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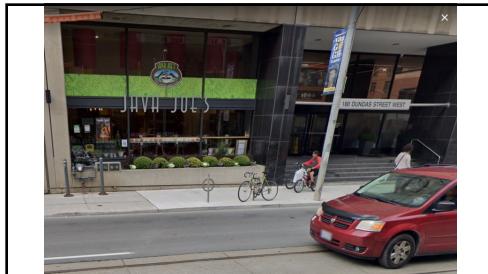
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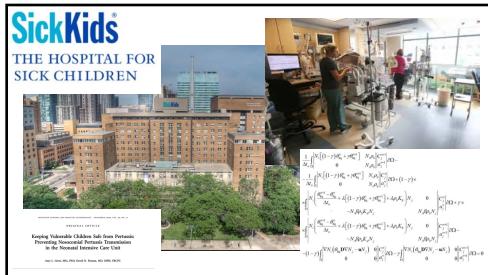
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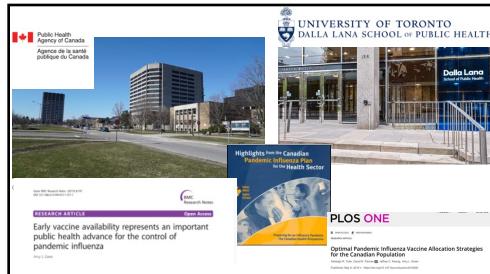
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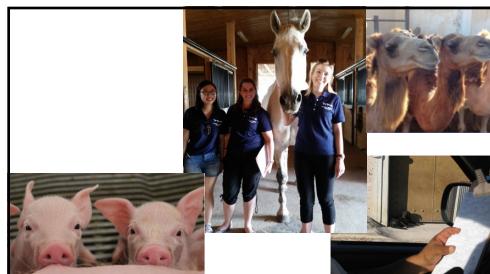
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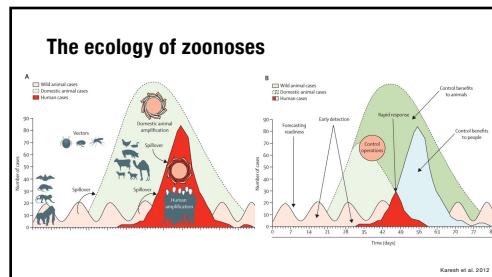
A presentation slide titled 'Zoonotic Spillover – Part 1' by 'Amy Greer, BSc MSc PhD'. The slide includes the text 'Canada Research Chair in Population Disease Modeling and Associate Professor' and the logos of the University of Guelph and Ontario Veterinary College. A circular diagram illustrates the concept of zoonotic spillover, showing overlapping circles for 'Human', 'Domestic animals', 'Wild animals', and 'Environment'.

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A slide titled 'What are you going to hear about today?' listing four topics:

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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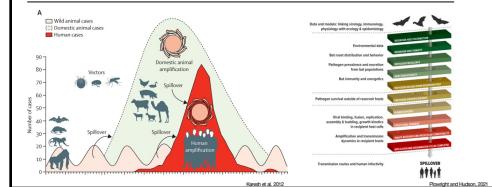
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How much do you think you know about spillover?

spillover.comQuiz

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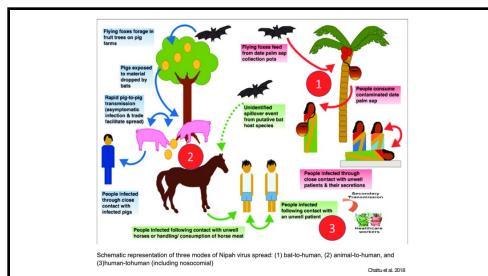
What factors are associated with disease spillover events?



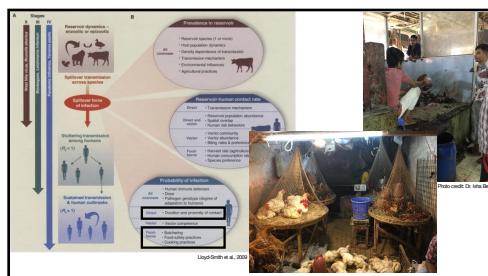
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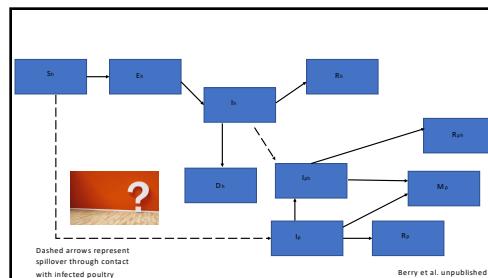
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Aim & Objectives

