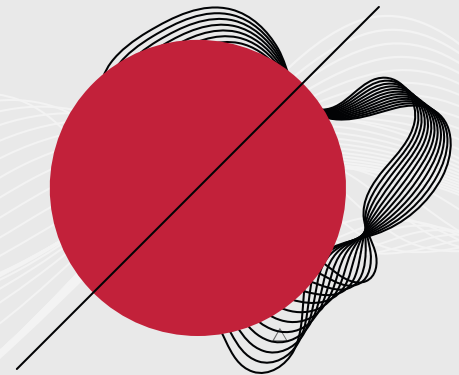
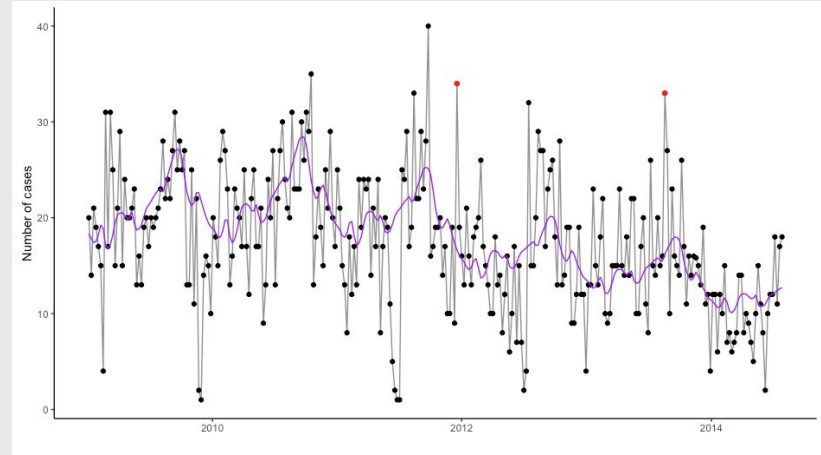
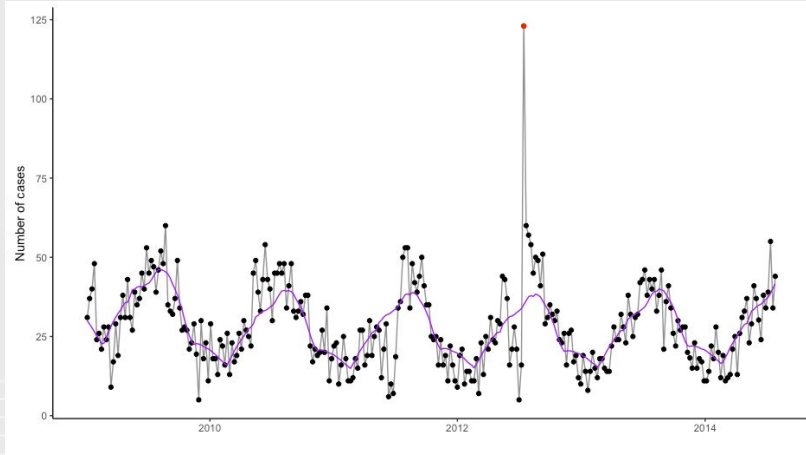


# **Disease Outbreak Radar:** **A Tool for Public Health Users**

UBC CPSC 547 Project  
Cloris Feng, Derek Tam, Tae Yoon (Harry) Lee  
Dec 10, 2020

