


## A career in disease dynamics: a marriage of biology and math

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 Department of Population Medicine, University of Guelph



Networks: EpiNet

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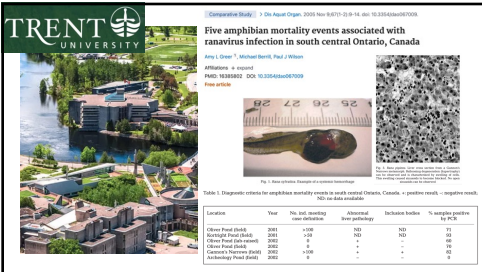
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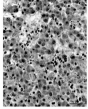
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**Five amphibian mortality events associated with ranavirus infection in south central Ontario, Canada**

Amy L. Greer<sup>1</sup>, Michael Berni, Paul J. Wilson  
 Affiliations: [a record](#)  
 PMID: 3618182, DOI: 10.3354/00007009

[Free article](#)



**Table 1. Diagnostic criteria for amphibian mortality events in south central Ontario, Canada.** <sup>1</sup> positive result, <sup>2</sup> positive result, <sup>3</sup> negative result, <sup>4</sup> not available

Location	Year	No. and mortality count (deaths)	Amphibian spp. mortality	Isolation positive for RV	% samples positive for RV
Chesapeake Park (Ontario)	2001	>100	ND	ND	0
Stony Brook (Ontario)	2002	20	100	ND	100
Chesapeake Park (Ontario)	2002	5	+	-	0
Chesapeake Park (Ontario)	2002	>100	+	+	100
Archibald Park (Ontario)	2002	0	-	-	0

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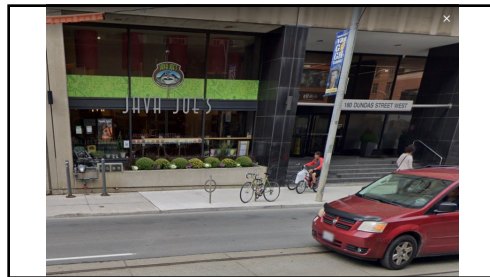
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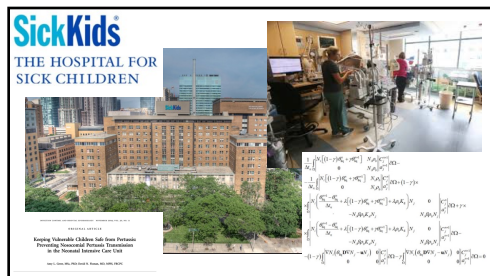
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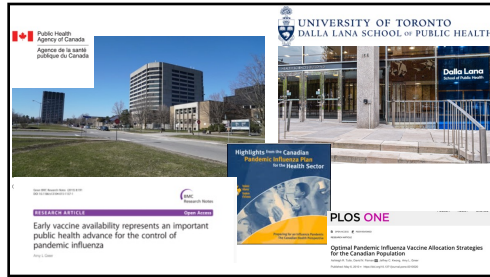
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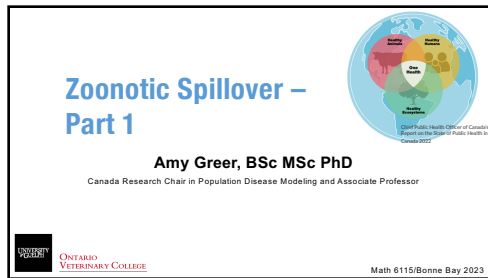
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**What are you going to hear about today?**

1. The ecology of zoonoses and spillover
2. What is disease spillover?
3. What factors are associated with disease spillover events?
4. Considerations for modelling spillover

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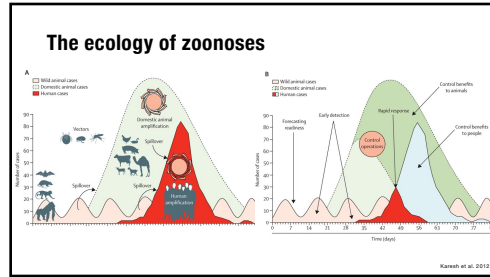
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**How much do you think you know about spillover?**

