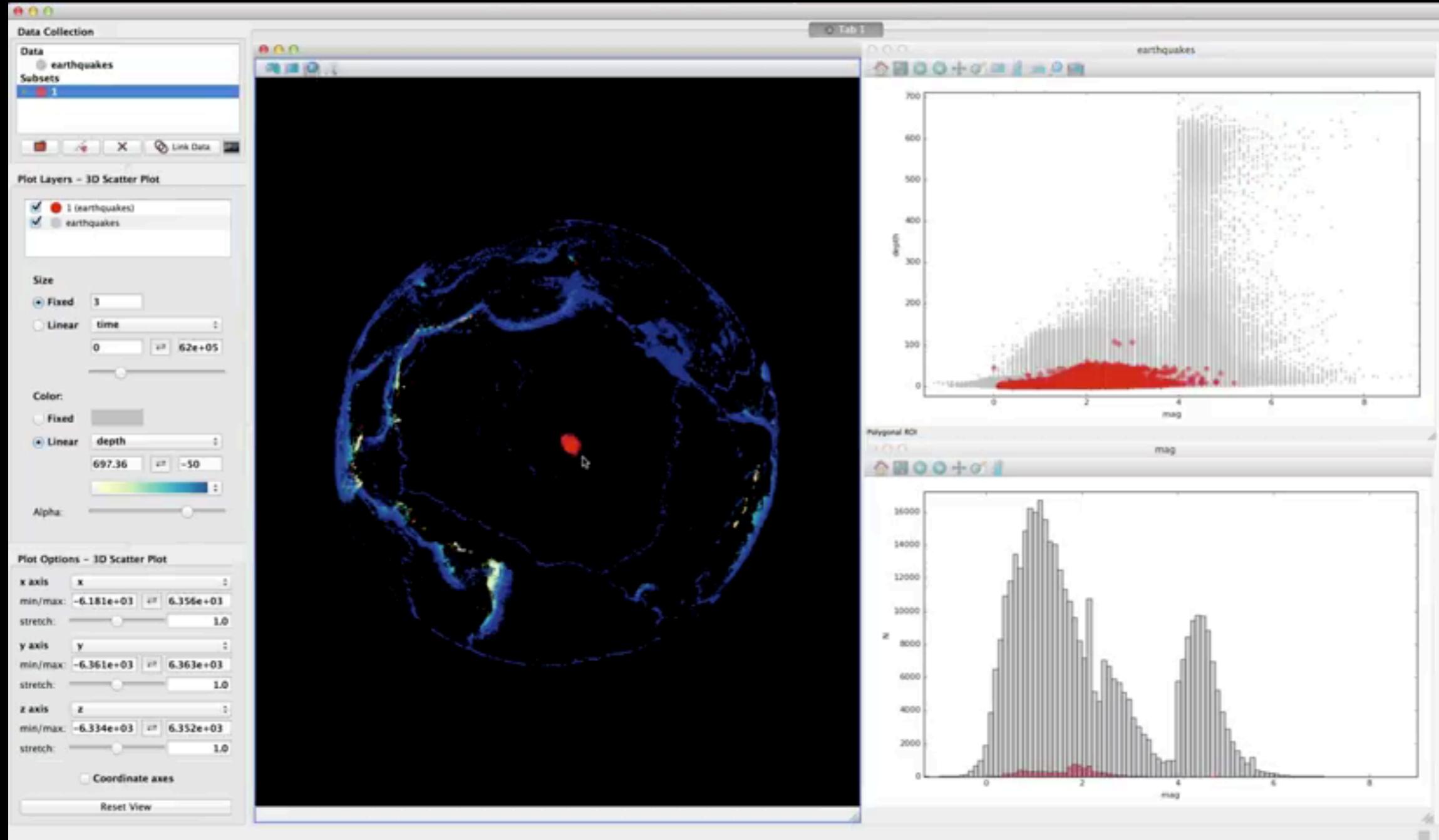
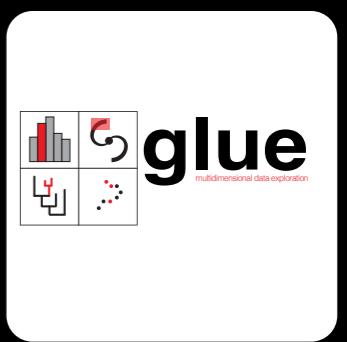
DIVERSE VIEWS

PRACTICALITY



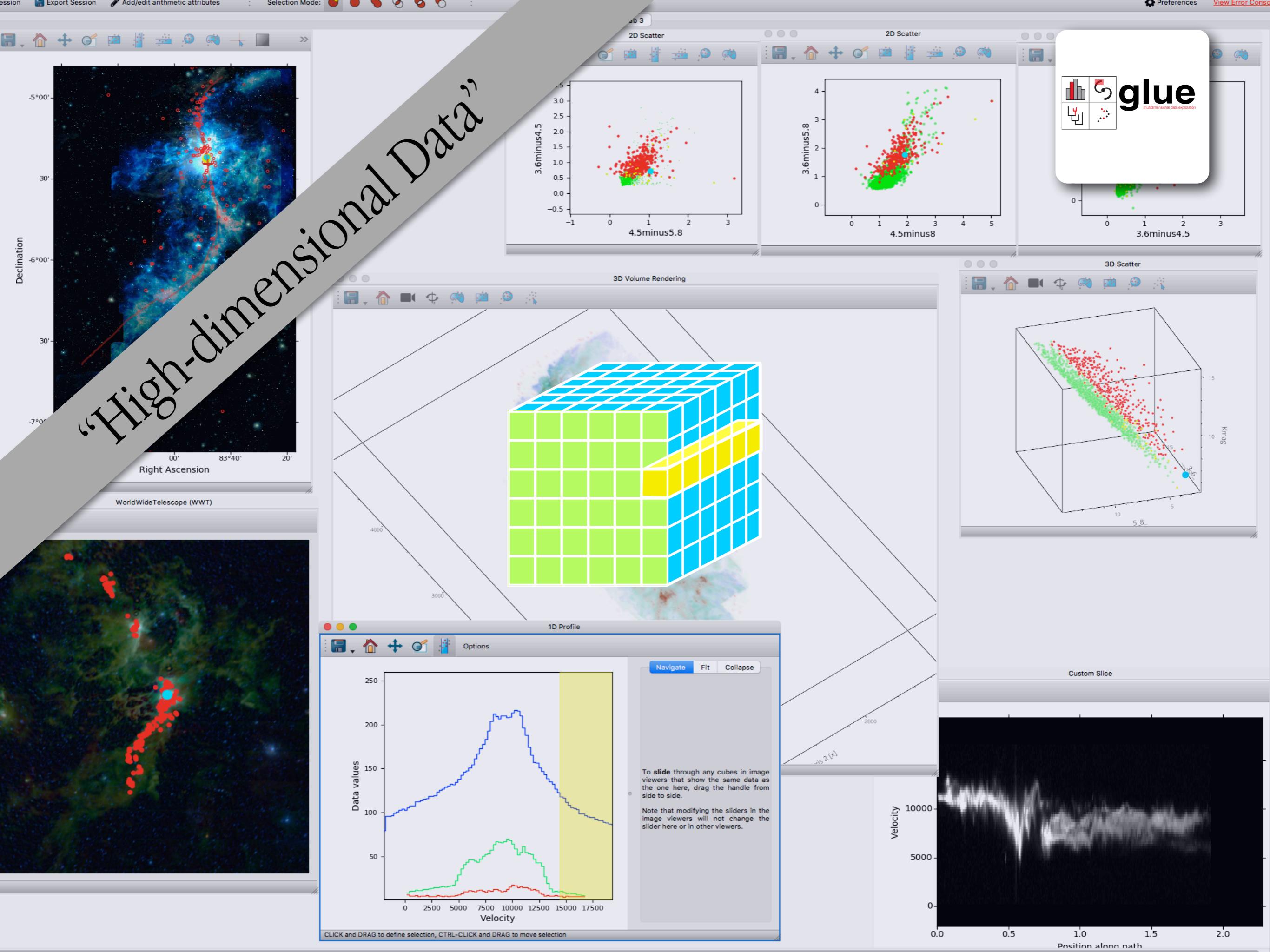
Linked Views of High-dimensional Data (in Python)

glue

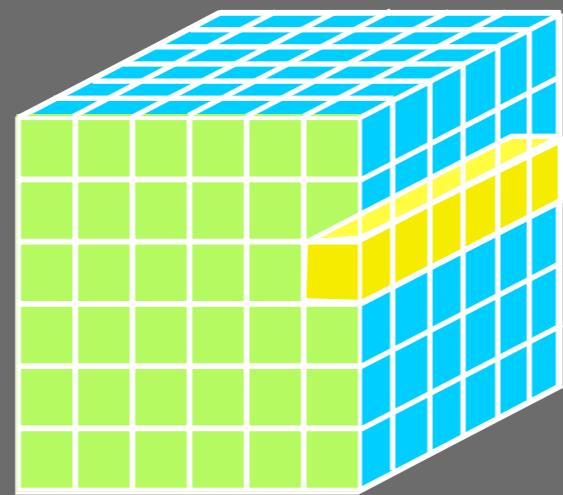


video by Tom Robitaille, lead glue developer

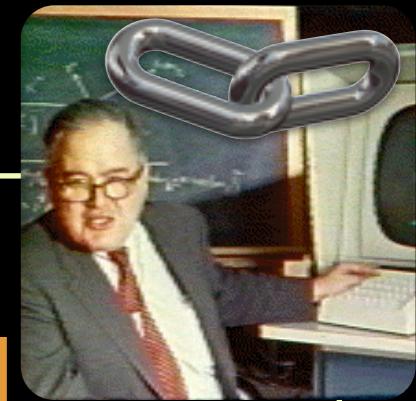
glue created by: C. Beaumont, M. Borkin, M. Breddels, T. Robitaille, C. Zucker, and A. Goodman, PI



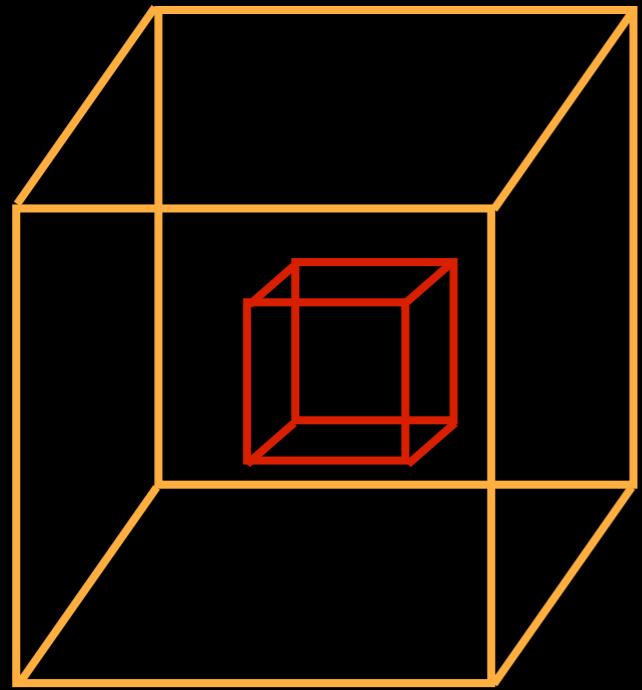
PRINCIPLES



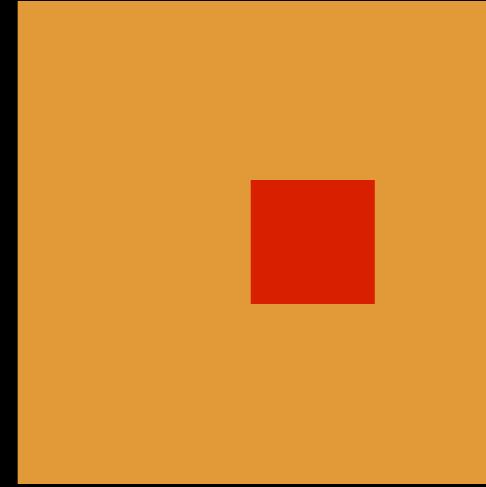
Linked Views of High-dimensional Data



John Tukey

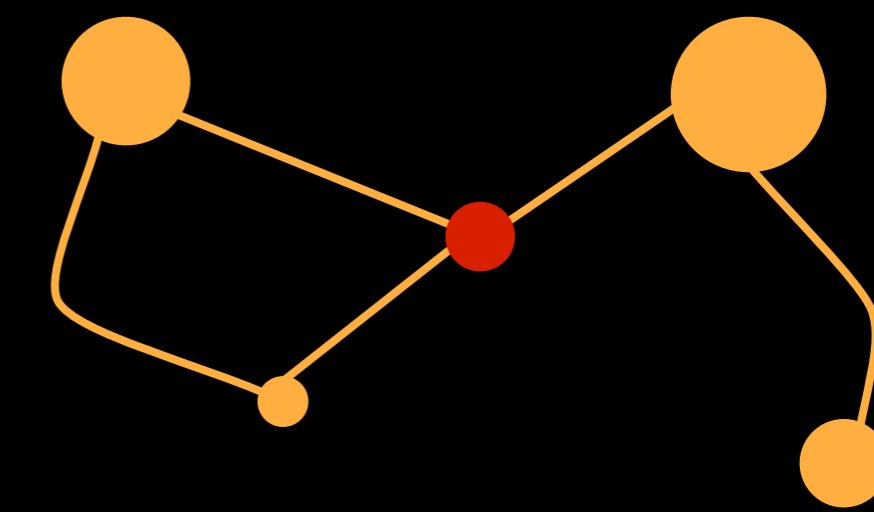
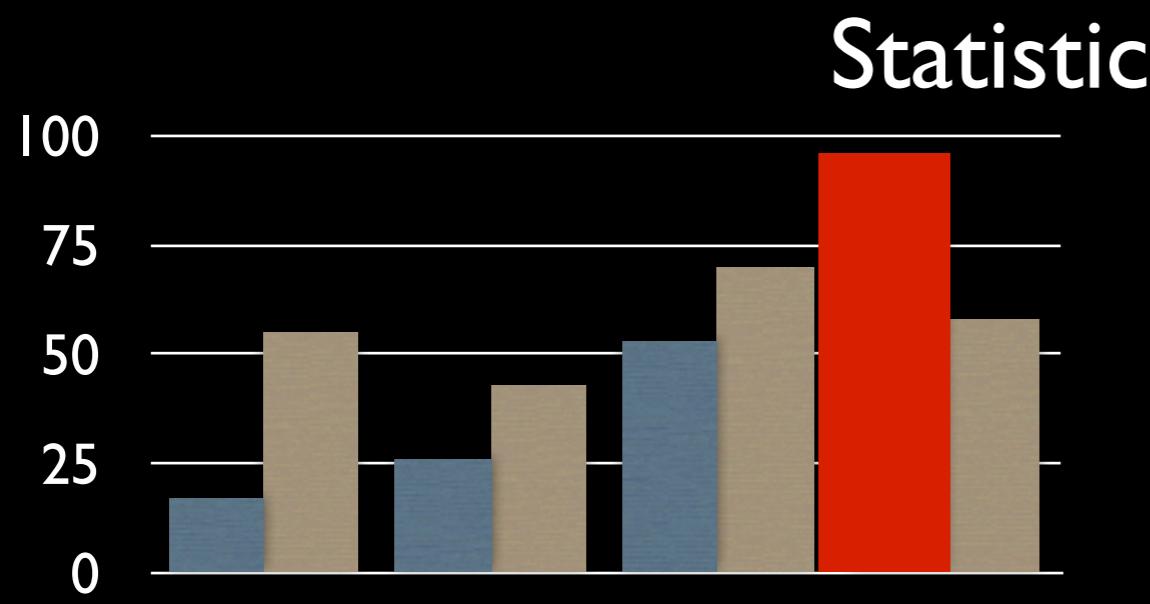


3D

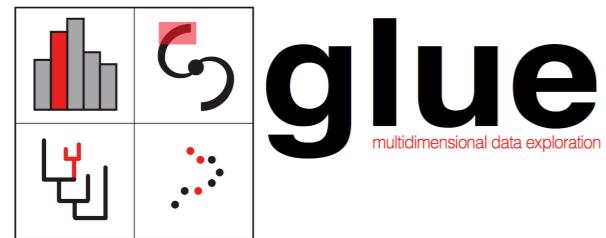


2D

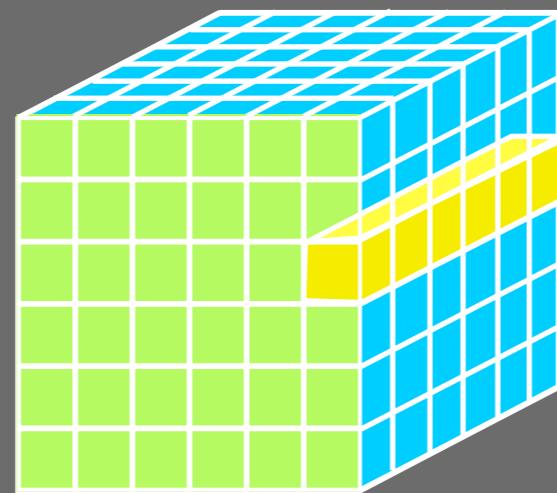
Data Abstraction

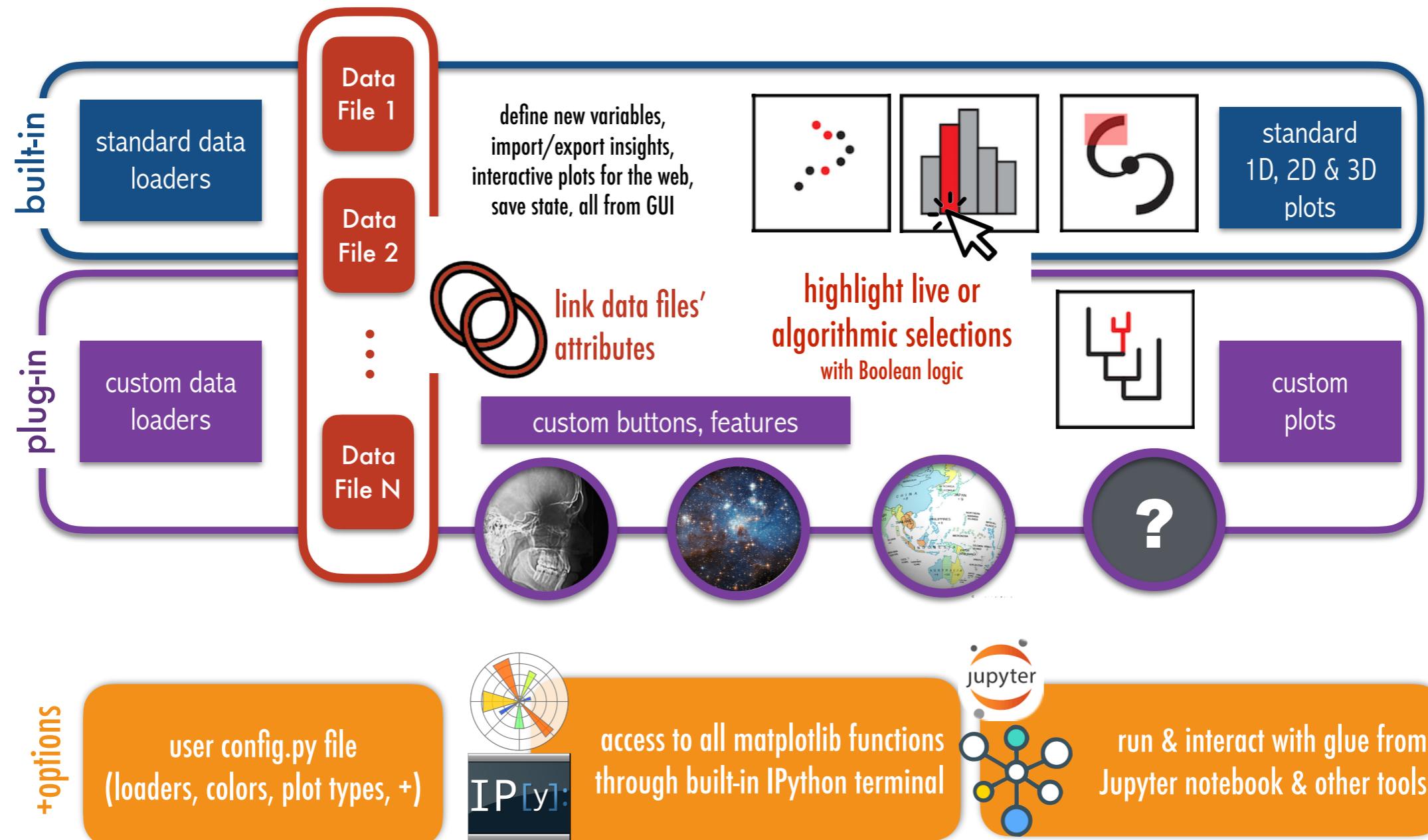
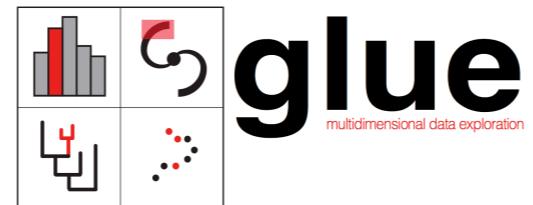


