

**Alexander Lex**

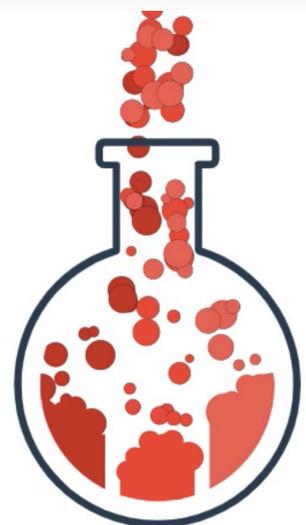
@alexander\_lex

<http://alexander-lex.net>



# Spatial Omics Visualizations:

## Lessons Learned from Networks and Maps



**visualization**  
**design lab**



# EVERYBODY IS TALKING ABOUT SPATIAL OMICS!

FOCUS | TECHNOLOGY FEATURE



## Method of the Year: spatially resolved transcriptomics

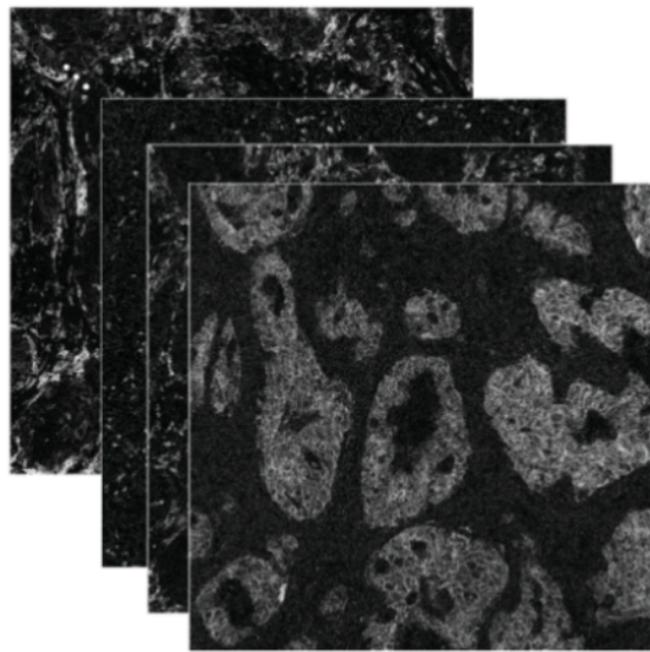
*Nature Methods* has crowned spatially resolved transcriptomics Method of the Year 2020.

Vivien Marx

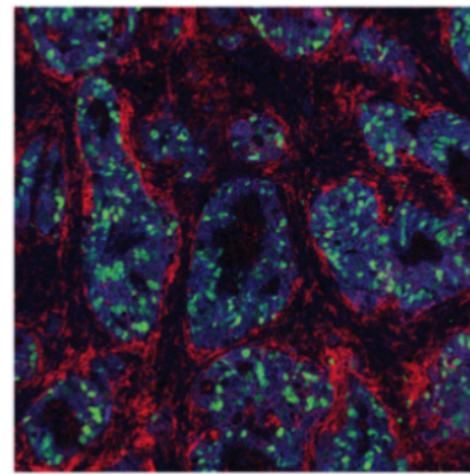
If a researcher is making a smoothie, it might be snack time. Or it could be the moment to prepare a sample for bulk RNA sequencing, in which tissue is homogenized and analyzed to yield



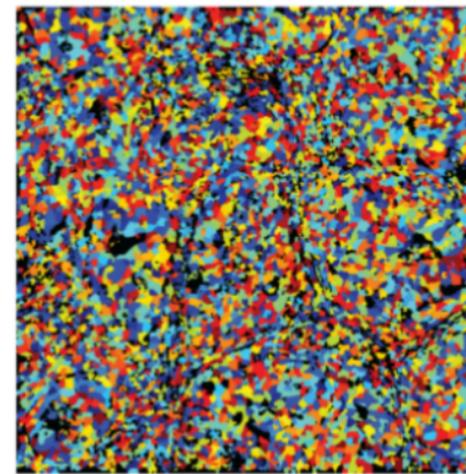
# BIOVIS CHALLENGE: FROM IMAGES TO ANALYSIS



Stack of images, each corresponding to one protein, i.e., one channel of the multi-dimensional measurement



Basic visualization e.g., overlaying 3 proteins as RGB



Segmentation mask.  
Cells indicated by random color



Downstream data analysis

# SPATIAL OMICS VISUALIZATION CHALLENGES

**High dimensional data**

**Similar to “classical” omics data (except for scale)**

**Spatial location / proximity is important**

**NOT encountered in “classical” omics data.**

# ANALOGY: MAPS!

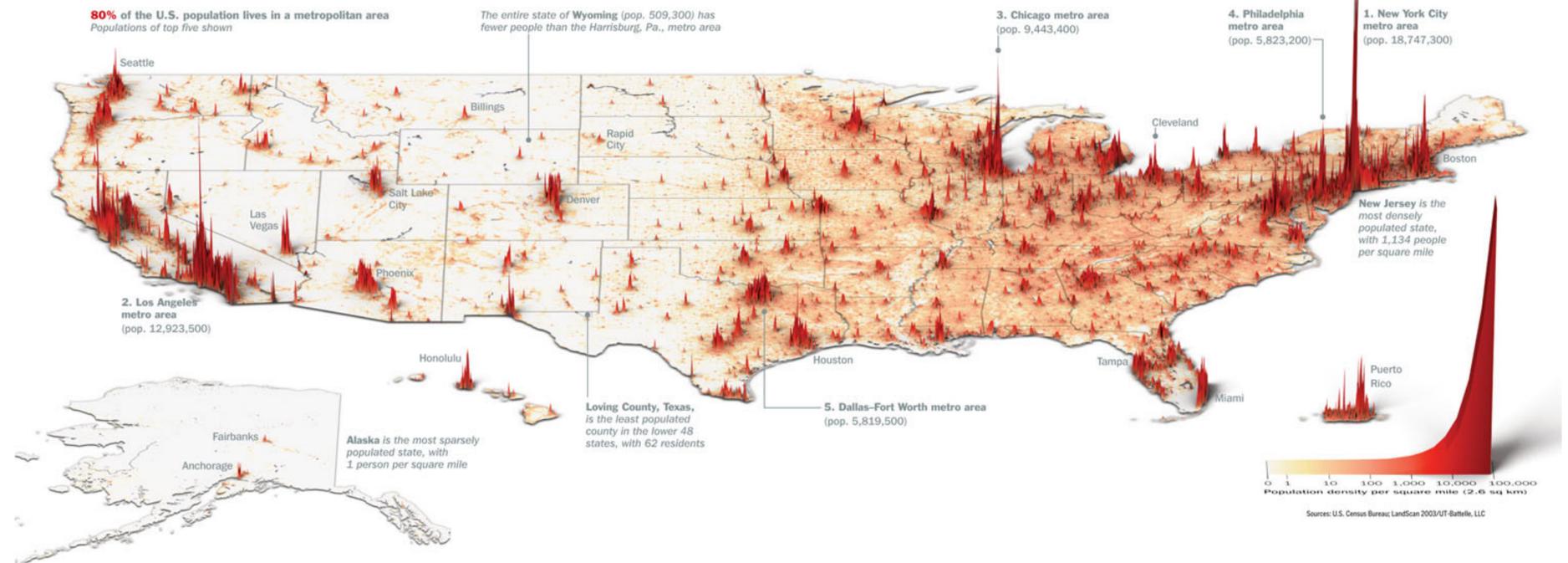
**Fixed location**  
**Potentially high-D**  
**vector for each**  
**location**

## Where We Live...

Unlike many developed countries, the U.S. keeps growing. We are also moving south and west. But compared with China or India, the nation is a vast prairie

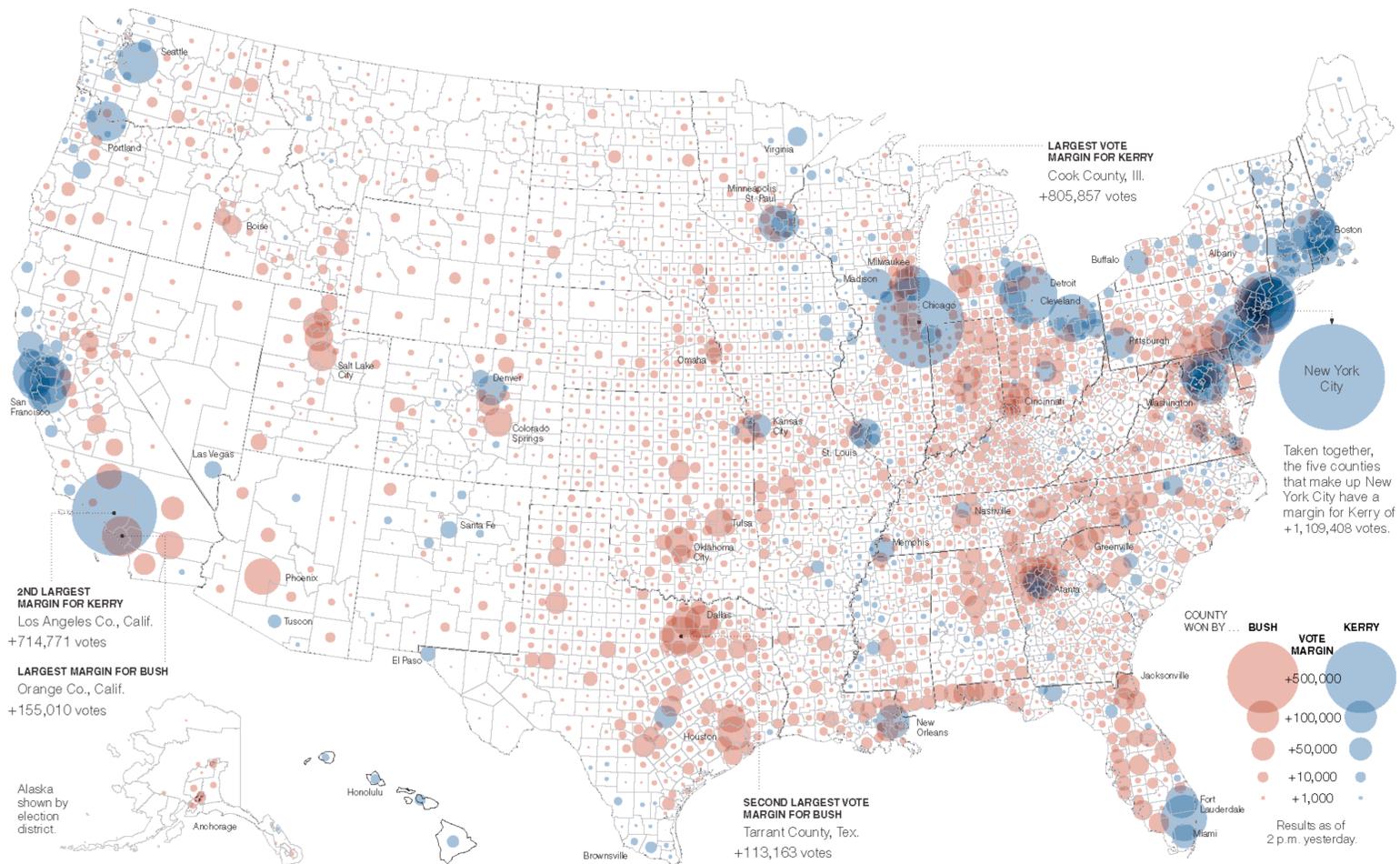
Our families are getting smaller—with one vital exception. Compared with those of Europe and Japan, the U.S. population is younger and more colorful because of the continued arrival of immigrants and their higher-than-average birthrates. Of the 100 million Americans who will join us in the next 37 years, half will be immigrants or their children. In the next few decades, 97% of the world's population growth will occur in the developing world; the U.S. is the largest developed country in the world that is still growing at a healthy clip. That matters, strategically, economical-

Ala.; Possum Trot, Ky.; or Lonelyville, N.Y. But they are all probably close to someone's idea of paradise. —By Nancy Gibbs

