

Mathematical models in outbreak response

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1 Background

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Background

Outbreaks:

Influenza, 2009

MERS-Cov (Middle-East Respiratory Syndrome)

Ebola, West Africa, DRC

Zika Virus, Brazil

Objective/Goal

during outbreak:

exploit all data

inform response team in real time

in general (also non outbreak situation)

allow evidence based decisions

compare/assess interventions

policy evaluation (before/after in), vaccine programmes

track of WHO targets (HIV, HCV)

Types of models

dynamic, mathematical (SIR)

statistical (e.g. Poisson regression)

Bayesian statistics

spacial stats/models

-> visualise outcome

Example of outbreak analytics workflow.

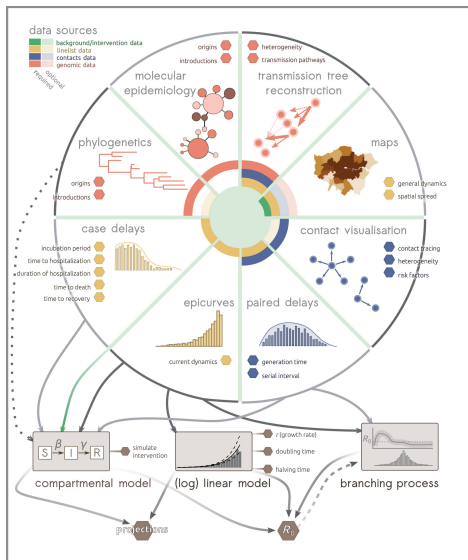
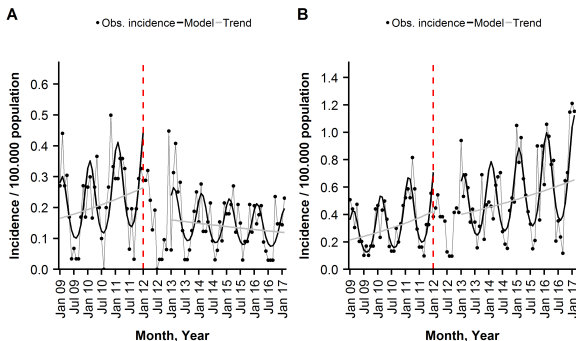


Figure: Monthly incidence of (A) PCV10 ST-IPD and (B) non-PCV10 ex ST 6A-/19A-IPD, among the ≥ 50 years old, observed and modelled by a segmented negative binominal regression, Austria, January 2009-February 2017, shown are overall and seasonal trends.



Model

$$\begin{aligned}\log(Y_t) = & \log(pop_t) + \beta_0 + \beta_1 t + \beta_2 \sin\left(\frac{2\pi t}{12}\right) \\ & + \beta_3 \cos\left(\frac{2\pi t}{12}\right) + \beta_5 (t - t_0)^+ \\ & + \mathbb{1}_{t-t_0>0} \left[\beta_4 + \beta_6 \sin\left(\frac{2\pi t}{12}\right) + \beta_7 \cos\left(\frac{2\pi t}{12}\right) \right]\end{aligned}$$

with

$$(x)^+ = \begin{cases} x, & \text{if } x > 0, \\ 0, & \text{otherwise.} \end{cases}$$

Zika [1]

Humans:

$$\frac{dS_h}{dt} = \mu_h N_h - \frac{\beta_h b}{N_h} S_h I_v - \mu_h S_h$$

$$\frac{dI_h}{dt} = \frac{\beta_h b}{N_h} S_h I_v - (\gamma_h + \mu_h) I_h$$

$$\frac{dR_h}{dt} = \gamma_h I_h - \mu_h I_h$$

Vectors:

$$\frac{dS_v}{dt} = \mu_v N_v - \frac{\beta_v b}{N_h} I_h S_v - \mu_v S_v$$

$$\frac{dE_v}{dt} = \frac{\beta_v b}{N_h} I_h S_v - (\delta + \mu_v) E_v$$

$$\frac{dI_v}{dt} = \delta E_v - \mu_v I_v$$

Zika2

```
Lv      <-      # life span of mosquitos (in days)
Lh      <-      # life span of humans (in days)
Iph     <-      # Infectious period in humans (in days)
IP      <-      # Infectious period in vectors (in days)
EIP     <-      # Extrinsic incubation period in adult mosquitos
muv     <-      # mortality of mosquitos
muh     <-      # mortality of humans
gamma   <-      # recovery rate in humans
delta   <-      # extrinsic incubation rate
b        <-      # Biting Rate
betah   <-      # Probability of transmission from vector to host
betav   <-      # Probability of transmission from host to vector
Nh      <-      # Number of humans (Population of Cali 2.4 million)
m       <-      # Vector to human ratio
Nv      <-      # Number of vectors
RO      <-      # Reproductive number
b       <-      sqrt((RO ^2 * muv*(muv+delta) * (muh+gamma)) /
                  (m * betah * betav * delta)) # biting rate

TIME    <-      # Number of years to run the simulation for
```

Zika3

S_h : Susceptible Humans

I_h : Infected/Infectious humans

R_h : Humans recovered from infection (with lifelong immunity)

S_v : Susceptible vectors

E_v : Exposed vectors

GO ??

a

Conclusion

Here comes the conclusion

Thank you!
Any questions?

- [1] Neil M. Ferguson et al. "Countering the Zika Epidemic in Latin America". In: *Science* 353.6297 (July 22, 2016), pp. 353–354. ISSN: 0036-8075, 1095-9203. DOI: 10.1126/science.aag0219. pmid: 27417493. URL: <https://science.sciencemag.org/content/353/6297/353> (visited on 06/19/2019).
- [2] Polonsky Jonathan A. et al. "Outbreak Analytics: A Developing Data Science for Informing the Response to Emerging Pathogens". In: *Philosophical Transactions of the Royal Society B: Biological Sciences* 374.1776 (July 8, 2019), p. 20180276. DOI: 10.1098/rstb.2018.0276. URL: <https://royalsocietypublishing.org/doi/10.1098/rstb.2018.0276> (visited on 06/18/2019).
- [3] Lukas Richter et al. "Invasive Pneumococcal Diseases in Children and Adults before and after Introduction of the 10-Valent Pneumococcal Conjugate Vaccine into the Austrian National Immunization Program". In: *PloS One* 14.1 (2019), e0210081. ISSN: 1932-6203. DOI: 10.1371/journal.pone.0210081. pmid: 30629620.