Declaration of Nina H. Fefferman, Ph.D.

1. I am a full professor at the National Institute for Mathematical and Biological Synthesis at the University of Tennessee, Knoxville. I am jointly appointed in both the Department of Mathematics and the Department of Ecology and Evolutionary Biology. I am also an Associate Director of the University of Tennessee One Health Initiative and the Director of the UT Mathematical Modeling Consulting Center. As a professor and academic administrator of research efforts, I have 16 years of experience in building, analyzing, and applying mathematical models of infectious diseases to help design public health policy and advise policy makers. My curriculum vitae is attached as Exhibit A.

2. For 9 years, I worked as a researcher for the United States Department of Homeland Security as part of the Command, Control, and Interoperability Center for Advanced Data Analytics, where my research focused on biosecurity, pandemic preparedness, and complex adaptive systems. I have served as a subject matter expert for the Los Alamos National Labs, the US Environmental Protection Agency, the Department of Defense, and the Centers for Disease Control and Prevention, all in the area of biodefense using mathematical models of outbreaks of infectious diseases. My work has been funded by grants from the National Institutes of Health, the National Science Foundation, the Department of Defense, the US Department of Agriculture, the US Fish and Wildlife Service, and the US Department of Homeland Security. In each of these roles, I have designed, implemented, and analyzed mathematical models and helped translate the insights derived from those models to shape policy to keep populations safe.

3. In 2006-2007, I served as a consultant to the New Jersey Department of Corrections, helping understand, anticipate, and plan mitigation for vulnerabilities of the NJ prison system to infectious disease outbreaks.

4. I have received federal funding for my work on pandemic preparedness and response plans for H1N1 2009, Ebola in 2014, Zika virus in 2016, and now for COVID-19. In each of the previous efforts, the results of my work have translated directly into policies implemented by municipal, state, federal, and international agencies (as appropriate).

The nature of COVID-19

5. The entire world is currently facing an ongoing pandemic of the SARS-nCoV-2 virus. The virus transmits very easily, with each infected person (estimated to the best of our current understanding) going on to infect between 2.4-3.8 other people over the average 10-14 days of infectiousness. This leads to exponential growth in the outbreak, meaning that there is rapid spread among individuals within a single population and simultaneous expansion to new populations.

6. The virus can be transmitted in the absence of symptoms, either before an individual develops signs and symptoms of illness, or even in the case that an individual never progresses to exhibit illness themselves. Clinical testing is the only way to identify who may be infectious. Currently, the United States does not have sufficient access to testing to adequately identify even a majority of current cases, much less the percentage of current active infections that would need to be identified and isolated to contain the spread of disease.

7. COVID-19 causes serious illness, with overall case fatality rates in the United States so far estimated at 5.8%. An estimated 20% of those who become infected and develop

symptoms require significant medical intervention. While certain medical conditions (including, but not limited to hypertension, asthma, COPD, diabetes, and AIDS) increase the probability of death from infection, otherwise perfectly healthy people are also vulnerable to death from COVID-19. Increasing age is a predictor of increased severity of illness and risk of death. Current data show that those between 55 and 64 years of age experience case fatality rates of between 1% and 3%, increasing with age until those between 65 and 84 years of age show case fatality rates between 3% and 11%, and those 85 years old or older show case fatality rates between 10% and 27%.

8. COVID-19 is now the leading cause of death in the United States.

9. Of the 20% of patients who develop severe symptoms in need of medical intervention, 30% are expected to require intensive care in order to survive. Given the infectiousness of the virus, this means that even if a relatively small percentage of a population is currently infected, the capacity of the medical support systems that serve them will be exhausted. When medical resources are exhausted, the mortality rates will increase even further above these already staggeringly high levels, including progression to death for more of the cases not currently considered necessarily fatal.

10. Although clinical trials to try to identify effective medicines and therapeutics to treat current infections, and research and trials in vaccine development to prevent future infections, are currently underway across the globe, as yet, no meaningful medical interventions have been shown to be effective against the virus. Our current best lines of defense are epidemiological rather than medical; interrupting the transmission of the disease rather than helping infected individuals to survive.

11. Individual practices such as the use of personal protective equipment, maintaining physical distance from others (also called "social distancing") and frequent washing (of hands, body, and environmental surroundings) with soaps and disinfectants are the only actions individuals can take that have been shown to be effective at self-protection from catching the virus.

The exponential rate of COVID-19 infection in FCI Fort Dix

12. FCI Fort Dix is already experiencing a serious outbreak of COVID-19, with an infection rate that far exceeds the state of New Jersey as a whole.

13. According to the BOP, there are 30 current, lab-result-confirmed cases of COVID-19 at FCI Fort Dix. This number was reported on April 28, 2020, and is undoubtedly much higher today, given the rate of spread and the failure to test patients who may be asymptomatic yet continuing to spread the virus.

14. With a population of 2,947, FCI Fort Dix therefore has a reported COVID-19 infection prevalence of 1,018 per 100,000. This is drastically higher than the surrounding community.

15. By comparison, the state of New Jersey as a whole reports a COVID-19 infection prevalence of 73 per 100,000.

16. As a result, at a minimum, the reported infection prevalence at Fort Dix is approximately 14 times that for New Jersey as a whole.

The implications of COVID-19 in detention facilities and proposed policies for mitigation

17. Based on my substantive expertise, long history of research, and as practice as a modeler of infectious outbreaks, I respectfully but strongly disagree with the stated premises and conclusions of the Bureau of Prisons recommendations (as most recently updated in BOP'S Home Confinement memorandum, April 22, 2020). These recommendations, by design, fail to reduce the population at FCI Fort Dix sufficiently to prevent the exponential spread of COVID-19, both within the facility and in the surrounding community. My disagreement is based on the following reasoning and logic, arrived at via my expertise in the field.

18. The goal of any action taken must be the increased safety and survival of the population served. In the case of a prison, the population under consideration consists of three separate groups, each at risk: the inmates themselves, the staff who serve at the prison facility in all capacities, and the general public into whose company the staff return after their work days and to into whose company the inmates within the prison would be returned in released from detention. Determination of inmate eligibility for release must serve to balance risks appropriately to best protect all of these groups. At the moment, some of the listed criteria for eligibility for release severely limit the potential population of inmates who may be considered. These limitations so drastically increase the epidemiological risk to inmates, staff, and public that they do not serve the greater goal of increasing the overall safety and survival of the total population.

19. This rationale is based on the understanding that, despite best efforts to increase personal hygiene and social distancing practices, and to reduce inmate movements and suspend access to members of the public (contractors, visitors, and legal professionals), prisons are inherently incapable of reducing the risks of transmission to those seen in the broader community. The needs for oversight over inmate populations by staff, physical limitations on space, housing,

and infrastructure required to maintain the incarcerated population, and even the common practices of employing inmates as laborers throughout the prison facilities in ways that require intermixing among cohorts all contribute to risks that individuals in the broader community do not face while practicing "stay-at-home" protocols.

20. Proposed efforts to screen inmates and staff are insufficient due to lack of clinical testing availability across the nation, meaning that only those who are currently showing symptoms are likely to be able to be tested, leaving room for significant transmission of infection prior to/in the absence of the development of any symptoms of illness.

21. There is already likely to be circulating infection that continues to go undetected due to the current CDC recommendations to test only those who show symptoms of infection. Current estimates, though based on incomplete data, suggest that as many as 40% of cases may be asymptomatic and these cases may still be capable of transmitting infection to others.

22. Incarceration is itself a source of physiological compromise — research has shown that incarcerated individuals have health outcomes that more closely resemble those described for patients 10-15 years older than their physiological age. This means that, even at the same level of effective social distancing and personal hygiene, a prison population becomes a greater reservoir for infection than would the same number of people behaving in the same way in the broader community.

23. While those at greatest risk of death from COVID-19 are medically vulnerable, severe health outcomes (including death) are regularly described in even young and otherwise healthy individuals. Actions that increase the spread of COVID-19 expose anyone infected to non-trivial risks. Staff who interact regularly with inmates must themselves be considered at risk

due to constant interaction with the population they oversee. As they return to the broader community at the end of their workday, those risks return with them.

24. Increasing the spread of COVID-19 also depletes valuable medical resources. Each new case not only incrementally increases the risk of death for COVID-19 patients, but also increases the risk of all-cause mortality in the community as the medical professionals/resources are insufficient to meet the baseline medical needs of the community (i.e. emergency rooms overwhelmed with COVID-19 patients will have increased numbers of deaths from heart attacks due to delays in access to medical attention). By keeping people in prisons and increasing the inmates' own likelihood of requiring medical attention (relative to the same numbers of cases requiring medical attention that would be needed for a population of the same size in the general population), we increase the burden to the medical staff and resources overall. This is true not only for medical facilities inside a prison as the number of seriously ill inmates are likely to exceed the capacity for attention within prison facilities, but also because each additional case caused in staff or staff-vectored community infection will also contribute to the broad burden on medical resources.

25. Epidemiologically, the only way to meaningfully reduce the risks posed to the entire population—inmates, staff, and public—is to drastically reduce the prison population. Due to exponential growth in outbreaks, each preventable infection that we fail to prevent directly impacts (on average) 3 people, who then also each impact 3 people, and so on. Every infection we can prevent saves lives directly and indirectly.