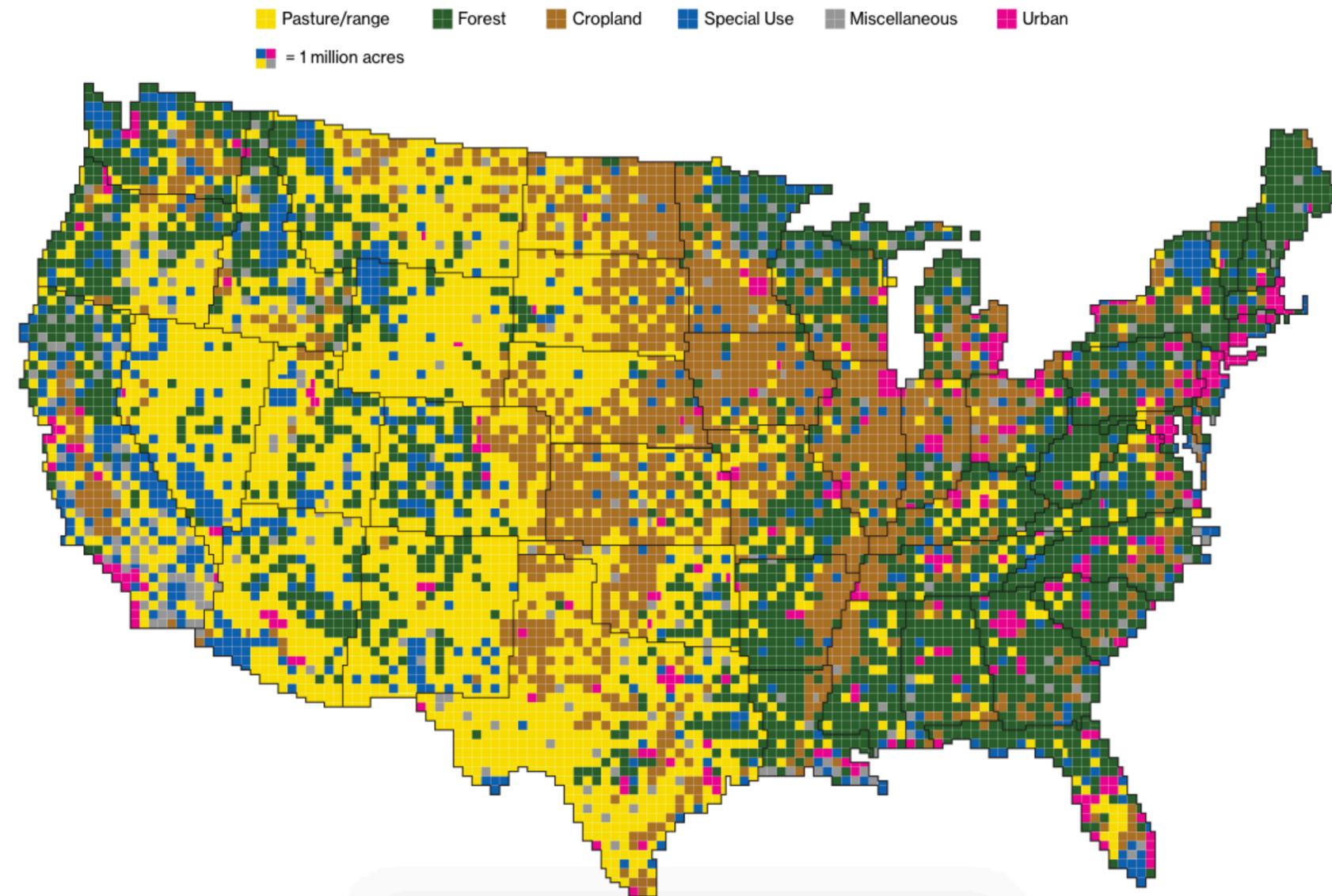


# SINGLE DATA VALUE + SPATIAL LOCATION

## Glyphs - Size

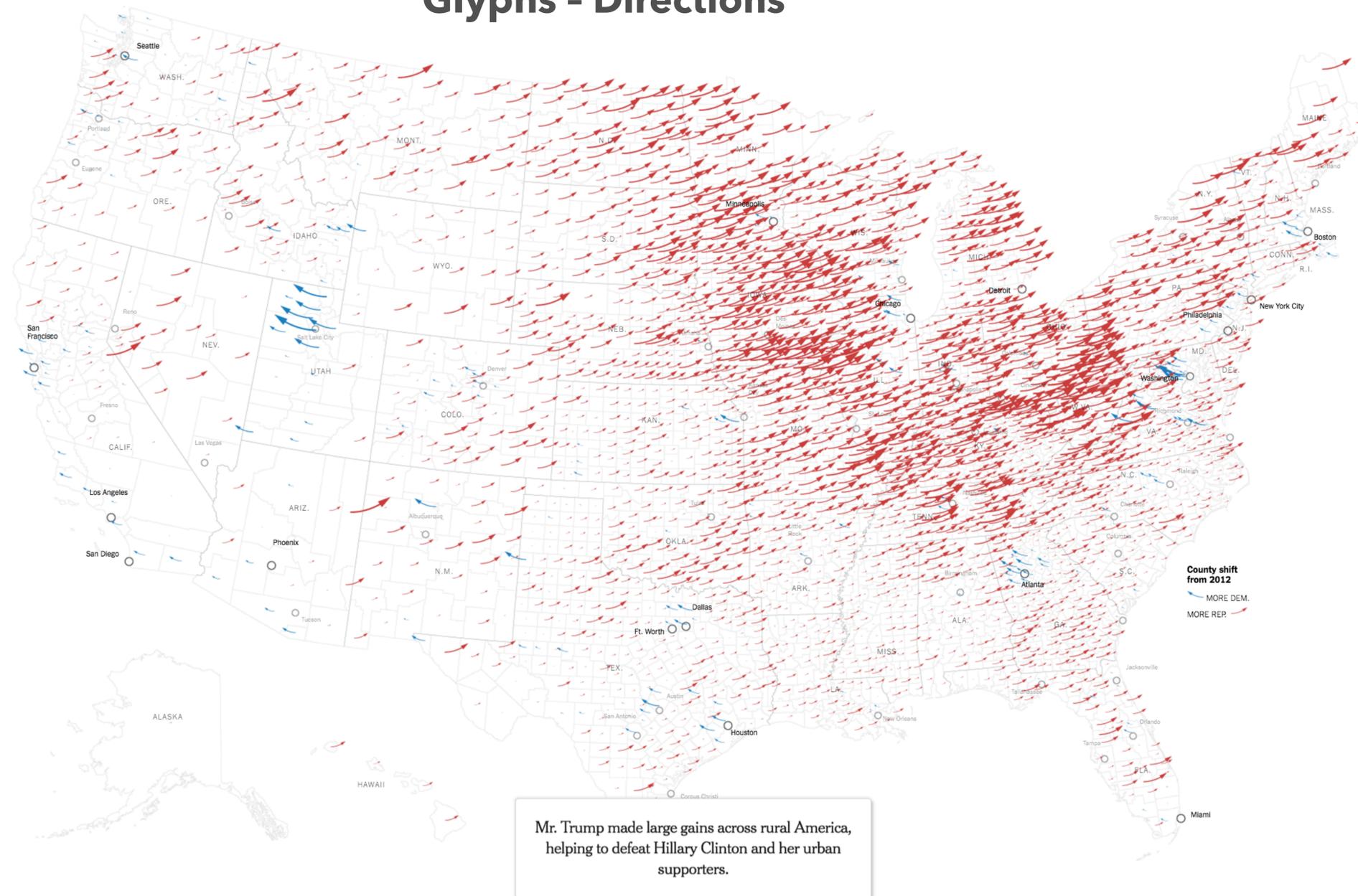


## Color



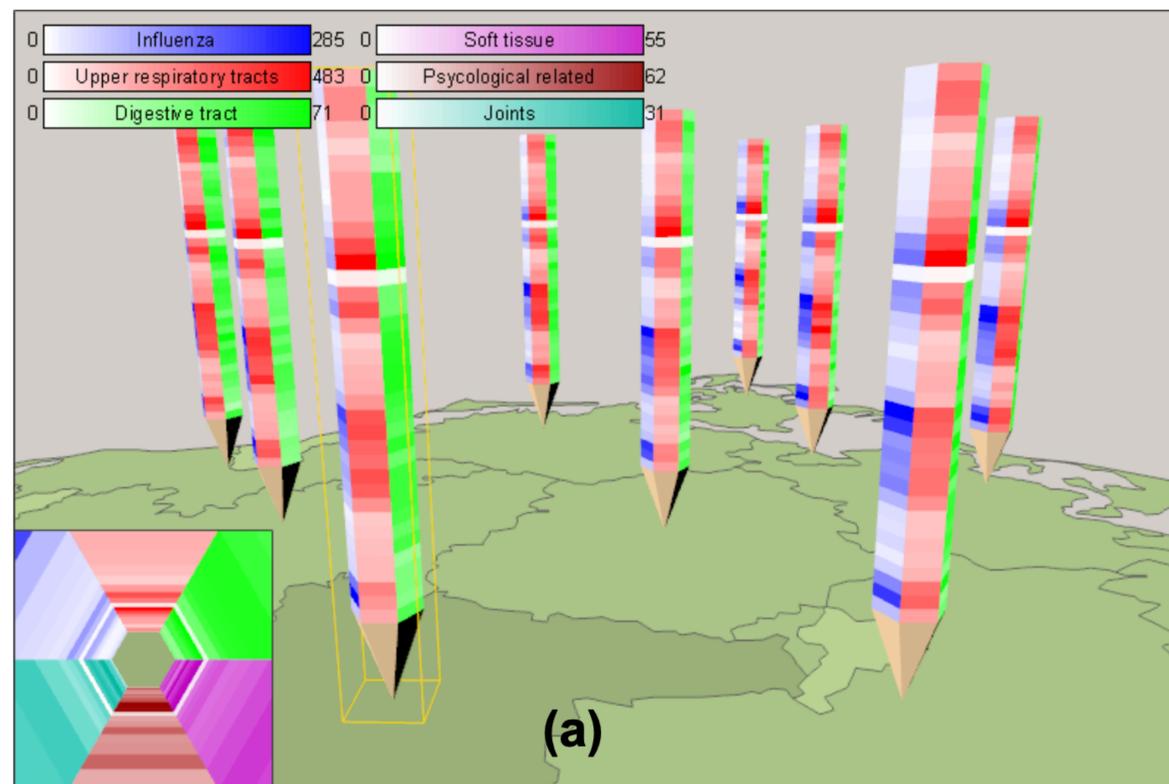
# SINGLE DATA VALUE + SPATIAL LOCATION

## Glyphs - Directions



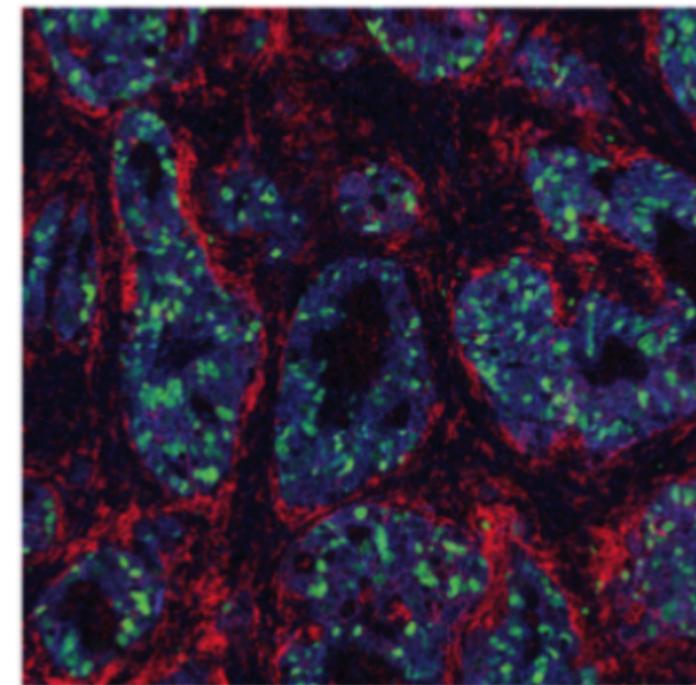
# BUT WHAT IF YOU HAVE MORE THAN ONE DATA VALUE?

