



Estimating the Exponential Growth Rate and \mathcal{R}_0

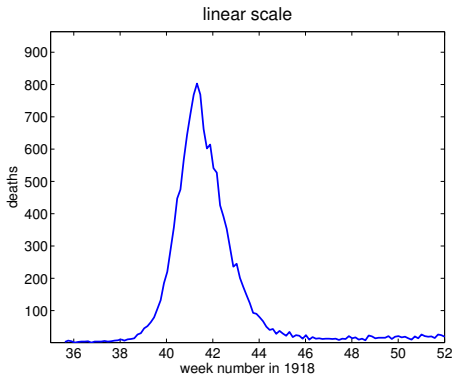
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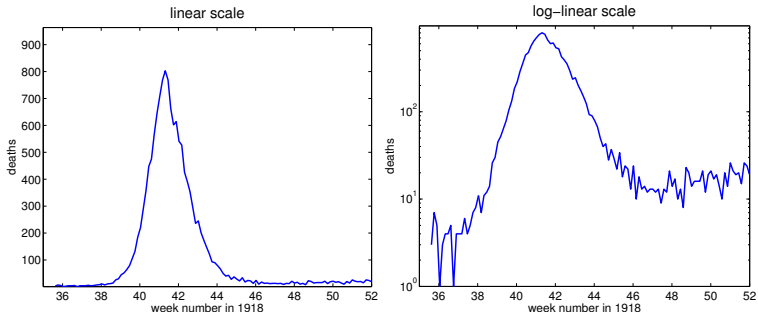
Introduction

Daily pneumonia and influenza (P&I) deaths of 1918 pandemic influenza in Philadelphia.





Re-examine 1918 Daily Philadelphia P&I Deaths



- ▶ An exponential growth phase
- ▶ Given infectious period and latent period, this rate implies \mathcal{R}_0





What We Will Learn

- ▶ Estimate the exponential growth rate
- ▶ Fit (phenomenological or mechanistic) models to data
- ▶ Estimate \mathcal{R}_0 from the exponential growth rate



The Exponential Growth Phase

- ▶ The 1918 pandemic epidemic curve, and most others, show an initial exponential growth phase,
- ▶ That is, during the initial growth phase, the epidemic curve can be modeled as

$$X(t) = X(0)e^{\lambda t},$$

where λ is the exponential growth rate, $X(0)$ is the initial condition.

- ▶ So, $\ln X(t)$ and the time t have a linear relationship during the initial growth phase

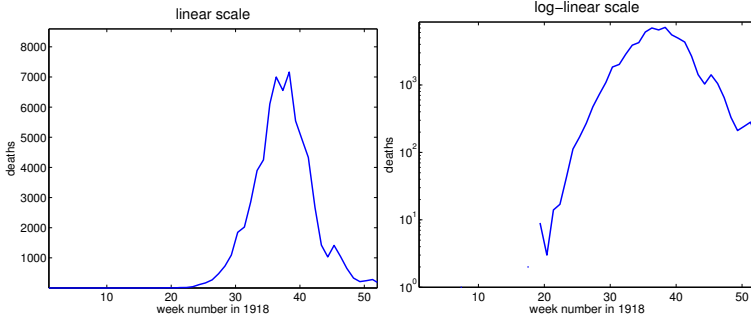
$$\ln X(t) = \ln X(0) + \lambda t.$$

- ▶ The exp growth rate measures how fast the disease spreads





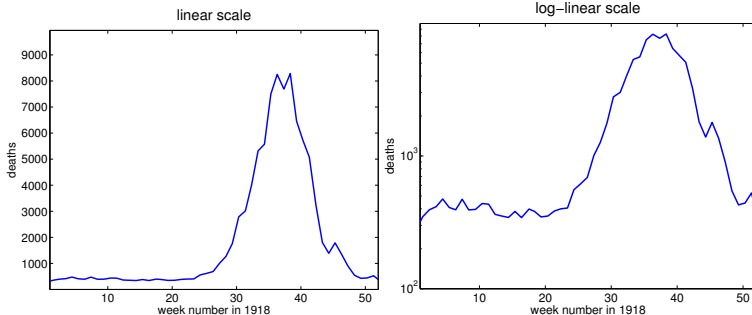
Example: 1665 Great Plague Deaths in London



- ▶ Exponential growth rate decreases around week 25?