26. For all of the reasons herein, in my expert opinion, the current efforts are epidemiologically insufficient to protect inmates, prison staff, or the general public surrounding the prison. It is my opinion that the current efforts are not sufficiently justified by the rationale of

Protecting the Public (as described in Attorney General's memorandum dated April 3rd, 2020). It is my opinion that the public interest is best served by relaxing the criteria for consideration for release until the point where epidemiological models of within-prison transmission, for populations whose health demographics incorporate the physiological compromise and physical restrictions inherent in incarceration, approach the same levels of risk of infection and transmission that would be seen in the same population were it to be released.

Pursuant to 28 U.S.C. 1746, I declare under penalty of perjury that the foregoing is true and correct.

Executed this 29th day of April, 2020, in Knox County, TN.

Nina H. Fefferman, Ph.D.

References

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2. Severe Outcomes Among Patients with Coronavirus Disease 2019 (COVID-19) — United States, February 12–March 16, 2020. MMWR Morb Mortal Wkly Rep 2020;69:343-346. DOI: http://dx.doi.org/10.15585/mmwr.mm6912e2

3. US Attorney General, and United States of America. "The First Step Act of 2018: Risk and Needs Assessment System." (2019).
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Nina H. Fefferman

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Education

2005	PhD in Mathematical Biology from the Department of Biology, Tufts University. Advisor: J. Michael Reed
2001	MS in Mathematics from the Department of Mathematics, Rutgers University. Advisor: J. Beck
1999	AB in Mathematics from Princeton University

Positions

2020-	Associate Director, UT One Health Initiative, University of Tennessee, Knoxville
2018-	Director, Mathematical Modeling Consulting Center, University of Tennessee, Knoxville
2018 -	Professor, Depts. of Mathematics & Ecology and Evolutionary Biology, University of Tennessee, Knoxville
2016 - 2018	Associate Professor, Depts. of Mathematics & Ecology and Evolutionary Biology, University of Tennessee, Knoxville
2015 - 2016	Program Director, Graduate Program in Ecology and Evolution, Rutgers University
2012 - 2016	Associate Professor, Dept. of Ecology, Evolution, and Natural Resources, Rutgers University
2011 - 2016	Assistant/Associate Professor, School of Public Health, University of Medicine and Dentistry of New Jersey
2008 - 2012	Assistant Professor, Dept. of Ecology, Evolution, and Natural Resources, Rutgers University
2007 - 2016	Research Assistant/Associate Professor, The Center for Discrete Mathematics and Theoretical Computer Science, Rutgers University
2005 - present	Co-Director, Tufts University Initiative for the Forecasting and Modeling of Infectious Disease (InForMID), Tufts University School of Medicine
2005 - 2007	Visiting Research Associate, Center for Discrete Math and Theoretical Computer Science (DIMACS), Rutgers University
2005	Short Term Visitor, School of Natural Sciences, Institute for Advanced Study

Honors/Awards

2019	Invited Participant of the 11th.Triennial Invitational Choice Symposium
2019	Invited Performer/Participant, Stand Up Science – a public performance featuring stand-up comics and scientists discussing their work
2017	Invited Research Team Leader: AWM Women in Mathematical Biology Workshop
2016	Invited Speaker at the National Academy of Sciences Sackler Colloquium
2015	Coauthored an article chosen for the cover of *Phil Trans Roy Soc B* (issue 370.1665)
2012	Invited to Health Foo 2012

2011 Shared the Virginia Governor's Technology Award in the category of 'Cross-Boundary Collaboration in Modeling & Simulation' for our study 'Strategic Default in the Context of a Social Network: An Epidemiological Approach'.
2010 Speaker at TEDx Midatlantic
2009 Rutgers University Packard Fellow Nominee
2007 Coauthored an article chosen for the cover of *The Lancet Infectious Diseases* (vol. 7)
- Invited to give 22 Keynote, Plenary, or Public Lectures (see Invited Talks for details), over three continents

Media Coverage (interviews and coverage):

Television/Online Video Broadcasts:

WBIR News, 2019
NJTV News, 2015
Discovery Channel "How Stuff Works" (Season 2: "Games Unboxed"), 2011
BBC World News Aug 21, 2007
CBS News Aug 22, 2007
Canada Television (CTV) Aug 21, 2007
AT&T Tech Channel Sept, 2007

Radio Broadcasts:

NPR Marketplace, Mar 2020
NPR WUOT Knoxville, Mar 2017
PRI Studio 360, Sept 2016
New Tech City, WNYC, Oct 2014
PRI Studio 360, Sept 2014
PRI Studio 360, Jan 2013
BBC UK News, Aug 2007
National Public Radio Podcast "Science Friday", Sept 2007
AM900 CHML, Sept 2007
National Public Radio "All Things Considered", Oct 2005

Print/Online Media (2005-present):

ABC News, ABS CBN News, ARS Technical, Canadian Press (via CBC), Cell, The Daily Mail (UK), The Daily Telegraph (Australia), The Economist, Forbes, Fox News, G1.com.br (Brazil), O Globo (Brazil), Gazet Van Antwerpen (Belgium), KevinMD, Knox News, NU.nl (Netherlands), Medical News Today, La Jornada (Mexico), New Scientist, PC Gamer, Reuters, TIME, The Washington Post, Science News, Slate.com, the South African Star, Tech News World, Wired, Yahoo! Entertainment *and many more*…

Research Support

Active

2020-2021	$198,932	NSF RAPID – DEB Coupled Social and Epidemiological Networks and COVID-19	PI
2020-2022	$359,849	DoD Minerva DECUR - The Topology of Interdependent Multi-Domain Behavioral Systems	PI
2017-2022	$138,964	NSF IOS - Melding Mathematical and Theoretical Models of Stress	UT-PI
2017-2021	$2,498,876	NSF EEID – Co-evolutionary Epidemiology of Avian	UT-PI

		Malaria	
Completed			
2018-2020	$196,628	SESYNC/NIMBioS Modeling Risk Perception, Vector-borne Diseases, and Environmental Integrity	PI
2016-2019	$99,938	NSF EAGER – CISE – Distributed Anomaly Detection	PI
2018-2019	$2,000	Haines Morris Grant – Internal UTK Competition	Co-PI
2016-2018	$50,000	US - Israel Binational Science Foundation (BSF)	Co-PI
2016-2018	$190,000	NSF RAPID – DEB – Modeling Zika Virus Control	PI
2015-2018	$292,804	USFWS – White-Nose Syndrome Open Grant	Co-PI
2015-2017	$21,003	NSF RAPID – Information & Intelligent Systems – Virtual Worlds and Experiential Learning	PI
2016-2017	$75,000	US START Center – Leadership in Social Networks	PI
2017	$30,000	Syngenta – Workshop Grant – Math of Agribusiness	Co-I
2016-2017	$100,000	National Academies Keck Futures Initiative	Co-PI
2015-2017	$130,000	NSF EAGER – DEB – Machine Learning for Co-Evolutionary Systems	Co-PI
2012-2016	$1,228,053	Dept. of Homeland Security – CyberSecurity	PI
2014-2016	$100,000	Dept. of Homeland Security – Next Generation Communications and Interoperability	Project PI
2009-2016	$275,000	Dept. of Homeland Security – BioSecurity	Project PI
2011-2014	$3,853,332	NSF EASM – Ocean Sciences – SocioEconomic Systems and Climate Change	Co-PI
2011-2012	$22,500	UCDPER – Emergency Preparedness	Co-PI
2010-2012	$384,000	Dept. of Homeland Security – Virtual Worlds and Experiential Education	Project PI
2010-2011	$99,944	Dept. of Homeland Security – Self-Organizing Surveillance Systems	Project PI
2010	$22,500	Dept. of Homeland Security – BioSecurity	Co-PI
2009-2012	$299,886	NSF – DEB – ULTRA-Ex	Co-PI
2009-2011	$89,318	UCDPER – Emergency Preparedness	PI
2009-2010	$10,000	USDA CSREES Multi-State Research Fund – Vector-borne Disease Control	Co-I
2008	$99,990	NIH NAID SBIR – Epidemiological Surveillance	PI
2008	$5,000	Rutgers Climate and Environmental Change Initiative	PI
2008	$75,000	Rutgers Academic Excellence Fellowship, Climate and Health Research Initiative	Co-I
2007	$22,500	Dept. of Homeland Security – BioSecurity	PI
2007	$22,500	Dept. of Homeland Security – BioSecurity	PI
2006	$5,000	Tufts Summer Scholars Award – Epidemiology	PI
2003-2004	$42,000	NIH R01 Supplement - Epidemiology	Co-PI
2003-2004	$1,500	Tufts Institute of the Environment	Co-I
2003	$500	MASI Student Travel Award	PI
2003	$1,500	TIES Student Travel Award	PI

Consultancies

2020	American Civil Liberties Union (ACLU)
2020	The State of Vermont, Department of Education
2018	Ogilvy

2017-present	Humane Society International
2009-present	US Centers for Disease Control
2011-2012	Research Institute for Housing America Trust Fund
2006-2007	New Jersey, Department of Corrections
2004-2009	NIH U19 (Center PI: Gorski) T-cell Mediated Immunity
2004	National Defense University
2004	DARPA