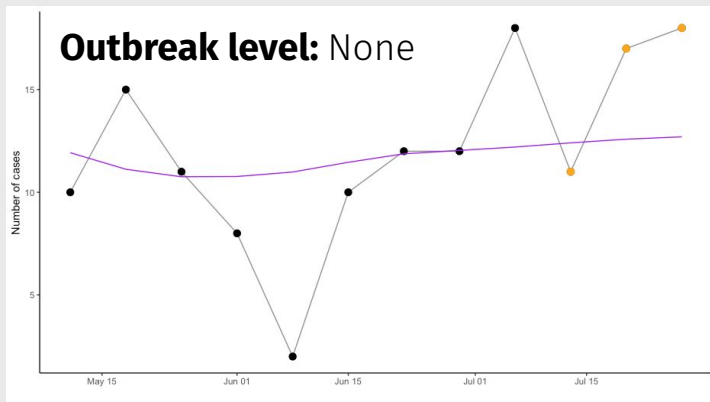
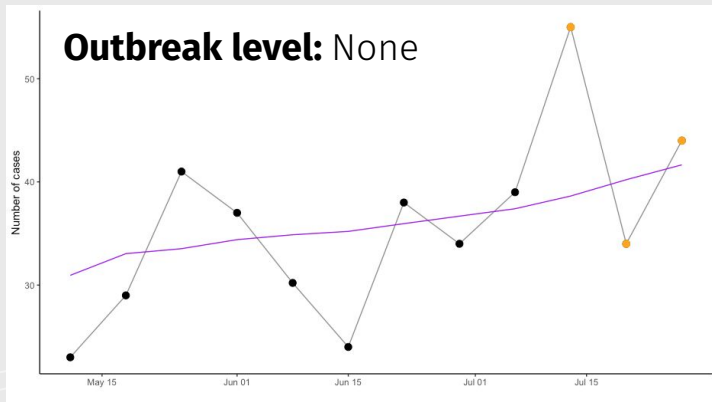
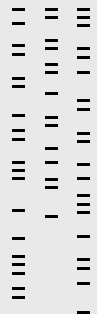


# Challenges in Disease Outbreak Detection



- “A disease outbreak is defined as the occurrence of disease cases in excess of normal expectancy.” [World Health Organization]
- Due to complex disease characteristics, it is difficult to define the “norm.”
- An automated method has been developed in collaboration with British Columbia Centre for Disease Control to reduce the burden and to help them identify disease outbreaks.

# Outbreak level



- Outbreak p-value: the probability of observing the **three** most recent numbers of cases.
- Outbreak level:
  - High: outbreak p-value  $\leq 0.1\%$
  - Medium:  $0.1\% < \text{outbreak p-value} \leq 1\%$
  - Low:  $1\% < \text{outbreak p-value} \leq 5\%$
  - $\blacktriangle$  None: outbreak p-value  $> 5\%$



# Motivation

BC CDC is monitoring around **60** diseases in each of the **16** regions of BC. They need to check whether there is an outbreak in  $60 \times 16 \approx$  **800** diseases **weekly**.

# WHAT: Multidimensional Table

Publicly available disease count data of the United States was used (Project Tycho).

Attribute name	Attribute type	Levels/range	Description
Disease	Categorical	6	△ Disease name
Week date	Sequential/ordered	2009-01-04 - 2014-07-27	Start and end dates of a week in which the number of cases is collected (~290 observations)
Region	Categorical	9	US census divisions
Number of cases	Quantitative	0 - 3,140 (integers)	Number of cases of a disease
Number of population	Quantitative	14,469,650 - 62,382,273	Number of population estimated by the US Census in the year of Week date
<b>Derived attributes</b>			
Rate of cases	Quantitative	0.16 - 1094.90 (per 10,000,000)	Number of cases divided by the number of population
Estimated number of cases	Quantitative	Non-negative real valued	Estimated by the method
Outbreak p-value	Quantitative	0-1 (real-valued)	Probability of observing the 3 most recent numbers of cases based on the method
Outbreak level	Categorical	User-defined	Level of outbreak specified by public health users based on the outbreak p-value