## Glyphs?



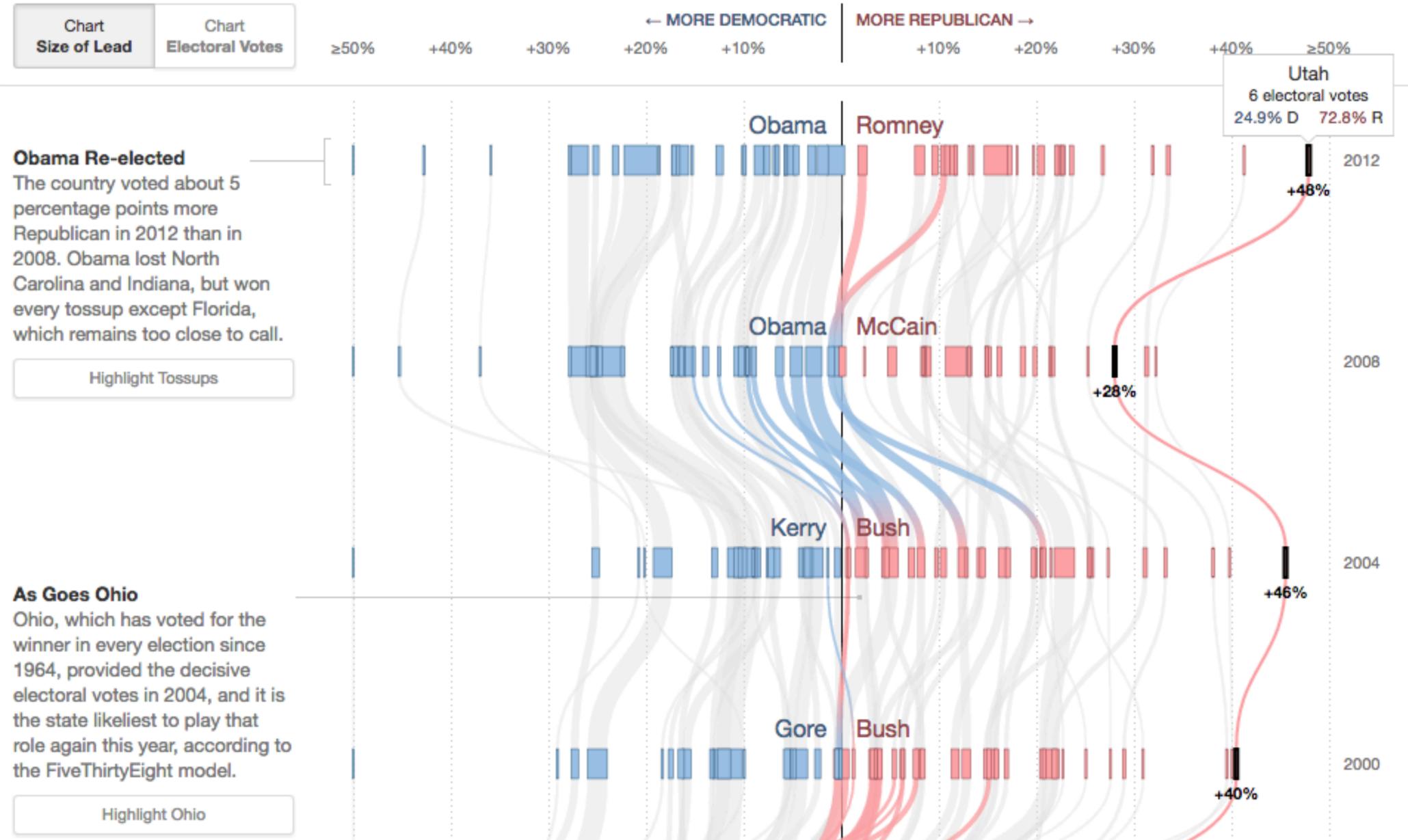
## Scalability? [Tominski]

Don't treat color channels as separate visual channels



Basic visualization e.g.,  
overlaying 3 proteins as RGB

# DO WE REALLY NEED A MAP?

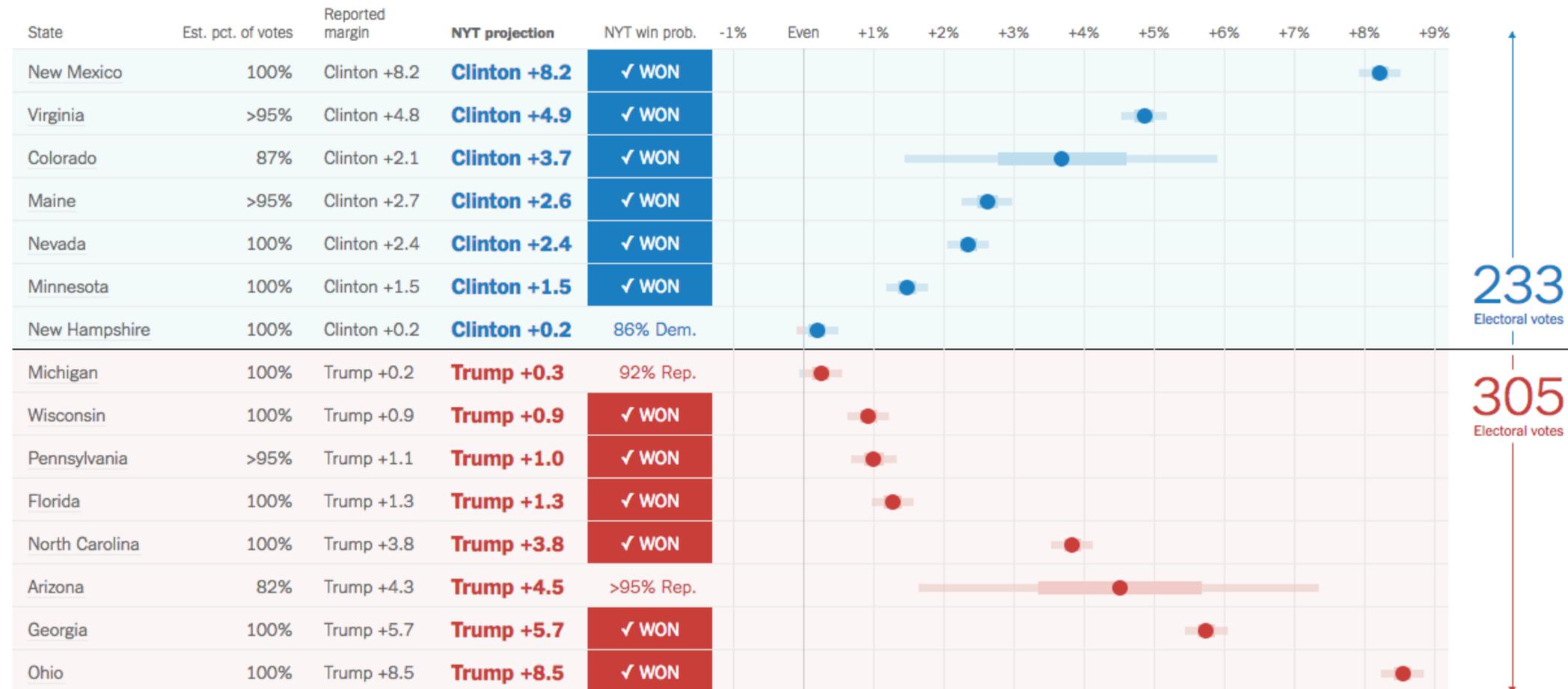


# DO WE REALLY NEED A MAP?

It's hard to do more complex things with maps

Is the spatial context paramount?

Is the spatial context a proxy for something?

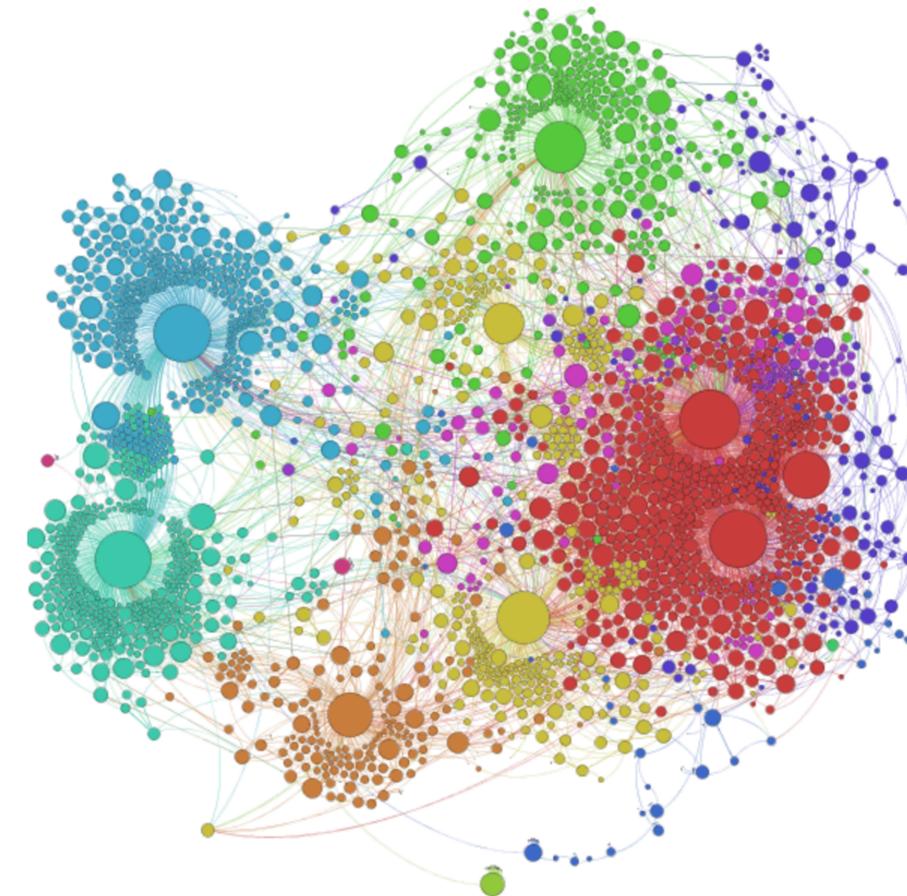


# ANALOGY: MULTIVARIATE NETWORKS

**Lots of attributes for nodes and edges**

**Location doesn't matter, but  
connectivity does**

**Can't choose location freely**



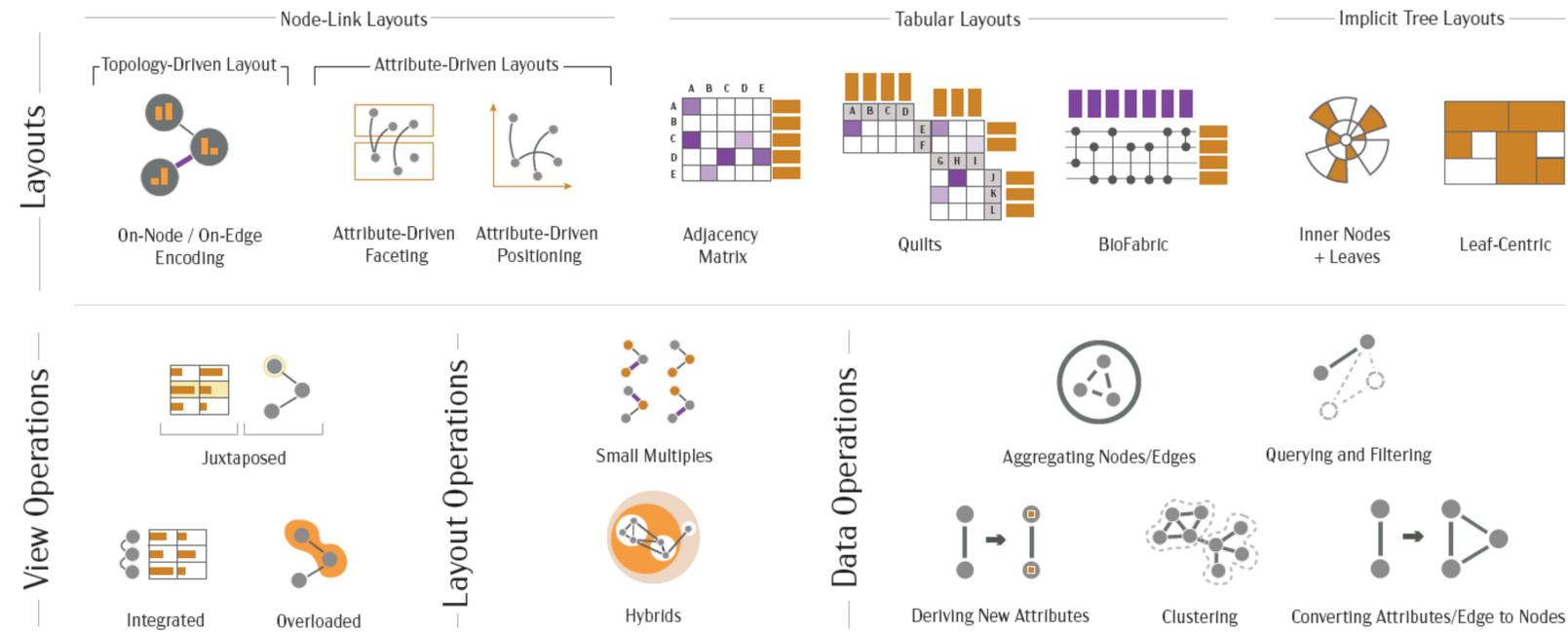


# The State of the Art in Visualizing Multivariate Networks

C. Nobre<sup>1</sup>, M. Meyer<sup>1</sup>, M. Streit<sup>2</sup>, and A. Lex<sup>1</sup>

<sup>1</sup>University of Utah, Utah, USA

<sup>2</sup>Johannes Kepler University Linz, Austria



**Figure 1:** A typology of operations and layouts used in multivariate network visualization. *Layouts* describe the fundamental choices for encoding multivariate networks. *View Operations* capture how topology and attribute focused visualizations can be combined. *Layout Operations* are applied to basic layouts to create specific visualization techniques. *Data Operations* are used to transform a network or derive attributes before visualizations. The colors reflect node attributes (orange), edge attributes (purple), and topology (grey).

