## Indian Buffet Epidemics A non-parametric Bayesian Approach to Modelling Heterogeneity

Ashley Ford, Gareth Roberts

Department of Statistics, University of Warwick

Inference For Epidemic-related Risk, 2011









Bipartite Graph Epidemic models

Inference for bipartite epidemic models

Indian Buffet Process

Indian Buffet Epidemics

- Need a model between
  - homogeneous mixing
  - over complex models with unknown parameters.
- Many have been proposed
  - household, spatial, multi-type
- Availability of data
  - contact surveys
    - RFID
    - POLYMOD
  - commuting data
- A non-parametric model for the heterogeneity that can represent a wide range of departures from homogeneity.

- Need a model between
  - homogeneous mixing
  - over complex models with unknown parameters.
- Many have been proposed
  - household, spatial, multi-type
- Availability of data
  - contact surveys
    - RFID
    - POLYMOD
  - commuting data
- A non-parametric model for the heterogeneity that can represent a wide range of departures from homogeneity.

・ロト ・ 個ト ・ ヨト ・ ヨト ・ 三日 ・ のへで

- Need a model between
  - homogeneous mixing
  - over complex models with unknown parameters.
- Many have been proposed
  - household, spatial, multi-type
- Availability of data
  - contact surveys
    - RFID
    - POLYMOD
  - commuting data
- A non-parametric model for the heterogeneity that can represent a wide range of departures from homogeneity.

- Need a model between
  - homogeneous mixing
  - over complex models with unknown parameters.
- Many have been proposed
  - household, spatial, multi-type
- Availability of data
  - contact surveys
    - RFID
    - POLYMOD
  - commuting data
- A non-parametric model for the heterogeneity that can represent a wide range of departures from homogeneity.

## Places and People

- Model heterogeneity in an epidemic amongst N people
- Each person belongs to 1 or more of many classes
  - e.g. households, schools, clubs, buses etcetera

- represented as
  - a bipartite graph
  - an  $N \times K$  binary matrix Z

## Places and People

- Model heterogeneity in an epidemic amongst N people
- Each person belongs to 1 or more of many classes
  - e.g. households, schools, clubs, buses etcetera

◆□▶ ◆□▶ ◆□▶ ◆□▶ 三回日 のの⊙

- represented as
  - a bipartite graph
  - an  $N \times K$  binary matrix Z

## Example bipartite graph



## Epidemics on bipartite graphs

Extend homogeneous mixing epidemic models to bipartite graph

▶ e.g. SIR, SEIR, SIS, Reed-Frost

Approaches to defining the infection rate

- a single infection rate could apply to all pairs of individuals connected through one or more locations
- Each class has an associated infection rate  $\lambda_k$
- ► Rate of infections on a susceptible individual j is  $\sum z_{jk} \lambda_k N_{k,t}^{l}$

•  $N_{k,t}^{l}$  is the number that are in classs k and infective at time t.

# Simulation

- ◆□▶ ◆□▶ ◆三▶ ◆三▶ 三目目 のへで

## Household epidemic models

### Global and within household infection rates $\lambda_g$ and $\lambda_h$

household size mnumber of housholds  $n_h$ a bipartite graph representation with adjacency matrix  $N \times (n_h + 1)$ where  $N = n_h m$ e.g. for 4 houses of sizes 2,3,3,4 the adjacency matrix is

| $\lambda_g$ | $\lambda_h$ | $\lambda_h$ | $\lambda_h$ | $\lambda_h$ |
|-------------|-------------|-------------|-------------|-------------|
| 1           | 1           |             |             |             |
| 1           | 1           |             |             |             |
| 1           |             | 1           |             |             |
| 1           |             | 1           |             |             |
| 1           |             | 1           |             |             |
| 1           |             |             | 1           |             |
| 1           |             |             | 1           |             |
| 1           |             |             | 1           |             |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |

## Household epidemic models

Global and within household infection rates  $\lambda_g$  and  $\lambda_h$ 

```
household size m
number of housholds n_h
a bipartite graph representation with
adjacency matrix N \times (n_h + 1)where
N = n_h m
e.g. for 4 houses of sizes 2,3,3,4 the
adjacency matrix is
```

| $\lambda_g$ | $\lambda_h$ | $\lambda_h$ | $\lambda_h$ | $\lambda_h$ |
|-------------|-------------|-------------|-------------|-------------|
| 1           | 1           |             |             |             |
| 1           | 1           |             |             |             |
| 1           |             | 1           |             |             |
| 1           |             | 1           |             |             |
| 1           |             | 1           |             |             |
| 1           |             |             | 1           |             |
| 1           |             |             | 1           |             |
| 1           |             |             | 1           |             |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |
| 1           |             |             |             | 1           |

## Multi-type model

A multi-type model with infection rate  $\gamma_{i,j}$  between an infective in group i and a susceptible in group j.

If  $\gamma_{i,j} = \gamma_{ji}$  and  $\sum_{j \neq i} \gamma_{i,j} \leq \gamma_{i,i} \forall i$  it can be represented as a bipartite model.

These conditions will usualy apply if the groups are geographically separate,

but may not if the groups are split by ages or varying susceptibility and infectiousness e.g.  $\gamma_{i,j}=lpha_ieta_j.$ 

With m types the bipartite representation has m columns for the within type infections and m(m-1)/2 for the between type infections.

