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Emily's Thesis Title

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A dissertation
submitted in partial fulfillment of the
requirements for the degree of

Doctor of Philosophy

University of Washington

2021?

Reading Committee:

Steven M. Goodreau, Chair

person 1

person 2

person 3

Program Authorized to Offer Degree:

Biological Anthropology

University of Washington

Abstract

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Emily Pollock

Chair of the Supervisory Committee:

Title of my chair Steven M. Goodreau

Biological Anthropology

“Here is my abstract”

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Acknowledgments

“My acknowledgments”

Dedication

“My dedication”

Introduction

- social sciency interest in networks & disease transmission
- dynamic networks require information about relationship duration
- transition to a history of the evolution of epidemic models (ie. from compartmental where everything is basically independent and exponential through to ERGMs where formation can be quite elaborate but we've never spent much time thinking about dissolution)
- why doing a bit better on dissolution/duration, especially by age, will be extra important for thinking about certain interventions for certain relatively short-lived infections (e.g. partner services in chlamydia).
- including age in dynamic models may sound straightforward but as we're going to see adds a surprising amount of complexity

Anthropologists have long recognized the importance of social connections and behavioral variation among humans and our nonhuman primate relatives. Indeed, the ability for us to participate in distinct but potentially interlocking complex social networks has fueled our evolution as a species and made our uniquely elaborate life possible. Network analysis has often been utilized as a way to visually and quantitatively represent these ties in order to understand their effects on those connected to each other, from kinship, social support and social capital, to the diffusion of information and transmission of disease. These latter networks are crucially important to our understanding of how human biosocial variation influences our health, where the oft-beneficial complex social networks we maintain and navigate every

day can also put us at risk of exposure to infection.

This work is guided by the theoretical framework of the human ecology of infectious disease, the investigation of how human behavior, social patterns, and built environments interact with the broader pathogen environment to influence our health. Of particular interest is not just aggregate behavior, but also how variation in individual behavior influences social patterns and alters the landscape through which diseases spread. In order to understand the complex patterns by which sexually transmitted infections (STIs) are transmitted throughout populations, we first need to understand the behavior of human relationships and how these behaviors generate the dynamic sexual network across which these types of infections can spread. Syndemic theory will also be used as a guide to understand how variation in behaviors and patterns act synergistically to increase vulnerability and exacerbate existing health disparities of certain population subgroups (Singer et al. 2006).(Singer et al., 2006)

from pinky textbook on stochastic modeling:

“Stochastic processes are ways of quantifying the dynamic relationships of sequences of random events. Stochastic models play an important role in elucidating many areas of the natural and engineering sciences. They can be used to analyze the variability inherent in biological and medical processes, to deal with uncertainties affecting managerial decisions and with the complexities of psychological and social interactions, and to provide new perspectives, methodology, models, and intuition to aid in other mathematical and statistical studies.”

1

Survival Analysis of Relationship Duration

- main goal here to understand where in the distribution we may not be capturing when we use models based on a memoryless process, and explore some ways to do better within the constraints of feasibility imposed by epidemic models
- so we primarily use the exponential distribution in the SA analyses
- discuss survey data and censoring / truncation issues
- And then flesh out the details of the analyses with a clear narrative and you have the core of a chapter!

2

Population Aging and Network Statistics

needs better title

3

Chlamydia, Acquired Immunity, & Expedited Partner Therapy?

some text from diss proposal:

C. trachomatis is an obligate intracellular bacterium transmitted through sexual contact among humans. Chlamydial infections are most often asymptomatic. Untreated infections in women are an additional public health concern because they can lead to a variety of sequelae including pelvic inflammatory disease, scarring of ovaries and fallopian tubes, ectopic pregnancies, chronic pain, and infertility. Repeat infections are common and are an additional risk factor for the development of the above sequelae (Brunham and Rey-Ladino 2005). There is a great deal of uncertainty regarding the natural history of chlamydia, but the

duration of infection for untreated individuals is generally thought to be up to 6 months for men and a year or more for women (Golden et al. 2000; Satterwhite et al. 2013). Chlamydia is usually treated with azithromycin or doxycycline, and unlike other common STIs like syphilis and gonorrhea, true antibiotic resistance is rare (Kong et al. 2015).