## Abstract

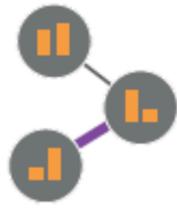
Multivariate networks are made up of nodes and their relationships (links), but also data about those nodes and links as attributes. Most real-world networks are associated with several attributes, and many analysis tasks depend on analyzing both, relationships and attributes. Visualization of multivariate networks, however, is challenging, especially when both the topology of the network and the attributes need to be considered concurrently. In this state-of-the-art report, we analyze current practices and classify techniques along four axes: layouts, view operations, layout operations, and data operations. We also provide an analysis of tasks specific to multivariate networks and give recommendations for which technique to use in which scenario. Finally, we survey application areas and evaluation methodologies.

Layouts

Node-Link Layouts

Topology-Driven Layout

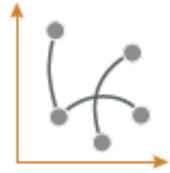
Attribute-Driven Layouts



On-Node / On-Edge Encoding

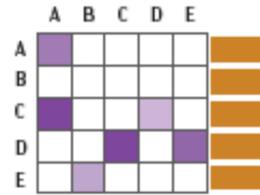


Attribute-Driven Faceting

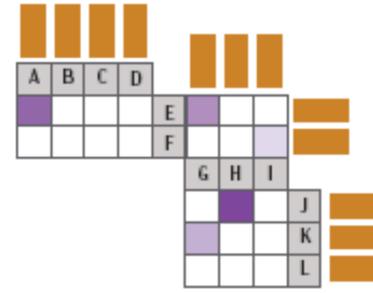


Attribute-Driven Positioning

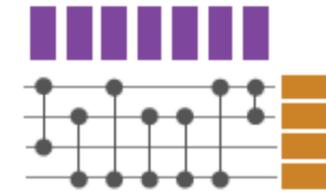
Tabular Layouts



Adjacency Matrix



Quilts

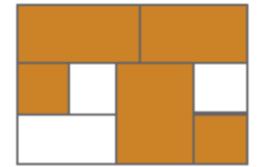


BioFabric

Implicit Tree Layouts



Inner Nodes + Leaves

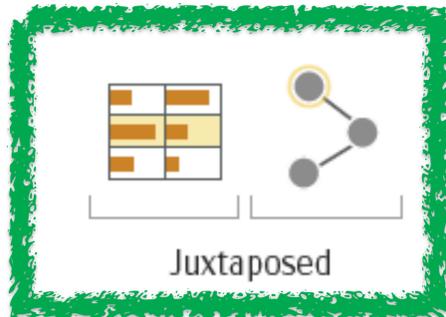


Leaf-Centric

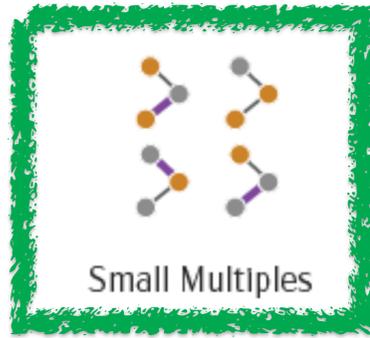
View Operations

Layout Operations

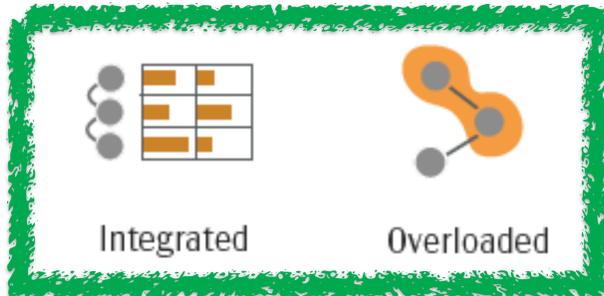
Data Operations



Juxtaposed



Small Multiples

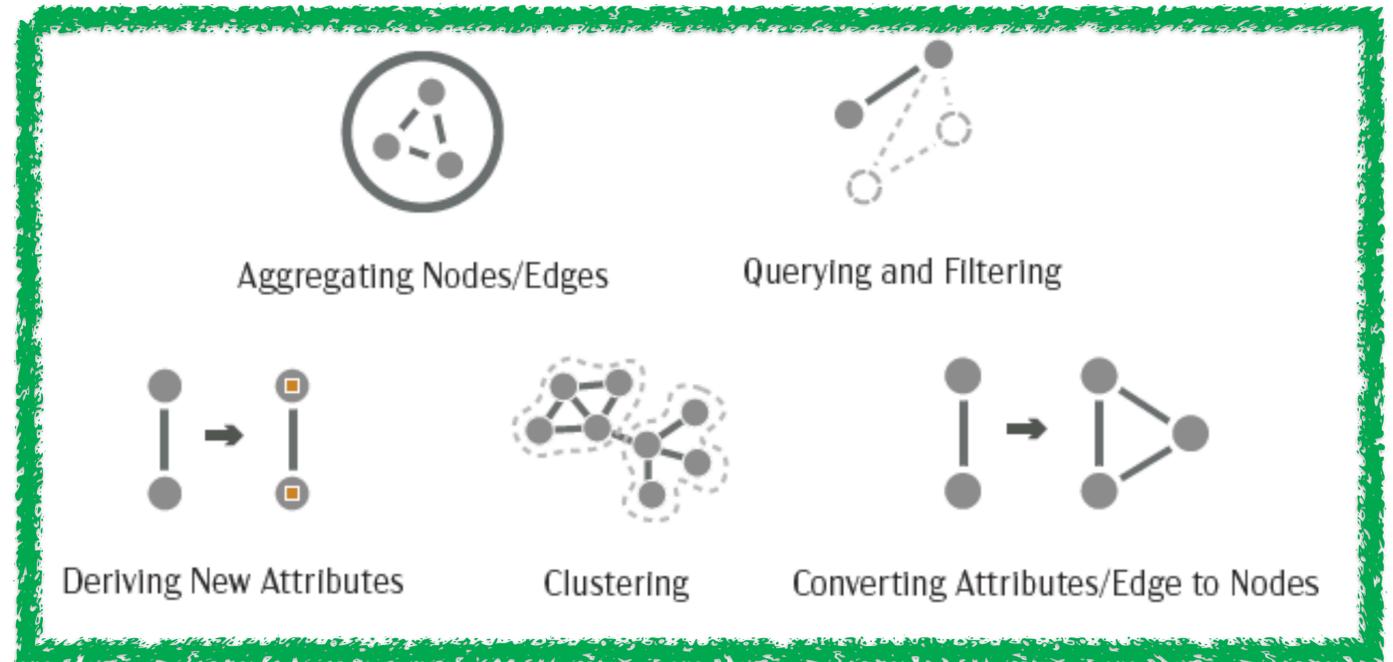


Integrated

Overloaded



Hybrids



Aggregating Nodes/Edges

Querying and Filtering

Deriving New Attributes

Clustering

Converting Attributes/Edge to Nodes

**SO WHAT ELSE CAN  
YOU DO?**

**INTERACTION!**

**Two Paths:**

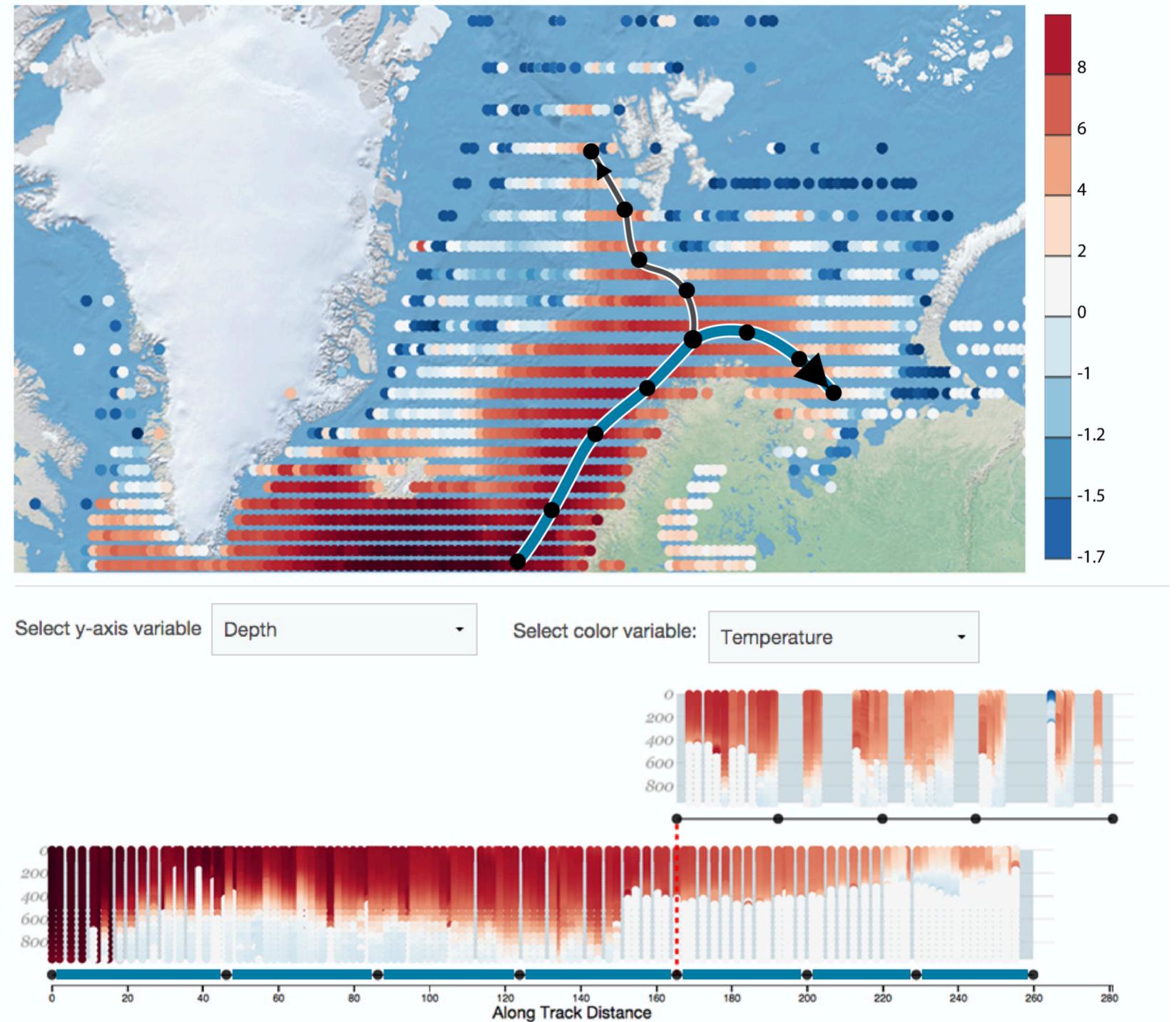
- 1. Select regions to show**
- 2. Select / derive data to show**

