

# A parameterized approach to the analysis of Otitis Media data

A Collaborative project between the PCRU and  
Menzies School of Health

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# Overview

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# Introduction

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# Motivation

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# Motivation

- ⊙ Otitis Media (Ear Infection) is a major health issue in NT AU.
- ⊙ 30% of Children are deaf. 99.7% of Indigenous inmates are deaf [5].
- ⊙ Currently vaccines and antibiotics not sufficiently effective
- ⊙ Multi-pathogen disease. See <http://www.ncbi.nlm.nih.gov/pubmed/23523773>
- ⊙ PCA and GLM limited to 1 : *many* interactions
- ⊙ Clinical researched requested Network Model analysis



## Research Direction

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PCRU has been developing a suit of tools for multivariate data analysis that utilizes FPT (Clique Centric) Algorithms

1. Fixed Parameter Tractable (FPT)  $T(n, k) = a^k + O(n^c)$
2. Microbiological [DNA, RNA, Protein-Protein interactions](#) [2, 6]
3. Ecological Data Silos [Northern Seas Ocean study](#) [2]
4. **Currently focusing on Clinical Data**
  - Type I Diabetes
  - Otitis Media (OM)
  - ALRI [4]



## Study Population and Design

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# Study Population and Design

- ⊙ Random collection trial between 1996-2001,2001-2004
- ⊙ swabs were collected from two months to 2 years of age
- ⊙ 740 nasopharyngeal swabs (these are sequenced)
- ⊙ to produce 29 variables
- ⊙ Correlations were computing between all pairs of variables
- ⊙ a threshold of 0.2 was used
- ⊙ Key variates are Otitis Media (OM) and ALRI.

# Unclustered Variables Hard to Understand

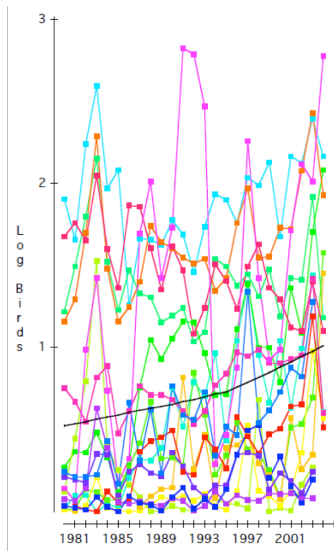


Figure: Unclustered variables are NP-Hard to unscramble.[2]

## Analysis Tools



# Clique Centric Pipeline

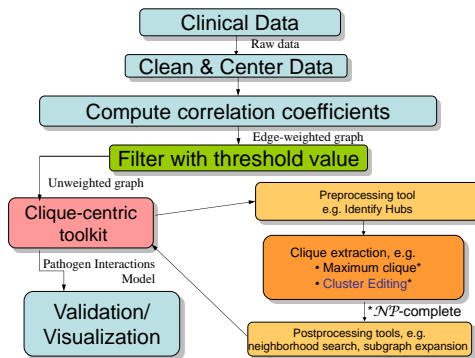


Figure: The role of CLUSTER EDITING in the clinical analysis pipeline. This figure and methodology are an adaptation of work by Langston et al. [3]

# Don't compound the error

- ⊙ Clinical, psychological and biological measurements can be fuzzy
- ⊙ Many unsupervised clustering problems are NP-Hard
- ⊙ FPT offers a way to compute **exact solutions**
- ⊙ Why use approximations when you can calculate the exact solution

## Computing a Network Model

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# CLUSTER EDITING Problem

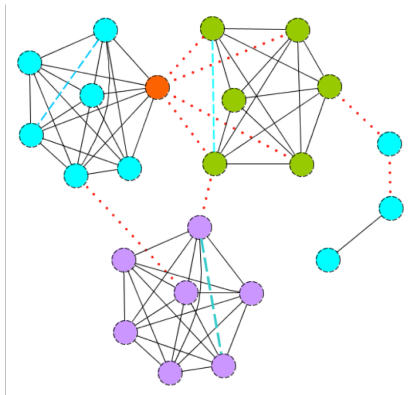


Figure: **Added Edges** colored blue. **Deleted Edges** Colored Red

- ⊙ Missing edges indicate False-negative correlations  
Perhaps the relationship is not linear or effected by outliers
- ⊙ Added edges indicate False-Positive correlations



Whats new about how we analyse this Clinical Data.