Example: 1665 Great Plague All-cause Deaths in London



- ▶ The decrease of the exponential growth rate in plague deaths may be caused by under reporting



Theoretical Exponential Growth Rate: SIR Model

$$S' = -\frac{\beta}{N}SI, \quad I' = \frac{\beta}{N}SI - \gamma I.$$

- ▶ Near the disease free equilibrium (DFE) $(N, 0)$

$$I' = (\beta - \gamma)I$$

- ▶ This is a linear ODE, with an exponential solutions

$$I(t) = I(0)e^{(\beta-\gamma)t}$$

- ▶ So, the exponential growth rate is $\lambda = \beta - \gamma$.
- ▶ What is the growth rate of the incidence curve $X(t) = \beta SI$?





Theoretical Growth Rate: SEIR Model

$$S' = -\frac{\beta}{N}SI, \quad E' = \frac{\beta}{N}SI - \sigma E, \quad I' = \sigma E - \gamma I.$$

- ▶ Near the disease free equilibrium (DFE) $(N, 0, 0)$

$$\frac{d}{dt} \begin{bmatrix} E \\ I \end{bmatrix} = \begin{bmatrix} -\sigma & \beta \\ \sigma & -\gamma \end{bmatrix} \begin{bmatrix} E \\ I \end{bmatrix} = J \begin{bmatrix} E \\ I \end{bmatrix}$$

- ▶ The exponential growth rate is

$$\lambda = \rho(J) = \frac{1}{2} \left(\lambda + \gamma + \sqrt{(\sigma - \gamma)^2 + 4\beta\gamma} \right)$$



Theoretical Growth Rate: General Case

Assume that a disease can be modeled with

- ▶ Susceptible classes $S \in \mathbb{R}^m$ and infected classes $I \in \mathbb{R}^n$
- ▶ parameters $\theta \in \mathbb{R}^p$.
- ▶ Assume a disease free equilibrium (DFE) ($S = S^*, I = 0$).

$$S' = f(S, I; \theta), \quad I' = g(S, I; \theta), \quad \text{where } \frac{\partial}{\partial S} g(S^*, 0) = 0$$

- ▶ Linearize about the DFE ($S^*, 0$):

$$I' = \frac{\partial g}{\partial I}(S^*, 0; \theta)I.$$

- ▶ The exponential growth rate

$$\lambda = \rho \left(\frac{\partial g}{\partial I}(S^*, 0; \theta) \right)$$





Fitting an Exponential Curve

- ▶ Model

$$x(t) = x(0)e^{\lambda t}$$

- ▶ Naive methods that have been widely used:

- ▶ Least square and linear regression
- ▶ Poisson regression
- ▶ Negative binomial regression

- ▶ These methods

- ▶ assume a mean that can be described by a deterministic model
- ▶ only consider observation errors around the deterministic model
- ▶ ignore the process errors are completely ignored



Point Estimates and Confidence Intervals

- ▶ The best estimate for (λ, x_0) is called a *point* estimate.
- ▶ A 95% confidence interval (CI) (a, b) for λ is an *interval* estimate that satisfies

$$\text{Prob}\{\lambda \in (a, b)\} = 95\%$$

- ▶ 95% is called the confidence level. Other examples, 99% CI
- ▶ Infinitely many CI with the same confidence level (95%)
- ▶ Wider CIs means the true parameter value may differ more widely from the point estimate
- ▶ E.g.: The 1918 pandemic influenza (fall wave) has $\mathcal{R}_0 = 1.86$ with 95% CI (1.82, 1.90) (Chowell et al Proc B 2008).