# SELECTING REGIONS TO SHOW

# PRINCIPLE

**Select one or multiple items/regions**  
**Show rich data about them in**  
**separate view**

# SELECT A PATH IN A MAP

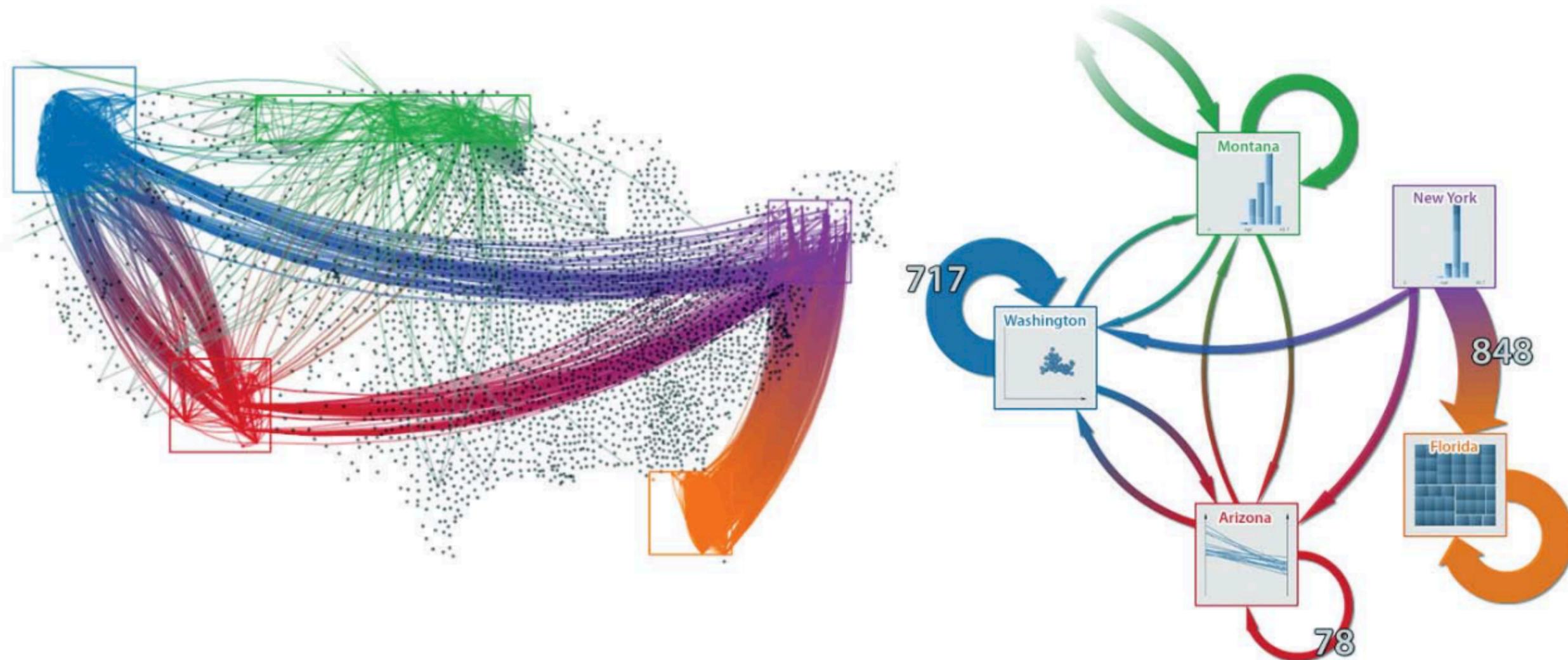


Carolina Nobre, Alexander Lex

OceanPaths: Visualizing Multivariate Oceanography Data

Proceedings of the Eurographics Conference on Visualization (EuroVis '15) - Short Papers, doi:10.2312/eurovisshort.20151124, 2015.

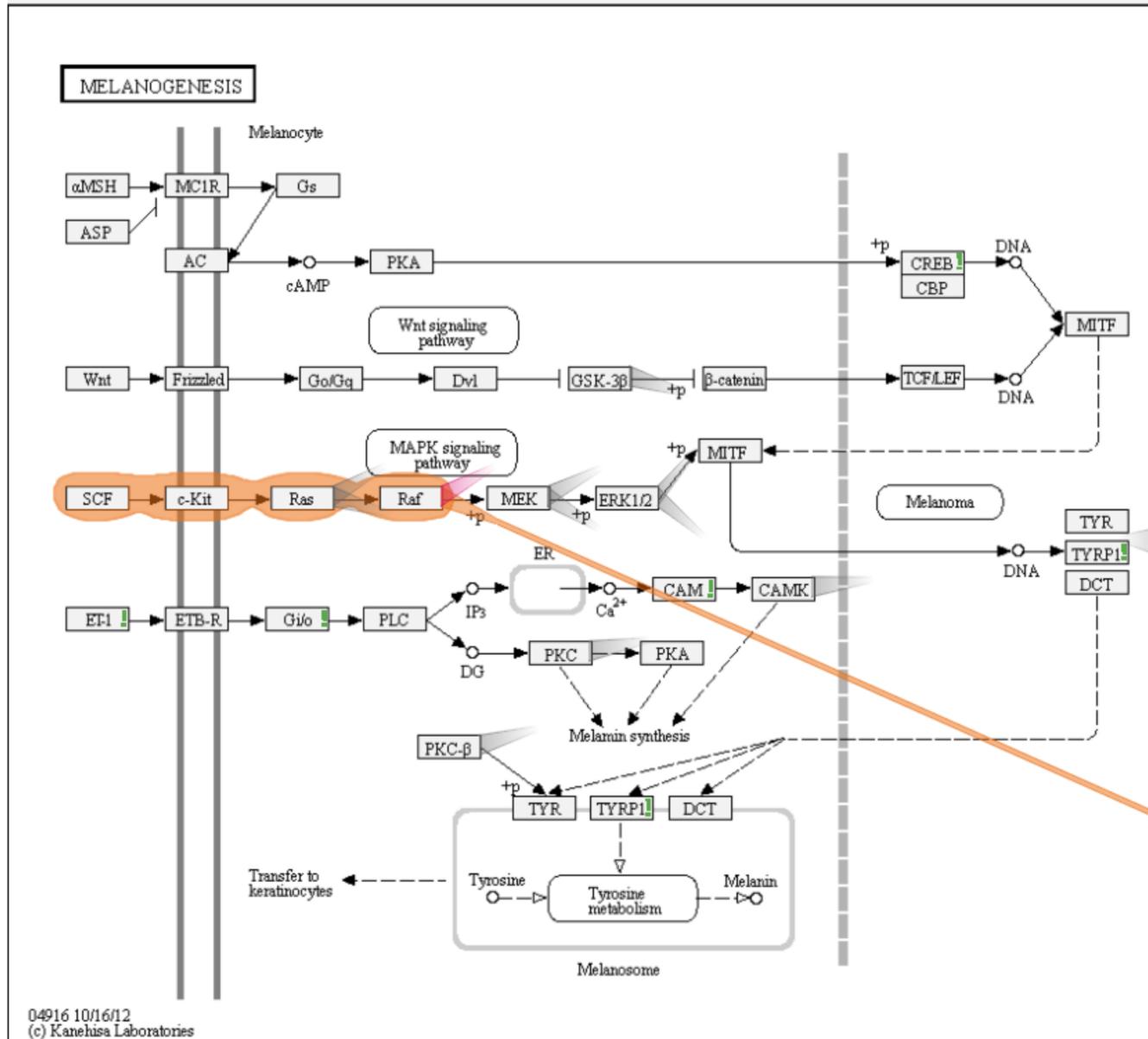
# SELECT A REGION AND AGGREGATE!



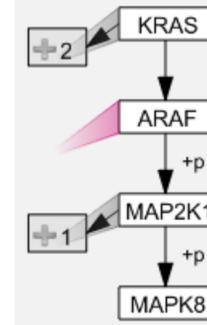
S. van den Elzen and J. J. van Wijk, "Multivariate Network Exploration and Presentation: From Detail to Overview via Selections and Aggregations," IEEE Transactions on Visualization and Computer Graphics (InfoVis '14), vol. 20, no. 12, pp. 2310-2319, 2014, doi: 10.1109/TVCG.2014.2346441.

