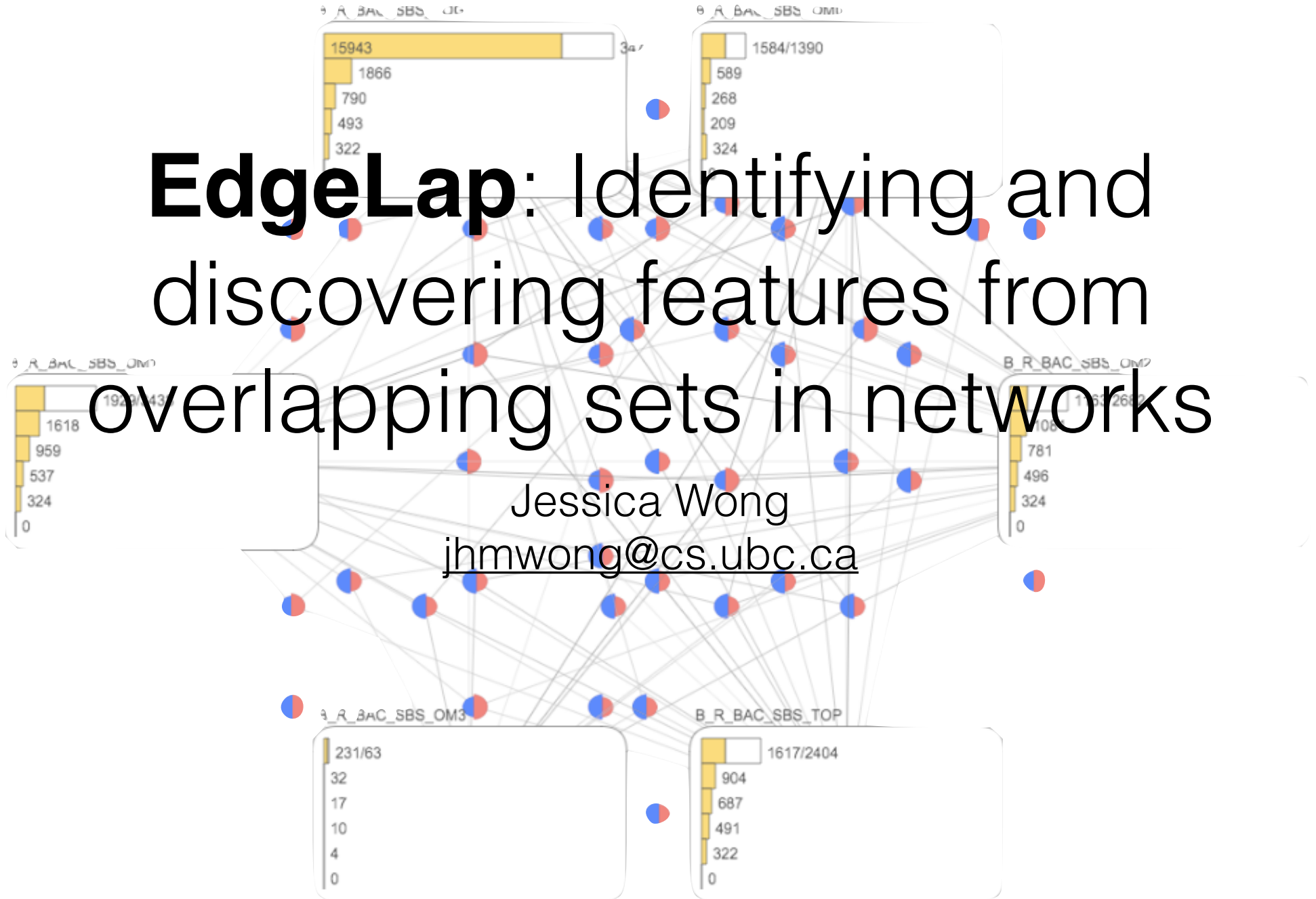


# EdgeLap: Identifying and discovering features from overlapping sets in networks

Jessica Wong

[jhmwong@cs.ubc.ca](mailto:jhmwong@cs.ubc.ca)