<http://spillover.nyu.edu>

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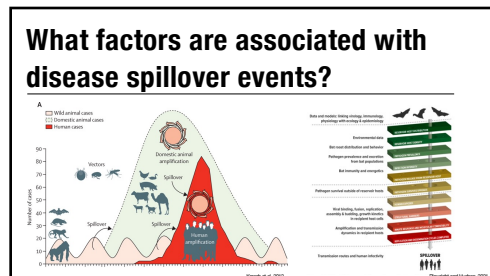
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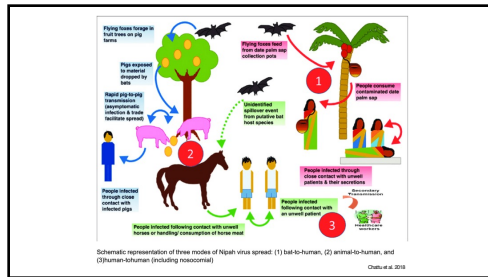
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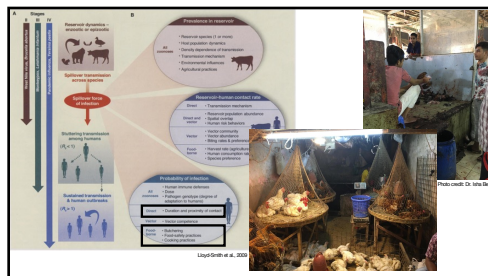
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### Aim & Objectives

- To develop a model that tracks influenza dynamics at the human-poultry interface to better understand risk of influenza reassortment and pandemic emergence in DCC, Bangladesh.
- To evaluate the potential effectiveness of intervention strategies on reducing avian influenza spillover and co-infections with seasonal influenza in humans in DCC, Bangladesh

Berry et al. unpublished

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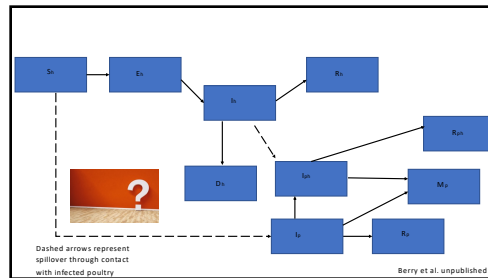
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### Spillover rate (Iacono 2016)

- Probability that K spillover events occur during time  $\tau$  is described by a stochastic Poisson process
- $P(k) = \frac{e^{-\lambda} \lambda^k}{k!}$
- Where  $\lambda$  is the expected number of zoonotic spillovers per unit time, which is specified as
  - $\lambda = N_h(t) Pr_p(N_p) \chi \rho_{hp} (N_p)$ 
    - $N_h(t)$  = Number of susceptible humans at time  $t$ , initial population size
    - $Pr_p(N_p)$  = Prevalence of infected poultry, calibrated using average prevalence of AIV in poultry from Sink Surveillance data in DCC = 52% in 2018
    - $\chi$  = infection response efficiency, estimated as 0.0091 (high uncertainty)
    - $\rho_{hp}(N_p)$  = Average human exposure to poultry in LBM, calibrated using DCC mobile phone survey  $\approx 30.2/365 = 0.082$

Berry et al. unpublished

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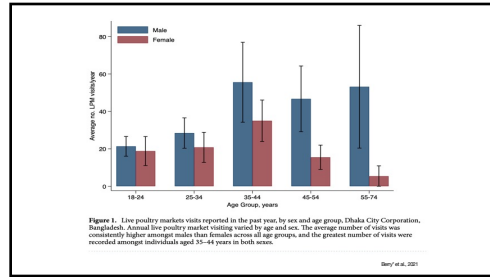
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	Male n = 656 95.05% CI	Female n = 309 96.05% CI	All n = 965 95.55% CI	p-value*
Washed hands with soap				<0.001
Always	18 (2.7)	18 (5.8)	36 (3.7)	
Not always	57 (8.7)	19 (6.2)	76 (7.8)	
Never	41 (6.3)	4 (1.3)	45 (4.6)	
Wore gloves				0.209
Always	0 (0)	0 (0)	0 (0)	
Not always	0 (0)	0 (0)	0 (0)	
Never	0 (0)	0 (0)	0 (0)	
Wore face mask				0.002
Always	0 (0)	0 (0)	0 (0)	
Not always	0 (0)	0 (0)	0 (0)	
Never	0 (0)	0 (0)	0 (0)	
Wore apron				0.182
Always	0 (0)	0 (0)	0 (0)	
Not always	0 (0)	0 (0)	0 (0)	
Never	0 (0)	0 (0)	0 (0)	
	95.05% CI	96.05% CI	95.55% CI	
Never	95.0 (94.8-95.0)	96.0 (95.8-96.3)	95.5 (95.3-95.6)	