# ENROUTE – PATH SELECTION

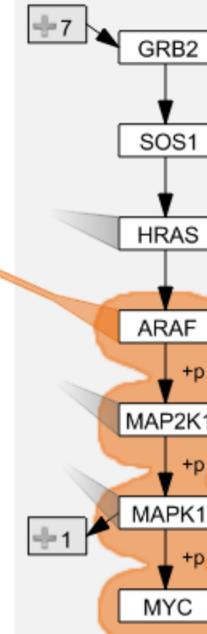
Melanogenesis



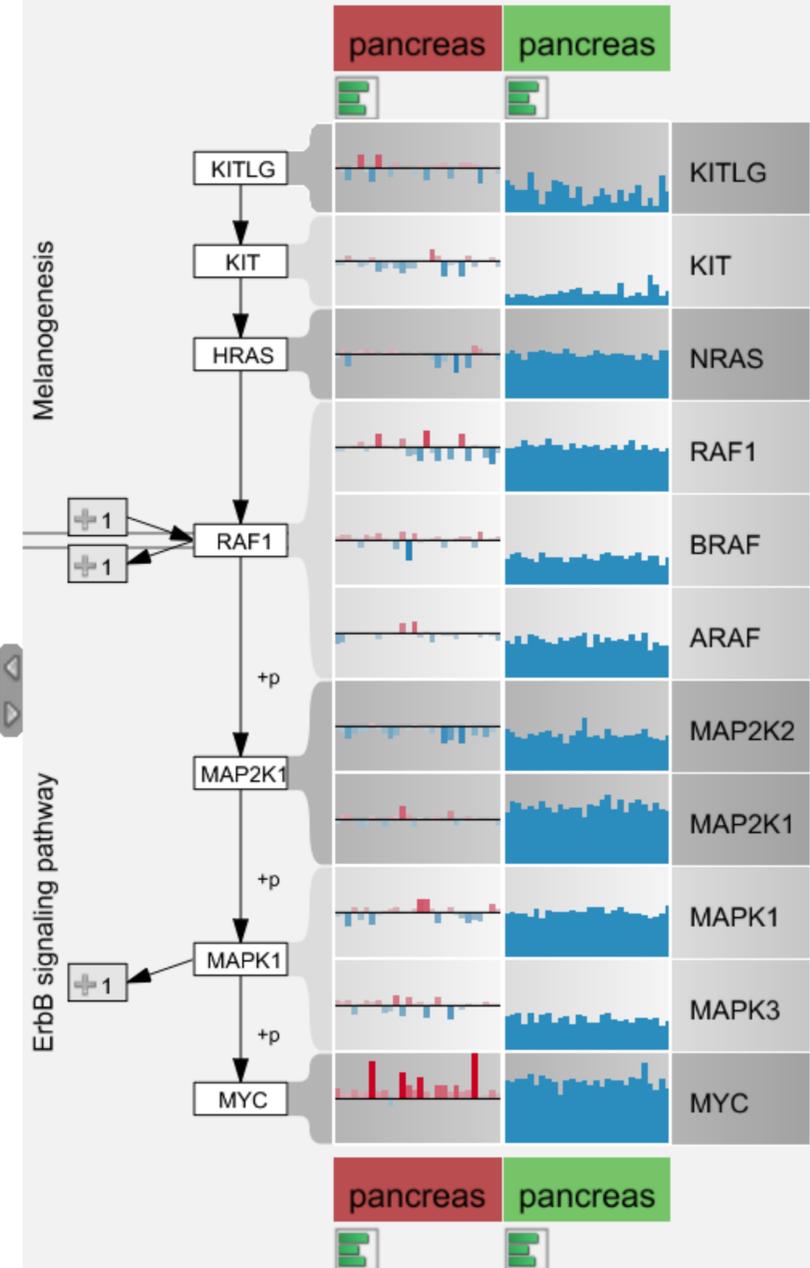
Pancreatic cancer



ErbB signaling pathway



Selected Path



**SELECT / DERIVE DATA  
TO SHOW**

**SHOWING RAW DATA  
IS HOPELESS!**

**But do you need to?**

**Show average expression (etc)**

**Show average expression for  
pathway of interest**

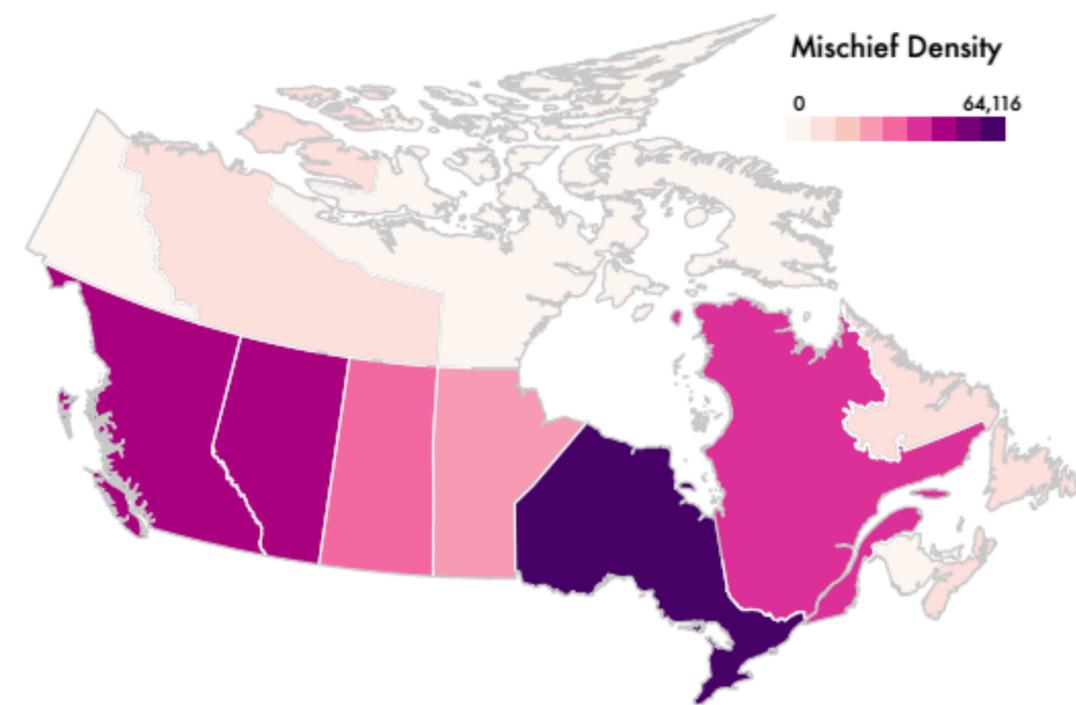
**Filter out uninteresting items**

**Create domain specific scores**

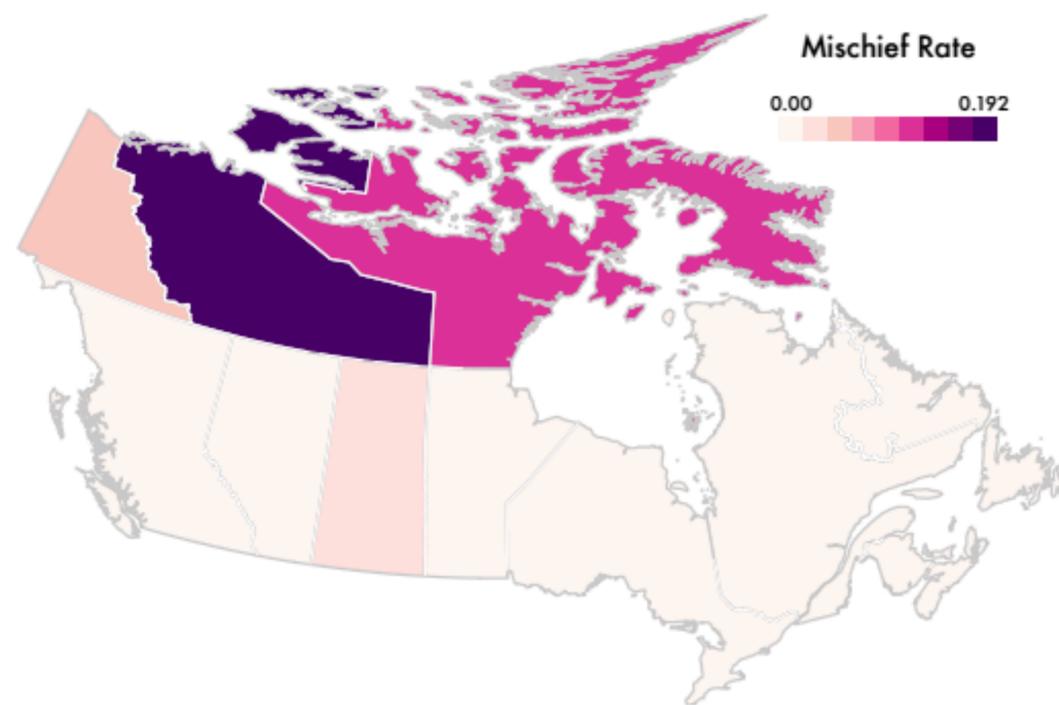
**Create clusters/archetypes**

**...**

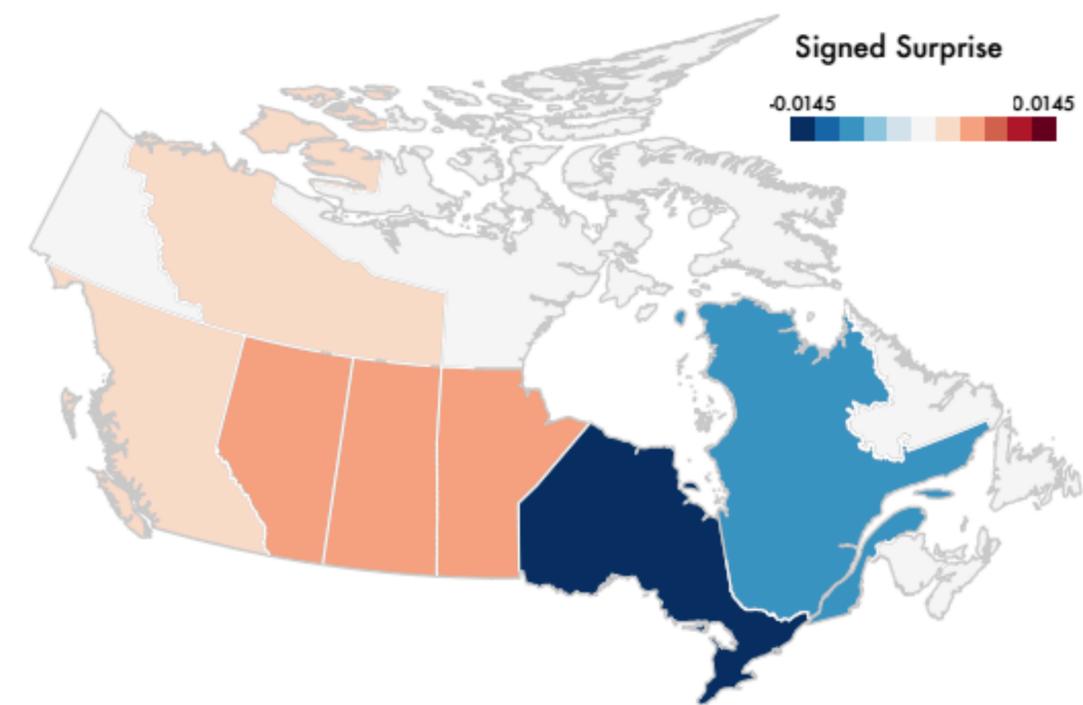
# APPROACH: USE A PRIOR, SHOW DIFFERENCE.



(a) The **Event Density** of “mischief” in Canada.



(b) The per-capita **Event Rate** of mischief.



(c) The **Surprise Map** of mischief.

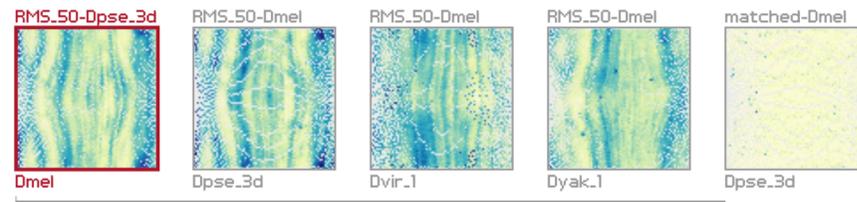
model of population density +  
accounting for variability when  
analyzing small numbers