PRACTICALITY



PRINCIPLES



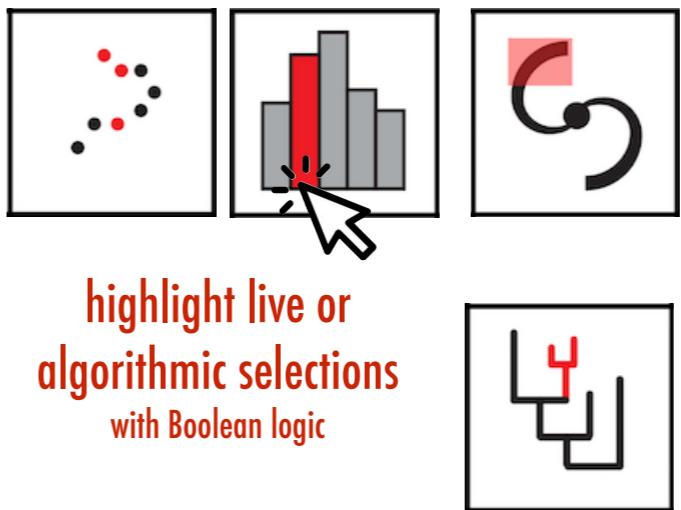
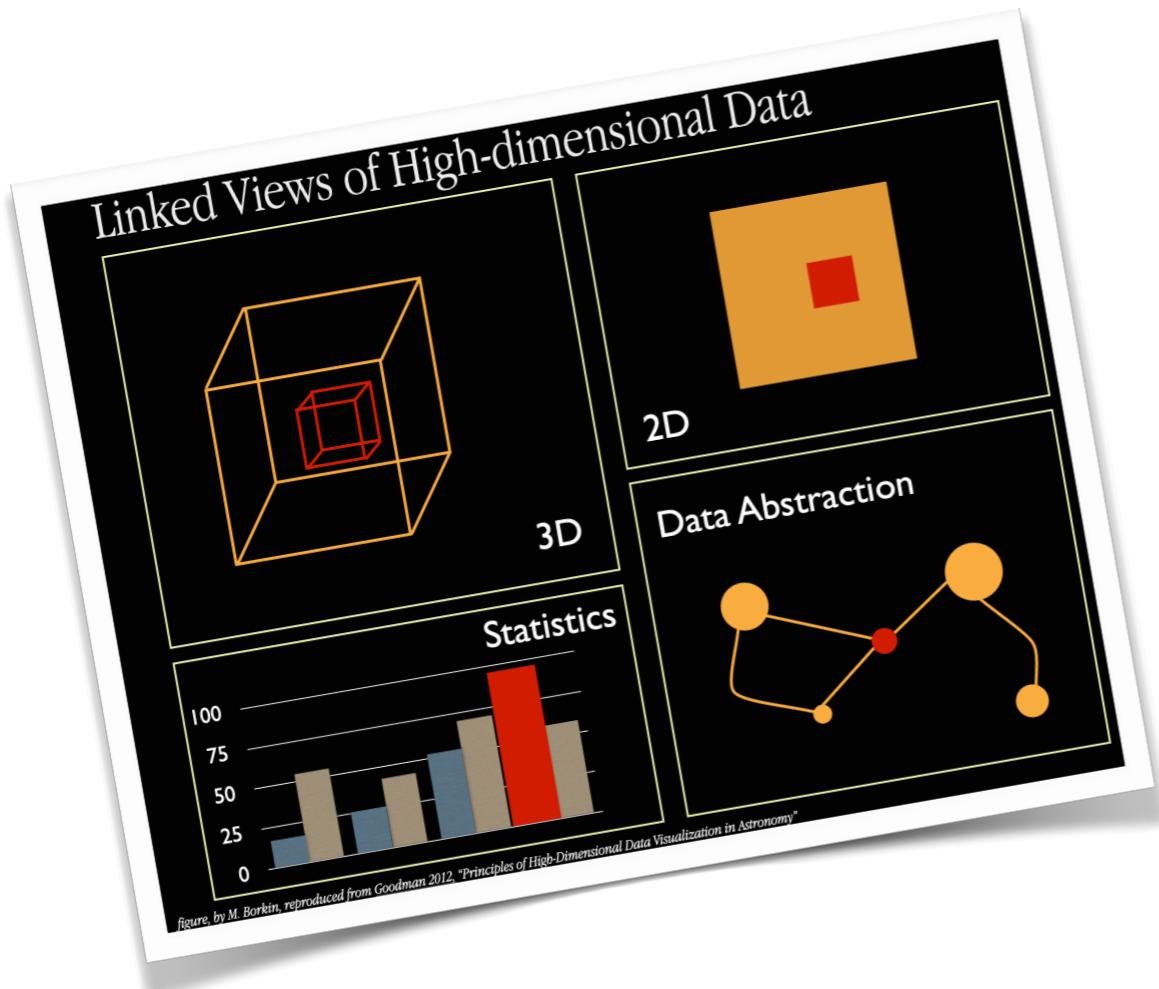
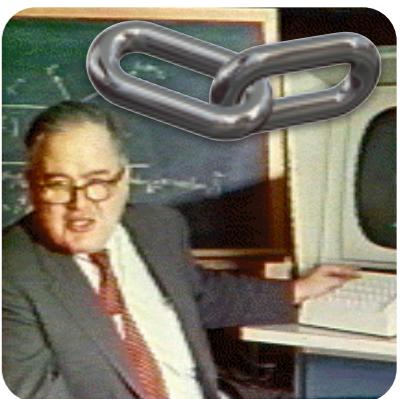


DIVERSE TOOLS

DIVERSE DATA

DIVERSE VIEWS

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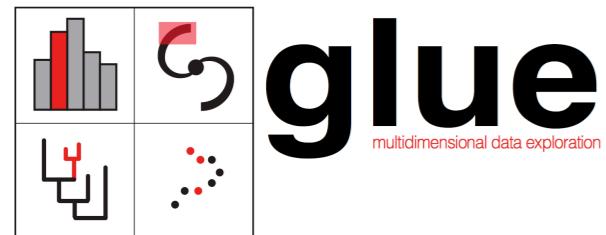


highlight live or
algorithmic selections
with Boolean logic

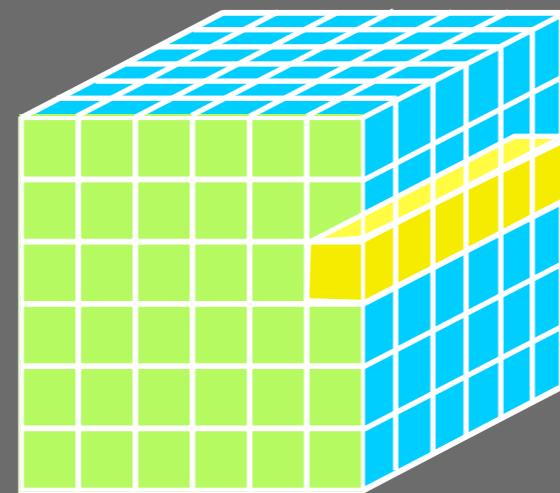
DIVERSE VIEWS

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PRACTICALITY



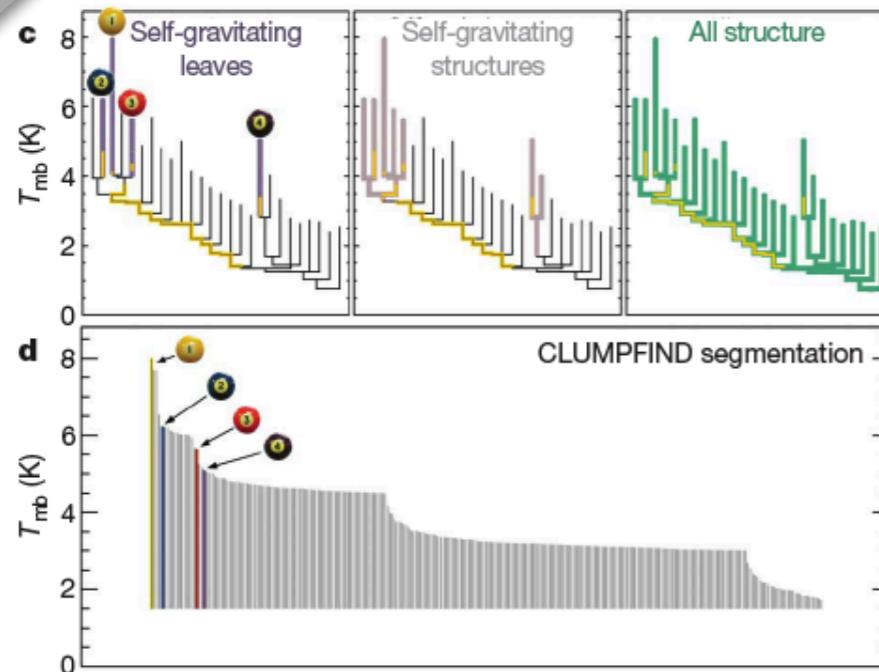
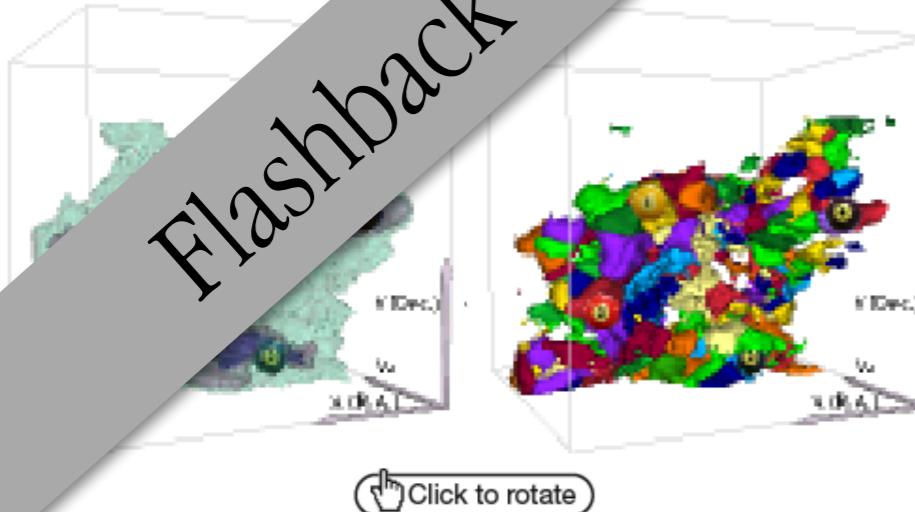
PRINCIPLES



“High-dimensional” or “Multivariate” Data (Astronomy=Biology)

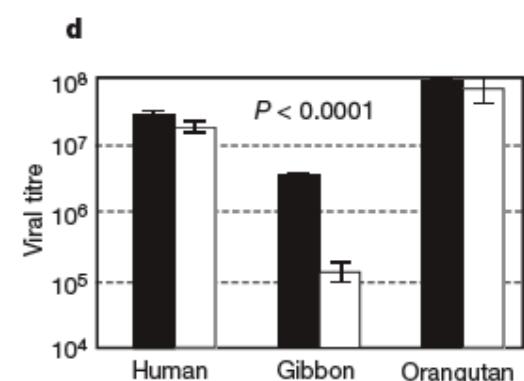
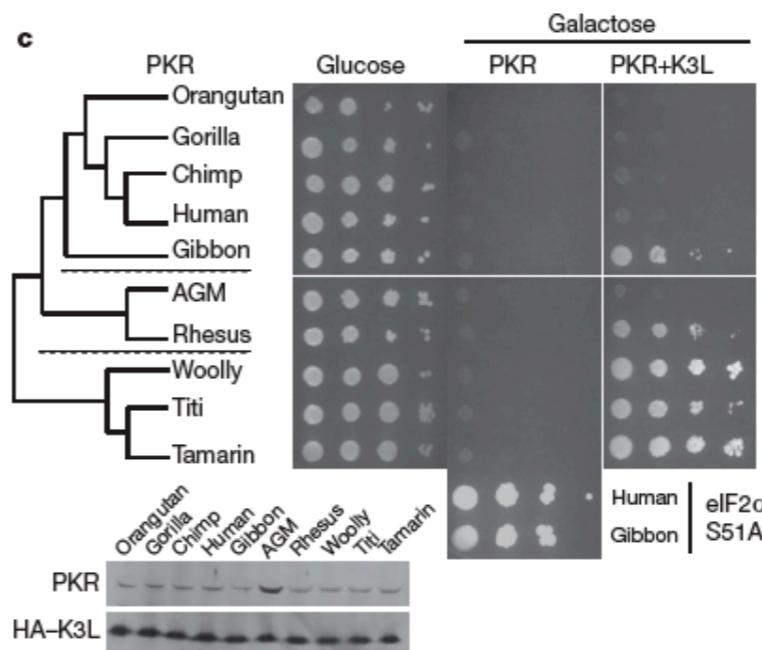
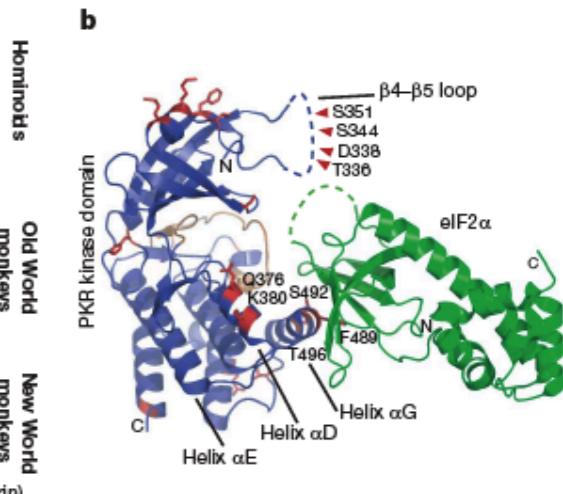
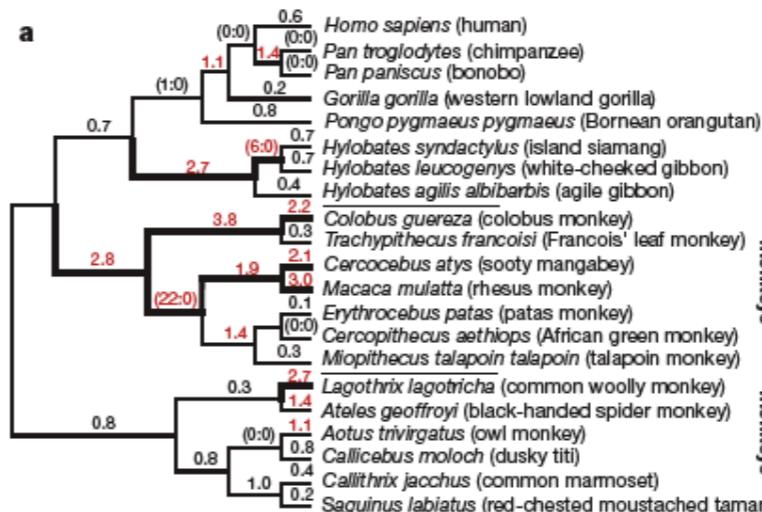
Flashback to 2009...