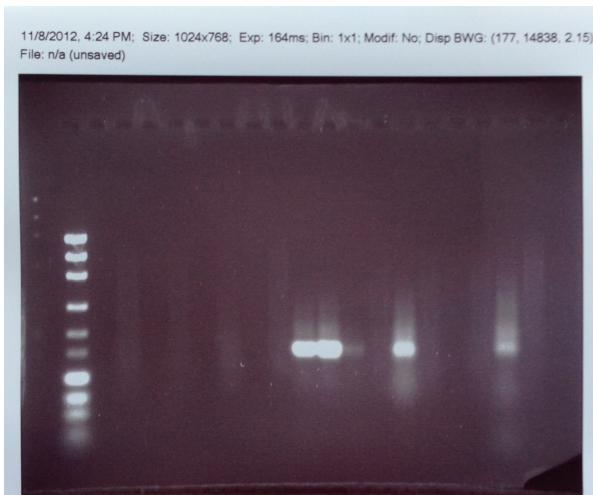
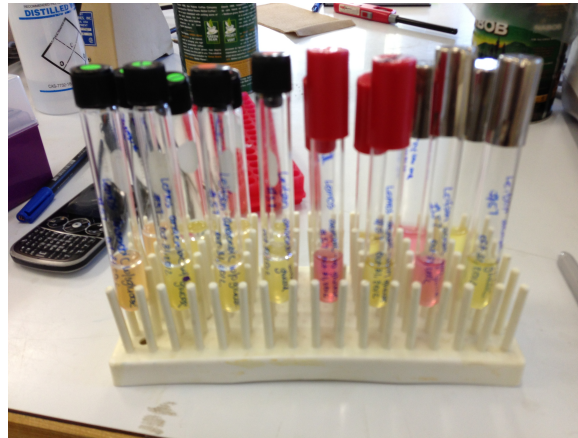
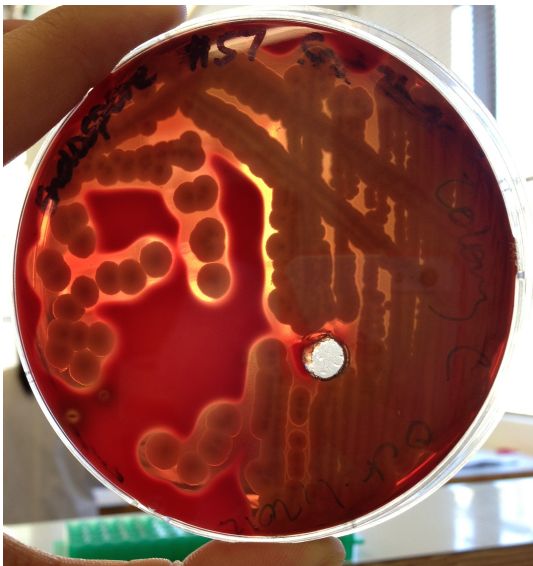


# Data Domain

- Microbes are **everywhere** and commonly work together to survive
- A group of microbes working together is called a **community**
- Common task is to identify microbes present in a community and try to find patterns in their interactions

# Data Domain

- You can identify microbes in many different ways



# Data Domain

- Soil samples were obtained from 18 different sites
- Each site had 4 different methods of harvesting DNA
- Microbes were identified using DNA

# Data Domain

- Each community has microbes and interactions between them
- Interactions can be positively correlated (copresence) or negatively correlated (mutually exclusive)
- Total of 605479 interactions

# Task Abstraction

- Identify and locate common microbial interactions that happen across different networks
- Goal: to generate a hypothesis about the types, and number of correlations found between microbes in different communities

# Complications

- We want to know common interactions that can occur in any combination of networks examined
  - $2^n$  combinations if there are  $n$  networks selected
- Also need to differentiate between positive/negative edges