Otitis Media (OM), Acute Lower Respiratory Infections (ALRI) analysis and Type I Diabetes

- ⊙ Heterogeneous Data
- ⊙ Categorical, Continuous, Logical Data means
- ⊙ New Measures of correlation
  - Spearmans Rank
  - Polychoric, Tetrachoric
  - Bi-Serial

# Higher levels of Noise

- ⦿ Lower thresholds for significant correlation
- ⦿ could use *eigen values* but also has to make sense in the domain
- ⦿ This means more Noise
- ⦿ Faster Algorithms needed


# Aggresize Parameterization

- ⊙ Multi-parameter CLUSTER EDITING

$$T(n, (k, a, d)) = \alpha^k + f(a) + f(d) + (n^c)$$

- ⊙ Bound the number of false +ve and -ve correlations that can be attributed to a single Variable.
- ⊙ We need different values to deal with hubs.
- ⊙ New (Hybrid) Graph Structure
- ⊙ New algorithms faster and also models the noise better.

## Results



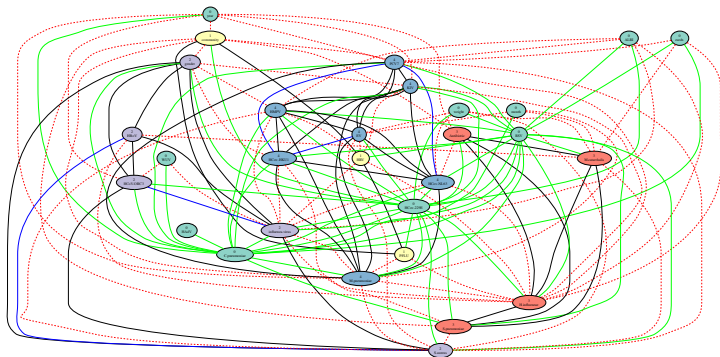


Figure: Pathway analysis of nasal pathogens reveals four distinct clusters using clique as a structural measure. (The diagram was produced using Graphvis by first weighting and coloring the graph based on the clusters found. [1]).

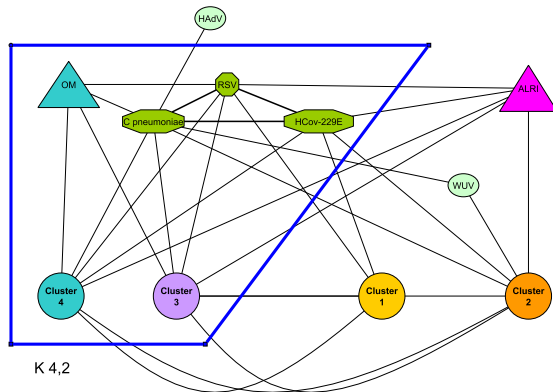


Figure: Pathway analysis of nasal pathogens reveals four distinct clusters using clique as a structural measure.

## Cluster 4

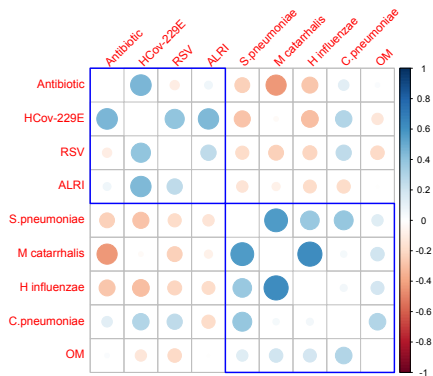


Figure: A heatmap of Cluster 4 with secondary hysterical clustering (produced using R:package corrplot [?])

With regard to the vaccination PCV7 we can see that this is

# Cluster 3

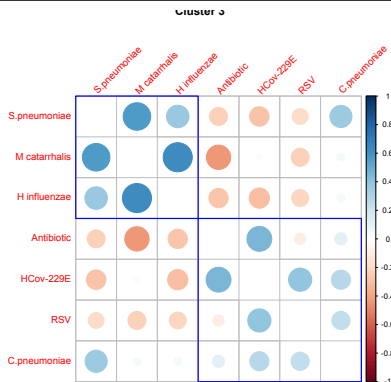


Figure: A heatmap of Cluster 3 with secondary hysterical clustering (produced using R:package corrplot)



- ⊙ Much faster >> 600 times
- ⊙ Hybrid Data Structure
- ⊙ Aggressive Multi-Parameter
- ⊙ Hubs

## Conclusion

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- ⊙ Use  $t(X)$  to cluster patient with similar symptoms
- ⊙ Data Linkage data sets  $50k - 100k$
- ⊙ Spectral pre-processing
- ⊙ Need good **heuristics** for these large data sets

\* **(1)** Does the Development of Hypotheses Stifle the Development of Knowledge? The Role of Cognitive Biases and the Benefits of Data Mining.

Simon Moss and Peter Shaw

\* **(2)** Effective Use of Multi-parameterized Correlation Clustering in Mining Nasopharyngeal Carriage and Disease data from Young Children

Peter Shaw, Faisal N. Abu-Khzam, Robyn Marsh, Heidi Smith-Vaughan (2017)

\* **(3)** Cluster Editing with Vertex Splitting

Faisal N. Abu-Khzam, Serge Gaspers, Alexis Shaw Peter Shaw

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