LETTERS



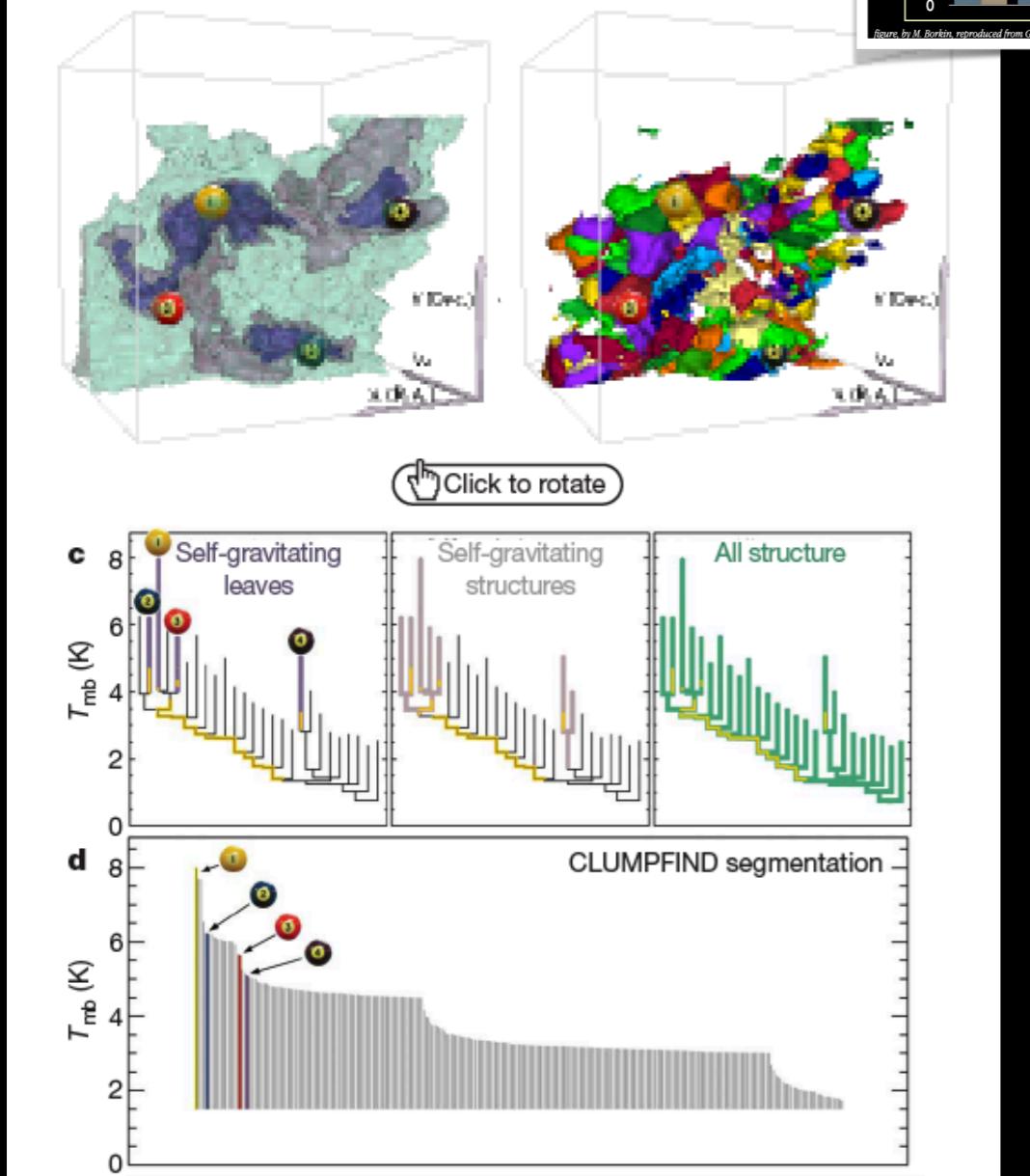
Goodman et al. *Nature*, 2009

LETTERS

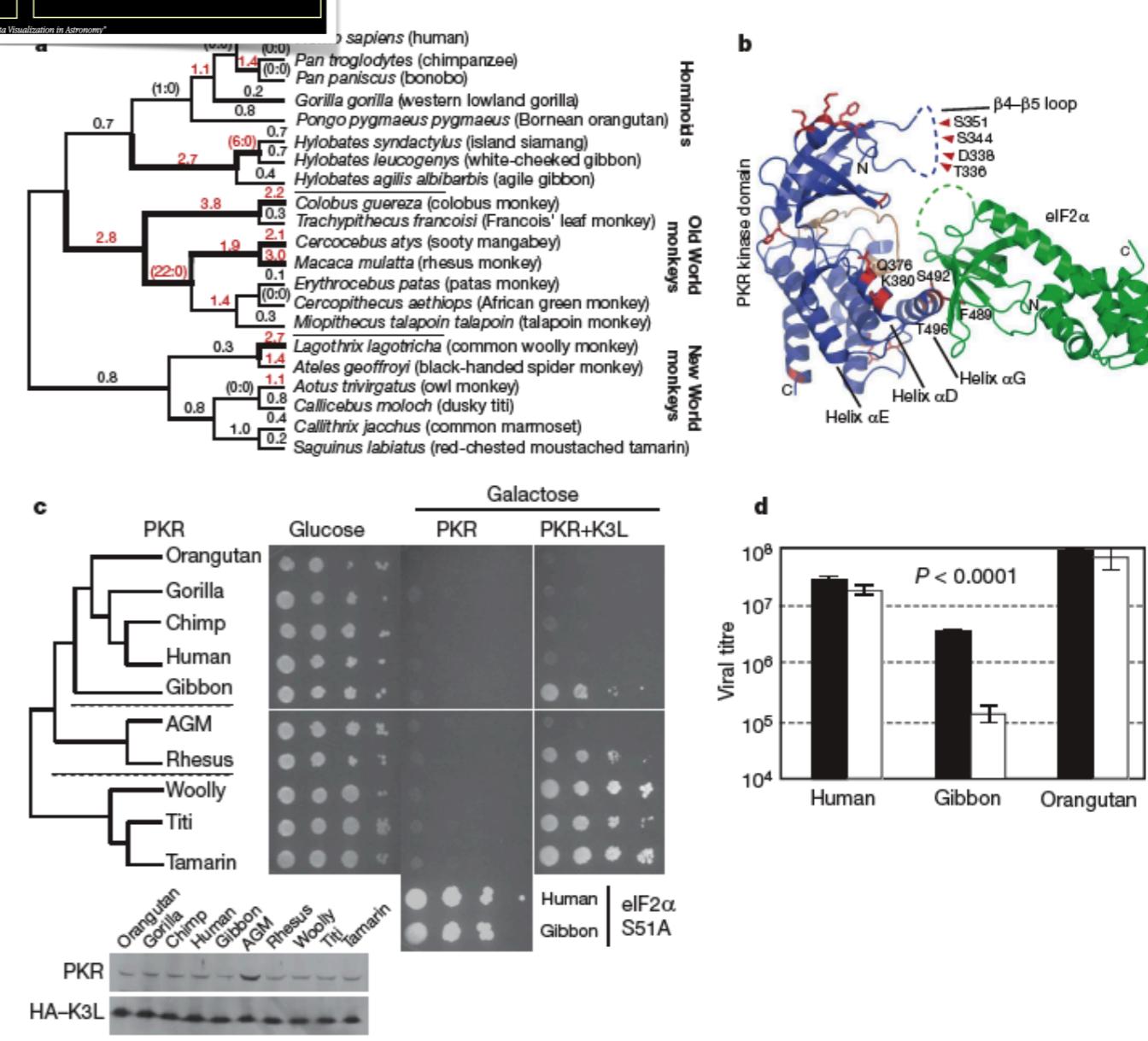


Elde et al. *Nature*, 2008

LETTERS

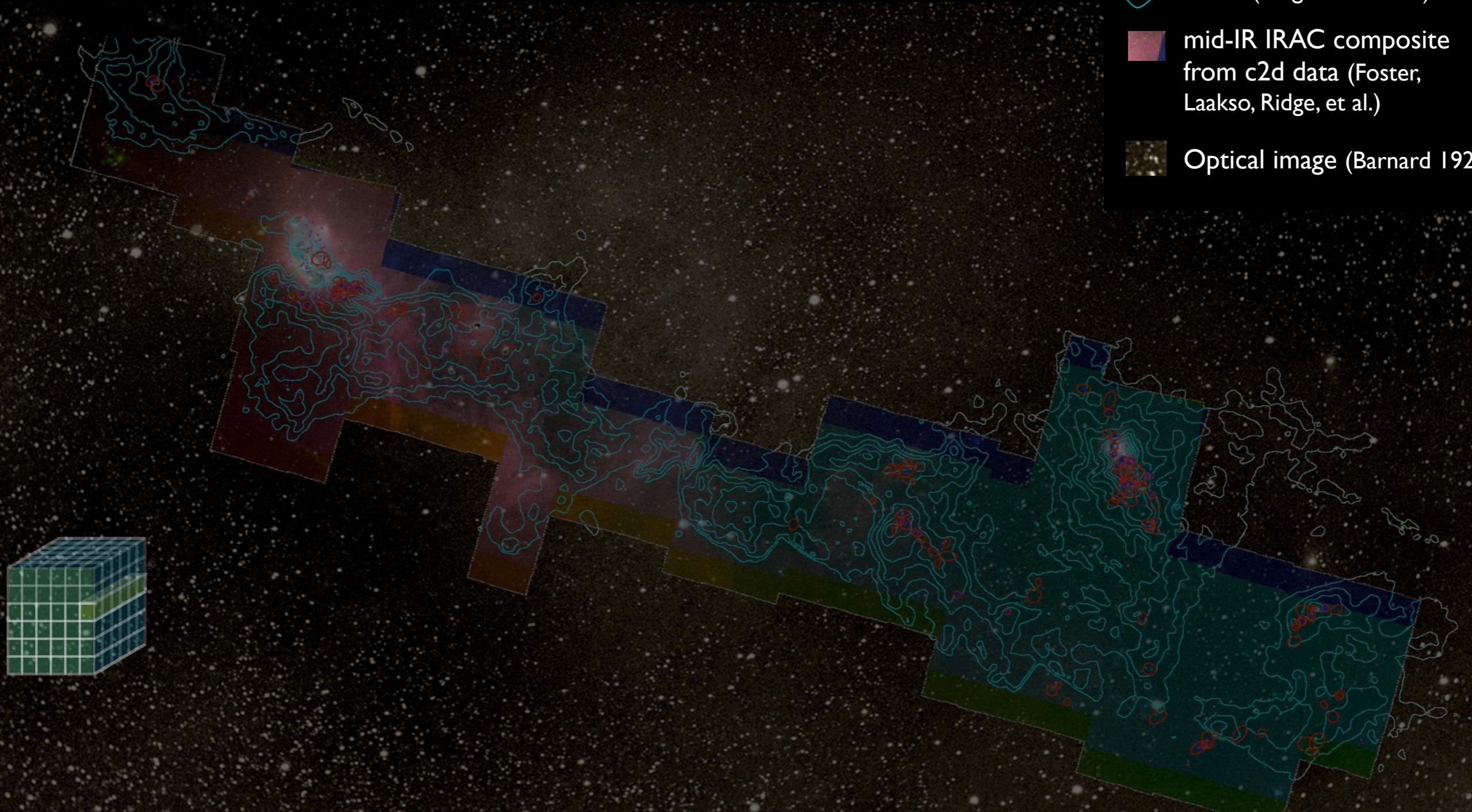


Goodman et al. *Nature*, 2009



DIVERSE HIGH-D DATA

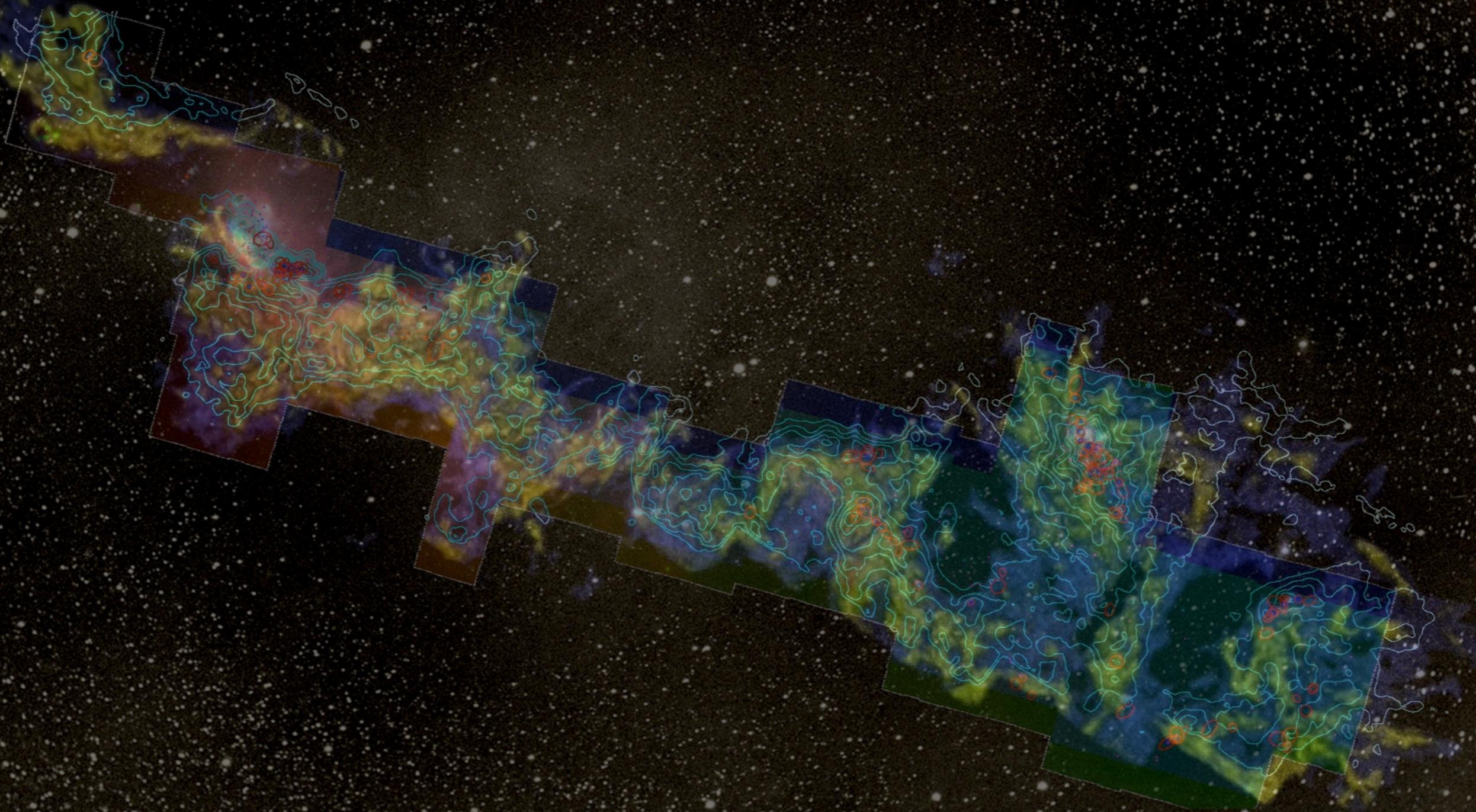
- mm peak (Enoch et al. 2006)
- sub-mm peak (Hatchell et al. 2005, Kirk et al. 2006)
- ^{13}CO (Ridge et al. 2006)
- mid-IR IRAC composite from c2d data (Foster, Laakso, Ridge, et al.)
- Optical image (Barnard 1927)



n: 1/249

Zoom: 227% Angle: 0

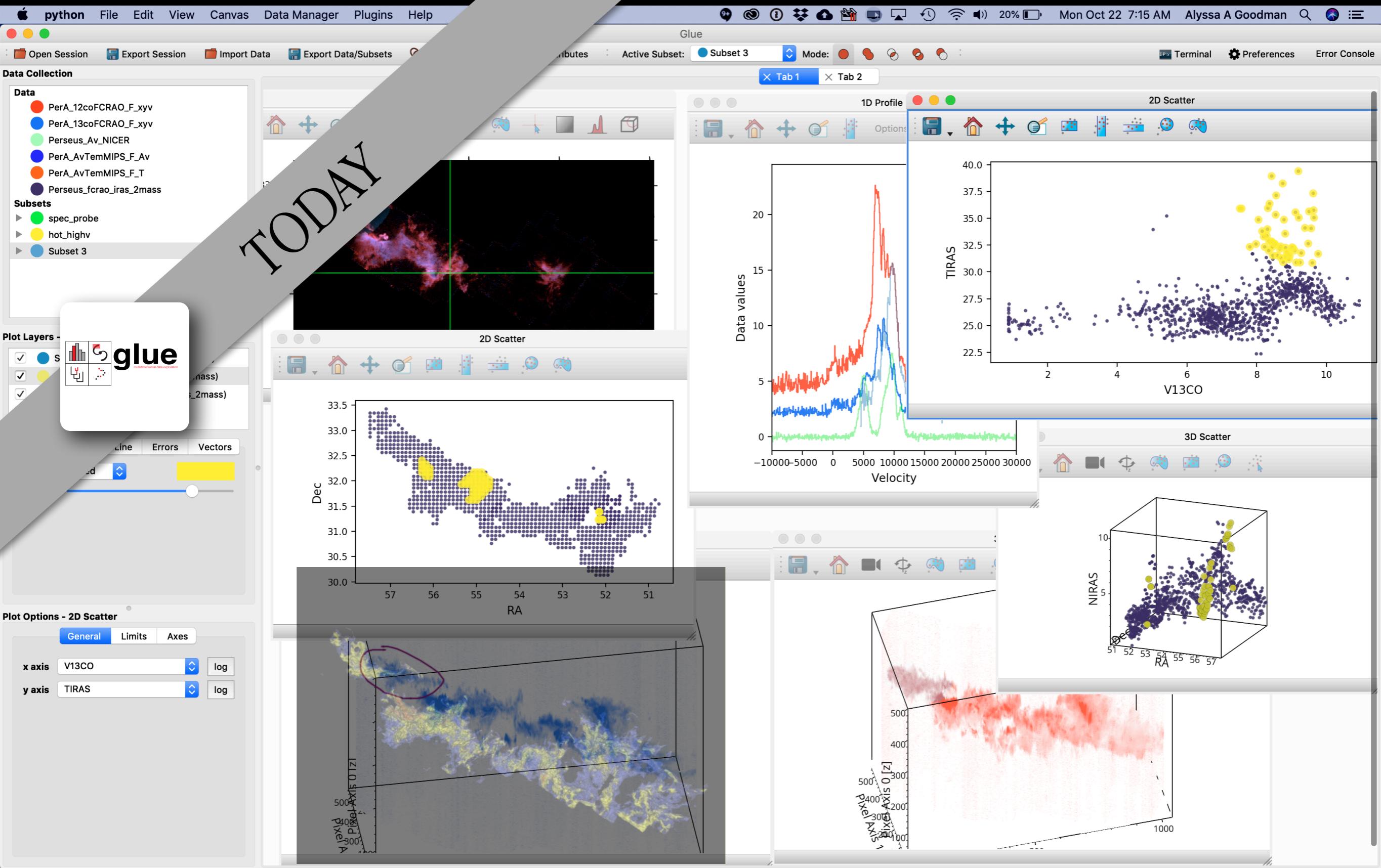




3D Viz made with VolView

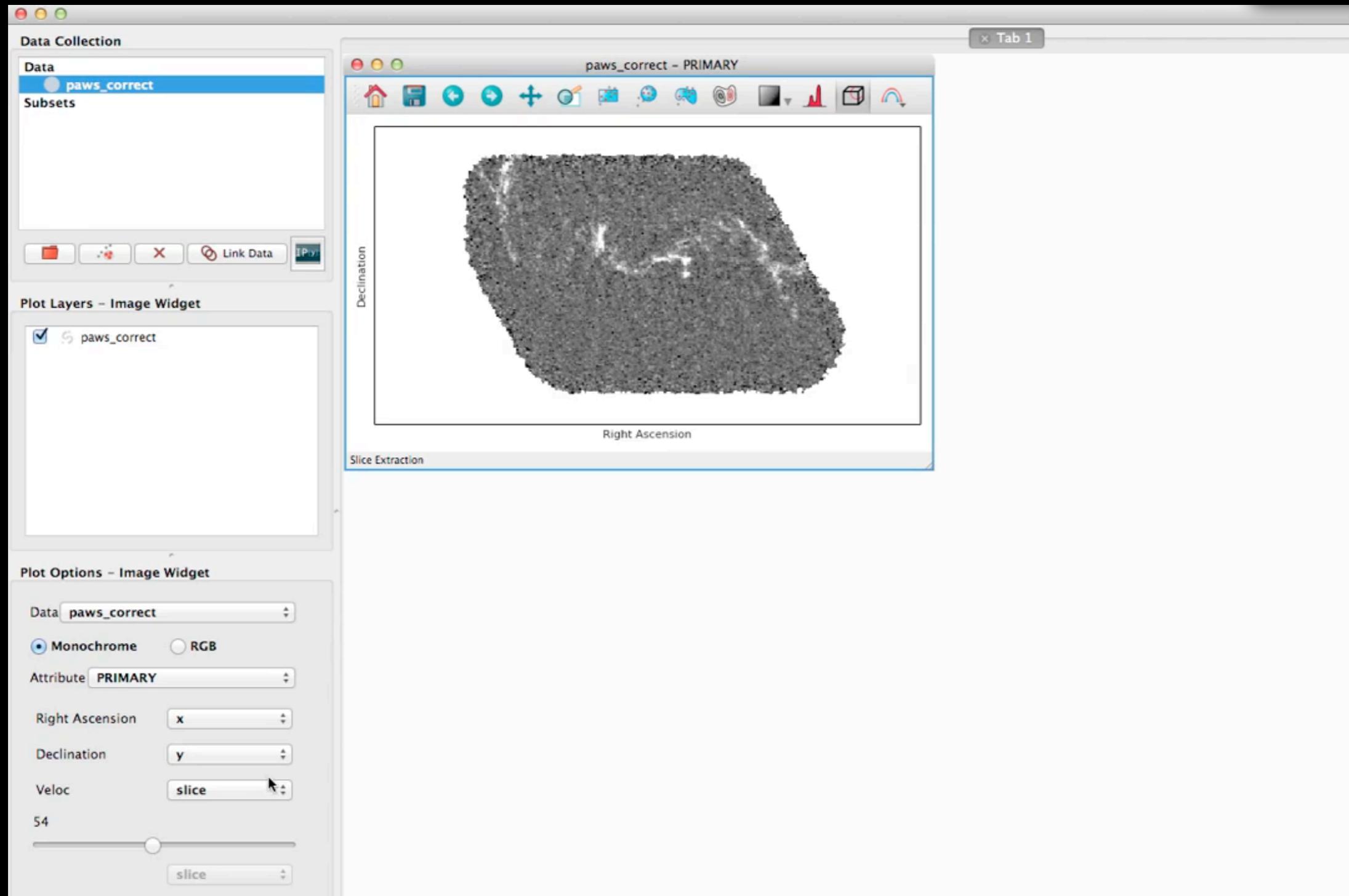
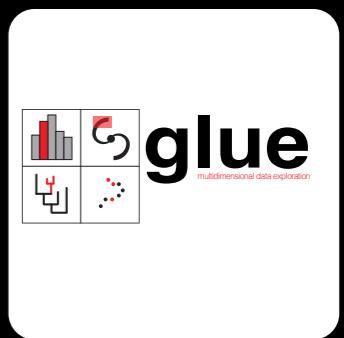
Astronomical Medicine @  IIG