- 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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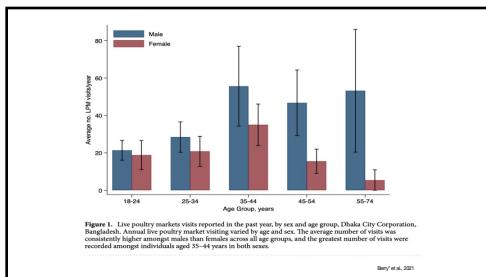
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Spillover rate (Iacono 2016)

- Probability that K spillover events occur during time τ is described by a stochastic Poisson process
- $$P(k) = \frac{\exp^{-\lambda\tau}(\lambda\tau)^k}{k!}$$
- Where λ is the expected number of zoonotic spillovers per unit time, which is specified as
 - $\lambda = N_h(t)P_{RH}(N_h)\chi\eta_{RH}(N_R)$
 - $N_h(t) = N$ total susceptible humans at time t , initial population size
 - $P_{RH}(t) =$ Prevalence of infected poultry, calibrated using average prevalence of AIV in poultry from Sork Surveillance data in DCC = 52% in 2018
 - $\chi =$ infection response efficiency, estimated as 0.0001 (high uncertainty)
 - $\eta_{RH}(N_R) =$ Average human exposure to poultry in LBM, calibrated using DCC mobile phone survey = 0.23/100-0.082

Berry et al. unpublished

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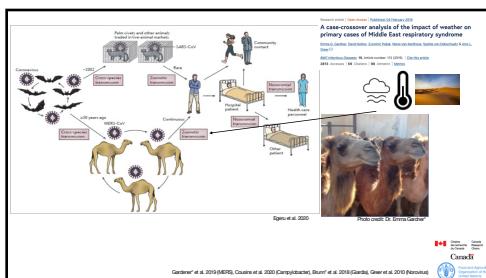
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	Male n = 400 95% CI n = 399-401	Female n = 359 95% CI n = 358-360	All n = 759 95% CI n = 758-760	p value*
Weighted sample?				< 0.001 *
Washed hands with soap				
Always				
Not always				
Never				
Wore gloves				
Always				0.239
Not always				-0.71
Never				-1.91
Wore apron*				0.082 *
Always				-1.11
Not always				-0.81
Never				-0.82
Time framework				
Always	0.0 [0.0, 0.0]	1.1 [0.1, 2.1]	1.1 [0.1, 2.1]	0.182
Not always				
Never				
Wear apron*				
Always	0.0 [0.0, 0.0]	1.1 [0.1, 2.1]	1.1 [0.1, 2.1]	0.182
Not always				
Never				

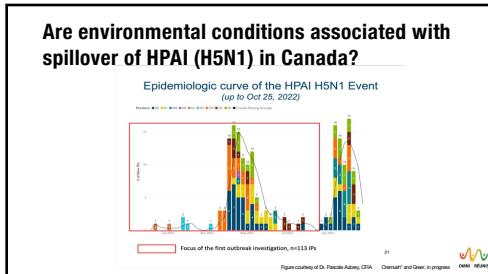
Table 3. Uptake of protective practices among those with poultry exposure in the past year, by sex, Dhaka City Corporation, Bangladesh. CI: confidence interval. *P-value obtained from chi-square test comparing males and females. Sample weighted by sex, sex and education to the Dhaka City Corporation demographic profile of the total Bangladeshi population. *Question only asked to those who report slaughtering, defeathering, eviscerating and/or cutting poultry; weighted denominator includes only those who report these exposures, n = 366.

Sany et al. 2021

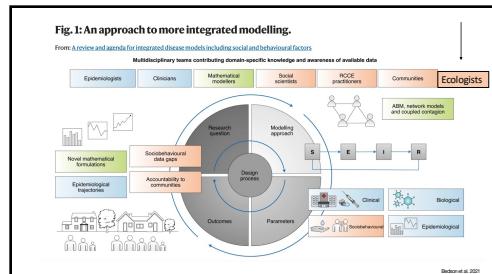
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Review
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Confronting models with data: the challenge of estimating disease spillover
Paul L. Doherty, Daniel J. Pascual, Andrew H. Dobson, Stephen H. Tulloch and Ben H. Peacor

Discussion
Get into 4 groups (birthdays: Jan-Mar, Apr-Jun, Jul-Sep, 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.

- What did you learn from this paper about how to confront models with data that you did not know before? (1 student)
- 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)
- 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?