**Table 1.** List of attributes and derived attributes.

# WHY: Search, Analyze, and Compare



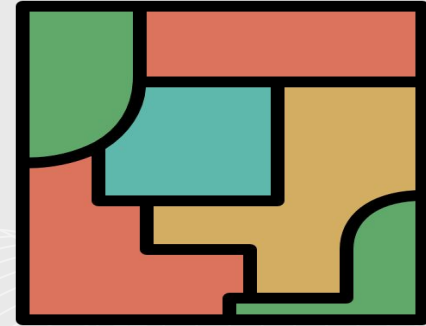
## Task 1

Efficiently **search** for an outbreak in diseases across regions



## Task 2

**Analyze** the trend in the number of cases and estimated number of cases for each disease with an outbreak

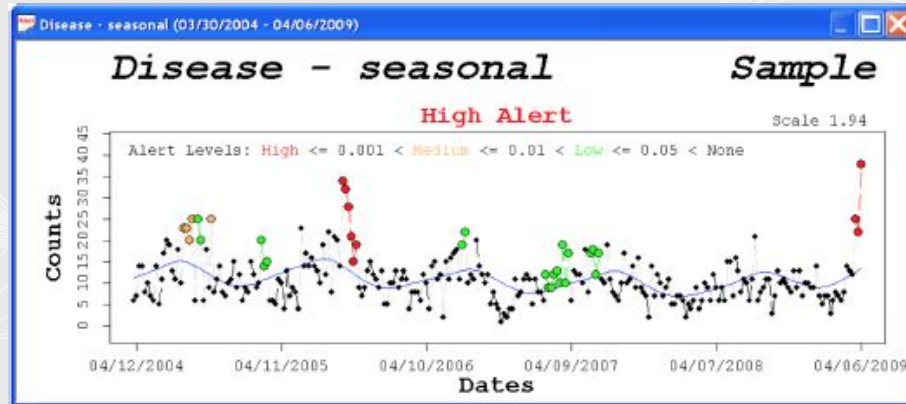


## Task 3

**Compare** the outbreak levels across regions for each disease with an outbreak

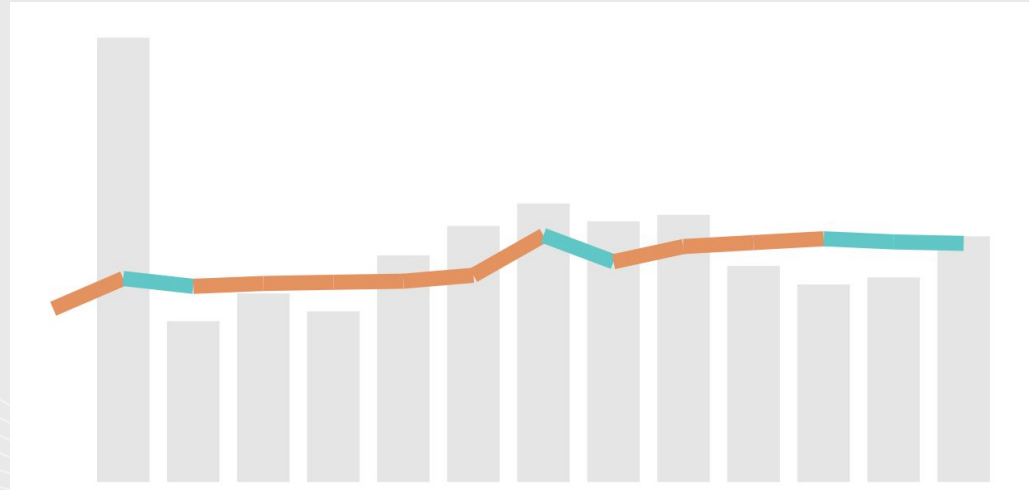
# Analysis of COVID Dashboards and Design Guidelines

- Analyzed 4 COVID dashboards & 8 design guidelines
  - Learned lessons specific to visualization of **case data** with **geographical context** and applied them to our prototype
  - Guidelines were used to specifically address Task 2 and Task 3



# Recommendation for Task 2

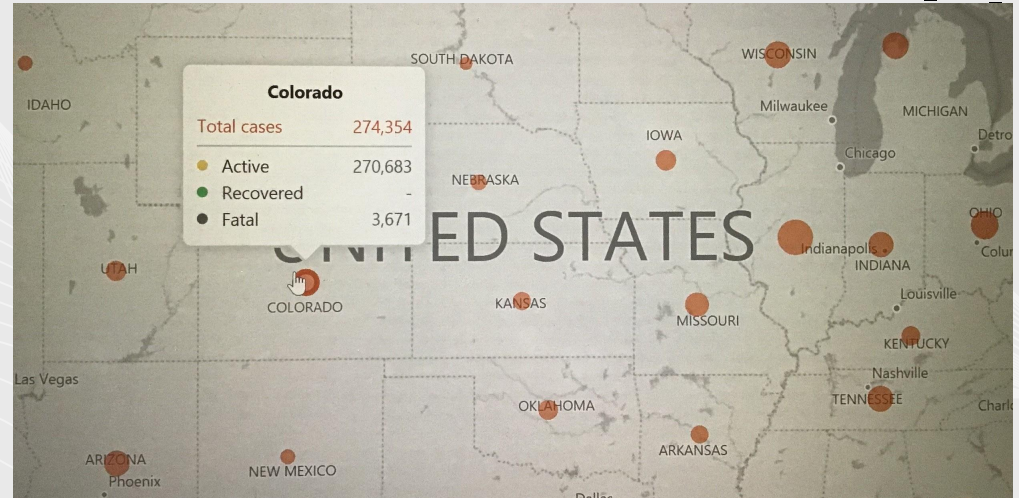
- **Task 2:**  
Analyze trends in the number of cases for a given region and disease
- **Recommendation:**  
A line-over-bar chart, with gain/loss trend-lines and annotated weeks of interest