COMPLETE



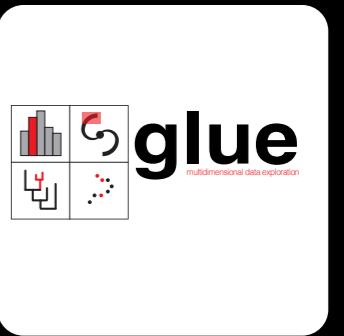
“Dimension” doesn’t always mean x , y , or z . . .

glue

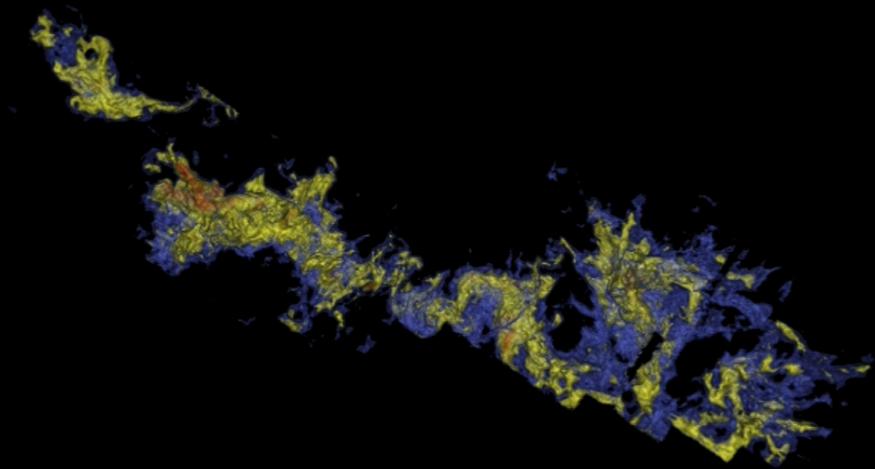


video by Chris Beaumont, glue developer
glue created by: C. Beaumont, M. Borkin, M. Breddels, T. Robitaille, C. Zucker, and A. Goodman, PI

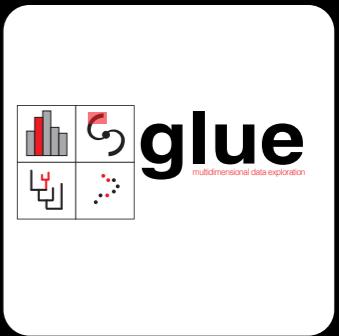
“Dimension” isn’t even always spatial . . .



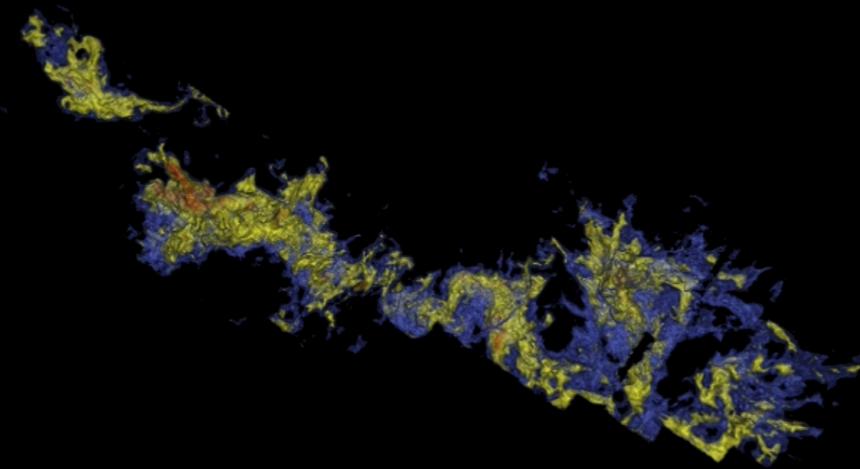
The “3rd” dimension in this 3D plot is “velocity” coming from Doppler Spectroscopy.



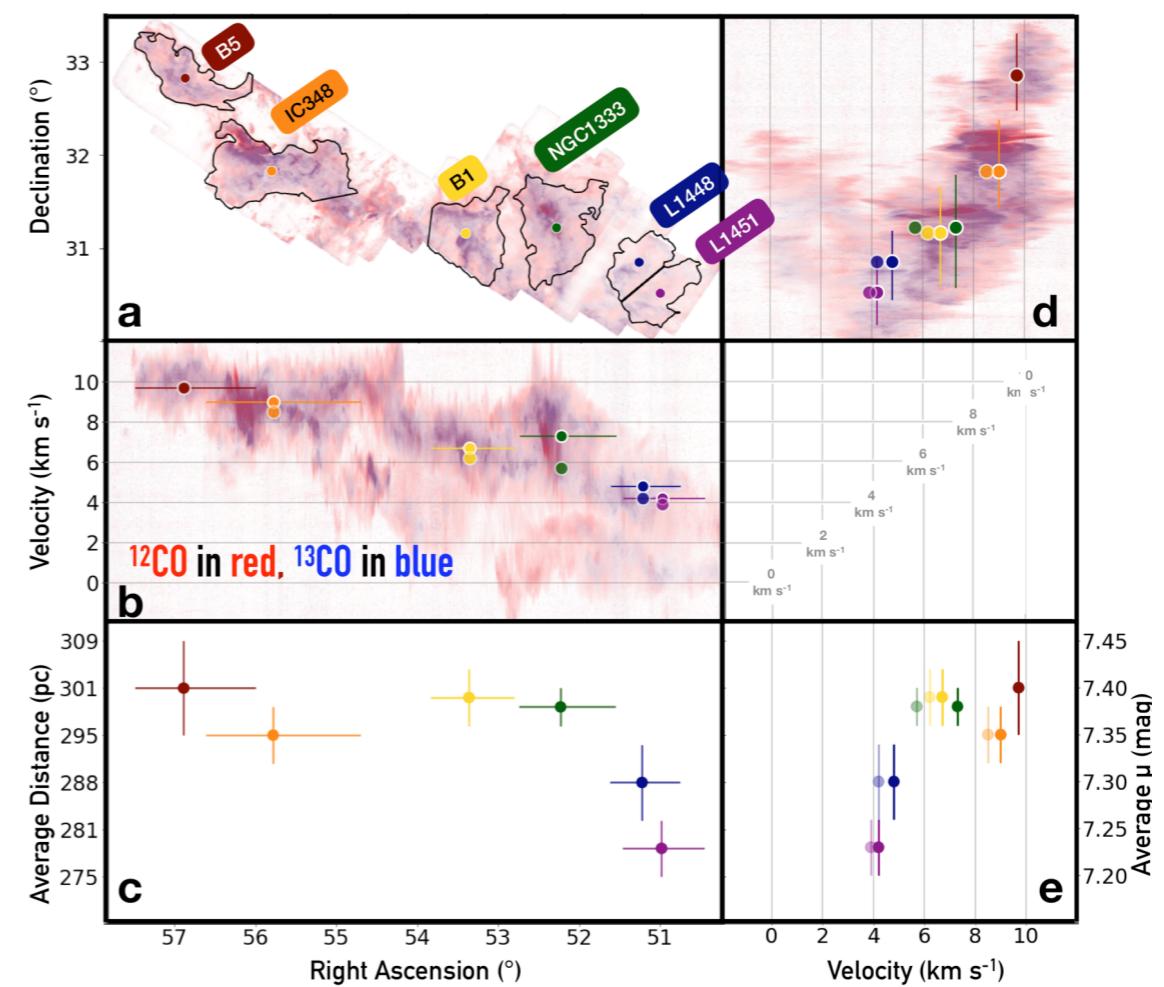
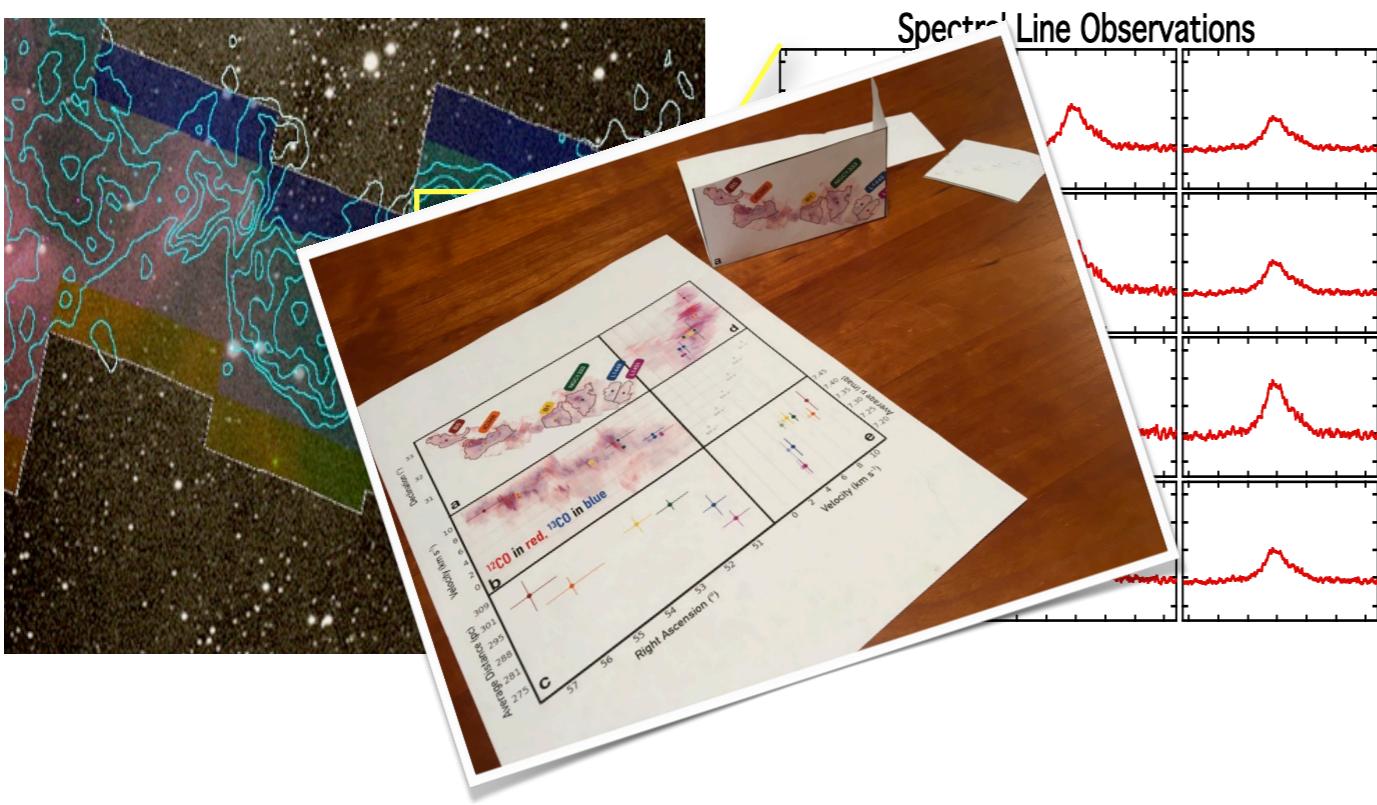
“Dimension” isn’t even always spatial . . .

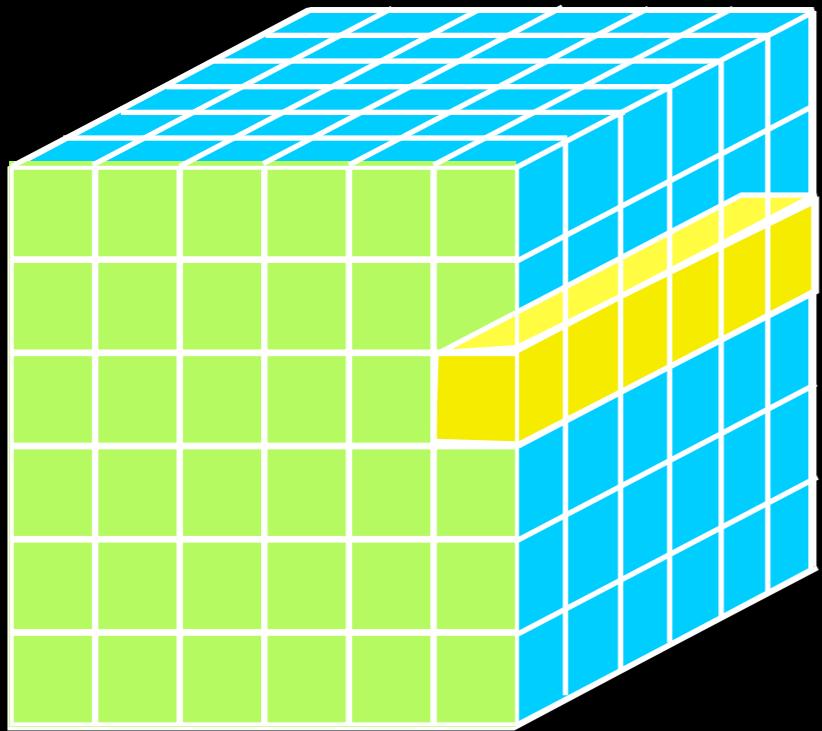


The “3rd” dimension in this 3D plot is “velocity” coming from Doppler Spectroscopy.



“Perseus” PROGRESS PREVIEW



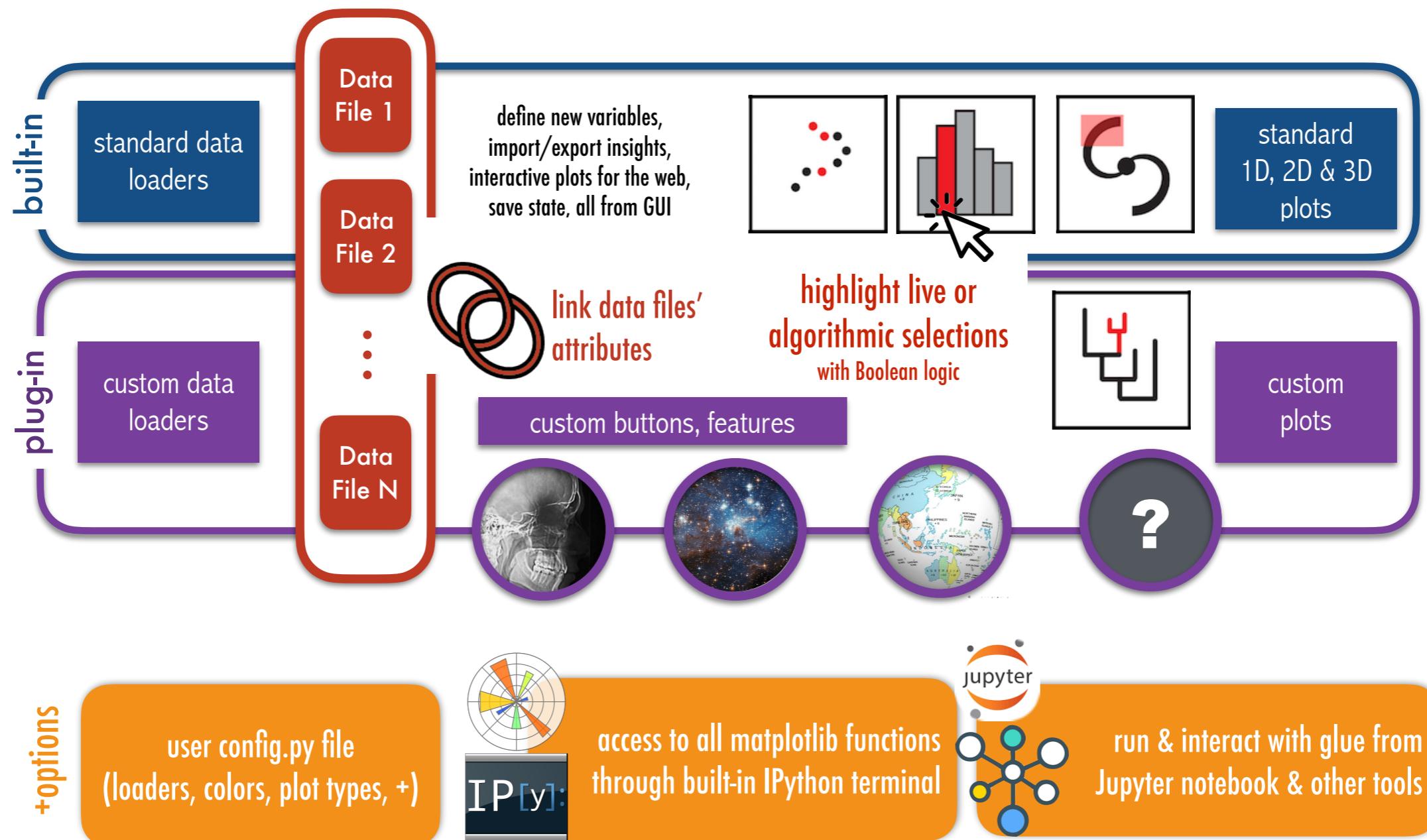
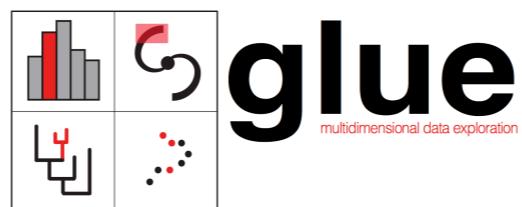


PRINCIPLE "DATA, DIMENSIONS, DISPLAY"