Linear Regression

$$x(t) = x(0)e^{\lambda t} \Rightarrow \ln x(t) = \ln x(0) + \lambda t$$

Commonly use the least square method:

- ▶ For a data set (t_i, x_i) , minimize

$$F(\lambda, x_0) = \sum_{t=1}^n (\ln x(t_i) - \ln x_0)^2$$

- ▶ Confidence intervals:
 - ▶ Assume that $\ln x_i$ are normally distributed,
 - ▶ i.e., x_i are log-normally distributed
 - ▶ Then (λ, x_0) are joint normal
 - ▶ The covariance matrix is $(D^2 F)^{-1}$.
 - ▶ If x_i is not log-normal, not an easy problem.



Poisson Regression

- ▶ For an epidemic curve (t_i, x_i) , x_i usually not \sim log-normal.
- ▶ If infection events have exponentially distributed waiting time, x_i are *Poisson* distributed.
- ▶ Poisson regression for these type of data, which is a maximum likelihood method.
 - ▶ A likelihood function is the probability of observing the data with a given set of parameters

$$\begin{aligned}
 L(\{x_i\}_{i=1}^n | \lambda, x_0) &= \prod_{i=1}^n \text{Prob}(x_i | \lambda, x_0) = \prod_{i=1}^n \frac{E[x_i]^{x_i} e^{-E[x_i]}}{x_i!} \\
 &= \prod_{i=1}^n \frac{x(t_i)^{x_i} e^{-x(t_i)}}{x_i!} = \prod_{i=1}^n \frac{x_0^{x_i} e^{\lambda t_i x_i - x_0} \exp(\lambda t_i)}{x_i!}.
 \end{aligned}$$

- ▶ Find the parameters λ, x_0 that maximize L



Poisson Regression: Maximize Log-likelihood

- ▶ Because L is a product, it is convenient to maximize $\ln L$, called the log likelihood

$$\ln L(\lambda, x_0) = \sum_{i=1}^n x_i \ln x_0 + \lambda t_i x_i - x_0 e^{\lambda t_i} - \ln(x_i!).$$

- ▶ Because x_i are constants, drop $\ln(x_i!)$ to maximize

$$\ln \tilde{L}(\lambda, x_0) = \sum_{i=1}^n x_i \ln x_0 + \lambda t_i x_i - x_0 e^{\lambda t_i}.$$

- ▶ This can only be maximized numerically.
- ▶ Covariance matrix of the parameters:

$$\text{Var}[\lambda, x_0] = \left(D^2 \ln \tilde{L} \right)^{-1}$$





Confidence Intervals: Likelihood Ratio Test

- ▶ To estimate the CI for λ , We use the likelihood ratio test
- ▶ Construct a likelihood profile for λ
 - ▶ λ is stepped to both directions of the point estimate $\hat{\lambda}$
 - ▶ At each step $k = \pm 1, \pm 2, \dots$
 - ▶ Find $L_k = \max L(x_0 | \lambda_k)$
 - ▶ Compute the likelihood ratio

$$D(\lambda_k) = 2 \ln \frac{L_0}{L_k} = 2 \ln L_0 - 2 \ln L_k$$

where L_0 is the likelihood at the point estimate.

- ▶ Best practice for step size is to use the standard deviation from the covariance matrix.
- ▶ Approx. $D_k \sim \chi_1^2$, find the 95% CI for $D(\lambda) : (D(\lambda_a), D(\lambda_b))$.
- ▶ The 95% for λ is (λ_a, λ_b) .



Negative Binomial Regression

- ▶ Poisson regression assumes $E[x_i] = \text{Var}[x_i]$.
- ▶ Over-dispersion: $\text{Var}[x_i] > E[x_i]$ because of
 - ▶ observation errors
 - ▶ non-exponentially distributed waiting times
- ▶ Solution: assume that x_i is Negative-Binomial with parameters r and $0 < p < 1$

$$\text{Prob}(x_i|r, p) = \frac{\Gamma(x_i + r)}{x_i! \Gamma(r)} p^r (1 - p)^{x_i}.$$

- ▶ Assume the same r for all x_i .

$$E[X_i] = r(1 - p)/p \Rightarrow p = r/(r + E[X_i]) = \frac{r}{r + x_0 e^{\lambda t_i}}$$