Participation in Research Centers

Center	**Position**	**Description of Role**
NIMBioS *(National Institute for Mathematical and Biological Synthesis)*	Leadership Team	Active participant in working group, organizer of multiple tutorials, mentor for summer research experience for undergraduates, and founding director of the Mathematical Modeling Consulting Center
InForMID *(Tufts University Initiative for the Forecasting and Modeling of Infectious Diseases)*	Center Co-Director	Researcher and Administrative lead in the area of mathematical modeling of infectious disease epidemiology
CCICADA *(US Dept of Homeland Security Command, Control, and Interoperability Center for Advanced Data Analysis)*	Project PI	Principle Investigator into data analysis relating to social behavior in virtual/technologically enable environments, bio-security, and bio-inspired algorithms in cyber-security
DIMACS *(The Center for Discrete Mathematics and Theoretical Computer Science)*	Member	Active participant in working groups, collaborations, and conferences (including acting as organizer for multiple workshops/conferences/tutorials) in all areas of mathematical macrobiology
START *(US Dept of Homeland Security Center for the Study of Terrorism and Responses to Terrorism)*	Project PI	Principle Investigator working on understanding social behavior and algorithms driving the emergence of extremism and leadership in

Publications (peer reviewed):

***** = *a student or post-doctoral researcher advised by Fefferman during the research effort reported*

Journal Articles:

Published or In Press

68. Lemanski*, N., S. Schwab, D. Fonseca, and N.H. **Fefferman**. (In press) Coordination Among Neighbors Improves the Efficacy of the Zika Control Despite Economic Costs. *PLoS Neglected Tropical Diseases.*
67. Wilson, S., S. Sindi, H. Brooks, M. Hohn, C. Price, A. Radunskaya, N. Williams, and N.H. **Fefferman**. 2020. How Emergent Social Patterns in Allogrooming Combat Parasitic Infections. *Frontiers in Ecology and Evolution*. 8:54.
66. DeNegre*, A., Myers*, K., and N.H. **Fefferman**. 2020. Impact of Strain Competition on Bacterial Resistance in Immunocompromised Populations. *Antibiotics*. 9(3):114
65. Myers*, K., A. Redere*, and N.H. **Fefferman**. 2020. How Resource Limitations and Household Economics May Compromise Efforts to Safeguard Children During Outbreaks. *BMC Public Health.* 20(1):1-14.
64. Suarez*, G., O. Udiani*, B. Allan, C. Price, S. Ryan, E. Lofgren, A. Coman, C. Stone*, L. Gallos*, and N.H. **Fefferman**. 2020. A Generic Arboviral Model Framework for Exploring Trade-offs Between Vector Control and Environmental Concern. *Journal of Theoretical Biology*. 490 (2020) 110161.
63. DeNegre*, A., Myers*, K., and N.H. **Fefferman**. 2020. Impact of Chemoprophylaxis Policy for AIDS-immunocompromised Patients on Emergence of Bacterial Resistance. *PLoS One*. 15(1): e0225861.
62. Gallos*, L., S. Havlin, G. Stanley, and N.H. **Fefferman**. 2019. Propinquity drives the emergence of network structure and density. *Proceedings of the National Academy of Sciences*. 116(41):20360-20365.
61. Stone*, C., S. Schwab*, D. Fonseca, and N.H. **Fefferman**. 2019. Contrasting the Value of Targeted vs. Area-Wide Mosquito Control Scenarios to Limit Arbovirus Transmission for Different Tropical Urban Population Centers. *PLoS Neglected Tropical Diseases.* 13.7: e0007479.
60. Myers*, K, A. DeNegre*, L.K. Gallos*, N. Lemanski*, A. Mayberry, A. Redere*, S. Schwab*, O. Stringham, & N.H. **Fefferman**. 2019. Dynamic Ad Hoc Social Networks in Improvised Intelligence / Counter-Intelligence Exercises: A Department of Homeland Security Red-Team Blue-Team Live-Action Roleplay. *Journal of Homeland Security and Emergency Management.* https://doi.org/10.1515/jhsem-2018-0027.
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57. Schwab*, S., C. Stone*, D. Fonseca, and N.H. **Fefferman**. 2019. (Meta)population Dynamics Determine Effective Spatial Distributions of Mosquito-Borne Disease Control. *Ecological Applications* 29(3): e01856.
56. Kebir*, A., N.H. **Fefferman**, and S.B. Miled. 2018. A general structured model of a hermaphrodite population. *Journal of Theoretical Biology*. 449:53-59.
55. Lemanski*, N.J. and N.H. **Fefferman**. 2018. Expanding the evolutionary theory of aging: honeybees as a test case for an optimal decision making model of senescence. *American Naturalist*. 191(6):756-766.
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51. Stone*, C.M., S.R. Schwab*, D.M. Fonseca, N.H. **Fefferman**. 2017. Human movement, cooperation, and the effectiveness of coordinated vector control strategies. *Journal of the Royal Society Interface.* 14(133):20170336.

50. Lemanski*, N.J. and N.H. **Fefferman**. 2017. Coordination Between the Sexes Constrains the Optimization of Reproductive Timing in Honey Bee Colonies *Nature Scientific Reports.* 7:2740.

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48. Greenbaum*, G. and N.H. **Fefferman**. 2017. Application of network methods for understanding evolutionary dynamics in discrete habitat. *Molecular Ecology*. DOI: 10.1111/mec.14059

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44. Maslo, B., S. Gignoux-Wolfsohn, and N.H. **Fefferman**. 2017. Success of Wildlife Disease Treatment Depends on Host Immune Response. *Frontiers in Ecology and Evolution*. 5(28).

43. Lofgren*, E., A. Egizi, and N.H. **Fefferman**. 2016. Patients as Patches: Ecology and Epidemiology in Healthcare Environments. *Infection Control and Hospital Epidemiology.* 37(12):1507-1512.

42. Korczynski*, M., A. Hamieh*, J. H. Huh, H. Holm, S. R. Rajagopalan, and N. H. **Fefferman**. 2016. Hive Oversight for Network Intrusion Early Warning Using DIAMoND: A Bee-Inspired Method for Fully Distributed Cyber Defense. *IEEE Communications Magazine* 54(6):60-67.

41. Gallos*, L. and N.H. **Fefferman.** 2015. Simple and efficient self-healing strategy for damaged complex networks. *Physical Reviews E*. 92(5):052806.

40. Kebir*, A., N.H. **Fefferman**, S. Ben Miled. 2015. Understanding hermaphrodite species through game theory. *Journal of Mathematical Biology*. 71(6-7):1505-1524.

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37. Burkhalter*, J.C., N.H. **Fefferman**, and J.L. Lockwood. 2015. The impact of personality on the success of prospecting behavior in changing landscapes. *Current Zoology*. 61:557-568.

36. Robinson*, O., J. Lockwood, O. Stringham*, and N.H. **Fefferman**. 2015. A Novel Tool for Making Policy Recommendations Based on PVA:Helping Theory Become Practice. *Conservation Letters*. 8(3):190-198.

35. **Fefferman**, N.H. and E.N. Naumova. 2015. Dangers of vaccine refusal near the herd immunity threshold: a modelling study. *Lancet Infectious Diseases*. S1473-3099(15)70130-1

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33. Parham, P E. J. Waldock, G.K. Christophides, D. Hemming, F. Agusto, K. J. Evans, N.H. **Fefferman**, H. Gaff, A. Gumel, S. LaDeau, S. Lenhart, R.E. Mickens, E. Naumova, R. Ostfeld, P. Ready, M. Thomas, J. Velasco-Hernandez, E. Michael. 2015. Climate, Environmental, and Socioeconomic Change – Weighing up the Balance in Vector-Borne Disease Transmission. *Philosophical Transactions of the Royal Society B.* 370.1665 (2015): 20130551.

32. Egizi, A., N.H. **Fefferman**, and D. M. Fonseca. 2015. Evidence that implicit assumptions of "no evolution" of disease vectors in changing environments can be violated on a rapid timescale. *Philosophical Transactions of the Royal Society B.* 370.1665 (2015): 20140136.

31. Greening*, B., N. Pinter-Wollman, and N.H. **Fefferman**. 2015. Higher-Order Analysis of Information Sharing and Knowledge Capacity in Animal Social Groups *Current Zoology*. 61(1): 114–127.

30. Gallos*, L. and N.H. **Fefferman.** 2014. Revealing effective classifiers through network comparison. *Europhysics Letters*. 108(3): 38001.

29. Lofgren*, E.T., R.W. Moehring, D.J. Anderson, D.J. Weber, and N.H. **Fefferman**. 2014. A Mathematical Model to Evaluate the Routine Use of Fecal Microbiota Transplantation to Prevent Incident and Recurrent *Clostridium difficile* Infection. *Infection Control and Hospital Epidemiology*. 35(1):18-27.

28. Greening*, B. and N.H. **Fefferman**. 2014. Evolutionary Significance of the Role of Family Units in a Broader Social System. *Nature Scientific Reports*. 4: 3608

27. Seiler, M.J., Collins, A.J., and N.H. **Fefferman**. 2013. Strategic Mortgage Default in the Context of a Social Network: An Epidemiological Approach. *Journal of Real Estate Research* 35(4).

26. Robinson*, O.J., N.H. **Fefferman**, and J.L. Lockwood. 2013. How to effectively manage invasive predators to protect their native prey. *Biological Conservation* 165: 146-153.

25. **Fefferman,** N.H., and L.M. Romero. 2013. Can physiological stress alter population persistence? A model with conservation implications. *Conservation Physiology*. 1(1): cot012. doi: 10.1093/conphys/cot012

24. Moorthy, M., D. Castronovo, A. Abraham, S. Bhattacharyya, S. Gradus, J. Gorski, Y.N. Naumov, N.H. **Fefferman**, and E.N. Naumova. 2012. Deviations in influenza seasonality: odd coincidence or obscure consequence? *Clinical Microbiology and Infection*. 18(10):955-962.

23. Hock*, K. and N.H. **Fefferman**. 2012. Social organization patterns can lower disease risk without associated disease avoidance or immunity. *Ecological Complexity*. 12:34–42.