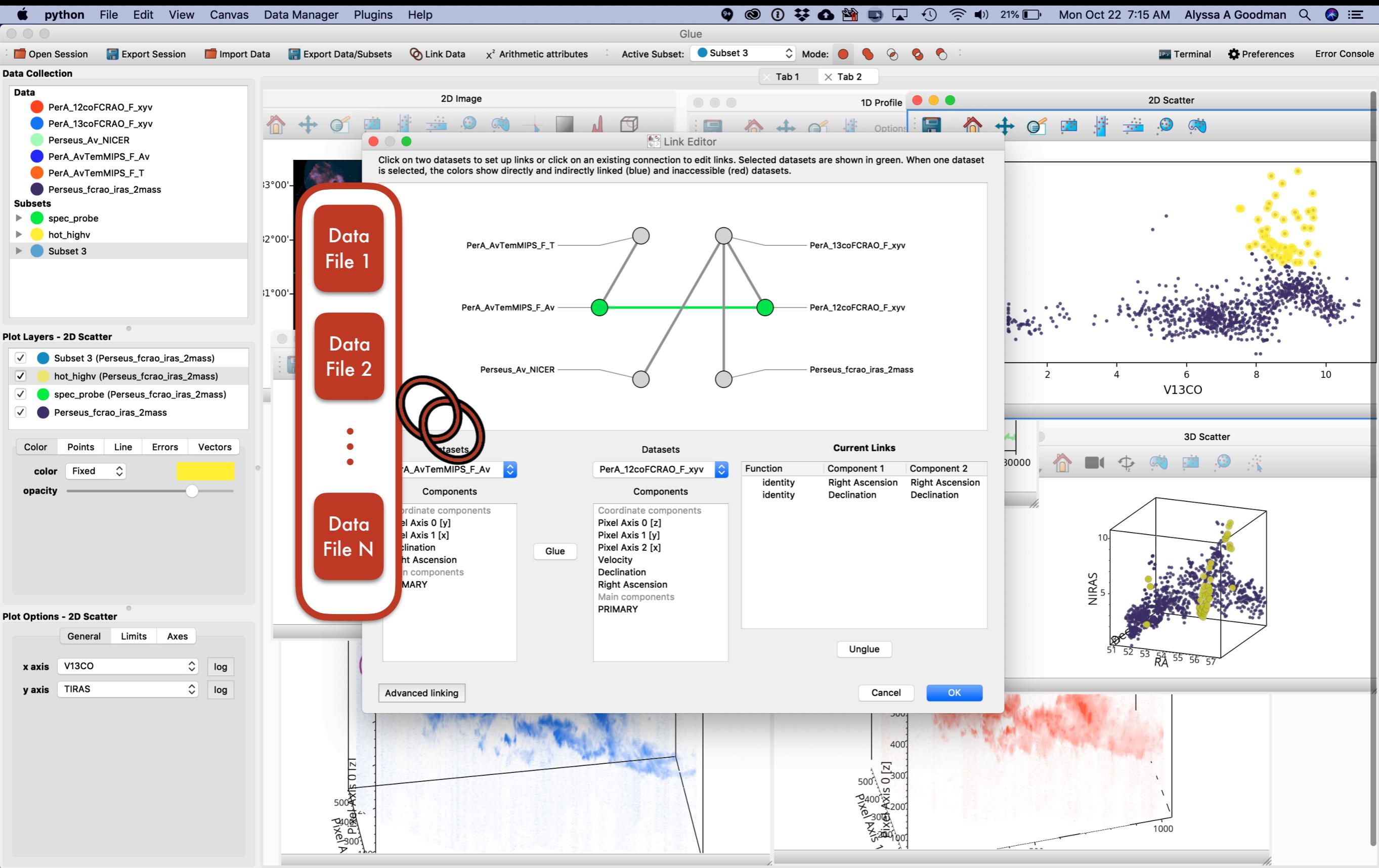
- 1D:** Columns = "Spectrum", "Time Series," "Sequence"
- 2D:** Faces or Slices = "Images," "Arrays"
- 3D:** Volumes = "3D Renderings", "2D Movies"
- 4D:** Time Series of Volumes = "3D Movies"

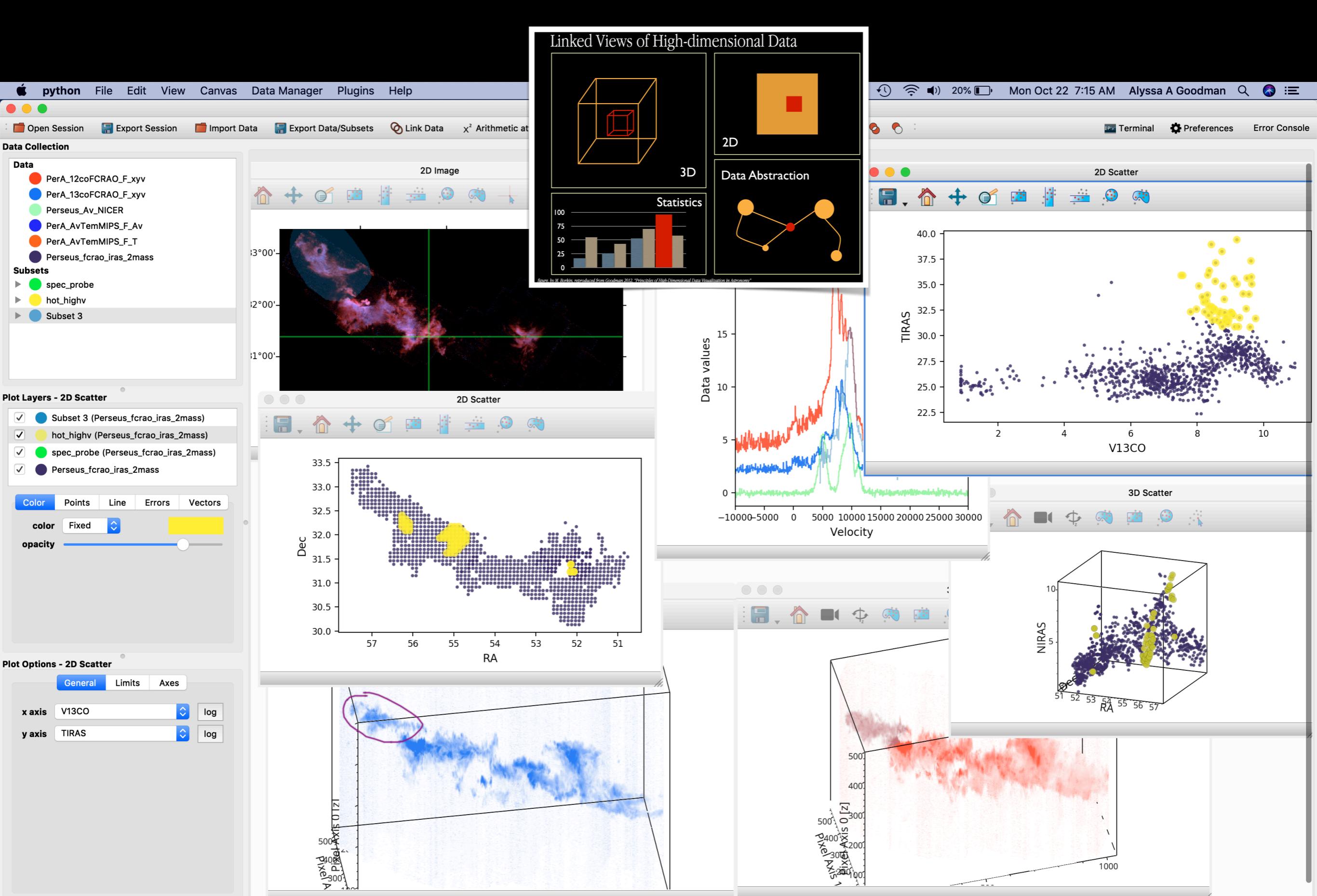
DIVERSE DATA

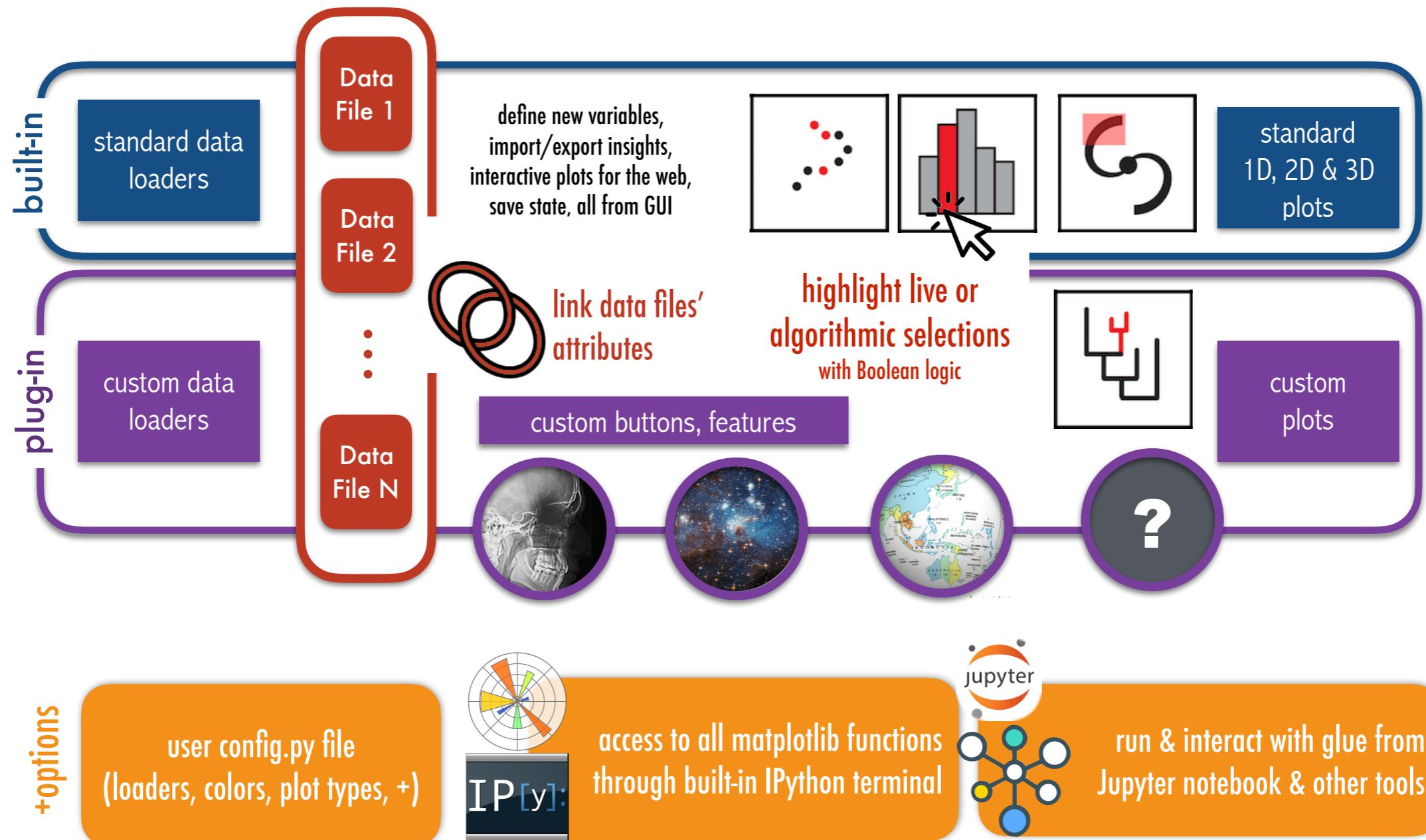
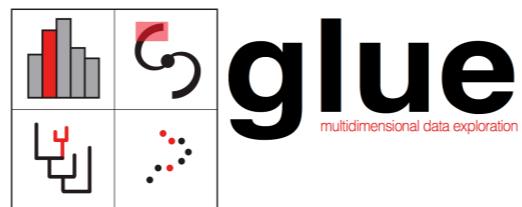
DIVERSE VIEWS



DIVERSE DATA
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DIVERSE TOOLS

DIVERSE DATA

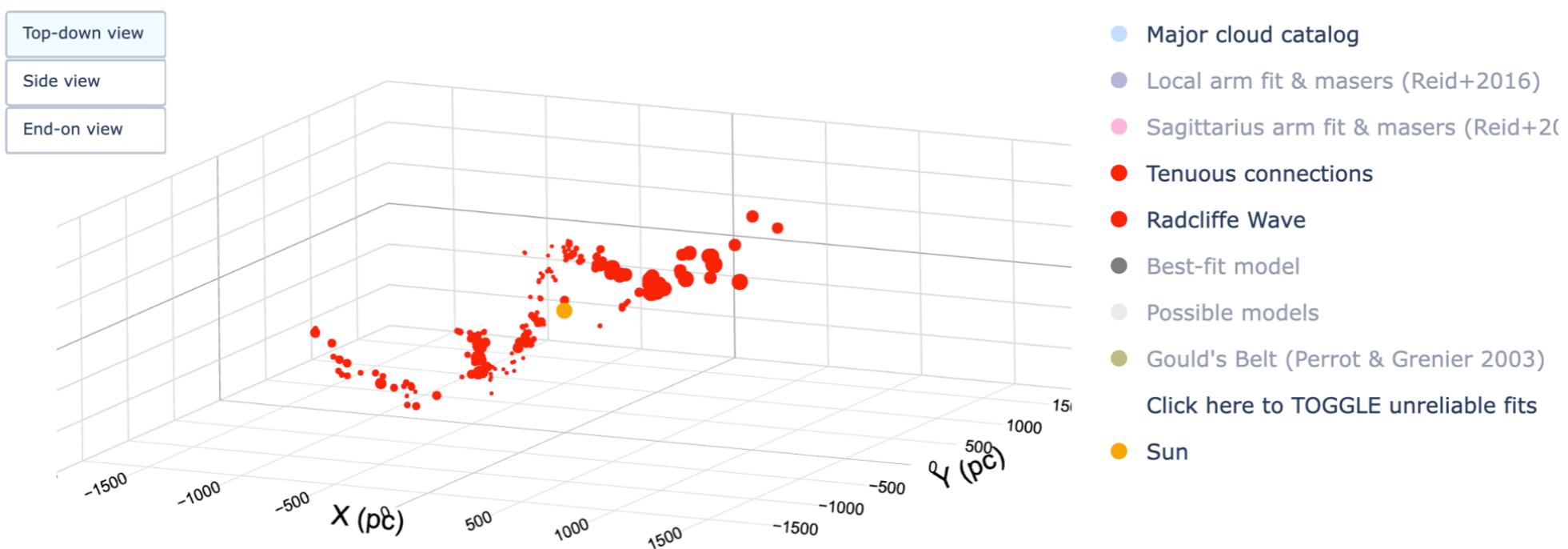
DIVERSE VIEWS

VISUALS

On this page: INTERACTIVES, FIGURES, VIDEOS -- scroll down to see it all.

INTERACTIVES

Explore the RadWave in 3D



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bioRxiv is receiving many new papers on coronavirus SARS-CoV-2. A reminder: these are preliminary reports that have not been peer-reviewed, practice/health-related behavior, or be reported in news media as established information.

New Results

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Cellular taxonomy and spatial organization of the ventral posterior hypothalamus reveals neuroanatomical parcellation of the mammillary bodies

Laura E. Mickelsen, William F. Flynn, Kristen Springer, Lydia Wilson, Eric J. Beltrami, Mohan Bolisetty, Paul Robson, Alexander C. Jackson

doi: <https://doi.org/10.1101/2020.05.14.096818>

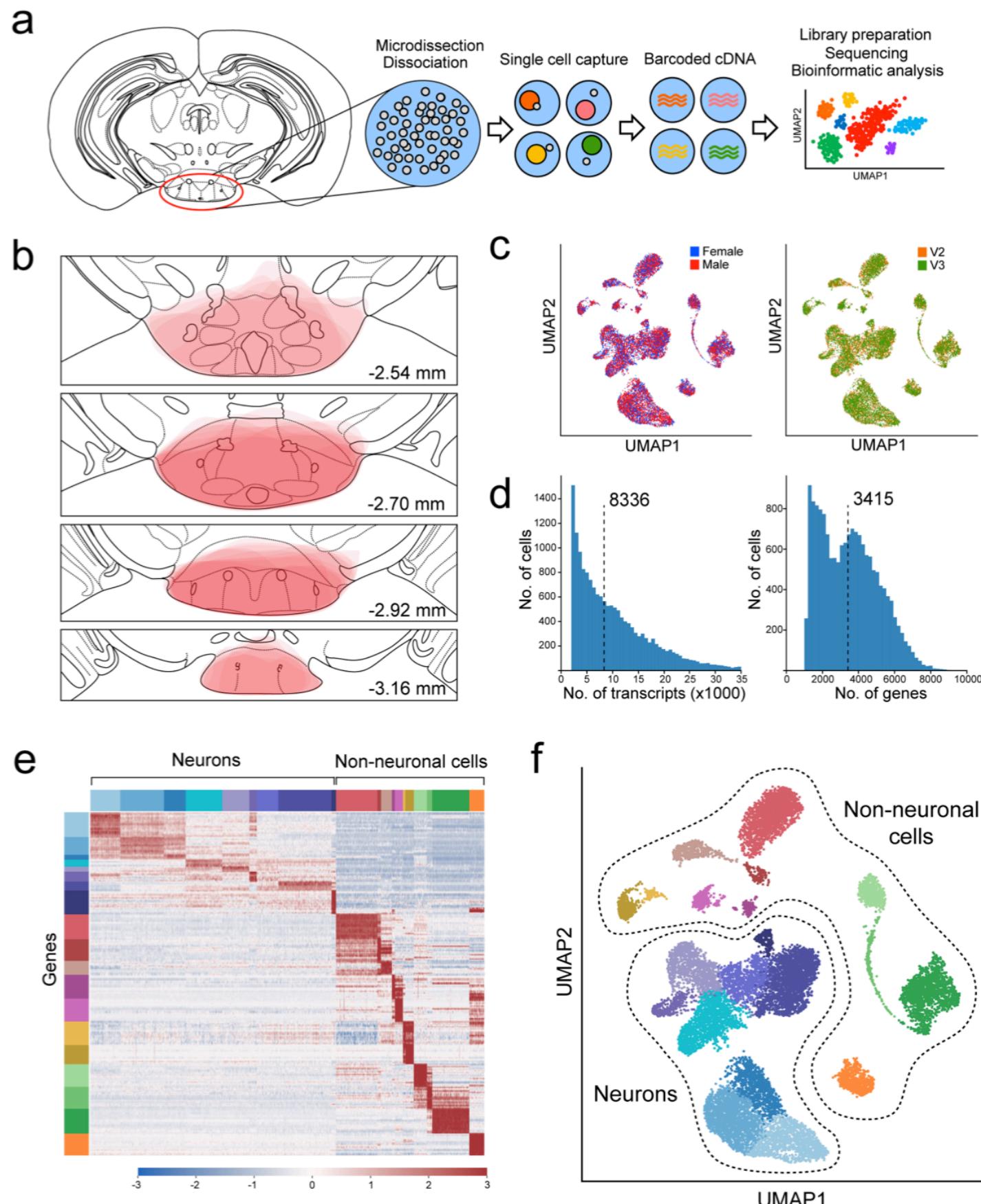
This article is a preprint and has not been certified by peer review [what does this mean?].