# EdgeLap

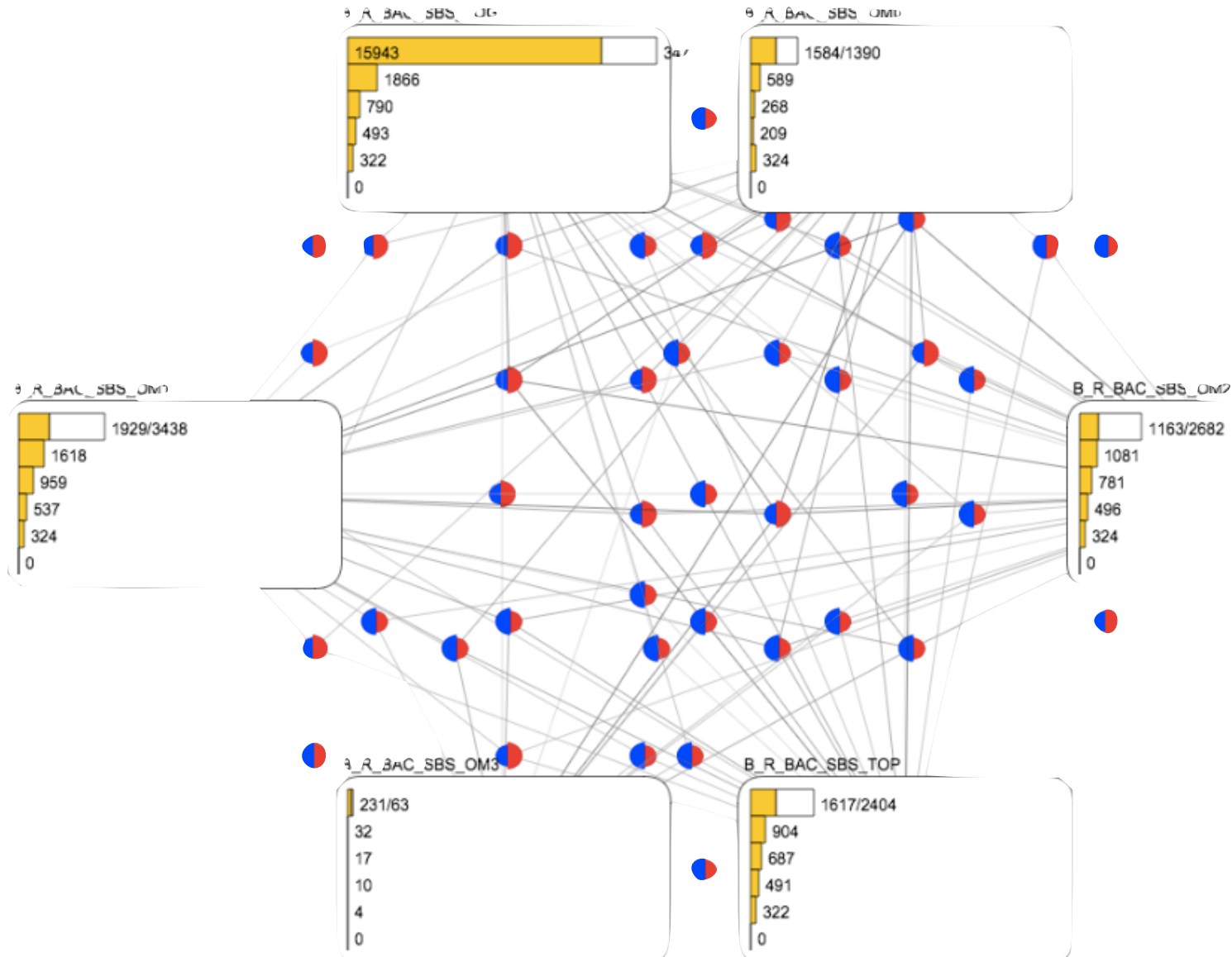
- Inspired from Radial Sets
- Created from a mix of Java, HTML, PHP, JavaScript, and Processing
- Finds common interactions between 2-7 networks



# Step One: Data

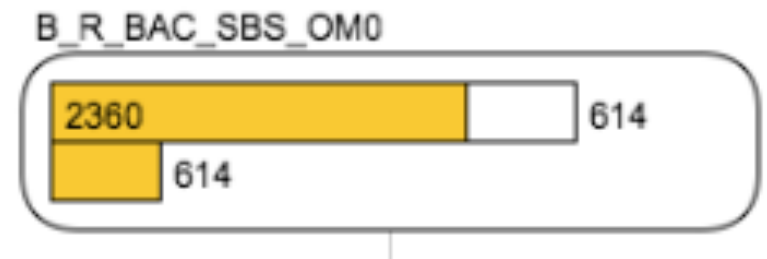
- Data needed to be processed and stored into a database
- Stored in Amazon RDS
- Sorted each interaction by name
- Created an index on the table to help speed up queries