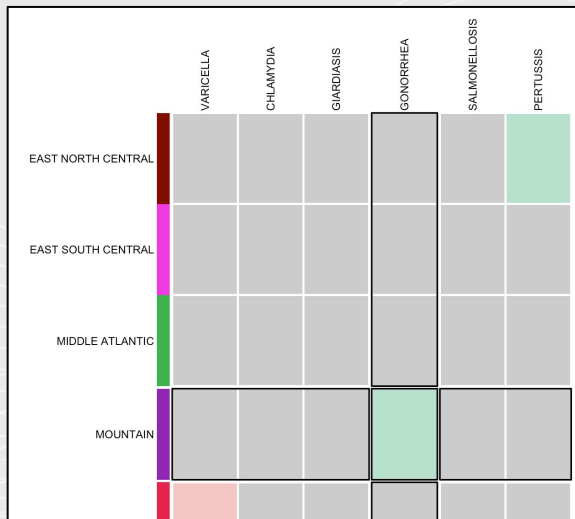


# Recommendation for Task 3

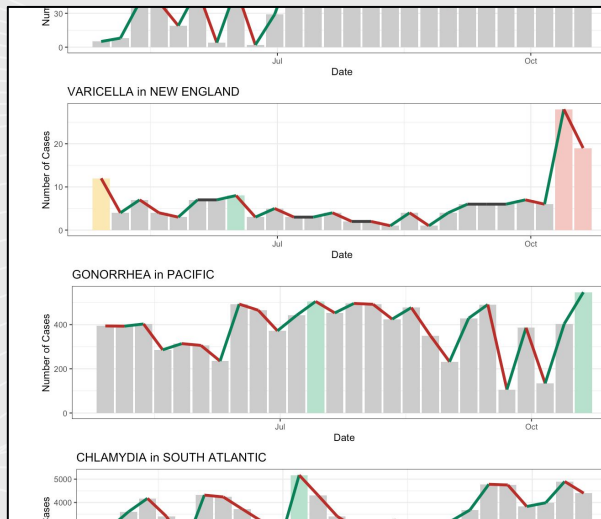
- **Task 3:**  
Compare the outbreak risk across regions for each disease
- **Recommendation:**  
A combination of bubble map and dot-density map to handle case number complexity



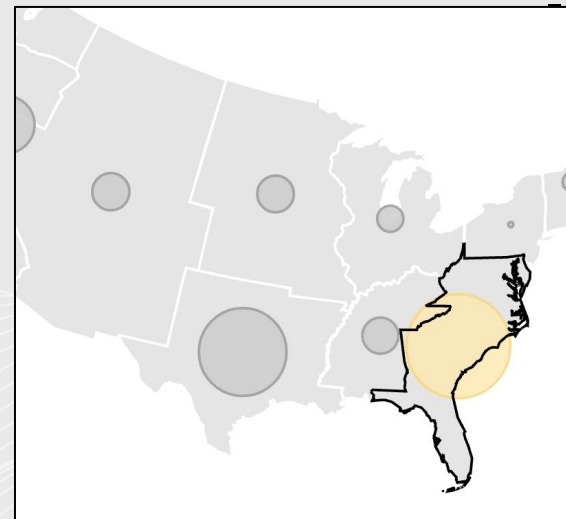
# HOW: Implementation Design



Task 1



Task 2



Task 3

# Implementation and Demo

- Outbreak detection algorithm was developed by Harry as part of his Master's thesis
  - Algorithm was applied to a subset of Tycho data
- Dashboard was implemented in **R**
  - Made interactive with the **ShinyApp package** and **plotly**
  - Web-hosting through **shinyapps.io**



# Limitations and Next Steps

- Tycho dataset is an **adequate but non-ideal** proxy for real data
- Dashboard could use additional quality-of-life features
  - Reordering and removal of individual timeseries
  - Region selection via map
- Reporting and export for diseases and regions of interest