Abstract Full Text Info/History Metrics

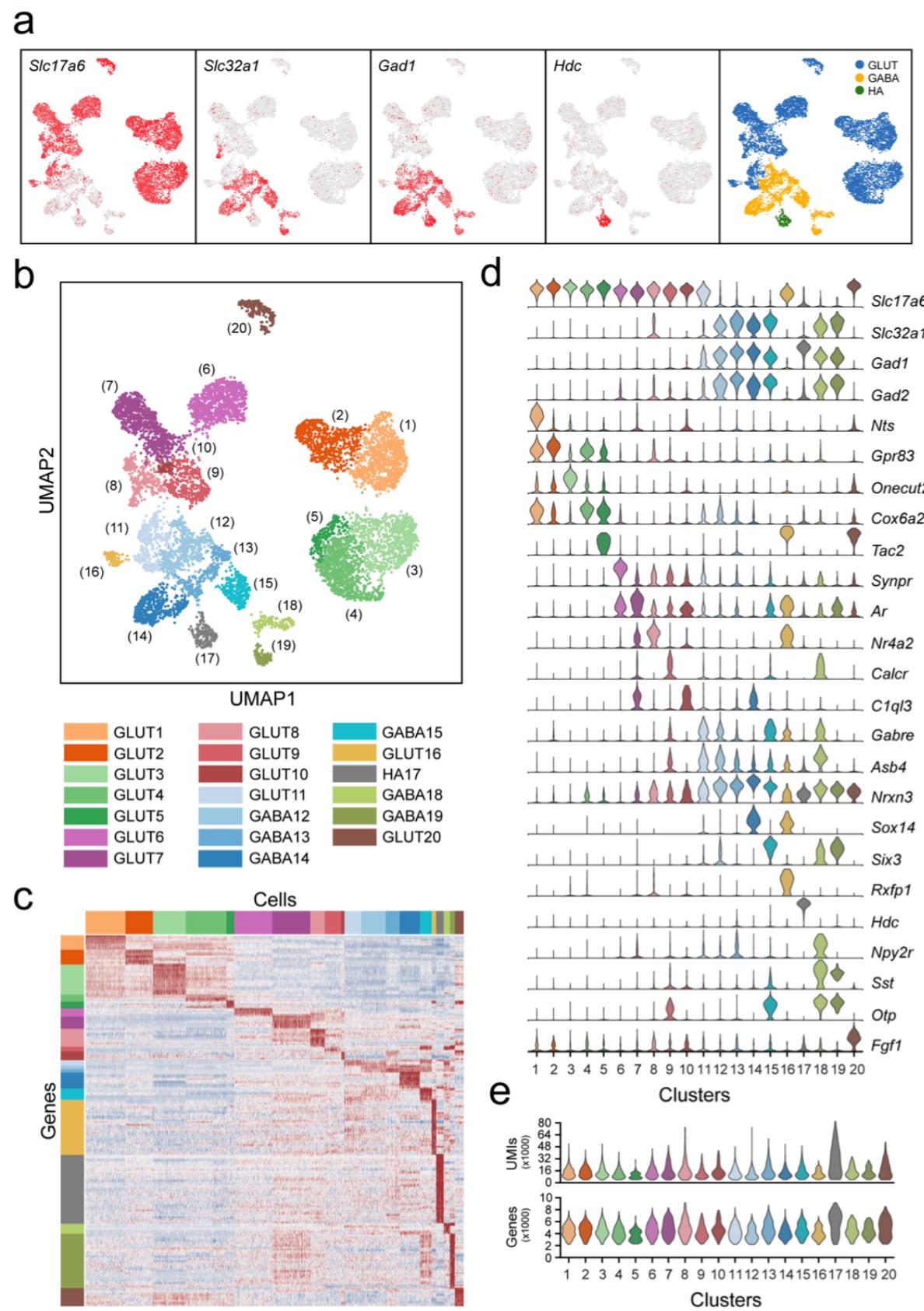
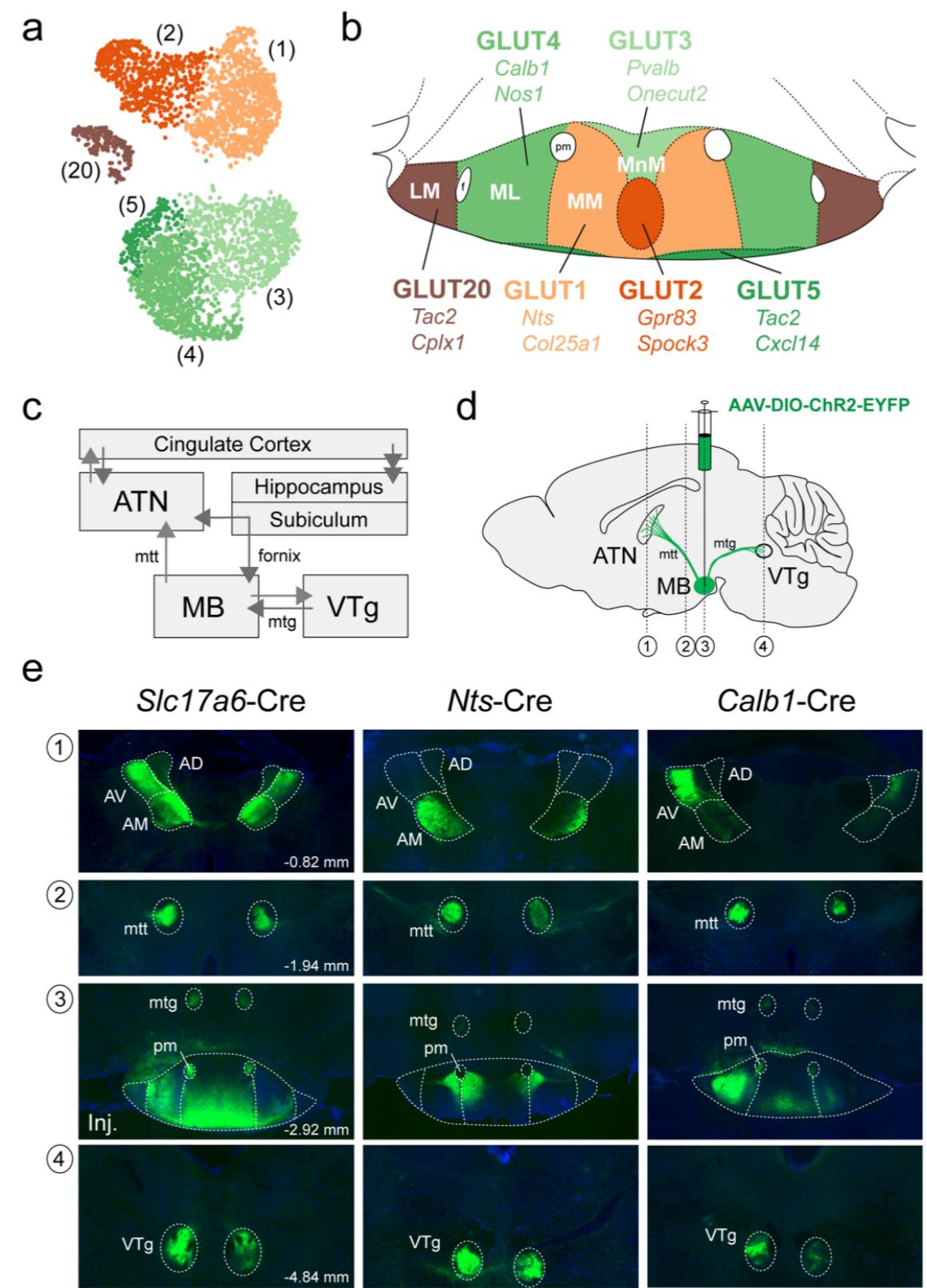
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ABSTRACT

The ventral posterior hypothalamus (VPH) is an anatomically complex brain region implicated in arousal, reproduction, energy balance and memory processing. However, neuronal cell type diversity within the VPH is poorly understood, an impediment to deconstructing the roles of distinct VPH circuits in physiology and behavior. To address this question, we employed a droplet-based single cell RNA sequencing (scRNA-seq) approach to systematically classify molecularly distinct cell types in the mouse VPH. Analysis of >16,000 single cells revealed 20 neuronal and 18 non-neuronal cell populations, defined by suites of discriminatory markers. We validated differentially expressed genes in a selection of neuronal populations through fluorescence *in situ* hybridization (FISH). Focusing on the mammillary bodies (MB), we discovered transcriptionally-distinct clusters that exhibit a surprising degree of segregation within neuroanatomical subdivisions of the MB, while genetically-defined MB cell types project topographically to the anterior thalamus. This single cell transcriptomic atlas of cell types in the VPH provides a detailed resource for interrogating the circuit-level mechanisms underlying the diverse functions of VPH circuits in health and disease.



accepted for publication at eLife (Mickelsen, Flynn, ... Robson, Jackson) 'Cellular taxonomy and spatial organization of the murine ventral posterior hypothalamus' doi: 10.7554/eLife.58901

Figure 2:**Figure 8:**



Genomes

Genome Browser

Tools

Mirrors

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My Data

View

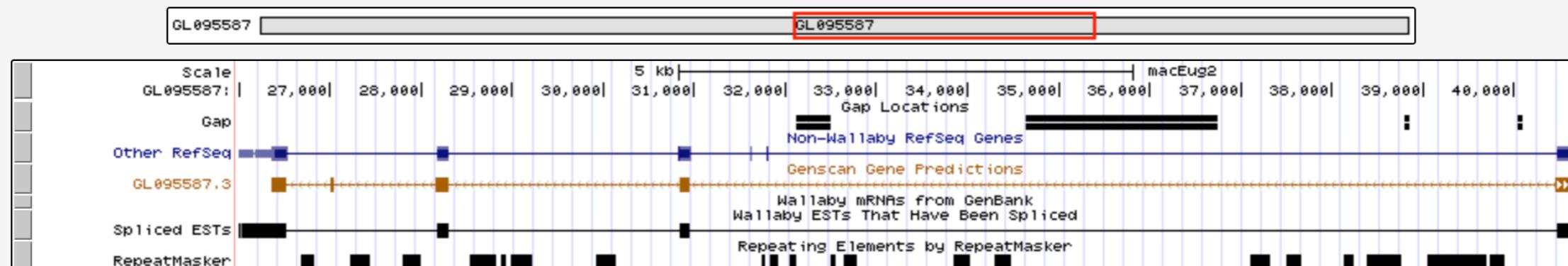
Help

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UCSC Genome Browser on Wallaby Sep. 2009 (TWGS Meug_1.1/macEug2) Assembly

move <<< << < > >>> zoom in 1.5x 3x 10x base zoom out 1.5x 3x 10x 100x

GL095587:25,998-40,635 14,638 bp. enter position or search terms go



move start

< 2.0 >

Click on a feature for details. Click or drag in the base position track to zoom in. Click side bars for track options. Drag side bars or labels up or down to reorder tracks. Drag tracks left or right to new position. Press "?" for keyboard shortcuts.

move end

< 2.0 >

track search

default tracks

default order

hide all

add custom tracks

track hubs

configure

multi-region

reverse

resize

refresh

collapse all

expand all

Use drop-down controls below and press refresh to alter tracks displayed.

Tracks with lots of items will automatically be displayed in more compact modes.



Mapping and Sequencing

refresh

[Base Position](#)

dense ▾

[Assembly](#)

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[Gap](#)

dense ▾

[GC Percent](#)

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[INSDC](#)

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[Restr Enzymes](#)

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[Short Match](#)

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Comparative Genomics

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Variation and Repeats

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[RepeatMasker](#)

dense ▾

[Interrupted Rpts](#)

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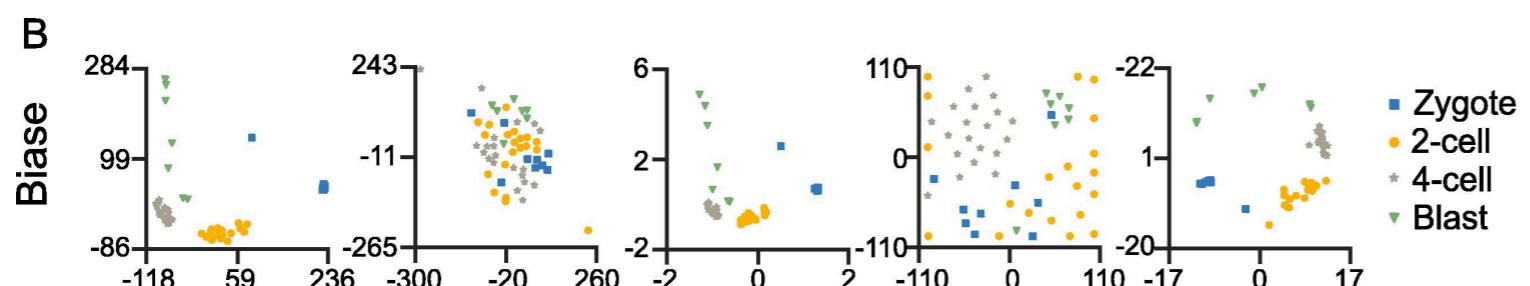
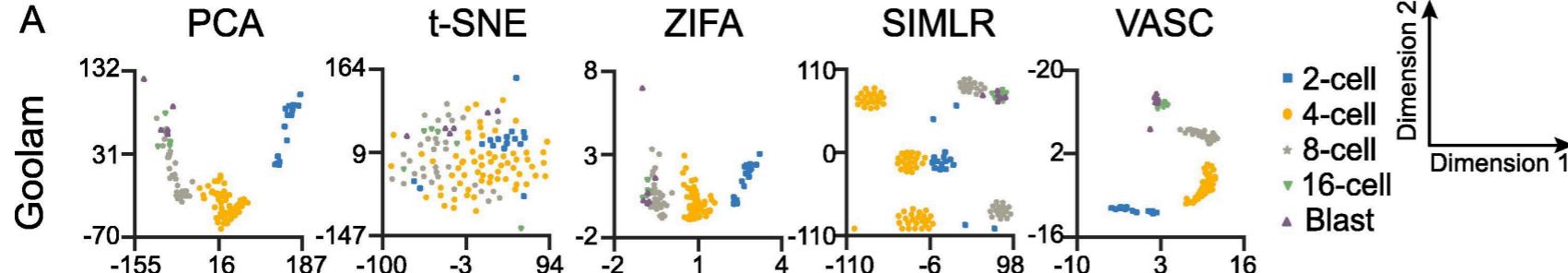
[Simple Repeats](#)

hide ▾

[WM + SDust](#)

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refresh



Method
VASC: Dimension Reduction and Visualization of Single-cell RNA-seq Data by Deep Variational Autoencoder

Dongfang Wang ^a, Jin Gu ^{a,b}✉

MOE Key Laboratory of Bioinformatics, BNRIST Bioinformatics Division & Center for Synthetic and Systems Biology, Department of Automation, Tsinghua University, Beijing 100084, China

Received 23 March 2018, Revised 9 July 2018, Accepted 8 August 2018, Available online 18 December 2018.

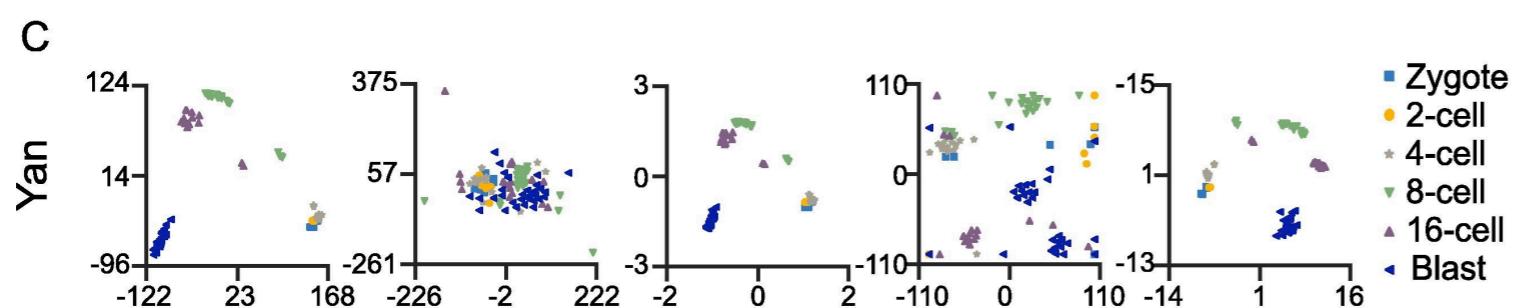
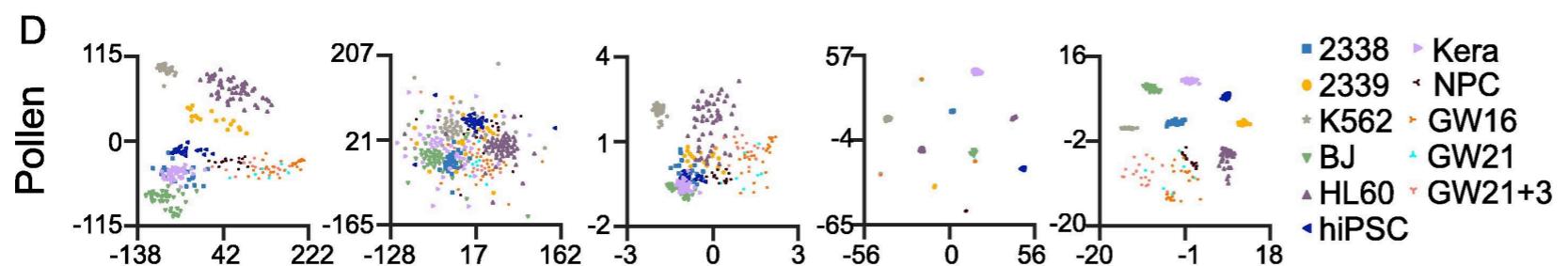
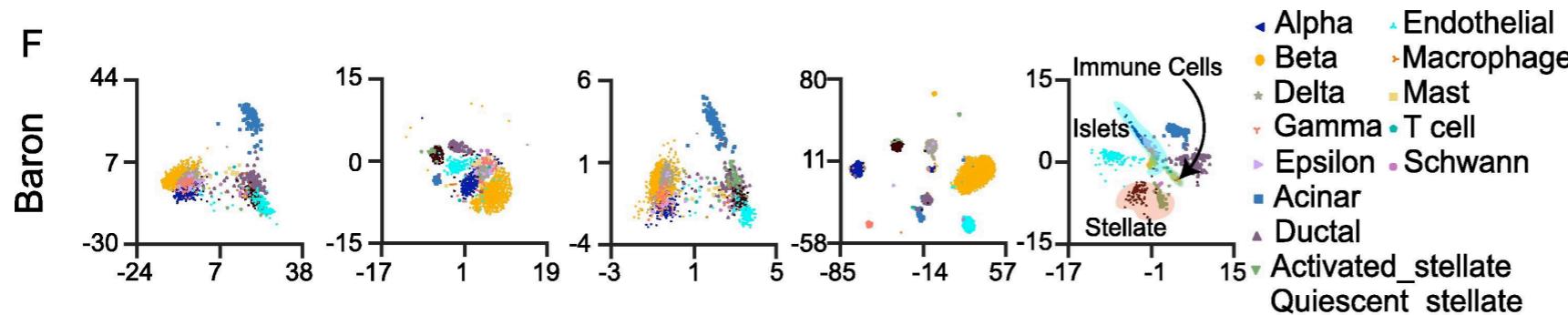
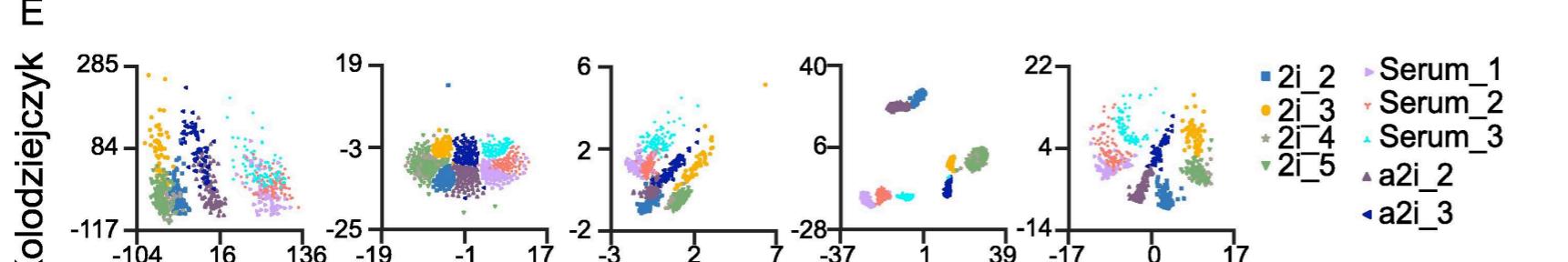