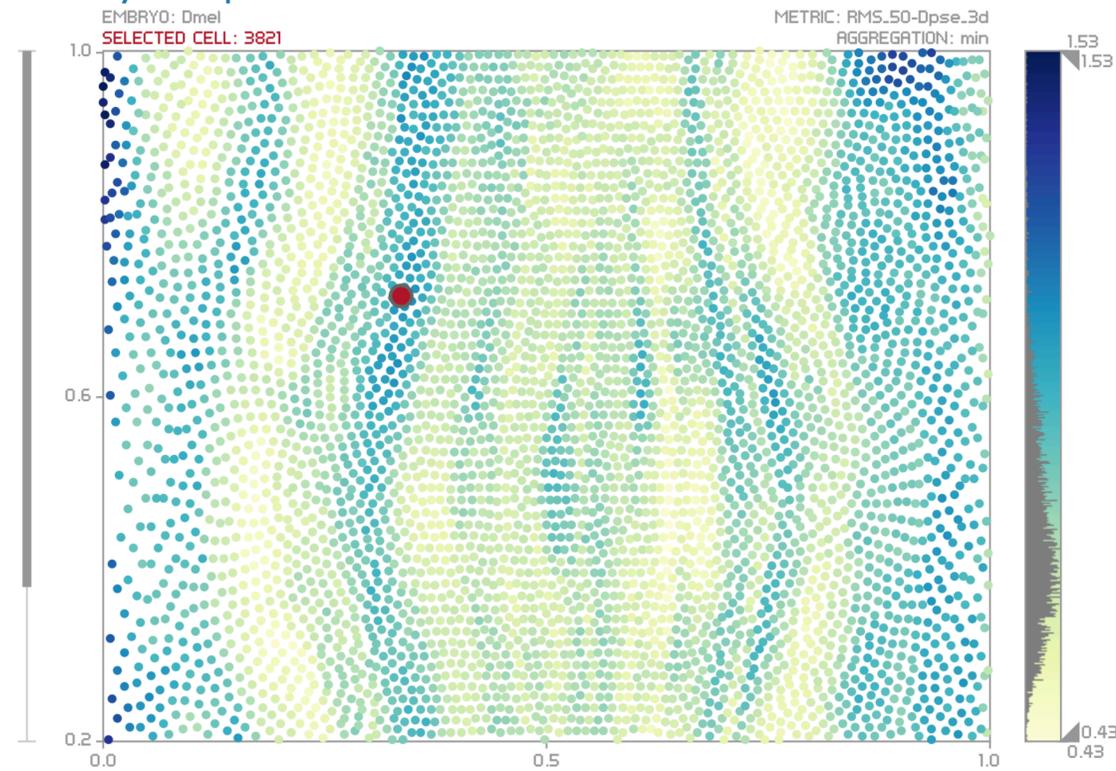


# EXAMPLE: DOMAIN SPECIFIC SCORE

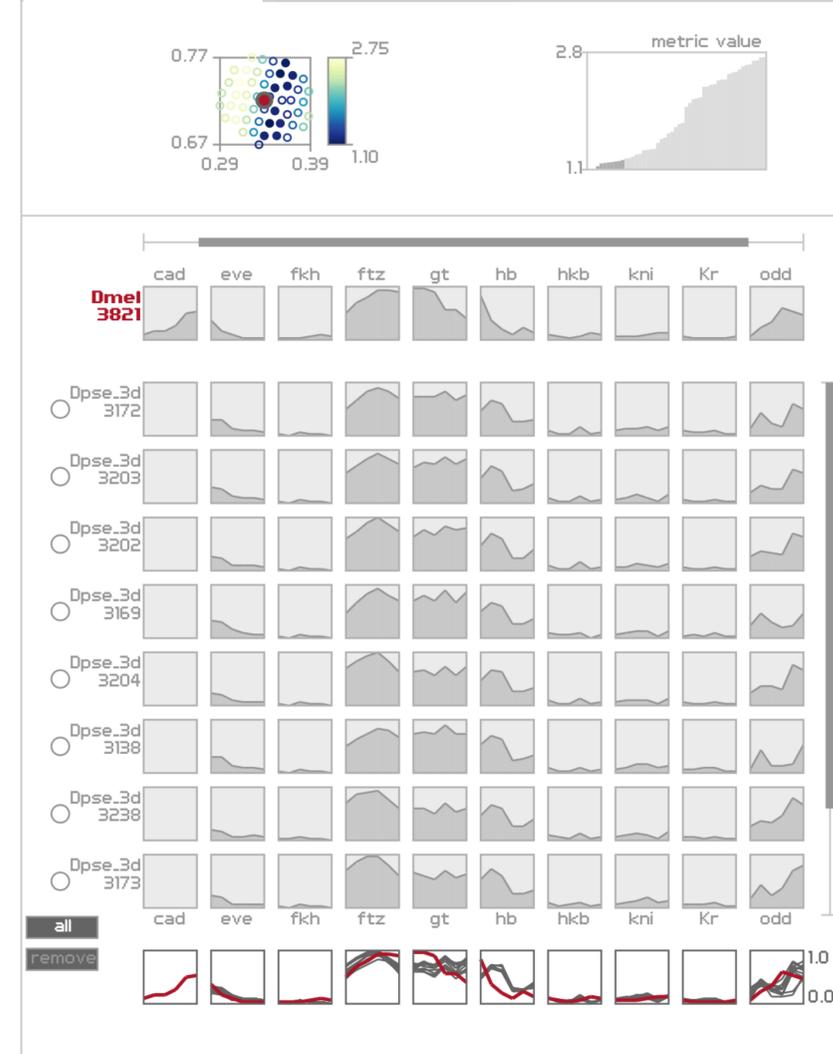
## Summaries



## Embryo Map



## Curvemaps



Miriah Meyer, Tamara Munzner, Angela DePace, Hanspeter Pfister

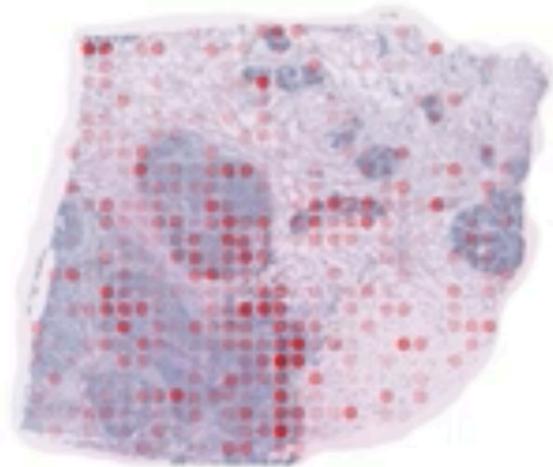
MulteeSum: A Tool for Comparative Spatial and Temporal Gene Expression Data

IEEE Transactions on Visualization and Computer Graphics (InfoVis), 16(6): 908--917, doi:10.1109/TVCG.2010.137, 2010.

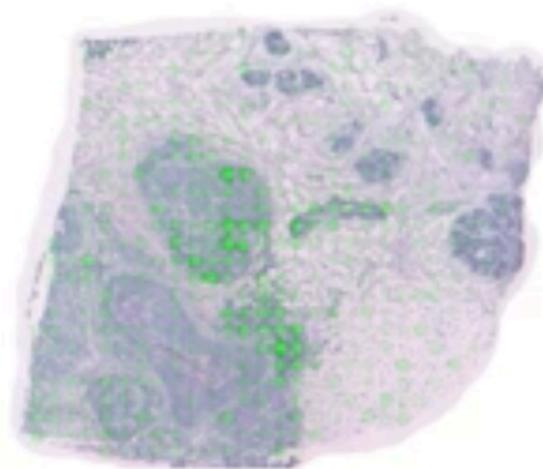
# USING SCORES @ BIOVIS BY ALMA ANDERSON

## ■ ■ ■ ■ Assessing co-localization

B-cells



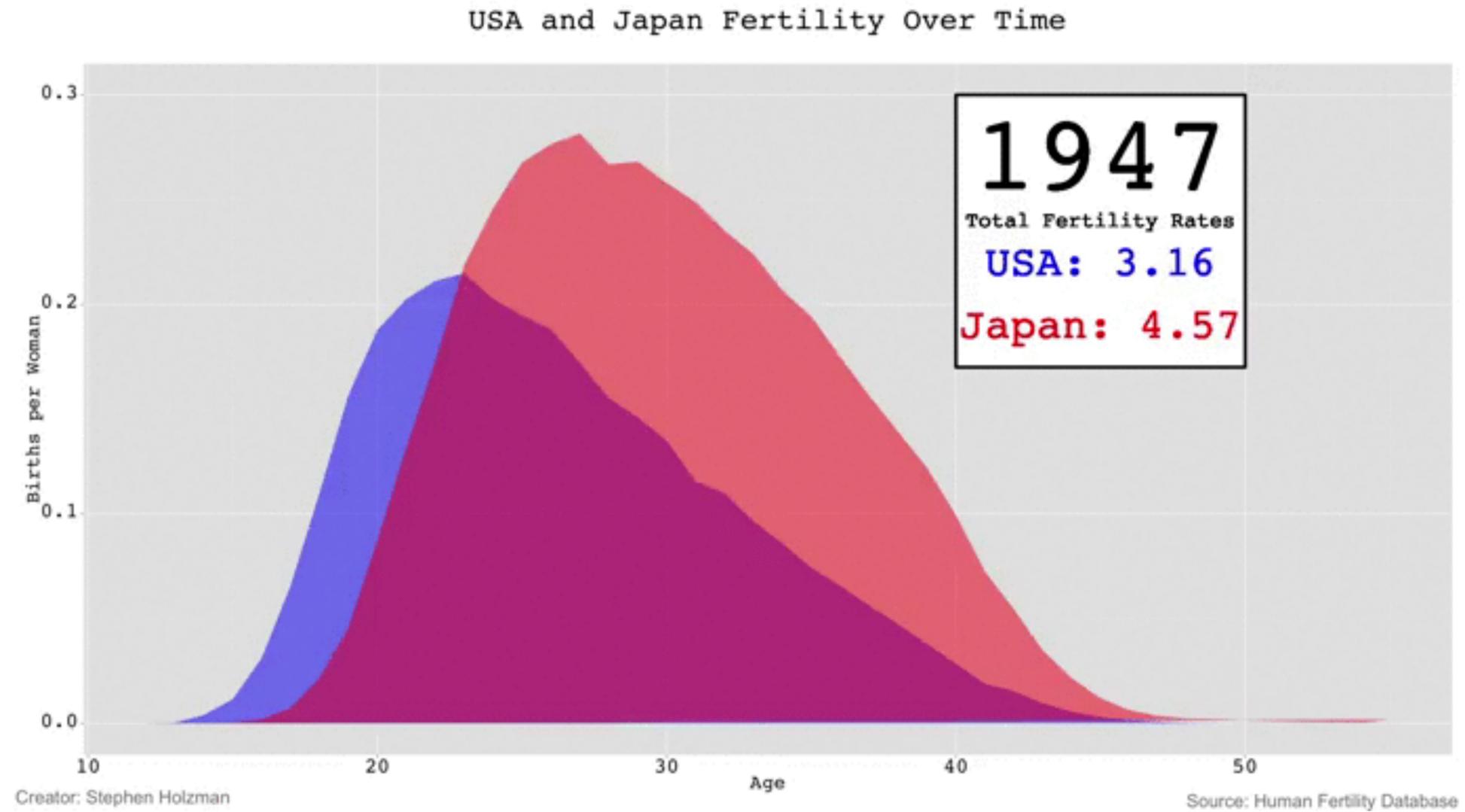
T-cells



Sample from breast cancer patient (HER2-positive)

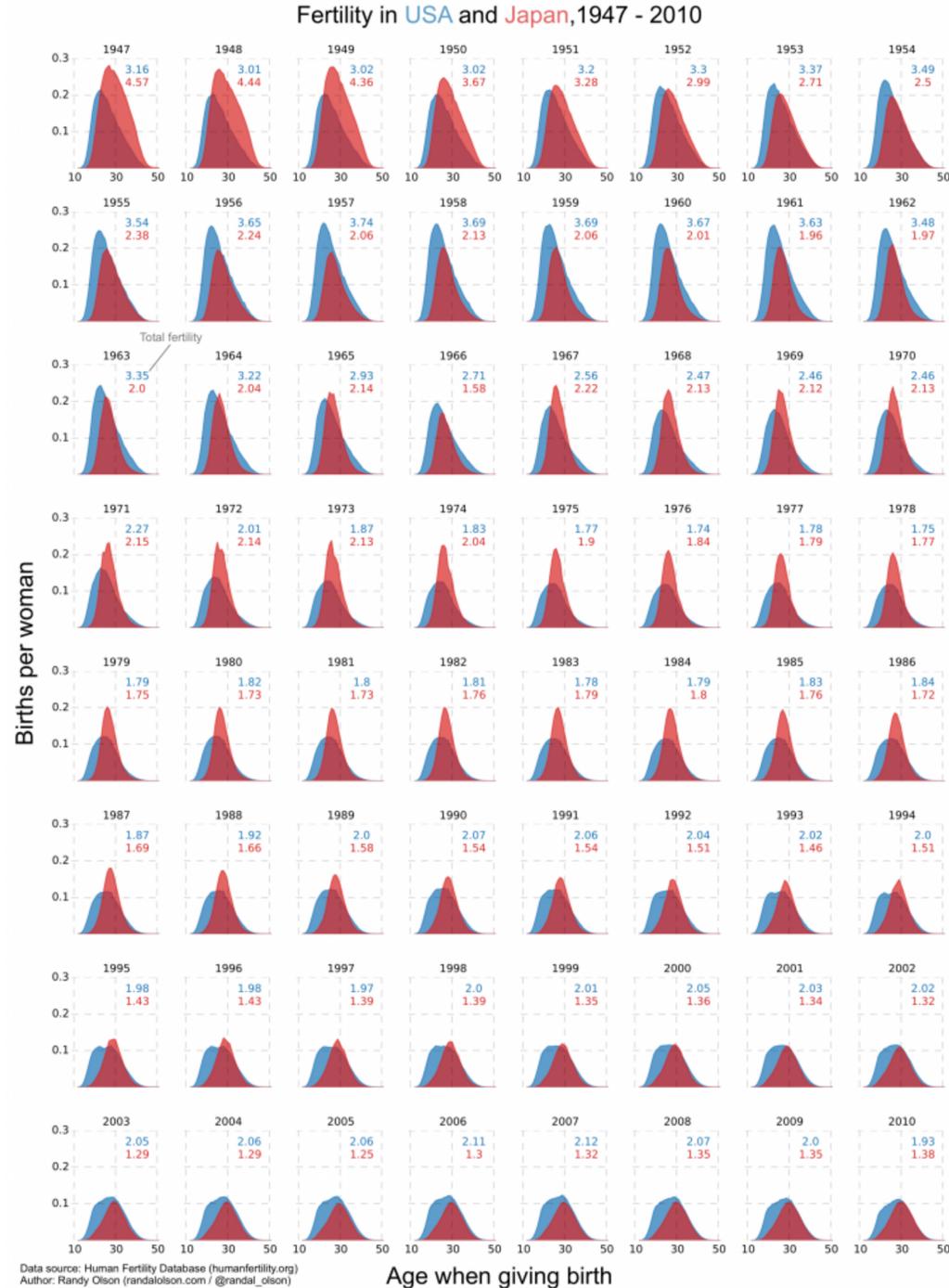
- We were interested in TLS-sites **enriched for both B and T-cells**; these have relevance for prognosis and understanding of the cancer
- Our immunologist wanted to relate this co-localization to the morphology
- **Solution:** transform to single metric (co-localization score) and show