22. Hock*, K. and N.H. **Fefferman**. 2011. Violating Social Norms when Choosing Friends: How Rule-Breakers Affect Social Networks. *PLoS One.* 2011; 6(10): e26652

21. Hock*, K. and N.H. **Fefferman**. 2011. Extending the role of social networks to study social organization and interaction structure of animal groups. *Annales Zoologici Fennici*. 48(6):365-370.

20. Kafai, Y.B. and N.H. **Fefferman**. 2010. Virtual Epidemics as Learning Laboratories in Virtual Worlds. *Journal of Virtual Worlds Research.* 3(2):2-15.

19. Hock*, K., K.L. Ng, and N.H. **Fefferman**. 2010. Systems approach to studying animal sociality: individual position versus group organization in dynamic social network models. *PLoS One.* 5(12): e15789.

18. **Fefferman,** N.H. and E.N. Naumova. 2010. Innovation in Observation: A Vision for Early Outbreak Detection. *Emerging Health Threats.* 3:e6. doi: 10.3134/ehtj.10.006

17. Lofgren*, E.T., J.B. Wenger, N.H. **Fefferman**, D. Bina, S Gradus, S. Bhattacharyya, Y.N. Naumov, J. Gorski, E.N. Naumova. 2010. Disproportional Effects in Populations of Concern for Pandemic Influenza: Insights from Seasonal Epidemics in Wisconsin, 1967-2004. *Influenza and Other Respiratory Diseases.* 4:205-212.

16. Phan, L., N.H. **Fefferman,** D. Hui, and D. Brugge. 2010. Impact of Street Crime on Boston Chinatown. *Local Environment*. 15(5):481-491.

15. Reed, J.M., N.H. **Fefferman**, and R.C. Averil-Murray. 2009. Vital Rate Sensitivity Analysis and Management Implications for Desert Tortoise. *Biological Conservation*. 14(12): 2813-3222.

14. Wilson-Rich, N., Spivak, M., **Fefferman**, N.H., Starks, P.T. 2009. Genetic, Individual, and Group Facilitation of Disease Resistance in Insect Societies. *Annual Reviews of Entomol*ogy. 54:405-23.

13. **Fefferman**. N.H. 2008. Biological Experimentation *in silico*. *Annales Zoologici Fennici*, 45: 367-368.

12. Lofgren*, E., M. Senese*, J. Rogers* and N.H. **Fefferman**. 2008. Pandemic Preparedness Strategies for School Systems: Is Closure Really the Only Way? *Annales Zoologici Fennici*, 45: 449-458.

11. **Fefferman**, N.H. and K.L. Ng*. 2007. How Disease Models on Static Graphs Fail to Approximate Epidemics in Shifting Social Networks. *Physical Review E*. 76:031919. (*This article was selected for reprinting by the Virtual Journal of Biological Physics Research 2007)*

10. Lofgren*, E. and N.H. **Fefferman**. 2007. The Untapped Potential of Virtual Game Worlds to Shed Light on Real World Epidemics. *The Lancet Infectious Diseases.* 7:625–629. *(article content was the cover of the journal)*

9. Lofgren*, E., N.H. **Fefferman**, Y.N. Naumov, J. Gorski and E.N. Naumova. 2007. Influenza Seasonality: Underlying Causes and Modeling Theories. *Journal of Virology*, 81(11):5429-5436.

8. Lofgren*, E., N.H. **Fefferman**, M. Doshi and E.N. Naumova. 2007. Assessing Seasonal Variation in Multisource Surveillance Data: Annual Harmonic Regression. *Lecture Notes in Computer Science.* BioSurveillance 2007. eds D. Zeng et al. 4506:114-123.

7. **Fefferman**, N.H. and K.L Ng*. 2007. The role of individual choice in the evolution of social complexity. *Annales Zoologici Fennici*, 44:58-69.

6. **Fefferman**, N.H., J.F.A. Traniello, R.B. Rosengaus and D.V. Calleri. 2007. Disease Prevention and Resistance in Social Insects: Modeling the Survival Consequences of Immunity, Hygienic Behavior and Colony Organization. *Behavioral Ecology and Sociobiology*, 61:565-577.

5. Starks, P.T.B. and N.H. **Fefferman**. 2006. Polistes Nest Founding Behavior: a Model for the Selective Maintenance of Alternative Behavioral Phenotypes. *Annales Zoologici Fennici*, 43:456-467.

4. **Fefferman**, N.H., and E.N. Naumova. 2006. Combinatorial Decomposition of an Outbreak Signature. *Mathematical Biosciences*, 202(2):269-287.

3. **Fefferman**, N.H. and J.M. Reed. 2006. A Vital Rate Sensitivity Analysis that is Valid for Non-Stable Age Distributions and for Short-Term Planning. *The Journal of Wildlife Management*, 70(3):649-656.

2. **Fefferman**, N.H., and P.T.B. Starks. 2006. A Modeling Approach to Swarming in Honey Bees. *Insectes Sociaux*, 53(1):37-45.

1. **Fefferman**, N.H., E.A. O′Neil, and E.N. Naumova. 2005. Confidentiality vs Confidence: The aggravation of aggregation as a remedy in public health. *Journal of Public Health Policy*, 26(4):430-449.

Under Review:

9. Chastain*, E. and N.H. **Fefferman**. The Evolution of Personality. (Under Review after Revision)
8. Feinberg, F., A. Patania, B. McShane, B. Falk, D. Larremore, E. Feit, J. Helveston, M. Small, M. Braun, N. **Fefferman**, and E. Bruch. A Framework for Studying Choices in Networks. (Under Review)
7. Beckage, B., K. Lacasse, J.M. Winter, N.H. **Fefferman**, F.M. Hoffman, L.J. Gross, S.S. Metcalf, T. Franck, E. Carr, A. Zia, and A. Kinzig. The Earth has humans, so why don't our climate models? (Under Review)
6. Udiani*, O., K. Lacasse, A. Zia, L. Gallos*, P. Zhong*, B. Beckage, E. Carr, T. Franck, L. Gross, F. Hoffman, P. Howe, A. Kinzig, S. Metcalf, J. Winter, and N.H. **Fefferman**. Recruitment and Mobilization for Social Movements: implications from network modeling. (Under Review)
5. Udiani*, O., and N.H. **Fefferman**. Could the Need for Rest Provide a Pathway for the Evolution of Division of Labor in Social Species? (Under Review)
4. Gignoux-Wolfsohn, S.A., Pinsky, M.L., Kerwin, K., Herzog, C., Hall, M., Bennett, A.B., **Fefferman**, N.H. and Maslo, B., Genomic signatures of evolutionary rescue in bats surviving white-nose syndrome. (Under Review)
3. Myers*, K., N.H. **Fefferman**, and J.M. Reed. Do Not Reject a Population Viability Analysis by Case Study: Observing an Unlikely Event Does Not Invalidate a Qualitative Model (Under Review)
2. Udiani*, O. and N.H. **Fefferman**. Has disease risk shaped the evolution of social complexity in insect societies? (Under Review)
1. Siewe*, N., B. Greening*, and N.H. **Fefferman**. The Potential Role of Asymptomatic Infection in Outbreaks of Emerging Pathogens (Under Review)

Book Chapters:

Published or In Press

10. **Fefferman**, N.H. When to Turn to Nature-Inspired Solutions for Cyber Systems. 2019. *in* Nature-Inspired Security and Resilience. eds. Eltoweissy, Elalfy, Fulp, and Mazurczyk. pp 29-50. The Institution of Engineering and Technology, London, UK.
9. Price, C.R. and N.H. **Fefferman**. 2019. A Preliminary Exploration of the Professional Support Networks the EDGE Program Creates. *in* A Celebration of the EDGE Program's Impact on the Mathematics Community and Beyond (pp. 317-325). Springer, Cham.
8. Brooks. H.Z., M.E. Hohn, C. Price, A.E. Radunskaya, S.S. Sindi, N.D. Williams, S.N. Wilson, N.H. **Fefferman**. 2018. Mathematical Analysis of the Impact of Social Structure on Ectoparasite Load in Allogrooming Populations. *in* Understanding Complex Biological Systems with Mathematics eds. A. Radunskaya, R. Segal, B. Shtylla. Association for Women in Mathematics Series, vol 14. pp 47-61. Springer
7. Williams, N.D., H.Z. Brooks, M.E. Hohn, C. R. Price, A.E. Radunskaya, S.S. Sindi, S.N. Wilson, and N. H. **Fefferman**. 2018. How Disease Risks Can Impact the Evolution of Social Behaviors and Emergent Population Organization. *in* Understanding Complex Biological Systems with

Mathematics eds. A. Radunskaya, R. Segal, B. Shtylla. Association for Women in Mathematics Series, vol 14. pp 31-46. Springer