Figure 2. Visualization of scRNA-seq datasets using different methods

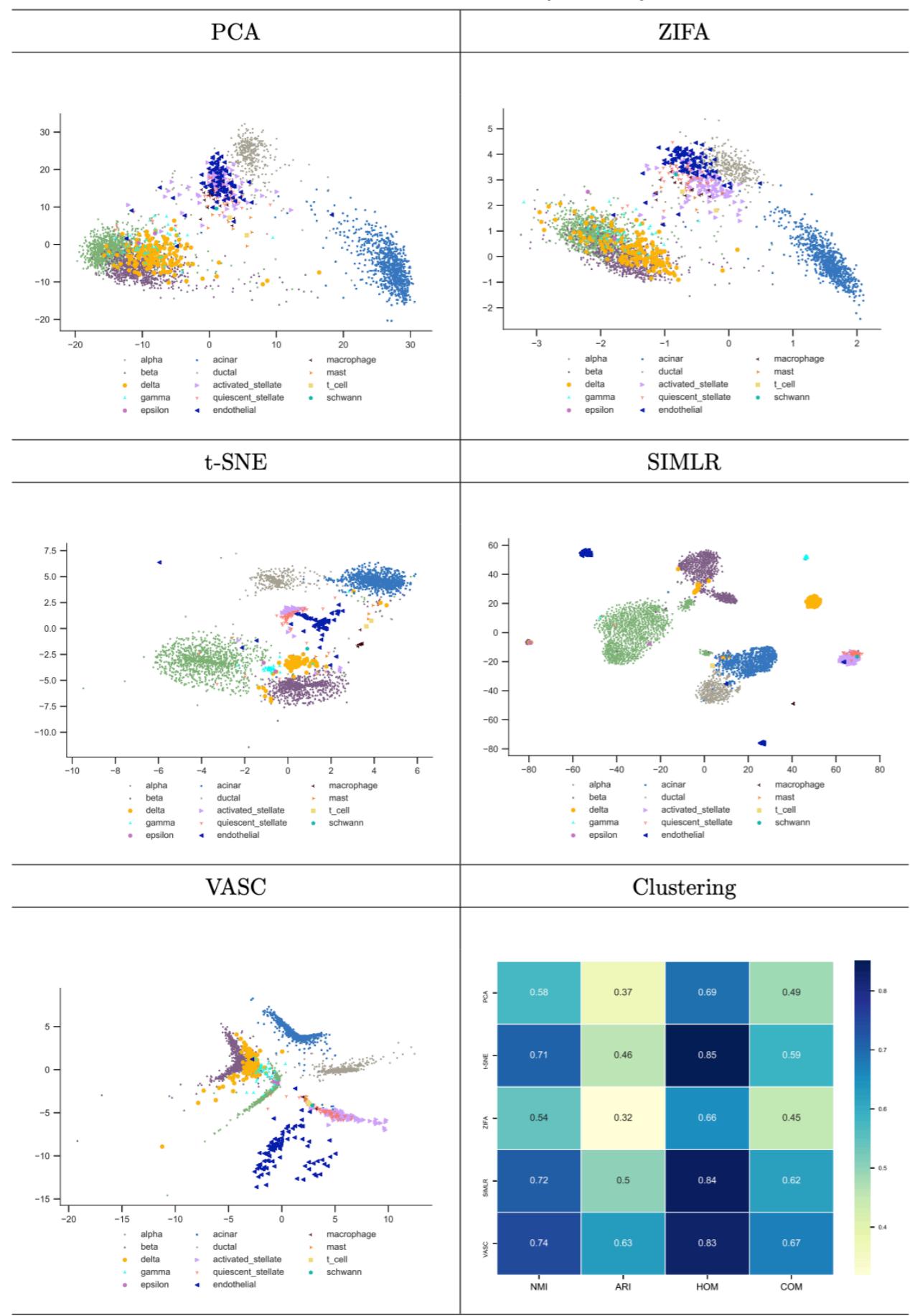


Each data point represents a cell. Different cell types are indicated in different colors and shapes. All datasets were run by PCA, t-SNE, ZIFA, SIMLR, and VASC respectively. Cell type information was retrieved from original studies. Shown in the figures are clustering output from the Goolam [22] (A), Biase [18] (B), Yan [30] (C), Pollen [27] (D), Kolodziejczyk [24] (E), and Baron_human-1 [17] (F) datasets. Visualization of other datasets is provided in the Section 4 of File S1.



PCA, principal components analysis; t-SNE, t-distributed stochastic neighbor embedding; ZIFA, zero-inflated factor analysis; SIMLR, single-cell interpretation via multiple kernel learning.

The “Baron” dataset contains a large number of cells from human and mouse pancreas. Totally, there are 4 human donors with 1937, 1724, 3605, and 1303 cells, and 2 mice with 822 and 1064 cells, respectively



Method
VASC: Dimension Reduction and
Visualization of Single-cell RNA-seq Data by
Deep Variational Autoencoder

Dongfang Wang ^a, Jin Gu ^{a,b}

MOE Key Laboratory of Bioinformatics, BNRIST Bioinformatics Division & Center for Synthetic and Systems Biology, Department of Automation, Tsinghua University, Beijing 100084, China

Received 23 March 2018, Revised 9 July 2018, Accepted 8 August 2018, Available online 18 December 2018.

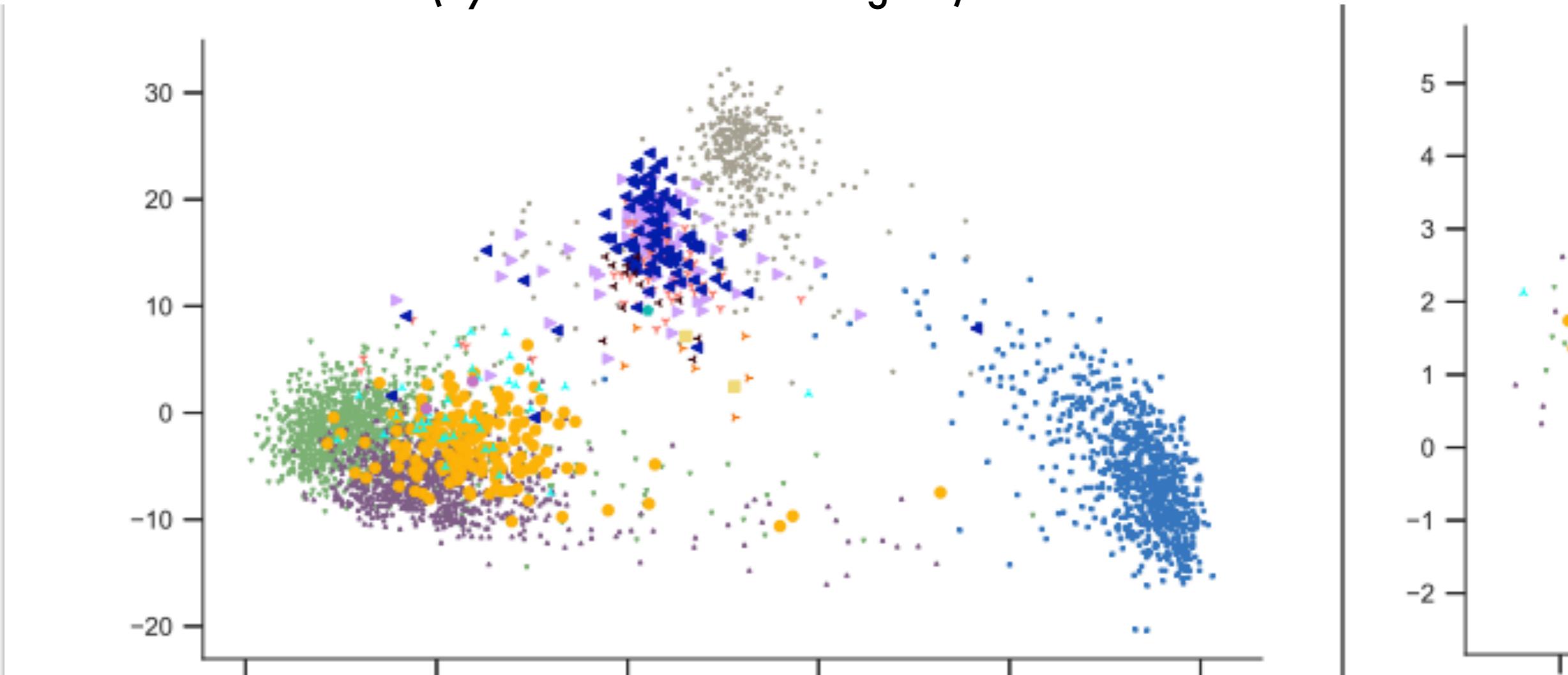
Figure 2. Visualization of scRNA-seq datasets using different methods

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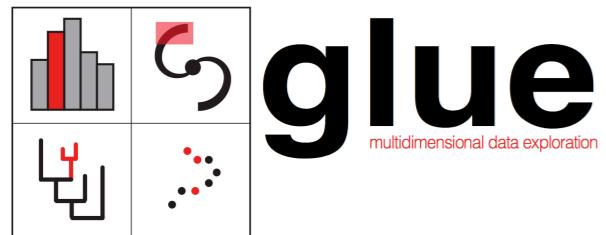
The “Baron” dataset contains a large number of cells from human and mouse pancreas. Totally, there are 4 human donors with 1937, 1724, 3605, and 1303 cells, and 2 mice with 822 and 1064 cells, respectively

PCA

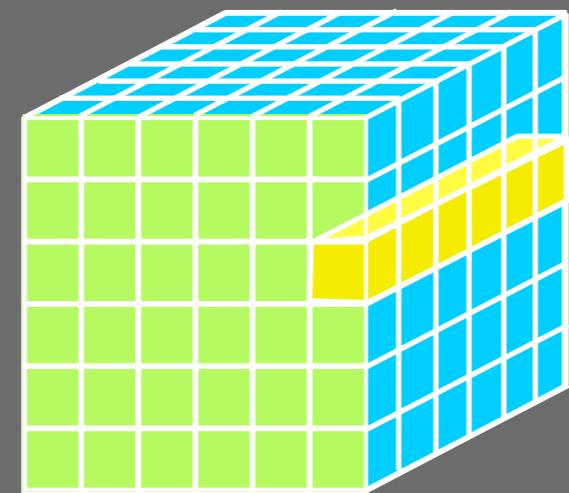
Tiny symbols show types of cells, color helps differentiate, as some symbol shapes are repeated.
(Symbol size not meaningful?)



PRACTICALITY



PRINCIPLES



PHILOSOPHY



DIVERSE TOOLS

DIVERSE DATA

DIVERSE VIEWS

MY WORLD & YOUR



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FACULTY ▾

EDUCATION &
LEARNING ▾

JAX MICE &
SERVICES ▾

PERSONALIZED
MEDICINE ▾

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MICE IN SPACE UPDATE

Researchers show that JAX Mice stayed mighty during their time in microgravity.

DIVERSE TOOLS

DIVERSE DATA

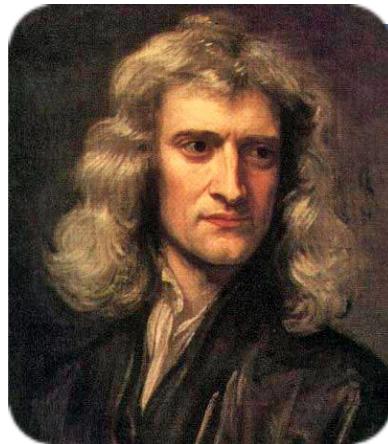
DIVERSE VIEWS

LEARN MORE +

PHILOSOPHY

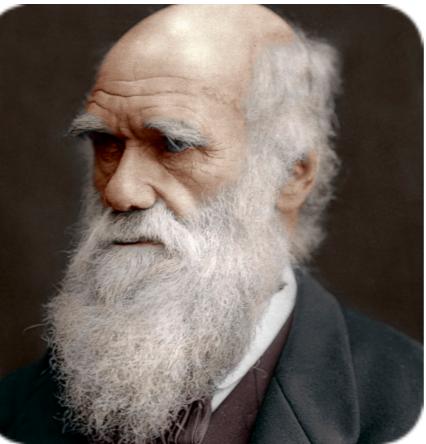




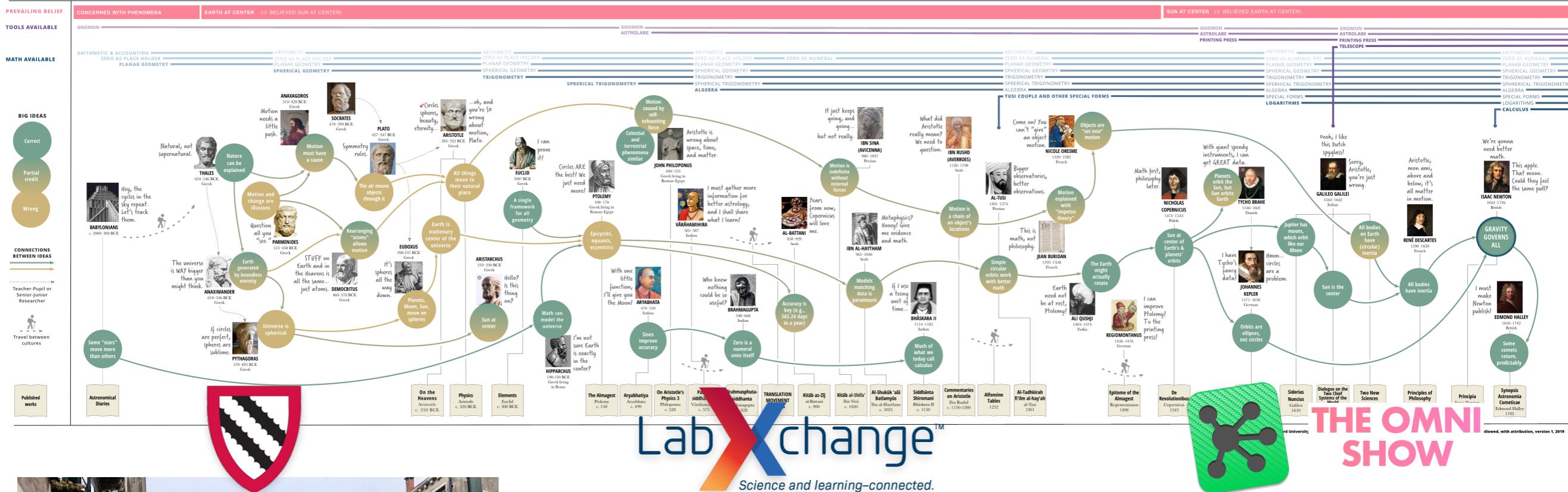


PATH TO

A project to track the evolution of science



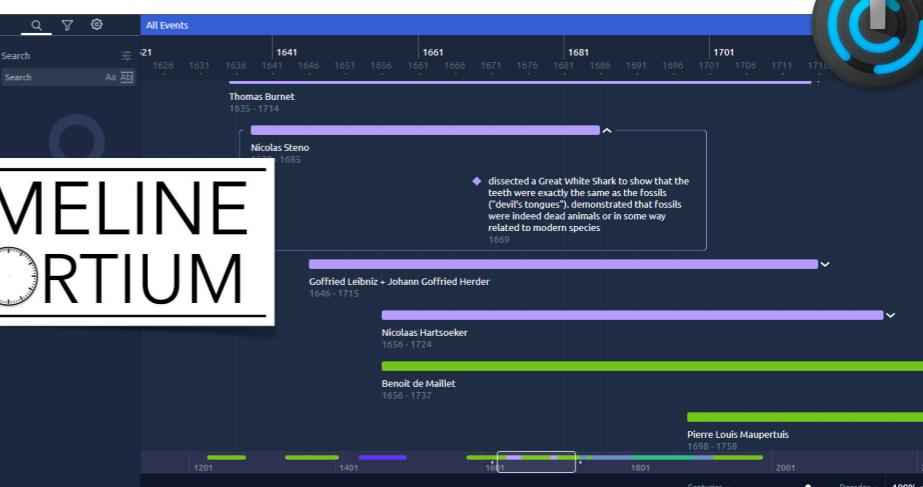
The Path to Newton



Phenomenon	Observation	Data	Rule	Theory	Explanation	Prediction
------------	-------------	------	------	--------	-------------	------------

Important Figures on the Path to Darwin

	Figure	Birth Date	Death Date	Time of Relevance	Rationale for Time of Relevance	idea/Discovery/Relevance
2	Anaximander of Miletus	-610	-546	-605	Around "On Nature" poem was published. No exact dates, I'm just guessing	proposed that the first ever animals lived in water and that the first land-dwelling ancestors of mankind must have been born in water and spent some time on land. Associated humans with fish bc they also required nru! Defended Essentialism -- that an entity's attributes are necessary to its identity and function. There have been simpler water-dwelling animals that preceded humans "most similar theory in antiquity to Darwinism.
3	goras	-570	-495	-550	Estimated provided by list for spermatogenesis/reproduction	"Kineness" and spermian: an obsolete biological theory that stated that sperm contained the preformed ger embryo -> Preformationism, that fathers contribute develop from miniature versions of themselves
4	itus	-535	-475	-500	PLACEHOLDER GUESSTIMATE	Believed in eternal change (subatomic particles). "I" birth and death in animals are just the mingling and Species don't change now, but species originated & adding more elements. Did not think of fixation of tr
5	Iodoces	-490	-430	-460	PLACEHOLDER GUESSTIMATE	
6	ritus	-460	-370	-400	PLACEHOLDER GUESSTIMATE	Atomism
Plato		-428	-347	-375	Publication of The Republic	"Theory of Form" all forms of life are imperfect copies of an eternal form. Early empiricism? Came up with Essentialism** Essentialism is the view that every entity has a set of attributes that are necessary to its function. Write about this in "The Republic." Perfection must already exist in another dimension. What we dimension, is a mere reflection of perfect realities in that other dimension (like how you can imagine a perfect God is in the other dimension. -> If something is changeable towards perfection, then it is not perfect to begin with. If it is perfect, then it cannot change)
8	Aristotle	-384	-322	-375	ESTIMATE. Generations of Animals was published in the "latter part of the fourth century B.C." (source)	Perfect forms do exist as extrapolations of our imagination. We create perfection in our own minds and we from nature, recognize a close connection between an organism's form and function (what we call "design" attributes to any divine cause. Inspired the dominant belief in the middle ages of the "Great Chain of Being" "Generations of Animals"
9	Zhuang Zhou	-369	-286	-299	Etc. On the Generation of Ani... list source	seed that biological species were not fixed and could change over time, in resp. regarding human nature and the heavens.





predictionx.org

PREDICTIONX: THE PAST & PRESENT OF THE FUTURE



ESSENTIALS

Predictive Systems Framework

Phenomena → Predictions

Understanding Uncertainty

Study Design

Timelines

Why predict?