# REDESIGN EXAMPLE

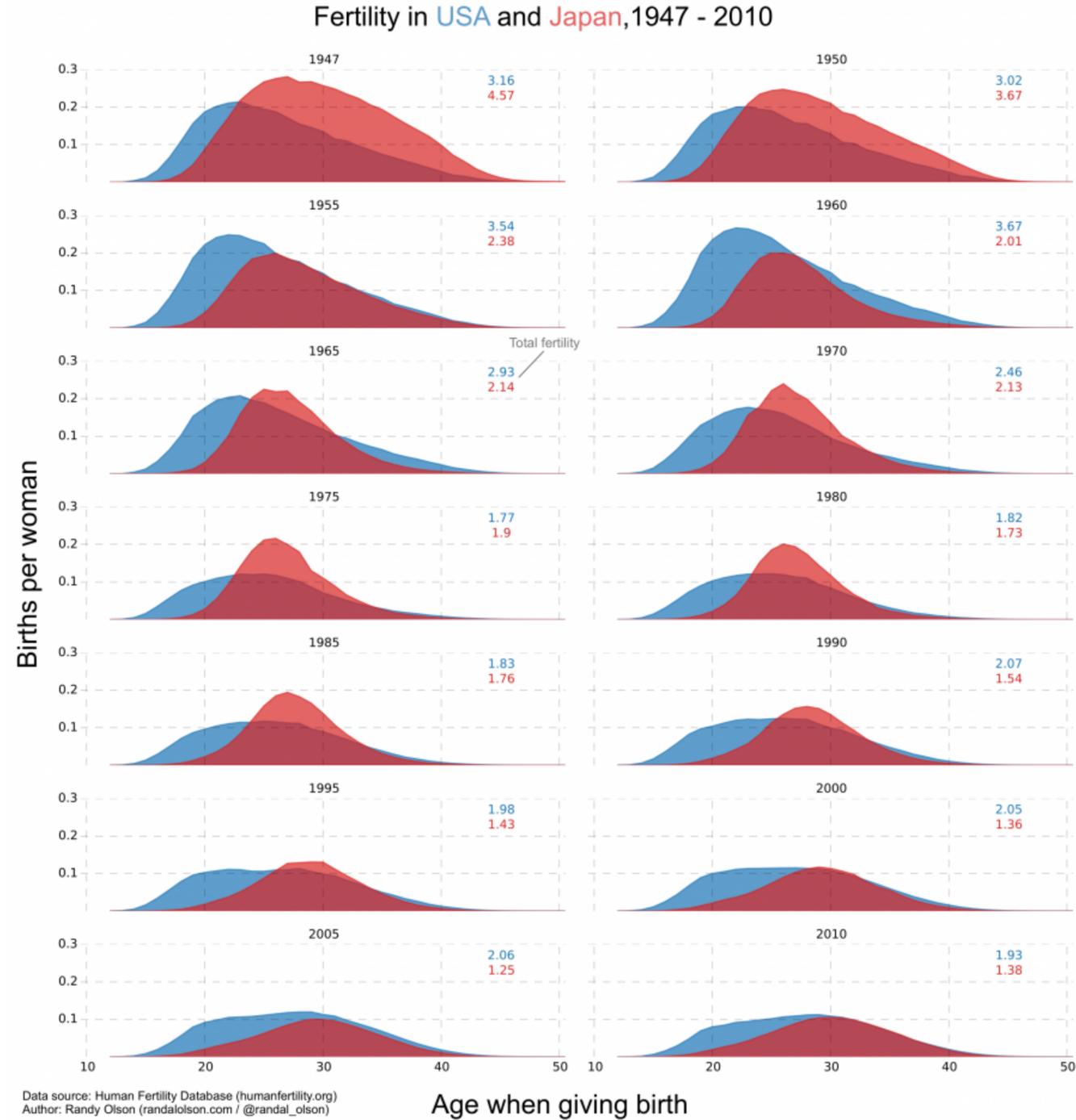


Randal S. Olson

# EYES BEAT MEMORY: SMALL MULTIPLES

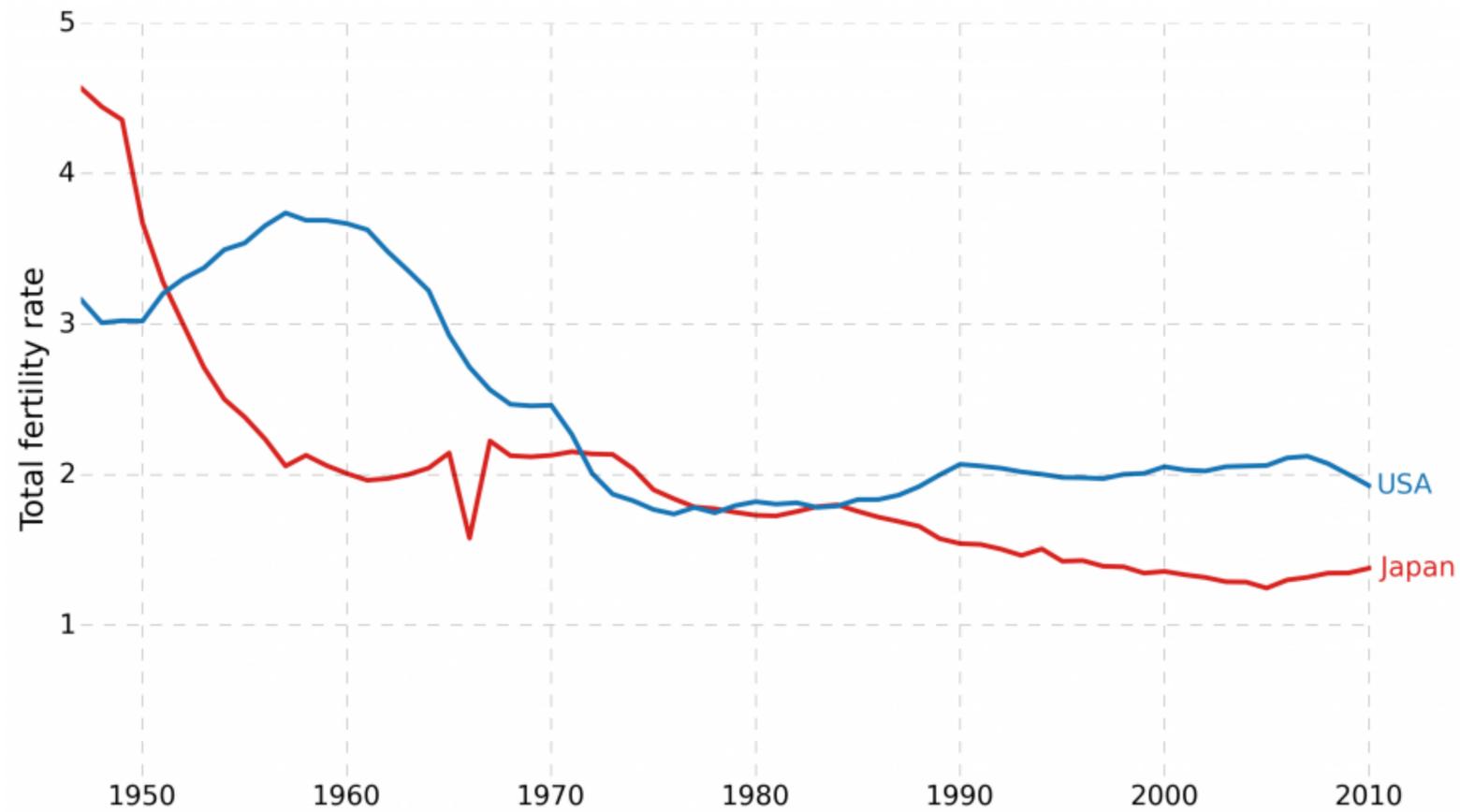


# EYES BEAT MEMORY: SMALL MULTIPLES



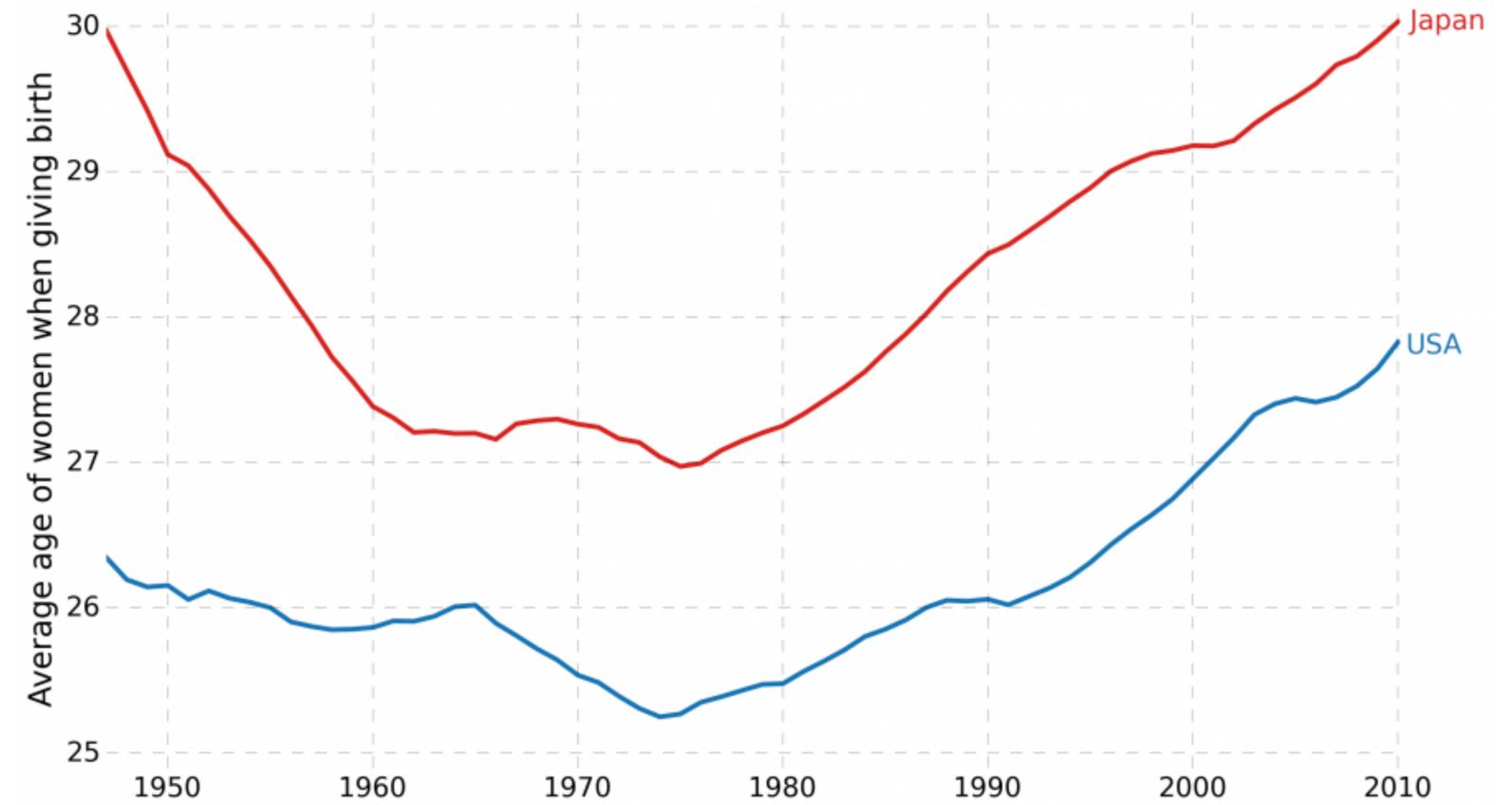
# SIMPLIFY!

Total fertility rate in USA and Japan, 1947 - 2010



Data source: Human Fertility Database (humanfertility.org)  
Author: Randy Olson (randalolson.com / @randal\_olson)

Average age when giving birth in USA and Japan, 1947 - 2010



Data source: Human Fertility Database (humanfertility.org)  
Author: Randy Olson (randalolson.com / @randal\_olson)

## CONCLUSION

**You can't show all the data!**

**But you can show what's important  
about the data.**

**Build tools that give analysts the  
ability to show that!**

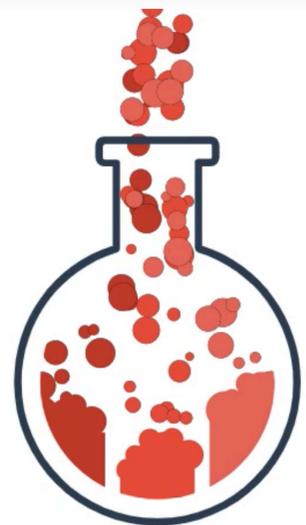
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Thanks to: Carolina Nobre, Kiran Gadhave, Jen Rogers, Haihan Lin, Dylan Wootton, Jochen Görtler, Oliver Deussen, Miriah Meyer, Jeff Phillips, Samuel Gratzl, Holger Stitz, Marc Streit, Nils Gehlenborg, Hilary Coon, Lane Harrison, Hendrik Strobel, Romain Vuillemot, Hanspeter Pfister, and many Others!



**visualization**  
**design lab**