6. Korczynski*, M., A. Hamieh*, J.H. Huh, H. Holm, S. R. Rajagopalan, and N.H. **Fefferman**. 2017. DIAMoND: Distributed Intrusion/Anomaly Monitoring for Nonparametric Detection (invited extended version). *in* Security, Privacy and Reliability in Computer Communications and Networks. eds. K. Sha, A Striegel, and M Song. River Publishers Series in Communications. River Publishers.
5. **Fefferman**, N.H. and L.M. Fefferman. 2011. Mathematical Macrobiology: An Unexploited Opportunity in High School Education. *in* Biomath in the Schools. eds. M.B. Cozzens, and F.S. Roberts. DIMACS Series in Discrete Mathematics and Theoretical Computer Science. Vol 76. American Mathematical Society.
4. Jagai, J., N.H. **Fefferman** and E.N. Naumova. 2011. Waterborne Disease Surveillance. *in* Encyclopedia of Environmental Health. eds. J. Nriagu, S. Kcew, T. Kawamoto, J. Patz, and D. Rennie. Elsevier Science. 1st edition
3. Ji, S., W.A. Chaovalitwongse, N.H. **Fefferman**, W. Yoo, and J.E. Perez-Ortin. 2009. Mechanism-based Clustering of Genome-wide RNA Levels: Roles of Transcription and Transcript-Degradation Rates. *in* Clustering Challenges in Biological Networks. eds. S. Butenko, P.M. Pardalos, and W.A. Chaovalitwongse. World Scientific Publishing Company.
2. **Fefferman**, N.H. and J.F.A. Traniello. 2008. Social Insects as Models in Epidemiology: Establishing the Foundation for an Interdisciplinary Approach to Disease and Sociality. *in* Organization of Insect Societies: From Genome to Sociocomplexity eds J. Gadau and J. Fewell. Harvard University Press
1. MacLeod, N., N. Ortiz, N.H. **Fefferman**, W. Clyde, C. Schulter, and J. MacLean. 2000. Phenotypic Response of Foraminifera to episodes of global environmental change. *in* Biotic Response to Global Change. eds S.J. Culver and P. Rawson. Cambridge University Press

Edited Volumes:

1. **Fefferman**, N.H. (Ed.) (2008) *Annales Zoologici Fennici* 45(5)

Peer Reviewed Contributed Conference Papers:

8. Suarez*, G.P., L.K. Gallos, and N.H. **Fefferman**. 2018. A Case Study in Tailoring a Bio-Inspired Cyber-Security Algorithm: designing anomaly detection for multilayer networks. *2018 IEEE Security and Privacy Workshops (SPW)*. IEEE, 2018.
7. Fields, D. A., Kafai, Y. B., Giang, M. T., **Fefferman**, N., & Wong, J. 2017. Plagues and people: Mass community participation in a virtual epidemic within a tween online world. *Proceedings of the 12th International Conference on the Foundations of Digital Games.* DOI: 10.1145/3102071.3102108
6. Kafai, Y. B., Fields, D. A., Giang, M. T., **Fefferman**, N., Sun, J., Kunka, D., & Wong, J. 2017. Designing for massive engagement in a tween community: Participation, prevention, and philanthropy in a virtual epidemic. In *Interaction Design & Children Conference*. New York: ACM, 365-370. ISBN: 978-1-4503-4921-5
5. Fields, D. A., Kafai, Y. B., Giang, M. T., **Fefferman**, N., & Wong, J. 2017. The Dragon Swooping Cough: Mass community participation in a virtual epidemic within a tween online world. *In* B. Smith, M. Borge, E. Mercier & K. Y. Lim (*Eds.*) *Proceedings of the 12th International Conference on Computer Supported Collaborative Learning,* Volume 2 (pp. 865-866). Philadelphia, PA: International Society of the Learning Sciences.

4. Fields, D. A., Kafai, Y. B., Sun, J., **Fefferman**, N., Ellis, E., DeVane, B., Giang, M. T., & Wong, J. 2016. The great dragon swooping cough: Stories about learning designs in promoting participation and engagement with a virtual epidemic. *In* Barany, A., Slater, S., & C. Steinkuehler *(Eds.), Proceedings of the Games + Learning + Society (GLS) 12.0 Conference* (pp. 419-424). Pittsburgh, PA: ETC Press.

3. Verma, S., A. Hamieh*, J. H. Huh, H. Holm, S. R. Rajagopalan, M. Korczynski*, and N. H. **Fefferman**. 2016. Stopping Amplified DNS DDoS Attacks Through Query Rate Sharing Between DNS Resolvers, to appear in the International Conference on Availability, Reliability and Security (ARES). (Note: this is the proceeding of a conference, not a journal, but is equivalent to journal publication for the field of computer science, however in keeping with the conventions of Biology, Fefferman is last author as PI on the sponsoring grant that funded the research.)

2. Korczynski*, M., A. Hamieh*, J.H. Huh, H. Holm, S. R. Rajagopalan, and N.H. **Fefferman**. 2015. DIAMoND: Distributed Intrusion/Anomaly Monitoring for Nonparametric Detection. *CCCN 2015: 24th International Conference on Computer Communications and Networks, IEEE, 2015*. (Note: this is the proceeding of a conference, not a journal, but is equivalent to journal publication for the field of computer science, however in keeping with the conventions of Biology, Fefferman is last author as PI on the sponsoring grant that funded the research.)

1. **Fefferman**, N.H., J. Jagai, and E.N. Naumova. 2004. Two - Stage Wavelet Analysis Assessment of Dependencies in Time Series of Disease Incidence. *Proceedings of the 2004 Conference of the International Environmetrics Society*

Research Mentoring

(**bold = current)**

Undergraduate Researchers:

Shyretha Brown, Danika Chari, Kaige Chen, Ian Clark, Liz Davis, Anne Eaton, Taylor Eisenstein, Brandon Grandison, Derek Hansen, David Haycraft, John Huffman, Ana Kilgore, John Kim, Edward Lee, Somair Malik, Andrew McConvey, Jeffrey Mandell, Zain Paracha, Luke Postle, Lauren Prince, Asya Pritsker, Cathy Reis, Jeremiah Rogers, Bolanle Salaam, Nicole Scholtz, Margaret Senese, Joshua Smith, Andrew Sohn, Kim Stanek, Johanna Tam, Colleen Thiersch, Elena Tsvetkova, Barton Willage, Immanuel Williams, Nakeya Williams, Barry Walker, Hannah Yin, Yi Ming Yu, Yongqing Yuan, Stefanie Yuen, James Xue, Bobby Zandstra

Graduate Researchers:

*(Committee Member, or Advisor for work on funded research projects – not primary dissertation advisor; * = special case)*

Kevin Aagard, Emma Bell, **Carissa Bleker**, Curtis Burkhalter, Jordan Bush, Huilan Chang, Erick Chastain, Fnu Eric Ngang Che, **Brittany Coppinger**, Ashley Crump, Kathyrn Fair, Alison Golinski, **Stephen Grady,** Gili Greenbaum, Candice JeanLouis, **Hwayoung Jung,** Ariel Kruger, Di Li, Eric Lofgren*, Nicholas Lorusso, Adam Marszalek, Benjamin Mcclendon, Anthony Ogbuka, Paul Raff, Orin Robinson, **Margaurete Romero,** Rajat Roy, Liliana Salvador, **Shelby Scott,** Tinevimbo Shiri, Brittany Stephenson**,** Alex Thorn, Rafael Valentine, Alex Villiard, Orion Weldon

(primary research advisor to)

Jessica Beck, **Kelly Buch**, Ashley DeNegre, **Jeff DeSalu**, Brad Greening, Natalie Lemanski, **Agnesa Redere**, Samantha Schwab, Oliver Stringham, Karen Wylie

Post-Doctoral Researchers:

Dr. Erick Chastain, Dr. Lazaros Gallos, Dr. Manuel Garcia-Quisimondo, Dr. Ali Hamieh, Dr. Karlo Hock, Dr. Cindy Hui, **Dr. Jing Jiao**, Dr. Amira Kebir, Dr. Maciej Korczynski, Dr. Natalie Lemanski, Dr. Kellen Myers, Dr. Kah Loon Ng, Dr. Chris Stone, Dr. Nourridine Siewe (co-advised by Prof. S. Lenhart), Dr. Gonzalo Suarez, **Dr. Oyita Udiani**, Dr. Peng Zhong

Courses Developed and Taught (all courses developed from scratch)

- Advanced Mathematical Ecology II (MAT/EEB 682 – University of Tennessee, Knoxville) Spring 2017 and 2019
- Evolution, Disease, and Medicine (ENR110 – Rutgers University / EEB 310 – UT, Knoxville) Fall each year 2009 – 2014, Spring 2018 and 2020
- Conversational Bio-Mathematical Modeling (ENR 428 – Rutgers University/ EEB 475 – UT, Knoxville) Spring 2011 – 2014, 2020
- Problems in Ecology: Academic Pedagogy (ENR 601 – Rutgers University) Fall 2015
- *(Co-Developed and Taught)* Ethics & Professional Development in Ecology and Evolution (ENR 602 01 – Rutgers University) Spring 2013-2016 (exception – sabbatical Fall 2014-Spring 2015)
- Introduction to Modeling Ecology, Evolution, and Epidemiology (ENR 604 – Rutgers University) Spring each year 2010 – 2016 (exception – sabbatical Fall 2014-Spring 2015)
- Introduction to Epidemiological Modeling (ENR 603 – Rutgers University) Fall each year 2009 – 2012
- Elements of Data Analysis and Epidemiology (CMPH 343 – Tufts University School of Medicine) Spring 2006

Professional Memberships

Association for Women in Mathematics (AWM)
Association for Women in Science (AWIS)
Complex Systems Society (CSS)
Institute of Electrical and Electronics Engineers (IEEE)
International Union for the Study of Social Insects (IUSSI)
Society for Industrial and Applied Mathematics (SIAM)
Society for Mathematical Biology (SMB)

Invited Presentations

*upcoming

2020

Public Interview: "Nina Fefferman," You Made it Weird podcast

Public Lecture: "The Role of Applied Math in Real-time Pandemic Response: How Basic Disease Models Work," NIMBioS Webinar Series, Knoxville, TN

Public Interview: "Math + Virus + Us," Here We Are podcast and YouTube video.