# Step Two: Display!

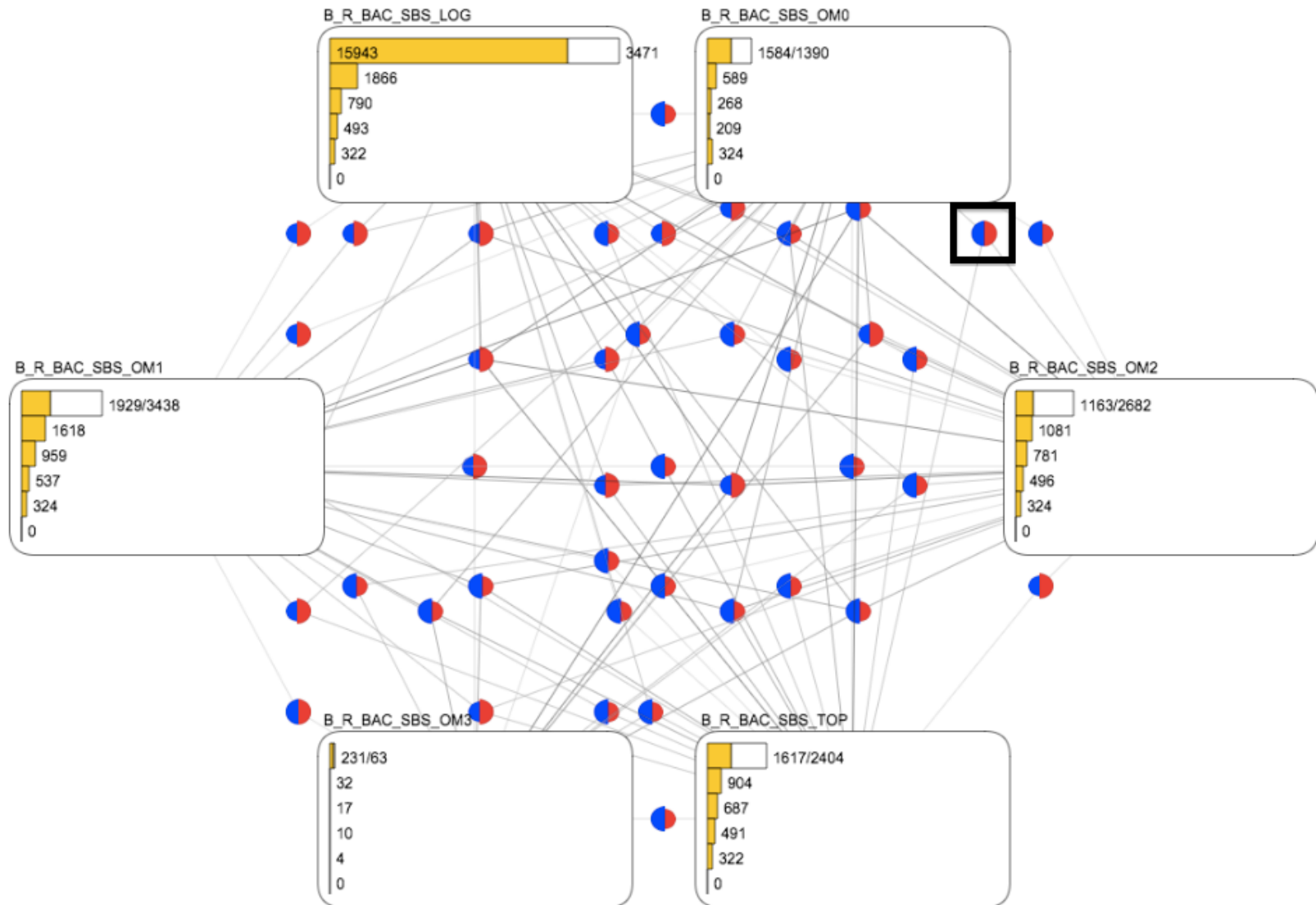


# Network Glyph

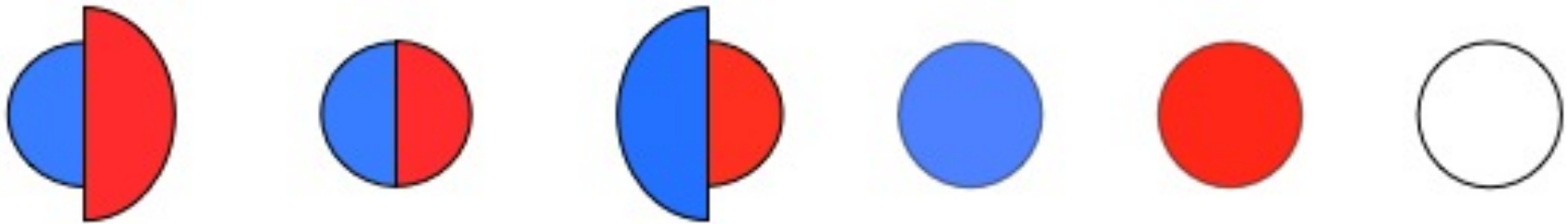
- Network glyphs are meant to show information about how many edges are shared between 0 to n networks
- Meant as secondary information about the network