Chlamydia is the most common reportable disease in the United States and incidence, particularly adolescents and young adults aged 15-29, is increasing nationwide. The Centers for Disease Control and Prevention (CDC) estimates that half of all new STI infections (including gonorrhea, syphilis, and others) occur in those aged 15-24 despite them making up only a quarter of the sexually active population. Even in places like King County, Washington, where overall rates have remained stable, longstanding acknowledged disparities in prevalence by race are marked and continue to increase (2015 SKCPH STD Report). These rates are particularly distressing in light of the fertility consequences of long-term infection and reinfection: it is estimated that in King County, over 60% of non-Hispanic Black women have had at least one chlamydia infection by age 34 (a rate 5x higher than non-Hispanic White women) and 1 in 500 of non-Hispanic Black women develops chlamydia-associated tubal factor infertility over their life-course (Chambers et al. 2018).

The United States has some of the highest STD rates in the industrialized world, and despite this, funding for public health programs dedicated to these issues has largely declined (CDC 2016 STD Report). As a result, few health departments are able to offer traditional partner notification services, where a patient who tests positive for an STI gives the contact information of their recent sex partners to the health department, and the department then contacts their partners with the hope that these partners will then get tested and, if necessary, treated. Expedited partner therapy (EPT) was developed with this scenario in mind (See figure 2). Under EPT, a patient who tests positive, upon receipt of their own treatment, receives either additional antibiotic pills for their recent sexual partners or prescriptions for treatment that their partners can fill. The patient then is expected to hand-deliver either the treatment or prescription to their partner(s), who take the medicine at their own discretion and without the need for a positive lab test. By using these actors to essentially leverage their sexual network in reverse, this system hopes to decrease the time to treatment for all possible infected partners and increase the total number of partners treated.

It can also reduce re-infection among the index patients if the partnerships are ongoing. There have been several clinical trials of EPT across the US (and Europe), including Washington State. These trials demonstrated that relative to traditional referral practices, EPT provision increased the proportion of partners who were ultimately treated, reduced the number of individuals who were re-infected at follow-up, and was less costly if at least 30% of partners were treated via EPT (CITE). Despite these results and a growing body of evidence in support, widespread implementation of EPT has been slow and there are still many questions to be answered.

Arrested Immunity Hypothesis One of the paradoxes in era of modern public health is that chlamydia incidence has actually increased overall in the presence of mass control programs. In Sweden, Norway, Finland and Canada the rates initially decreased but then resumed increasing, and in Australia, United States, and the United Kingdom the rates never stopped increasing even after program initiation, although this second pattern has been attributed to the challenges of implementing control programs consistently throughout a large population (Brunham and Rekart 2008). These areas now experience incidence rates higher than rates prior to introduction of control programs. Additionally, a regression analysis using data from family planning clinics in Region X of the United States (Alaska, Washington, Idaho, and Oregon) found that, after controlling for any changes in demographics, sexual behaviors, and increased sensitivity of clinical tests, there was a remaining 5% ‘true’ and unexplained annual increase in chlamydia positivity from 1997-2004 (Fine et al. 2008). In response to these and other examples of unabated chlamydia infection in the presence of control programs, Brunham and Rekart have proposed the arrested immunity hypothesis (Brunham and Rekart 2008). Under this hypothesis, early detection and treatment of chlamydia interrupts the development of acquired immunity, making treated individuals particularly vulnerable to reinfection almost immediately after treatment. While we have no natural history studies of chlamydia infection in humans that address the development, duration, and extent of immunity, there is growing evidence beyond rodent models and trends in incidence that partial immunity can develop and play a role. Rodent models of chlamydial infection suggest that a high proportion are able to resolve their primary infection and are temporarily resistant to infection. Rodents that then eventually become reinfected with chlamydia have a shorter duration of disease, lower pathogen load and decreased inflammatory response