Omens, Oracles & Prophecies

Mesopotamian Haruspicy	Egyptian Priests	Yoruba Ifa
Roman Augury	Tarot	Casting Lots
Chinese Oracle Bones	The Diviner's Guide	Greek Astronomy
Oracle of Delphi	Turkish Tasseography	Astrology
Aztec Rituals	Maya Spacetime	Comets of Doom

cross-cultural conversations



THE RISE OF THEORY

Ancient Mesopotamia, Egypt, Greece & Rome
Islamic Science
The Path to Newton
► Indian Mathematics
European Renaissance

The Royal Society



MODERN SIMULATION

John Snow & Cholera
Cholera Map

- Epidemiology
- Personal Genomics
- Population Genetics

- Climate & Wealth
- Behavioral Economics

The Future of the Future
► AI, Derek's Day
► Philosophy
► Uncertainty

- Climate & Energy
- Climate Policy
- Tent Tarot
- Earthquakes

- Futures of our Universe
- SETI

Coming Soon

Interactive Resource

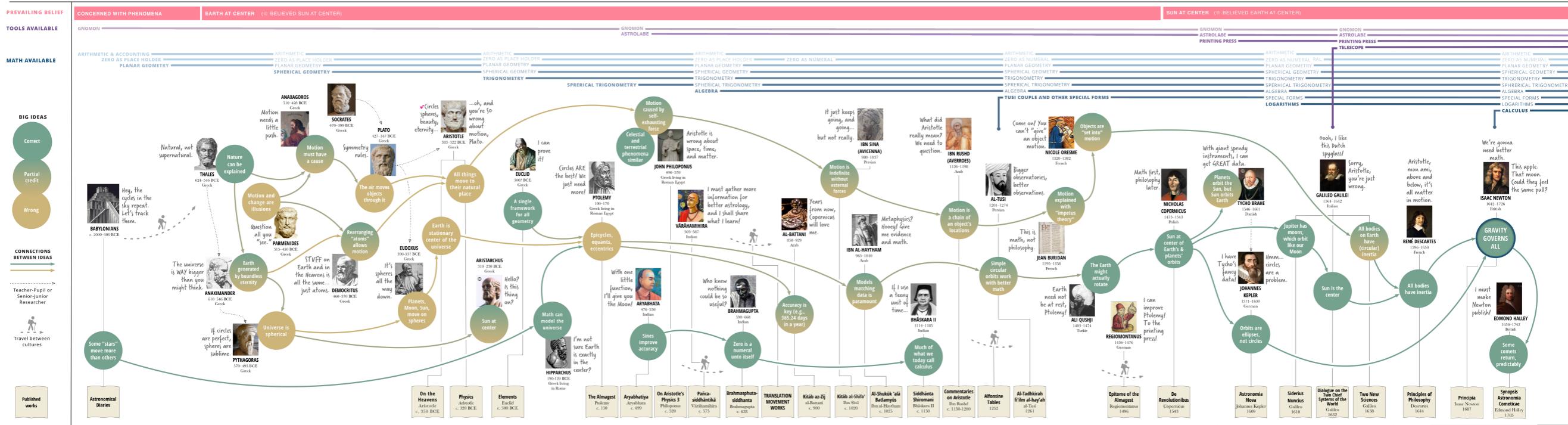
► video(s)

edX on edX as mini-course, otherwise find material via predictionx.org

THE "PADUA" RAINBOW



The Path to Newton



or, Experiment

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Mendel

Phenomenon

Observation

Data

Rule

Prediction



Darwin

Phenomenon

Observation

Data

Theory

Explanation

Prediction

BIOLOGY

PHYSICS



Kepler

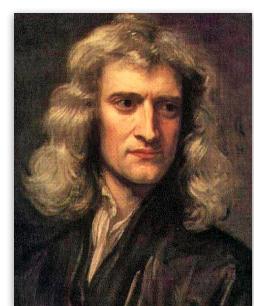
Phenomenon

Observation

Data

Rule

Prediction



Newton

Phenomenon

Observation

Data

Rule

Theory

Explanation

Prediction



Mendel 1865



Darwin 1859

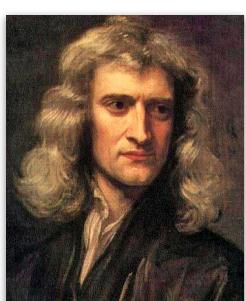


NO FULLY PREDICTIVE GENERAL THEORY

ONLY PREDICTIVE GENERAL THEORY



Kepler 1609



Newton 1687



THE FUTURE OF THE FUTURE

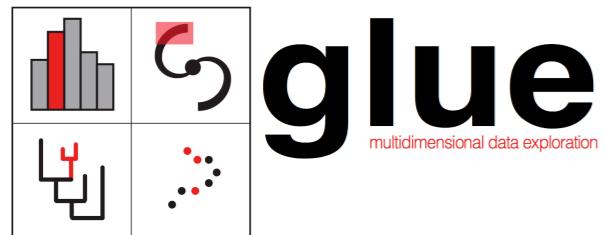
20th century



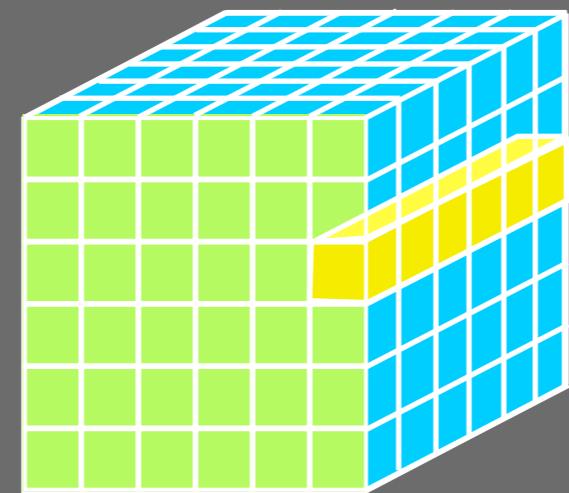
21st century?



PRACTICALITY



PRINCIPLES



PHILOSOPHY



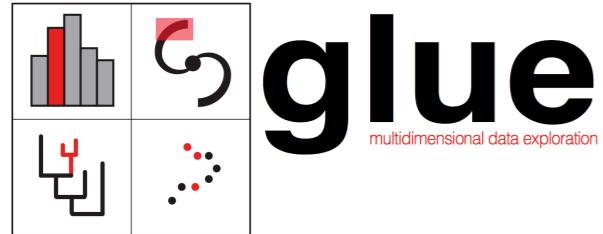
DIVERSE TOOLS

DIVERSE DATA

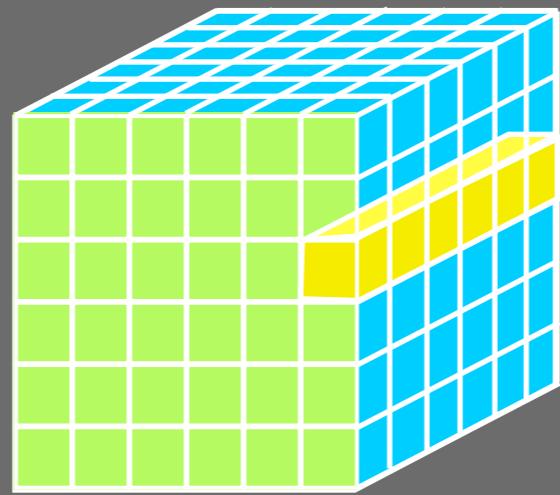
DIVERSE VIEWS

THE FUTURE OF PUBLISHING

PRACTICALITY



PRINCIPLES



PHILOSOPHY

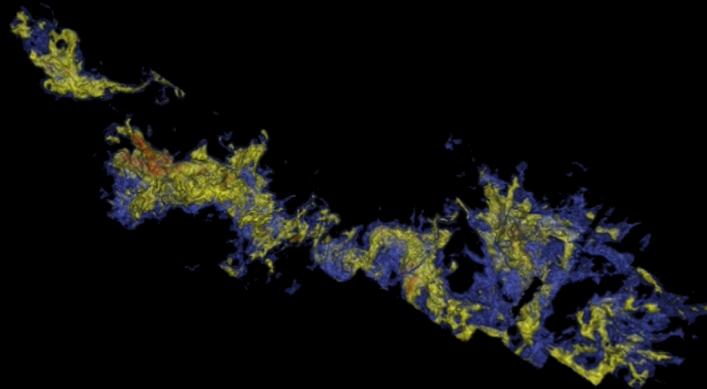


DIVERSE TOOLS

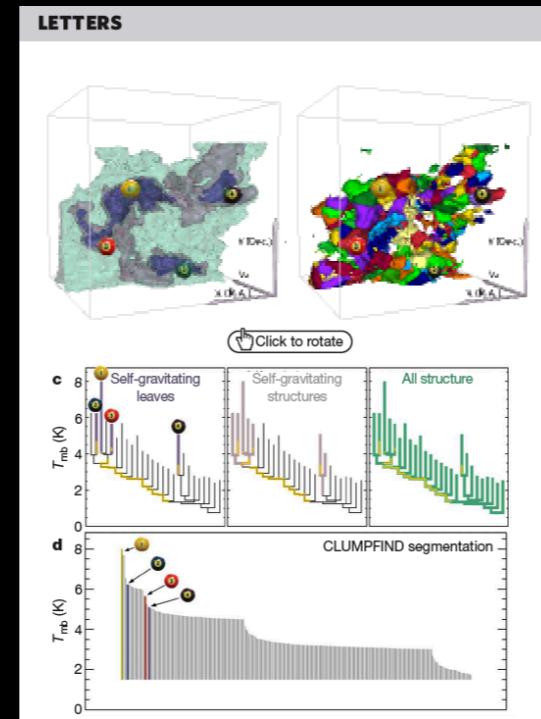
DIVERSE DATA

DIVERSE VIEWS

2008



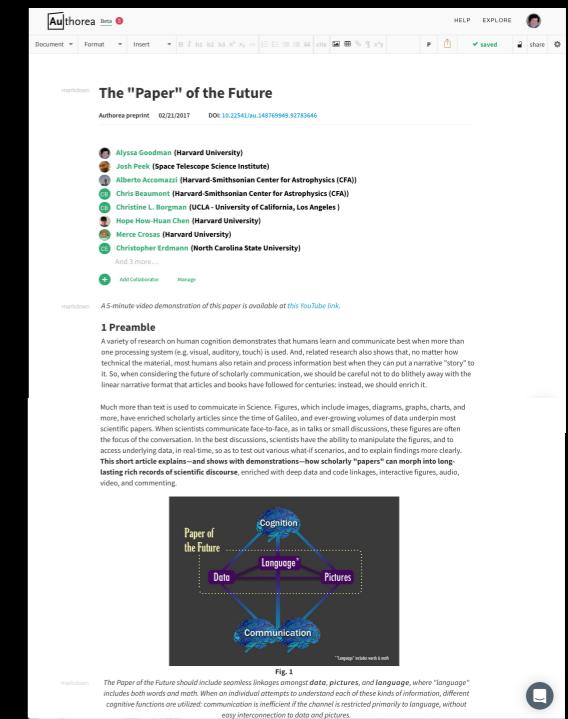
2009



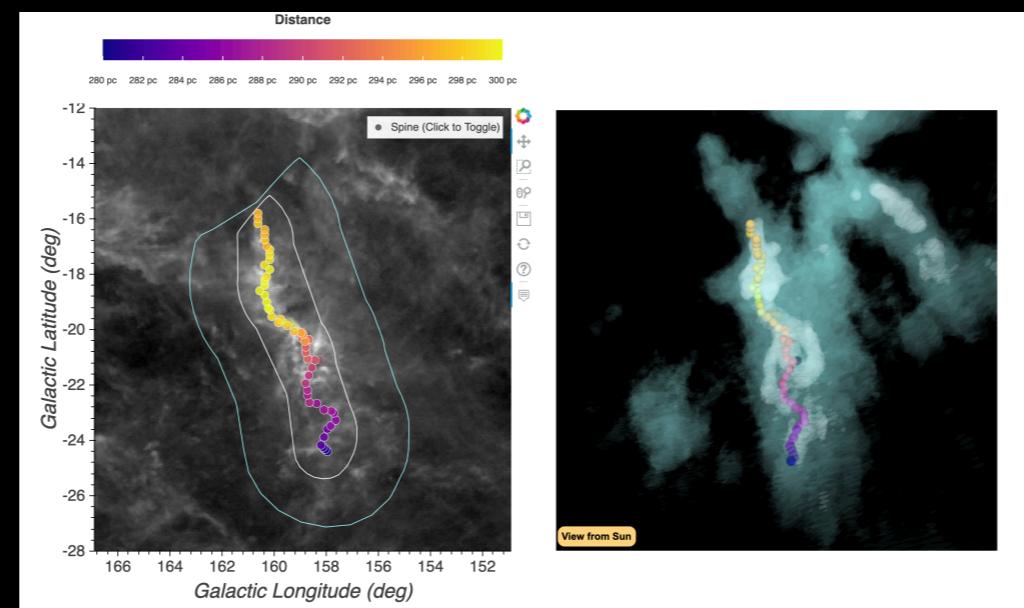
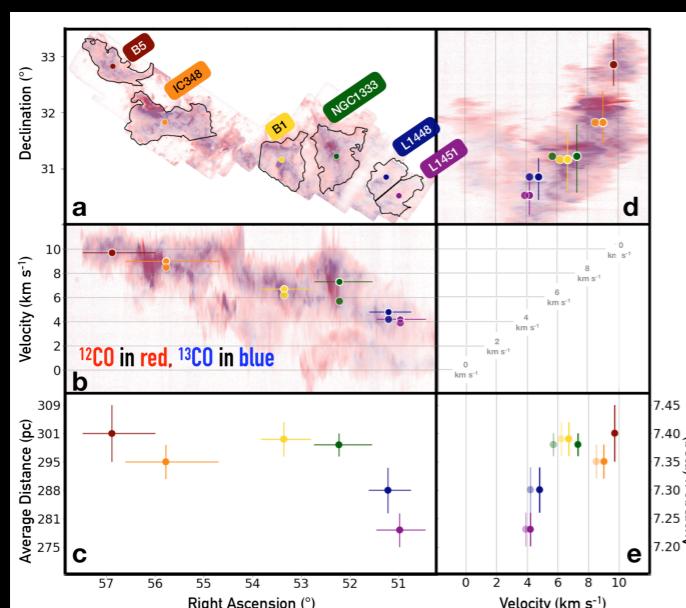
2012



2015



"Perseus" Progress



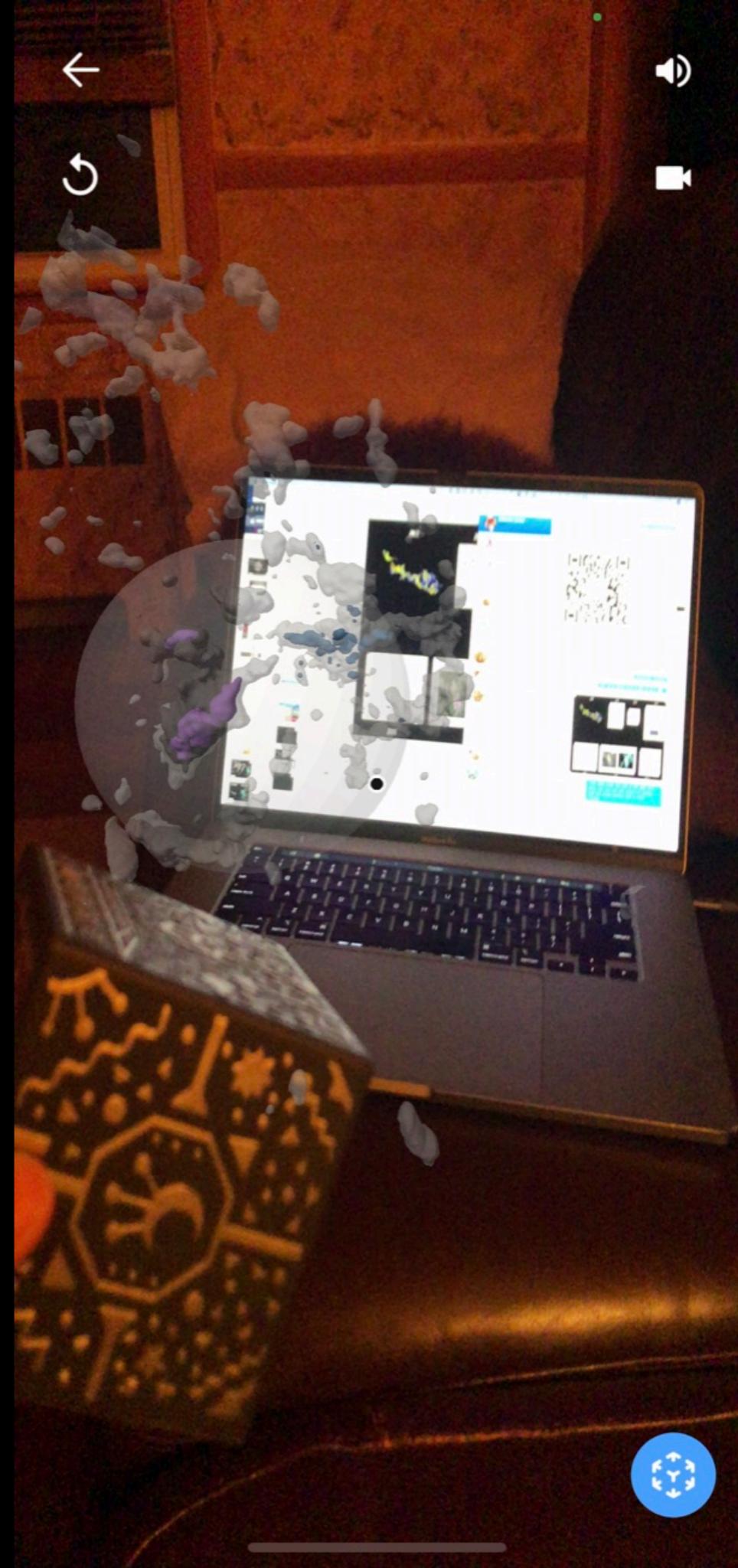
2018

2020

2020+



THE very near FUTURE



PERSEUS IN ACTUAL 3D





TEN QUESTIONS TO ASK WHEN CREATING A VISUALIZATION

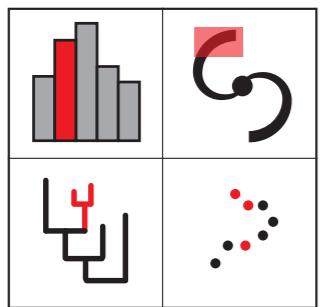
The 10 Questions

1. **Who** | Who is your audience? How expert will they be about the subject and/or display conventions?
 2. **Explore-Explain** | Is your goal to explore, document, or explain your data or ideas, or a combination of these?
 3. **Categories** | Do you want to show or explore pre-existing, known, human-interpretable, categories?
 4. **Patterns** | Do you want to identify new, previously unknown or undefined patterns?
 5. **Predictions & Uncertainty** | Are you making a comparison between data and/or predictions? Is representing uncertainty a concern?
 6. **Dimensions** | What is the intrinsic number of dimensions (not necessarily spatial) in your data, and how many do you want to show at once?
 7. **Abstraction & Accuracy** | Do you need to show all the data, or is summary or abstraction OK?
 8. **Context & Scale** | Can you, and do you want to, put the data into a standard frame of reference, coordinate system, or show scale(s)?
 9. **Metadata** | Do you need to display or link to non-quantitative metadata? (including captions, labels, etc.)
 10. **Display Modes** | What display modes might be used in experiencing your display?
-

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To read an in-process manuscript giving the scholarship behind the recommendations on this site, see [Coltekin & Goodman 2018](#).



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multidimensional data exploration

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