

9 Epidemic Data II

10 Epidemic Data III

11 Epidemic Data Tools



Mathematics and Statistics

$$\int_{M} d\omega = \int_{\partial M} \omega$$

Mathematics 4MB3/6MB3 Mathematical Biology

Instructor: David Earn

Lecture 8 Epidemic Data Wednesday 24 January 2018

Announcements

- Thanks everyone for doing the contributions survey for Assignment 1.
- Don't stress about the ratings about each other's contributions. The issue is whether some group members did not pull their weight. If somebody didn't try and others had to pick up the slack, that person should be penalized. I will not penalize somebody because they tried but felt they didn't contribute as much to the final document as they could have. Do try to even out the work across the assignments.
- Make sure everyone in your group gets a chance to be in control of the LATEX for one assignment.

More Announcements!

Assignment 2:

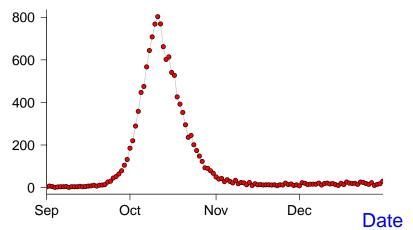
Due Monday 5 February 2018 in class (and by e-mail) at 11:30am.

Midterm test:

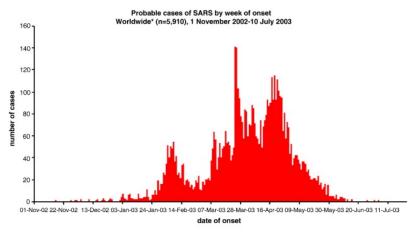
- Date: week of 5–9 March? or 12–16 March?
- Time: TBA
- Location: TBA

P&I Mortality, Philadelphia, 1918

P&I Deaths



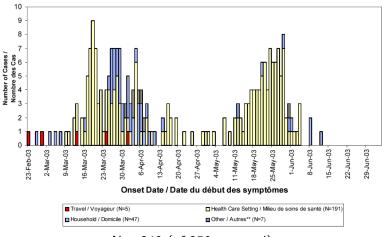
SARS in 2003 (Worldwide)



*This graph does not include 2,527 probable cases of SARS (2,521 from Beijing, China), for whom no dates of onset are currently available.

Motivating Data – Single epidemics

SARS in 2003 (Toronto)



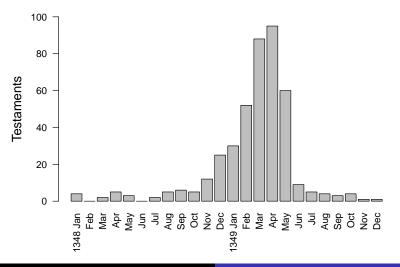
N = 249 (of 250 reported)

Some SARS Facts

High case fatality

- 1918 flu < 3%
- SARS > 10%
- Long hospital stays
 - Mean time from admission to discharge or death: ~ 25 days in Hong Kong
- 8098 probable cases, 774 deaths
- How bad would it have been if it had not been controlled?

The Black Death in London, England, 1348–1349



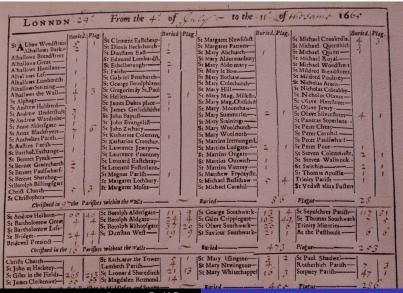
Instructor: David Earn Mathematics 4MB3/6MB3 Mathematical Biology

London Bill of Mortality, 26 Sept to 3 Oct 1665

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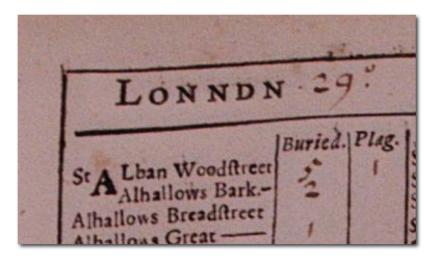
Instructor: David Earn

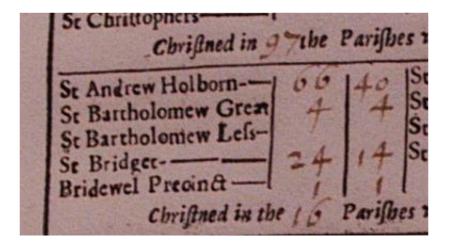
Mortality Bills are typically handwritten



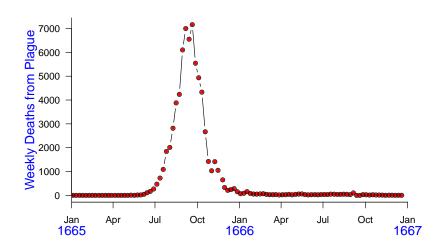
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But handwriting is usually very clear