2019

Public Lecture: "Vaccine Acceptance and Epidemic Risks," Infinite Futures Event Series, Museum of Science and Industry, Chicago, IL.

"When to Turn to Biology for Inspiration in Systems Design," DIMACS 30th Anniversary Conference, New Brunswick, NJ.

"Patients as patches: Ecological challenges from the epidemiology of healthcare environments," ESA 2019, Louisville, KY.

"Math and Disease," Possibilities in Postsecondary Education and Science (PIPES), UTK, Knoxville, TN.

Keynote Address: "Evolving Efficient Solutions: How simple natural systems solve the most complicated problems," MBI Capstone Conference 2019, Columbus, OH (virtual)

Plenary Talk: "How AIDS prevalence impacts the emergence of antibiotic resistance in bacterial infections," SIAM BAMM 2019, Richmond, VA.

Public Lecture: "Math and Disease," Stand Up Science, Farragut, TN.

"Biosurveillance and Homeland Security," Princeton University, NJ.

"Understanding Social Communication Systems with Homology Theory," Complex Systems Seminar, University of Michigan, Ann Arbor, MI.

"Going Against the Grain," Women Empowered in STEM (WeSTEM) 2019, Champaign, IL.

"You're Worth It: Job Negotiations," Women Empowered in STEM (WeSTEM) 2019, Champaign, IL.

2018

"Math: A Critical, Treacherous Bridge Between Scientific Disciplines," American Geophyiscal Union (AGU 2018), Washington DC.

"The Evolution of Social Complexity as Multi-Scale Feedback Control on Networks," Systems Theory Lunch Colloquium, Harvard Medical School, Boston, MA.

"Saving Bats from Fungal Diseases with Linear Algebra," Claremont Center for Mathematical Sciences Colloquium, Claremont, CA.

Plenary Talk: "Evolving Efficient Solutions: How simple natural systems solve the most complicated problems," NIMBioS Undergraduate Research Conference 2018, Knoxville, TN.

Plenary Talk: "Linking Local Decisions with Global Outcomes in Networks: Case Studies in Behavior and Population Health" SIAM Life Sciences 2018, Minneapolis, MN.

"The mathematical biology of networks: from disease outbreaks to cyber-attacks," TN Governor's School, University of Tennessee, Knoxville, TN.

"Trans-disciplinary adventures in the mathematical biology of networks: from disease outbreaks to cyber attacks," DIMACS REU, Rutgers University, Piscataway, NJ.

Public Webinar: "Social and Biological Networks: The Evolution of Social Systems," US National Academies of Sciences, Engineering, and Medicine: Math Frontiers Webinar Series

2017

"Self-Diagnosing Networks," Data Institute San Francisco Conference (DSCO17), San Francisco, CA.

Keynote: "Evolving Efficient Solutions: How simple natural systems solve the most complicated problems," Workshop on Bio-Inspired Security, Trust Assurance, and Resilience (BioSTAR 2017), San Jose, CA.

"Wildlife Disease Management Outcomes May Depend on the Mechanism of Host Immune Response," Distinguished Lecture Series in Immunology and Infectious Diseases, Center for Emerging & Re-emerging Infectious Diseases, School of Medicine, University of Washington, Pullman, WA.

2016

"Evolving Healthy Populations," International Symposium on Biomathematics and Ecology Education and Research 2016, Charlseton, SC.

"Individuals, Societies, and Climate: Modeling motivations to change," Oak Ridge National Laboratory Workshop on Human Activity at Scale in Earth System Models, Oak Ridge, TN.

"Network Models in Epidemiology," US-Canadian Institutes Epidemiology Summer School: Mathematical Modeling of Infectious Disease Spread, MBI, Columbus, OH.

"The Invasion Ecology of Diseases in a Human Environment," Arthur M. Sackler Colloquia of the National Academy of Sciences, Coupled Human and Environmental Systems, Washington DC.

"Global Feedback Control on Centrality in Self-Organizing Systems", Mathematical Biosciences Institute Workshop on the Control and Observability of Network Dynamics, MBI, Columbus, OH.

"Zika Control: More Complicated than Hoped?" Next Einstein Forum, Dakar, Senegal.

2015

"Linear Algebraic Tools in Conservation Ecology," Simon A. Levin Mathematical, Computational and Modeling Sciences Center Seminar, Tempe, AZ.

"Applications of Homology Theory to Animal Communication Systems," Mathematics and Statistics Colloquium, Arizona State Univ., Tempe, AZ.

"Trade-offs Between Collaboration and Infection Risk: Can 'social distancing' improve colony function?" Conference on Complex Systems 2015, Tempe, AZ.

"The Benefits of Ongoing Dynamics in Self-Organizing Social Systems," Conference on Collective Dynamics and Evolving Networks, Bath, UK.

Plenary Talk: Exploiting the Complexity of Identity to Infiltrate Clandestine Groups – Lessons from a LARP, CyDentity Conference, CCICADA, New Brunswick, NJ.

"Incorporating Evolutionary Rescue into Population Viability Models," Mathematics of Planet Earth: Workshop on Management of Natural Resources, Washington D.C.

"Distributed Detection Algorithms for Real-Time Maritime CyberSecurity," Joint CCICADA & AMU Conference on Maritime CyberSecurity, New Brunswick, NJ.

"The Definition of Communication: One way biology and math people accidentally talk past each other and what we might be able to do to fix it," Annual Meeting, Society for Integrative and Comparative Biology, West Palm Beach, FL.

2014

"BioInspired Anomaly Detection: Social Insects and Network Security," Dept. of Homeland Security Science and Technology HSARPA CyberSecurity Division Research and Development Showcase and Technical Workshop, Washington D.C.

"n-TANGLE: a new method for comparing networks across scales" Workshop on Advances in Discrete Networks, Dept. of Mathematics, Univ. of Pittsburgh, Pittsburgh, PA.

Keynote Address: "Virtual Worlds Helping Public Health Preparedness," New Jersey Health Care Quality Institute Annual Meeting, Trenton, NJ.

"A Mathematician's Role in Fighting Ebola," Saint Ann's School, Brooklyn, NY.

"Provable Boundaries on Disease Outbreaks in Self-Organizing Social Networks," The Duke University Mathematical Biology Colloquium, Durham, NC.

Keynote Address: "Designing your own role: Women in STEM," Tufts University Graduate Student Luncheon for Women in Science, Medford, MA.

"Division of Labor as an Adaptation to Combat Disease Risks?" The Seventh International Symposium on Biomathematics and Ecology: Education and Research (BEER), Claremont, CA.

"How dynamic networks affect disease transmission," The BioCircuits Institute, UCSD, San Diego, CA.

"The Evolution of Social Complexity," Plant Biology Dept. Seminar, Univ. of Vermont, Burlington, VT.

"Provable Boundaries on Disease Outbreaks in Self-Organizing Social Networks," Math Dept. Seminar, Univ. of Tennessee at Knoxville, TN.

"Mathematics, Optimization, and the Evolution and Behavior of Social Insects," Math Dept. Junior Colloquium, Univ. of Tennessee at Knoxville, TN.

"The Life of a Mathematical Researcher," Saint Ann's School, Brooklyn, NY.

"Mathematics, Optimization, and the Evolution and Behavior of Social Insects," Social Insect Research Group Seminar, School of Life Sciences, Arizona State Univ., AZ.

"N-tangle: A Network Comparison Method," Workshop on Animal Social Networks, NIMBioS, TN

2013

"Evolutionary pressures, Infectious Diseases, and Self-Organizing Social Systems," Evolutionary Studies Seminar, Co-Sponsored by the Collective Dynamics of Complex Systems Research Group, the Undergraduate Math Club, Upsilon Pi Epsilon, and Pi Mu Epsilon, SUNY Binghamton, NY.

"BioInspired Anomaly Detection," DHS CyberSecurity PI Meeting, Arlington, VA.

"Mathematics, Evolutionary Biology, Epidemiology, and National Security", Saint Ann's School, Brooklyn, NY.

"Evolution of Reproductive Timing and Social Organization in Honey Bees," Scientific Learning Forum at FMC, Ewing, NJ.

"Crowd Sourcing WoW: A Case Study in Improving Pandemic Preparedness," Annual George M. Sideris Biology Conference, LIU, Brooklyn, NY.

2012

Public Lecture: "Math, Complexity, and Social Groups: Using math to understand the nature of society," Campus Life Enrichment Committee (CLEC) Lecture, Georgia Southern Univ., GA.

"How and Why Static Approximations Can Fail to Give Adequate Insight into Processes on Dynamic Networks," Math Dept. Colloquium, Georgia Southern Univ., GA.

"Theoretical Worlds: An Exploration of Models and Model Systems," Tufts Univ, Dept. of Civil and Environmental Engineering Seminar Series, Medford, MA.

"Help, my avatar is sick!" Panel Talk, SXSW, Austin, TX.

"WISE – Women, Ignore Silly Expectations!" 2012 WISE Conference, Texas A&M, TX.

2011

"The Evolution of Social Complexity," CUNY Initiative for the Theoretical Sciences Workshop on A Unified Theory of Evolution, CUNY, NY.

"Balancing Workforce Productivity Against Disease Risks for Environmental and Infectious Epidemics," Math Dept. Seminar, Univ. of Ghana, Legon, Ghana.

"Selective Pressures from Disease on Social Behavior in Hosts," DIMACS/MBI US - African BioMathematics Initiative: Workshop on Genetics and Disease Control, Elmina, Ghana.