## Multi-type model

A multi-type model with infection rate  $\gamma_{i,j}$  between an infective in group i and a susceptible in group j.

If  $\gamma_{i,j} = \gamma_{ji}$  and  $\sum_{j \neq i} \gamma_{i,j} \leq \gamma_{i,i} \forall i$  it can be represented as a bipartite model.

These conditions will usualy apply if the groups are geographically separate,

but may not if the groups are split by ages or varying susceptibility and infectiousness e.g.  $\gamma_{i,j} = \alpha_i \beta_j$ .

With m types the bipartite representation has m columns for the within type infections and m(m-1)/2 for the between type infections.

# Multi-type model

| $\gamma_{1,1}-\gamma_{1,2}-\gamma_{1,3}$ | $\gamma_{2,2} - \gamma_{1,2} - \gamma_{2,3}$ | $\gamma_{3,3}-\gamma_{1,3}-\gamma_{2,3}$ | <b>Ŷ</b> 1,2 | <b>γ</b> 1,3 | <b>γ</b> 2,3 |
|--|--|--|--------------|--------------|--------------|
| 1  |  |  | 1            | 1            |              |
| 1  |  |  | 1            | 1            |              |
| 1  |  |  | 1            | 1            |              |
|  | 1  |  | 1            |              | 1            |
|  | 1  |  | 1            |              | 1            |
|  |  | 1  |              | 1            | 1            |
|  |  | 1  |              | 1            | 1            |
|  |  | 1  |              | 1            | 1            |
|  |  | 1  |              | 1            | 1            |

# Examples



≣I≡ ୬**୯**୯

## Other Models

- three level
- commuting
- ► known labels

### MCMC inference for known contact matrix

log likelihood when observed on  $[0, T_{max}]$ with *n* infections at  $T_i^I$  and removals at  $T_i^R$ 

$$\sum_{j} \log \eta_j(T_j^I) - \int_0^{T_{\max}} \sum_j \eta_j(t) dt + \sum_{j} \log g(T_j^R - T_j^I) + \sum_{j} \log \{1 - G(T_{\max} - T_j^I)\}$$

where  $\eta_j(t) = \sum_k z_{jk} \lambda_k N_{k,t-}^l$  is the instantaneous rate of infections on individual *j*, *g* and *G* are the pdf and cdf of time to recovery. A simulation with *Z* 1000 × 21 and  $\lambda_k = .5/n_k$  has 626 infections

ヨト イヨト ヨヨ のへの



## MCMC inference for known contact matrix

### RW - Metropolis-Hastings for 21 parameters



true values 0.00058, 0.00150, 0.17000

## non-parametric inference

- Parametric distributions
  - ► Gaussian, Cauchy
  - exponential, gamma, log-normal, Weibull

- Non-parametric
  - histograms
  - kernel density estimate

## Indian Buffet Process- a culinary metaphor



Introduced by Griffiths and Ghahramani (2005).

- ► N customers enter a restaurant one after another.
- The first customer selects Poisson(α)
- The *j*th customer selects each dish with probability  $m_k/j$ 
  - ▶ where m<sub>k</sub> is the number of previous customers who have chosen k.
- and then tries  $Poisson(\alpha/j)$  new dishes.
- A distribution over all binary matrices with N rows
  - $\blacktriangleright$  Expected number in each row is lpha
  - Expected number of non zero columns  $\alpha \sum_{j=1}^{N} 1_{j}$ , where  $\beta \in \mathbb{R}^{n}$

### Indian Buffet Process - example



Classes

IBP Z generated with  $N = 260, \alpha = 15$ 

Individuals

## Limit of Finite K

- $\psi_k$  is the probability that an individual is in class k
- $\psi_k \sim Beta(lpha/\kappa, 1)$  or from stick breaking
- ▶ The model for Z is:  $z_{ik}|\psi_k \sim Bernoulli(\psi_k)$  independently

• The Indian Buffet process is obtained as  $K \to \infty$ 

### Indian Buffet Epidemic

Combine the bipartite graph model with the Indian Buffet Process as a prior for the contact graph.



 $lpha=6,\ N=600,\ K_{+}=31$  , different initial infectives in 8,4,1 . It is also seen

## MCMC for Indian Buffet Epidemic

- A challenging MCMC problem
  - Very high dimensional
  - multi-modal
- Proposals
  - non centered, using fixed K
  - independence, using sequential IBP

◆□▶ ◆□▶ ◆□▶ ◆□▶ ヨヨ のへ⊙







## Conclusions

- Bi-partite graph epidemics provide a generic formulation for modeling and inference.
- The Indian Buffet Epidemic provides a non-parametric model for heterogeneity in contact processes.
- MCMC inference is possible on small epidemics
  - ▶ Work continues on extending the size which can be handled

◆□▶ ◆□▶ ◆目≯ ◆目≯ ◆□▶

◆□▶ <舂▶ <差▶ <差▶ 差目 のへの</p>

## For Further Reading I

#### 

T.L. Griffiths and Z. Ghahramani.

Infinite latent feature models and the Indian buffet process (tech. rep. no. 2005-001). Gatsby Computational Neuroscience Unit, 2005.

◆□▶ ◆□▶ ◆□▶ ◆□▶ ヨヨ のへ⊙