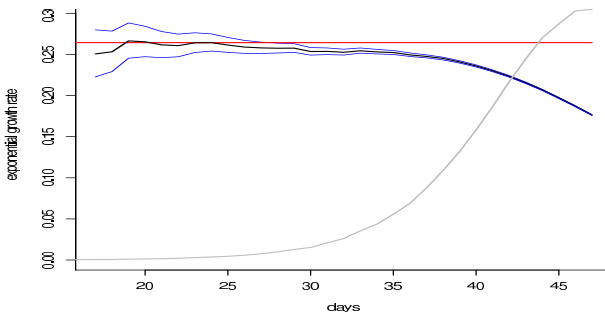
- ▶ The parameters are λ , x_0 , and r .
- ▶ As $r \rightarrow \infty$, the Negative Binomial approaches Poisson.



Fitting an Exponential Curve

Application to Simulated Epidemics

The trend of estimated exponential growth rate when using more data points towards the peak of epidemic:



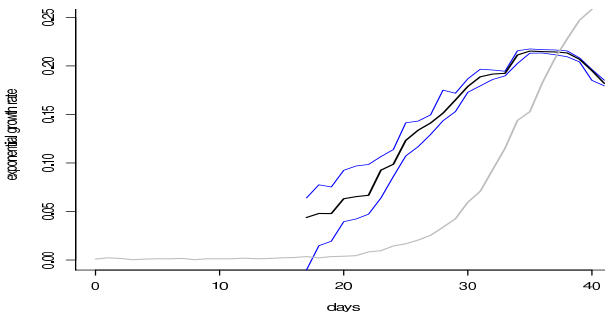
Red: theoretical rate; Black: estimation; blue: 95% CI;





1918 Pandemic Influenza in Philadelphia

The trend of estimated exponential growth rate when using more data points towards the peak of epidemic:



Black: estimation; blue: 95% CI; grey: epidemic curve



Baseline

- ▶ The early flat phase are non-flu deaths, such deaths are called the baseline P&I deaths
- ▶ In a pandemic, most P&I deaths are flu deaths. We can thus ignore the variation in the baseline
- ▶ So, we can use a new model for the mean P&I deaths

$$x(t) = b + x_0 e^{\lambda t}$$

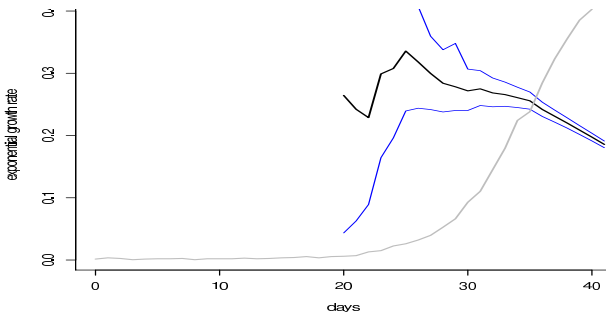
where b is the baseline.

- ▶ We use Poisson regression.



1918 Pandemic Influenza in Philadelphia with Baseline

The trend of estimated exponential growth rate when using more data points towards the peak of epidemic:



Black: estimation; blue: 95% CI; grey: epidemic curve



Taking Account of Decreasing Growth Rate

- ▶ Exponential growth rate decreases because of the depletion of the susceptibles.
- ▶ Use the exponential model,
 - ▶ Find the best fitting window by testing goodness of fit.
- ▶ Use more sophisticated phenomenological models
 - ▶ Logistic model for cumulative cases
 - ▶ Richards model for cumulative cases
- ▶ Use a mechanistic model, e.g., SIR, SEIR, ...



Single Epidemic Phenomenological Models

- ▶ Logistic model:
 - ▶ The cumulative cases $C(t)$ initially grow exponentially, then approach the final size
 - ▶ The same shape as the logistic model.

$$C'(t) = \lambda C[1 - C/K]$$

This model introduces one more parameter K .