**Table 3.** Uptake of protective practices among those with poultry exposure in the past year, by sex, Dhaka City Corporation, Bangladesh. CI confidence interval. \*P-value obtained from chi-square test comparing males and females. \*Sample weighted by age, sex and education to the Dhaka City Corporation demographic profile of the 2011 Bangladesh census. Weighted denominator is those who report any exposure to live poultry in the past year. \*Question was only asked to those who report slaughtering, defeathering, eviscerating and/or cutting poultry; weighted denominator includes only those who report these exposures, n = 566.

*Beer et al., 2021*

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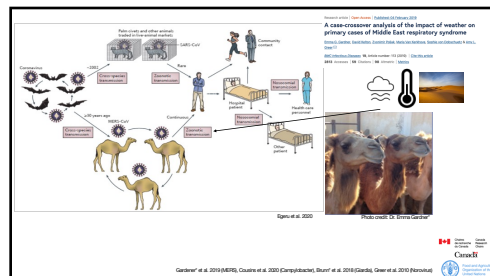
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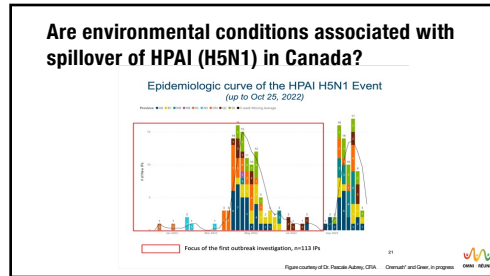
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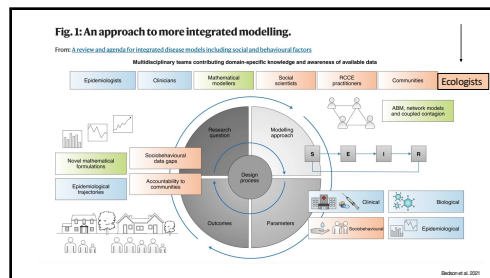
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**PHILOSOPHICAL TRANSACTIONS B**

**Confronting models with data: the challenges of estimating disease spillover**

Paul E. Lead, Ryan J. Pebody, Andrew H. Ross, Graham H. Smith, and Ben R. Hunt

**Abstract**

**Notes**

**Keywords**

**Supplemental Material**

**Discussion**

Get into 4 groups (birthdays: Jan-Mar, Apr-Jun, Jul-Sept, Oct-Dec). Identify a person from your group who will be able to speak to your group response to each of the following questions when we come back together.

1. What did you learn from this paper about modelling disease spillover that you did not know before? (1 student)
2. There are 2 case studies in this paper (HPAI and brucellosis). Briefly summarize the case studies and identify some of the challenges the authors describe in relation to modelling disease spillover for each. (1 student/case study)
3. Identify 2 questions/ideas that came up when you were discussing the paper that you would like to discuss further with the broader class.

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**Problem set 1.7**

1. Think of a simple example of a zoonotic pathogen where the transmission of the pathogen requires both a human and animal host. Describe the natural history of the host-pathogen system in a short paragraph with a link to a reference.
2. Translate the biology from part 1 into a compartment diagram and write the basic corresponding equations to describe the dynamics of the disease in the populations of interest.
3. If you were going to move forward with modelling this zoonotic host-pathogen system, what components of the system would be the most difficult aspects to parameterize? Why?

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