# Step Two: Display!

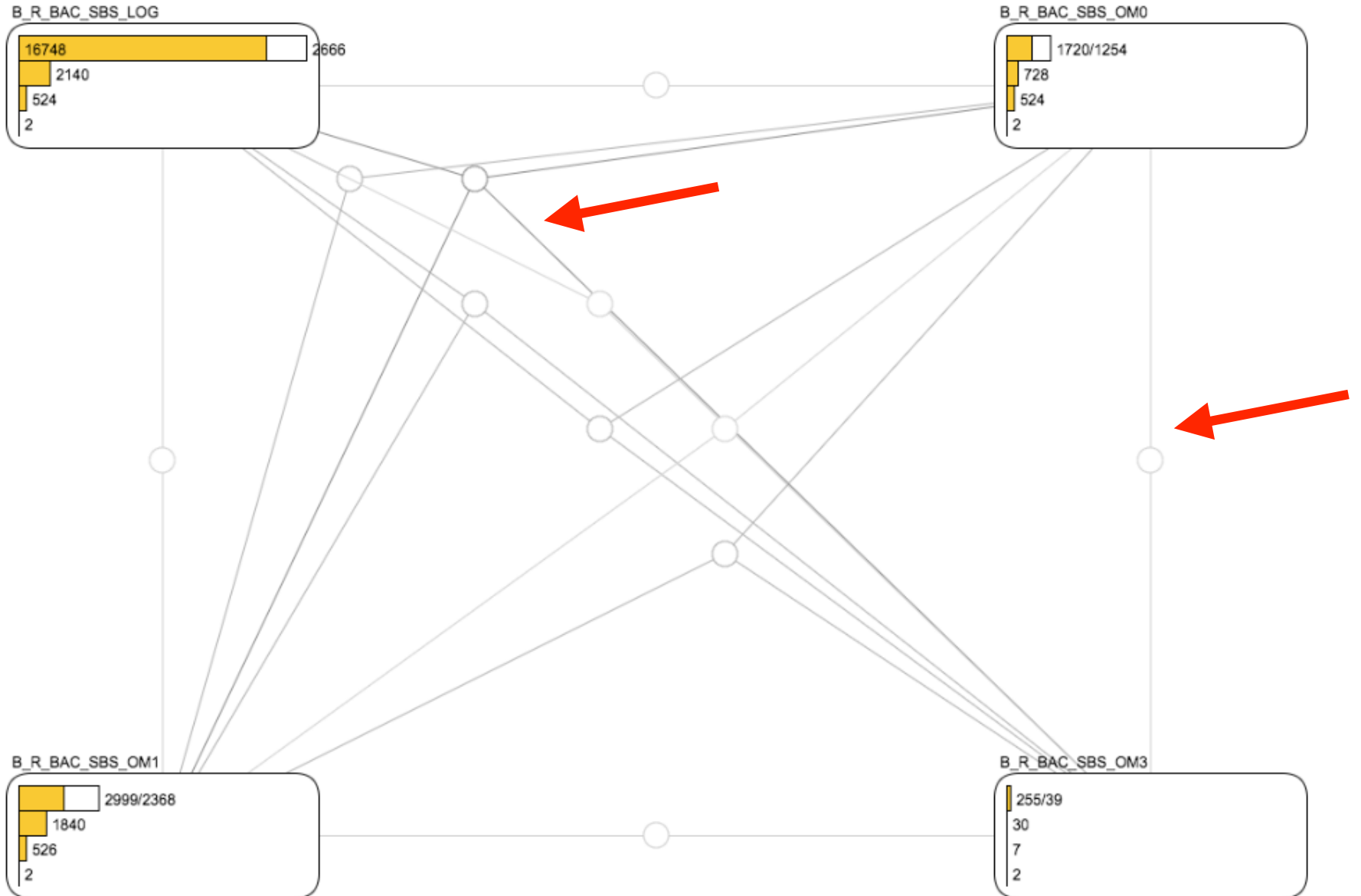


# Step Two: Display!



- Used to show properties of set we are looking at
- Blue = copresence (positive correlation)
- Red = mutual exclusion (negative correlation)
- White doesn't take the type of interaction into account

# Step Two: Display!



# Step Two: Display!

EdgeLap x

localhost/~jhmwong/CPSC547/Processing/

**Networks**

- B\_R\_BAC\_SBS\_LOG
- B\_R\_BAC\_SBS\_OM0
- B\_R\_BAC\_SBS\_OM1
- B\_R\_BAC\_SBS\_OM2
- B\_R\_BAC\_SBS\_OM3
- B\_R\_BAC\_SBS\_TOP
- R\_BAC\_IDF\_OL
- R\_BAC\_IDF\_OM1
- R\_BAC\_IDF\_OM2
- R\_BAC\_IDF\_OM3

Draw Visualization

Undirectional Relationships Only

Copresence Relationships Only

Mutually Exclusive Relationships Only

Clear Selection

To SVG

# Future Work

- Filtering option
- Hovering over the circle glyphs will highlight associated lines, display summary information
- Better method to draw lines to avoid too much overlap
- Improve performance



# Q&A

- Questions?