Plenary Address: "The Future of Technology and Knowledge," Next-Generation Communications Interoperability Workshop, Chicago, IL.

"Virtual Worlds and Real Epidemics - Insights from WoW's Corrupted Blood Plague," E-Virtuoses International Conference on Serious Games, Valenciennes, France.

Plenary Address: "Disease Robustness and Evolutionary Selective Pressures on Social Organization in Eusocial Insects," Mathematical Biosciences Institute Workshop on Insect Self-Organization and Swarming, Ohio State Univ., OH.

"Hakkar's Corrupted Blood Plague: How an Outbreak in WoW is Helping Epidemiologists Create Better Disease Models," Game Developer's Conference 2011, San Francisco, CA

"Exploring the Role of Behavior in Infectious Disease Dynamics: Mathematical Insights from World of Warcraft and other Virtual Worlds," DIMACS/CCICADA Student Workshop on Where the Mathematical and Computational Sciences Meet Society, Rutgers University, NJ

"Multi-Dimensional Data and the Influence of Human Behavior in Biosurveillance for Infectious Disease Outbreaks," Global Biosurveillance Conference: Enabling Science and Technology – 2nd Meeting in the Biological Threat Non-Proliferation Conference Series, Santa Fe, NM

2010

"Distributed Algorithms for Collective Visualization of Data," Visualanalytics Workshop 2010, Imperial College London, UK

"The Importance of Behavioral Dynamics on Disease Burden," Southern African Wildlife College, South Africa

"The Impact of Stress on Populations," DIMACS Advanced Study Institute on Conservation Biology, Limpopo, South Africa

"Social Behavior in Virtual Worlds," Panel Discussant – InPlay 2010, Toronto, Canada

"Self-Organizing Networks, Social Complexity, and Disease Dynamics," Rensselaer Polytechnic Institute, NY

"Playing with Plague: Exploring Disease Dynamics from Within," 2010 AAAS Annual Meeting, San Diego, CA

"Epidemiological Pressures on the Evolution of Social Complexity," Mathematical Methods in Systems Biology, Tel Aviv, Israel

2009

"Information Theoretic Tool for Biosurveillance," CCICADA Kickoff Meeting, Rutgers Univ., NJ

"Perspectives, Challenges, and Creativity in Understanding Behavioral Epidemiology," Workshop on Behavioral Epidemiology, Rutgers Univ., NJ

"Evolutionary Implications of Epidemics on Social Behavior," Evolutionary Genetics and Genomics at Rutgers, Rutgers Univ., NJ

Panel participant and Speaker on Popular Culture and Science, Sheffield Documentary Film Festival '09, Sheffield, United Kingdom

Keynote Address: "Epidemiological Insights from Virtual Worlds," Life Science Dialogue Heidelberg, - Inaugural Conference, Germany

"Social Stability and Success: A new concept in self-organizing systems and preferential attachment," Office of Naval Research Workshop on Complex Systems, Institute for Pure and Applied Mathematics, Los Angeles, CA

"The Impact of Household Capital Models on Targeted Epidemiological Control Strategies for Diseases with Age-Based Etiologies," Makerere Univ., Kampala, Uganda

Keynote Address: "Hakkar's Corrupted Blood Plague: How an Outbreak in World of Warcraft is Helping Epidemiologists Create Better Disease Models," Games for Health – Virtual Worlds, Boston, MA

"Network Representations and the Evolution of Social Complexity," Frontiers in Applied and Computational Mathematics, New Jersey Institute of Technology, NJ

"Mathematical Optimization, Evolutionary Sociobiology, and Eusocial Insects," Conference on The Power of Analysis, Princeton Univ., NJ

"Mathematical Insights into Behavioral Epidemiology," Univ. of Texas Health Science Center, Houston, TX

"Basics of Mathematical Modeling," Mosquito Modeling Made Easy Day, Center for Vector Biology, Rutgers Univ., NJ

"Mathematical and Computational Methods in Epidemiology and BioSurveillance," Jackson State University, MS

"Mathematics, Optimization, and the Evolution and Behavior of Social Insects," UNC, Chapel Hill, Applied Math, NC

"Network models in Epidemiology and Sociobiology: Introduction, Overview, and Recent Advances," Mathematical Sciences, RPI, NY

2008

"Social Behavior and the Dynamics of Corrupted Blood," Rice University/Games for Health, Houston, TX

"Possible Selective Mechanisms for the Evolution of Disease-defensive Social Organizations," Ecology and Evolution Seminar, Boston Univ., MA

"Behavioral Epidemiology in Virtual Worlds: Exploiting the virtual experience," Advanced Technology Applications for Combat Casualty Care 08; Telemedicine and Advanced Technologies Research Center Medical Simulation & Training Technology

"Recent Advances in the What, How and When of Network Models in Infectious Disease Epidemiology," SIAM 2008, CA

"World of Warcraft Corrupted Blood Disease: Epidemiological Observations and Findings," Games for Health, Baltimore, MD

"Computational Ecology: The Evolution of Sociality," Frontiers in Applied and Computational Mathematics, New Jersey Institute of Technology, NJ

Plenary Talk: "Self-organizing social behavior and disease-defensive organizational strategies in social species," Complexity 2008, Univ. Illinois Urbana, IL

"From the Individual to the Population: Modeling the many levels of evolutionary fitness in social species," Dept. of Ecology and Evolution and Natural Resources, Rutgers Univ., NJ

"Individual Decisions, Group Efficiency," ExxonMobil, Clinton, N.J.

2007

Public Lecture: "Virtual Games, Real Epidemics: Can We Learn Real-Life Lessons in BioDefense from Online Games?" Biosecurity, Biotechnology and Global Health Seminar Series, Program on Science and Global Security, Princeton Univ., NJ

"Disease on Networks: Can Static Representations Capture the Full Complexity of a Dynamic Process?" NDSSL Seminar Series, Virginia Bioinformatics Institute, Virginia Tech, VA

Public Lecture: "Real People, Virtual Worlds: Watching a Plague Unfold," Institute for Mathematical Sciences, National Univ. of Singapore

"The Continued Mystery of Regular, Old, Annual Flu," Workshop on Mathematical models for the Study of the Infection Dynamics of Emergent and Re-emergent Diseases in Humans, Institute for Mathematical Sciences, National Univ. of Singapore

"Epidemics and the Evolution of Social Complexity," Program in Ecology and Evolution Seminar Series, Rutgers Univ., NJ

"Playing Games at School: Parents, Public Schools, and Children's Health," DIMACS Workshop on Game Theory in Epidemiology and Ecology, Rutgers Univ., NJ

"Analyzing Entropy in Biosurveillance," U.S. Dept. of Homeland Security research briefing, Washington D.C.

"Fantastic Problems in Mathematical Ecology," DIMACS Bio-Math Connection Field Testers Workshop, Rutgers Univ., NJ

"Does Securing Infrastructure Against Workforce-Depletion Depend on Whether the Risk is Environmental or Infectious?" DIMACS Workshop on Mathematical Modeling of Infectious Diseases in Africa, Univ. of Stellenbosch, South Africa

"Social interaction and disease dynamics," Workshop on Analysis of Time Series Data in Epidemiology, Tufts Univ. School of Medicine, Boston, MA

"The Behaviors of Individuals and Populations," Working Group on Spatio-Temporal and Network Modeling of Diseases, ICMS, Edinburgh, Scotland

"The Evolution of Complexity in Already Social Groups," Dept. of Ecology and Evolutionary Biology, Princeton Univ., NJ

"Disease as a Selective Pressure and the Evolution of Social Complexity," Applied Biomathematics, Stony Brook, NY

"Vital Rate Sensitivity Analysis: A new method for population viability analysis - Two examples of its use," Applied Biomathematics, Stony Brook, NY

"Disease as a Selective Pressure and the Evolution of Social Complexity," Morin Lab, Dept. of Ecology, Evolution and Natural Resources, Rutgers Univ., NJ

2006

"The Role of Individual Choice in the Evolution of Social Complexity and its Implications Towards the Emergence of Zoonotic Infections," DIMACS Computational and Mathematical Epidemiology Seminar, Rutgers Univ., NJ

"Preparing Societal Infrastructure Against Disease-Related Workforce Depletion," DIMACS Workshop on Facing the Challenge of Infectious Diseases in Africa, University of the Witswatersrand, South Africa

"Fantastic Problems in Mathematical Ecology," DIMACS Bio-Math Connect Institute for High School Teachers, Denver, CO

"Societal Bio-defense - How Can we Accomplish Safety, Stability and Efficiency?" SIAM Annual Meeting, Boston, MA

"When females should stop supporting lazy males: mathematics and honey bees?" DIMACS REU Seminar Series, Rutgers Univ., NJ

"Selected Problems in Epidemiology." DIMACS Tutorial on Data Mining and Epidemiology, NJ

"How Would Termites Prepare for Pandemic Bird Flu and What Should We Learn From Them?" Joint Dept. of Entomology and Center for Infectious Disease Dynamics Seminar, Penn State Univ., PA

"Different Scales of BioDefense - Can societies be both safe and efficient?" DIMACS Computational and Mathematical Epidemiology Seminar, Rutgers Univ., NJ

2005

"Termites in the Nation′s Service," DIMACS Computational and Mathematical Epidemiology Seminar, Rutgers Univ., NJ

"Applications of Self-Organizing Systems to Epidemiology." DIMACS Mixer Series, Rutgers Univ., NJ

"Disease Signatures: A New Combinatorial Method for Epidemiology," DIMACS Computational and Mathematical Epidemiology Seminar, Rutgers Univ., NJ

"Fantastic Problems in Mathematical Ecology," DIMACS Bio-Math Connect Institute for High School Teachers, Rutgers Univ., NJ

"How Complex Systems Can Simplify a Complex Problem: What Epidemiologists Can Learn From Insects," Institute for Advanced Study, Center for Systems Biology Seminar Series, NJ

2004

"Incorporating Behavior and Social Structure into Pathogen Defense Strategies. Conference on Innate Immunity for Biodefense," National Defense University's Center for Technology and National Security Policy (CTNSP) & the Department of Defense, Washington D.C.