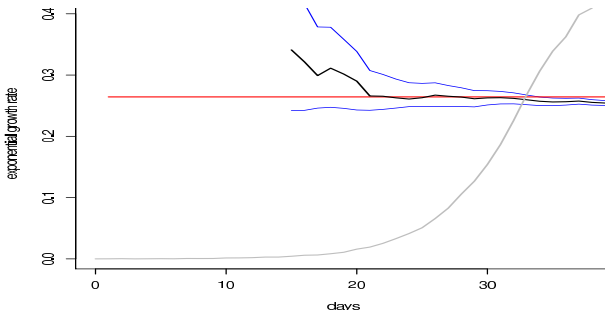
- ▶ But we should not directly fit the cumulative cases data $c_k \sum_{i=0}^k x_k$ to this model, because c_k are not independent.
- ▶ Instead, we compute the interval cases $x(t) = c(t+1) - c(t)$, and fit $x(t)$ to the data x_i .
- ▶ Richards model: cumulative cases has a mean

$$C'(t) = \lambda C[1 - C/K]^\alpha$$



Fit Logistic Model to Simulated Epidemics

Allows the use of more data points:



Red: theoretical rate; black: estimation; blue: 95% confidence interval; grey: epidemic curve

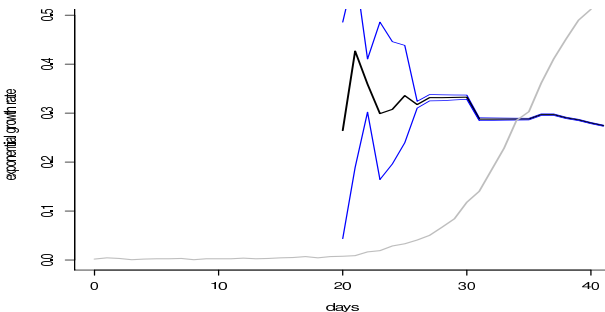




Decreasing Growth Rate

Philadelphia 1918 Pandemic w/ Baseline + Logistic Model

The trend of estimated exponential growth rate when using more data points towards the peak of epidemic:



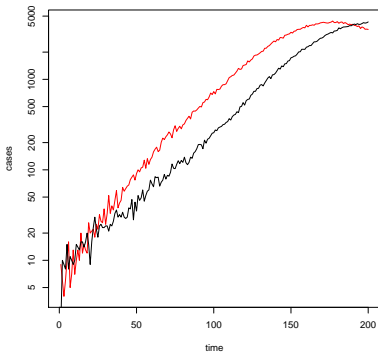
Black: estimation; blue: 95% confidence interval; grey: epidemic





Process Errors

Same disease parameters may produce different epidemic curves.





Coverage Probability

- ▶ Coverage probability of a CI is the probability that CI contains the true parameter value.
- ▶ A 95% CI should have 95% coverage probability.
- ▶ Because we ignored process errors, this method generally produces narrower confidence intervals
- ▶ Simulations can verify that the coverage probability for incidence cases is poor.
- ▶ Larger observation errors, for example, small reporting rates, improve coverage.

Methods that can handle process errors include: one-step ahead, particle filters, MCMC, ...



Estimate \mathcal{R}_0 : SIR Model

First, as an example, we look at an SIR model

$$S' = -\frac{\beta}{N}SI, \quad I' = \frac{\beta}{N}SI - \gamma I.$$

- ▶ Recall that $\lambda = \beta - \gamma$, so $\beta = \lambda + \gamma$

$$\mathcal{R}_0 = \frac{\beta}{\gamma} = \frac{\lambda + \gamma}{\gamma} = 1 + \frac{\lambda}{\gamma}.$$

- ▶ What if λ is the exponential growth rate of the incidence curve $X(t) = \beta SI$?



Estimate \mathcal{R}_0 : SEIR Model

$$S' = -\frac{\beta}{N}SI, \quad E' = \frac{\beta}{N}SI - \sigma E, \quad I' = \sigma E - \gamma I.$$

- ▶ Recall that

$$\lambda = \rho(J) = \frac{1}{2} \left(\lambda + \gamma + \sqrt{(\sigma - \gamma)^2 + 4\beta\gamma} \right)$$

- ▶ Isolate β ,

$$\beta = \sigma + \frac{\lambda}{\gamma}(\lambda + \gamma + \sigma)$$

$$\mathcal{R}_0 = \frac{\beta}{\gamma} = 1 + \lambda \left(\frac{\lambda}{\sigma\gamma} + \frac{1}{\gamma} + \frac{1}{\sigma} \right).$$



Estimate \mathcal{R}_0 with a Model: in General

$$\begin{aligned} S' &= f(S, I; \theta), \\ I' &= g(S, I; \theta). \end{aligned}$$

where $S \in \mathbb{R}^m$, $I \in \mathbb{R}^n$, θ is the vector of parameters.