(Rank et al. 2003). However, it has also been shown that treatment early in the course of infection interrupts the development of this protective immunity (Su et al. 2002). There is also some indirect evidence in humans. A 2010 review article acknowledged that in several studies of infection status among couples, the rates of discordance (i.e. one partner is infected while the other is not), are higher for chlamydia than for gonorrhea and that this discordance increases with age, providing indirect evidence for some level of protective immunity to chlamydia that increases with age, likely due to exposure over time. There is little immunity that develops to gonorrheal infection due to high levels of antigenic variation (Batteiger et al. 2010). Recent modeling using data from both the UK and United States has demonstrated that at least some immunity to chlamydia following natural clearance is necessary to generate observed patterns in incidence (Omori, Chemaitelly, and Abu-Raddad 2019). These questions are particularly relevant in the context of expedited partner therapy, where the goal is to interrupt transmission by treated individuals and their partners as quickly as possible. However, due to the arrested immunity of those treated quickly, if the timing of delivery and uptake of partners is not sufficient, the initially treated is likely at higher risk of reinfection than under the standard referral scenario. If sufficient numbers of partners are treated effectively and quickly and transmission throughout the network is greatly diminished, then EPT may be able to overcome the effects of this arrested immunity.

Conclusion

We conclude.



The First Appendix

additional figures?

B

The Second Appendix

more technical stuff in here?

Colophon

This document is set in **EB Garamond**, **Source Code Pro** and **Lato**. The body text is set at `upt` with *EBGaramond(3)*.

It was written in R Markdown and \LaTeX , and rendered into PDF using **huskydown** and **bookdown**.

This document was typeset using the XeTeX typesetting system, and the **University of Washington Thesis class** class created by Jim Fox. Under the hood, the **University of Washington Thesis LaTeX template** is used to ensure that documents conform precisely to submission standards. Other elements of the document formatting source code have been taken from the **Latex, Knitr, and RMarkdown templates for UC Berkeley's graduate thesis**, and **Dissertate: a LaTeX dissertation template to support the production and typesetting of a PhD dissertation at Harvard, Princeton, and NYU**

The source files for this thesis, along with all the data files, have been organised into an R package, `xxx`, which is available at <https://github.com/xxx/xxx>. A hard copy of the thesis can be found in the University of Washington library.

This version of the thesis was generated on 2020-08-10 14:22:53. The repository is currently at this commit:

The computational environment that was used to generate this version is as follows:

```
- Session info -----
setting  value
version  R version 3.6.1 (2019-07-05)
```

```

os      macOS Catalina 10.15.3
system  x86_64, darwin15.6.0
ui      X11
language (EN)
collate en_US.UTF-8
ctype   en_US.UTF-8
tz      America/Los_Angeles
date    2020-08-10

```

- Packages -----

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| backports | 1.1.8 | 2020-06-17 | [1] | CRAN (R 3.6.2) |
| bookdown | 0.20.2 | 2020-08-06 | [1] | Github (rstudio/bookdown@f9cf1ac) |
| callr | 3.4.3 | 2020-03-28 | [1] | CRAN (R 3.6.2) |
| cli | 2.0.2 | 2020-02-28 | [1] | CRAN (R 3.6.0) |
| colorspace | 1.4-1 | 2019-03-18 | [1] | CRAN (R 3.6.0) |
| crayon | 1.3.4 | 2017-09-16 | [1] | CRAN (R 3.6.0) |
| desc | 1.2.0 | 2018-05-01 | [1] | CRAN (R 3.6.0) |
| devtools | * 2.3.1 | 2020-07-21 | [1] | CRAN (R 3.6.2) |
| digest | 0.6.25 | 2020-02-23 | [1] | CRAN (R 3.6.0) |
| dplyr | 1.0.1 | 2020-07-31 | [1] | CRAN (R 3.6.2) |
| ellipsis | 0.3.1 | 2020-05-15 | [1] | CRAN (R 3.6.2) |
| evaluate | 0.14 | 2019-05-28 | [1] | CRAN (R 3.6.0) |
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| knitr | 1.29 | 2020-06-23 | [1] | CRAN | (R 3.6.2) |
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| stringr | 1.4.0 | 2019-02-10 | [1] | CRAN | (R 3.6.0) |

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| xfun | 0.16 | 2020-07-24 | [1] | CRAN | (R 3.6.2) |
| yaml | 2.2.1 | 2020-02-01 | [1] | CRAN | (R 3.6.0) |

[1] /Library/Frameworks/R.framework/Versions/3.6/Resources/library

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