Keynote Address: "Social Insects, Immunocompetence and Epidemiology: A Model System for Systems Modelers," Vanderbilt Medical School, Dept. of Microbiology and Immunology Annual Retreat, TN

"Disease and Immunocompetence in Group-Living Animals: Implications for Human Epidemiology," DARPA/DSO Workshop on Endogenous Defense, VA

Contributed Presentations

2008. "An Interdisciplinary Framework for Defining and Distinguishing Security Desiderata for Personally Sensitive Information," DIMACS/DyDAn Workshop on Internet Privacy: Facilitating Seamless Data Movement with Appropriate Controls

2006. "A Vital Rate Sensitivity Analysis (VRSA) for Non-stable Age Distributions and Short-term Planning," North American Ornithological Conference

2004. "A Mathematical Analysis of Reproductive Fission," North American Section of the International Union for the Study of Social Insects (with published abstract)

2004. "Two-stage Wavelet Analysis Assessment of Dependencies in Time Series of Disease Incidence," The 2004 Conference of the International Environmetrics Society (with published abstract)

2004. "Mathematical Modeling of Behavior and Ecology in Social Insects: Social mechanisms of pathogen control in termite colonies," Departmental Research Seminar, Tufts Univ.

2003. "Modeling Waterborne Infectious Outbreaks: When, where and how bad will they be?" The 2003 Conference of the International Environmetrics Society (with published abstract)

2003. "Modeling Disease Resistance through Social Interactions in Termites," The 2nd Conference on the Mathematics and Algorithms of Social Insects (with published abstract)

Service (external to Home Institution)

Ongoing	Referee of papers for *American Naturalist, Annales Zoologici Fennici, Behavioral Ecology and Sociobiology, Biological Conservation, BMC Evolutionary Biology, Bulletin for Mathematical Biology, Canadian Biosystems Engineering, Conservation Letters, IMA Journal of Applied Mathematics, Journal of Biological Dynamics, Journal of Infectious Diseases, Journal of Insect Science, Journal of Nonlinear Dynamics, Mathematical Biosciences, Journal of Medical Internet Research, Journal of the Royal Society Interface, Malaria Journal, Nature, Nature Scientific Reports, Parasites and Vecotrs, PeerJ, Phyiscal Reviews X, PLoS Computational Biology, PLoSOne, PloS Medicine, PNAS, Vaccine, Vector-Borne and Zoonotic Diseases*
2020	Deputy Editor *PLOS Computational Biology*
2019-2021	Director of Development, Enhancing Diversity in Graduate Education (EDGE) Foundation
2019	Guest Editor *PLOS Computational Biology*
2019	Co-Organizer SIAM Network Science Annual Meeting (NS 19)
2018	NSF ad hoc proposal reviewer
2018	Burroughs Wellcome Fund grant proposal reviewer
2018	Co-Organizer IEEE Symposium on Security and Privacy, entitled: 3rd Workshop on Bio-inspired Security, Trust, Assurance and Resilience (BioSTAR 2018)

2017-cont.	Member of the Leadership Team of the National Institute for Mathematical and Biological Synthesis
2017	Co-Organizer NIMBioS Workshop on Applying Optimization Techniques to Agricultural Problems
2017	ARO grant proposal reviewer
2016	Co-Organizer MBI (the Mathematical Biosciences Institute at Ohio State) Workshop on Generalized Network Structures and Dynamics
2016	Co-Organizer MBI (the Mathematical Biosciences Institute at Ohio State) Emphasis Semester on Dynamics of Biologically Inspired Networks
2014	ARO grant proposal reviewer
2013- 2016	Member of Scientific Advisory Board for MBI (the Mathematical Biosciences Institute at Ohio State)
2013	NIH grant proposal reviewer
2013-2016	Co-Organizer NIMBioS Working Group on Climate Change and Vector-borne Diseases
2013-2019	Invited Participant Joint NIMBioS-SESYNC Working Group on Human Risk Perception and Climate Change
2012	Invited Grant Proposal Reviewer for the United States – Israel Binational Science Foundation
2012	US Environmental Protection Agency FIFRA Scientific Advisory Panel (SAP) on Pollinator Risk Assessment Framework
2011	Invited Participant - External Expert Review Panel for Bioscience Research and Development at Los Alamos National Laboratory
2011	Program Committee Member, The Third International UKVAC Workshop on Visual Analytics (VAW 2011)
2011	NSF grant proposal reviewer
2011	Co-Organizer DIMACS/MBI US - African BioMathematics Initiative: Advanced Study Institute and Workshop on Genetics and Disease Control
2010	Organizer of the DIMACS Mini-Workshop on ‘Emergent Properties of Dynamic Biological Networks’
2010	Lecturer at DIMACS/MBI US - African BioMathematics Initiative: Workshop and Advanced Study Institute on Conservation Biology
2010	Organizer of the DIMACS Mini-Workshop on ‘Game-theoretic Approaches to Medical Prognosis’
2010	NSF grant reviewer/panel participant
2010	Invited International Reviewer for Centre of Excellence Grants for the Australian Research Council
2010	Co-Organizer of the DIMACS Workshop on Modeling and Mitigation of the Impacts of Extreme Weather Events to Human Health Risks
2009	Co-Organizer DIMACS Workshop on Economic Epidemiology, Makerere Univ., Kampala, Uganda
2009	NSF grant reviewer/panel participant
2009	Co-Organizer/ Program Co-Chair Workshop on Economic Epidemiology, Makerere Univ., Kampala, Uganda
2009	Co-Organizer Mosquito Modeling Made Easy Day at the N.J. Center for Vector Biology
2008-2010	Member Chief Editorial Committee for the DIMACS Book Series
2008-2010	Member Editorial Board of DIMACS Educational Modules Series
2008	Invited organizer SIAM mini-symposium on Network Models of Infectious Disease
2008	Ran the Reconnect Program on Biosurveillance at DIMACS – a week long short course for teaching faculty at liberal arts institutions on an advanced topic to expand their

	own and their students research opportunities
2007	Mentor to two teams of researchers for Department of Homeland Security funded Research Experience for those at Minority Serving Institutions
2006-2016	Advisory/Editorial Board Member for the journal *Annales Zoologici Fennici*
2004	Subject Matter Expert on Innate Immunity and Biodefense, National Defense University
2004	Research Consultant, DARPA (via Strategic Analysis, INC.)
2003	Developed algorithm for Managing Endangered Species Habitat in Hawaii - MESHH software package (Reed, J.M., N.H. Fefferman, C.S. Elphick, and M. Silbernagle. 2004)
2000-2002	Technical Editor (Cryptography) to MacMillan Press
1999	Invited Reviewer of AES submission to the National Institute of Standards and Technology, later published as The Twofish Encryption Algorithm, Schneier, et al, 1999, John Wiley & Sons Inc.

Service (internal to Home Institution)

2020	Advisor to the COVID-19 Re-Imagining Fall Task Force
2019-cont.	Head of Graduate Admissions, Program in Ecology and Evolutionary Biology
2019	Research Mentor for the NIMBioS Summer Research Experiences (SRE) for Undergraduates
2019	Co-Organizer Tutorial on Networks at NIMBioS
2018	Serve on departmental Promotion and Tenure Committee for Prof. O'Meara
2018-cont.	Serve on Faculty Mentoring Committee for Prof. Kivlin
2017-cont.	Served as Departmental Coordinator for University Future Faculty Program
2017	Research Mentor for the NIMBioS Summer Research Experiences (SRE) for Undergraduates
2017	Lecturer for Joint 2017 MBI-NIMBioS-CAMBAM Summer Graduate Program
2016-2017	University of Tennessee, Knoxville Department of Ecology and Evolutionary Biology Search Committee Member and Diversity Advocate (Ecosystem Ecology Search)
2016-2017	University of Tennessee, Knoxville Department of Mathematics Search Committee Member (Mathematical Biology Search)
2016-cont.	University of Tennessee, Knoxville Program in Ecology and Evolutionary Biology Graduate Affairs Committee Member
2015-2016	Rutgers University Biological Sciences Area Committee Member
2014	Rutgers University EENR Department Wildlife Biology Faculty Search Committee Member
2010	Co-Mentor to a team of researchers for Department of Homeland Security funded Research Experience for those at Minority Serving Institutions
2009-2010	Organizer of the EENR seminar series
2009	Organizer of the DIMACS Workshop on Behavioral Epidemiology
2009-2010	Member E&E Executive Committee
2008-2012	Member of EENR Curriculum Committee
2008-2010	Member Chief Editorial Committee for the DIMACS Book Series
2008-2010	Member Editorial Board of DIMACS Educational Modules Series
2007-2009	Member of the Rutgers University Advisory Board to the Office for the Promotion of Women in Science, Engineering and Mathematics
2006-2015	Research Advisor for Rutgers Univ. DIMACS REU
2005-2007	Co-organizer DIMACS seminar series Mathematical and Computational Epidemiology