- ▶ Recall that the exponential growth rate is the largest eigenvalue of

$$\frac{\partial g}{\partial I}(S_0, 0; \theta).$$

- ▶ This relationship usually gives us an estimate of the transmission rate given all the other disease parameters.
- ▶ \mathcal{R}_0 can be computed using the inferred transmission rate and all other given parameter values.



Estimate \mathcal{R}_0 using Generation Interval

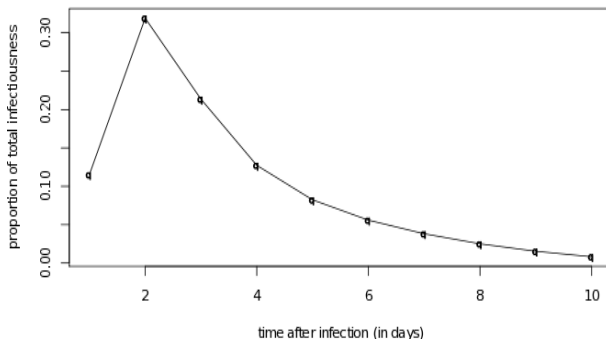
See Wallinga and Lipsitch (Proc B 2007, 274:599604)

- ▶ Generation interval (serial interval): the waiting time from being infected to secondary infections
 - ▶ The generation interval distribution $w(t)$ can be estimated (e.g., from contact tracing) without a mechanistic model.
- ▶ Let $n(t)$ is the transmission rate at age of infection τ .
 - ▶ $\mathcal{R}_0 = \int_0^\infty n(\tau) d\tau$, and $w(\tau) = n(\tau)/\mathcal{R}_0$.
 - ▶ The incidence curve $x(t) = x(t) * n(t)$
- ▶ Assume $x(t) = x(0)e^{\lambda t}$,

$$\mathcal{R}_0 = \frac{1}{\int_0^\infty e^{-\lambda\tau} w(\tau) d\tau}$$



The Influenza Generation Interval Distribution



Taken from N.M. Ferguson, et al., (Nature 2005, 437:209-214)



The Basic Reproduction Number of 1918 Pandemic Influenza in Philadelphia

Given that we have estimated the exponential growth rate to be

$$\lambda = 0.288, \quad \text{with 95\% confidence interval: } (0.286, 0.290)$$

and with the above generation interval distribution, we can compute that

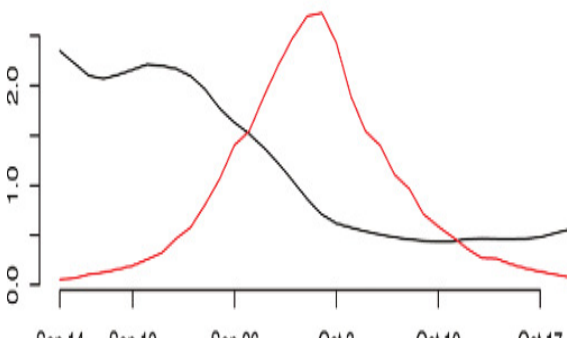
$$\mathcal{R}_0 = 2.16, \quad \text{with 95\% confidence interval: } (2.15, 2.17)$$

This is consistent with other estimations such as Mills et al. (Nature 2004) and Goldstein et al. (PNAS 2009)



Basic Reproduction Number Estimated by Goldstein et al (2009)

Daily reproductive numbers, Philadelphia (above) and NY State (below)





Some Considerations

- ▶ Deaths v.s. incidences
- ▶ Temporal aggregation (e.g., weekly incidences)
- ▶ Spatial aggregation (e.g., overall Canada v.s. city level curves)



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