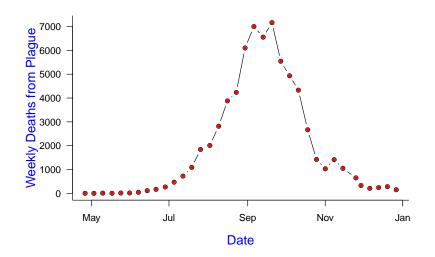




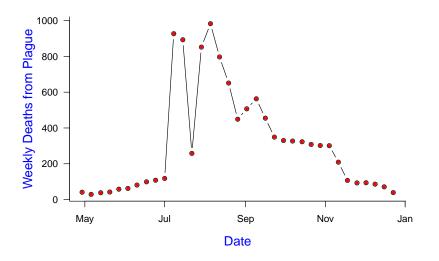
The Great Plague of London, 1665



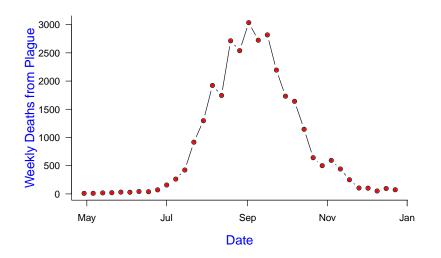
The Great Plague of London, 1665



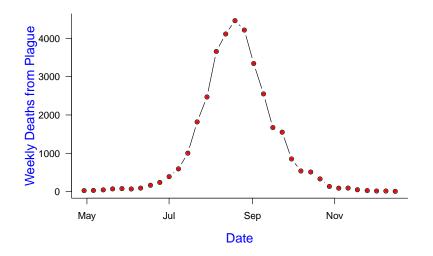
London Plague of 1593



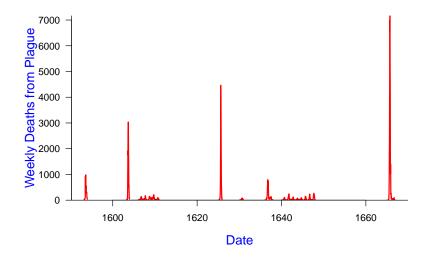
London Plague of 1603



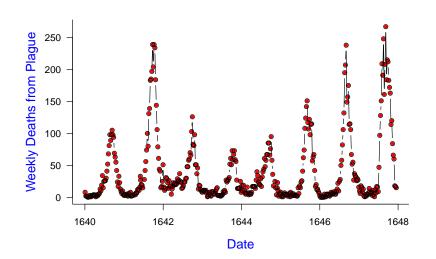
London Plague of 1625



Weekly Deaths from Plague in London, 1592–1666



Weekly Plague in London, 1640–1648



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Some Plague Facts

- Plague epidemics recorded from Roman times to early 1900s.
- $\label{eq:last} \mathbf{I}/3 \mbox{ Europe's population died in "Black Death" of 1348} \\ \mathbf{I} \sim 300 \mbox{ years for the population to reach the same level.}$
- Recently (2011) established (at McMaster!) that the pathogen that caused The Black Death was Yersinia pestis

[Bos et al. 2011, Nature 478, 506-510]

 More recently (2014) established (again at McMaster!) that the pathogen that caused The Plague of Justinian (541–543 AD) was Yersinia pestis

[Wagner et al. 2014, Lancet Infectious Diseases 14, 319-326]

■ *Y. pestis* still a concern?

Yes: Rodent reservoir, antibiotic-resistant strains, bioterrorism

Spatial data for any plagues? Yes, for London in 1665...

Visualization of spatial structure of Great Plague

- GIS encoding of parish boundaries
- Overlay parish boundaries on more modern map for reference
- Colour parishes as they become infected
- Is there evidence for spatial spread or was the spatial pattern random?
- DE low-tech animation...
- CBC high-tech animation...
 - The Nature of Things, 21 August 2014. http://www.cbc.ca/natureofthings/episodes/ secrets-in-the-bones-the-hunt-for-the-black-death-killer



Mathematics and Statistics

$$\int_{M} d\omega = \int_{\partial M} \omega$$

Mathematics 4MB3/6MB3 Mathematical Biology

Instructor: David Earn

Lecture 9 Epidemic Data II Friday 26 Jan 2018

Announcements

Assignment 2:

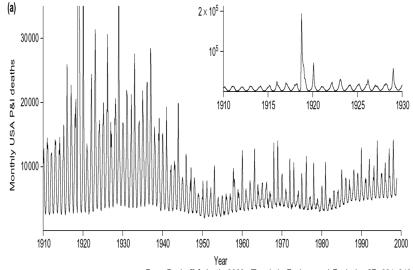
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 - Time: 7:00pm to 9:00pm
 - Location: TBA

Visualization of entire course of the Great Plague

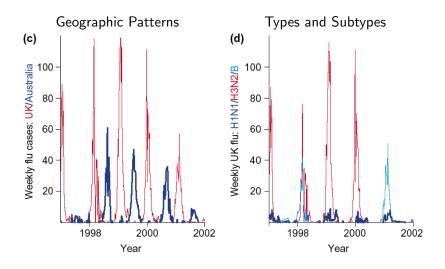
- What happenned after initial spatial spread?
- Visualize full spatial epidemic structure
- Show magnitude of epidemic in each parish with cylinder.
- Epidemic Visualization (EpiVis) software by Junling Ma.

P&I mortality in U.S.A., 1910–1998



Earn, Dushoff & Levin 2002, Trends in Ecology and Evolution 17, 334-340

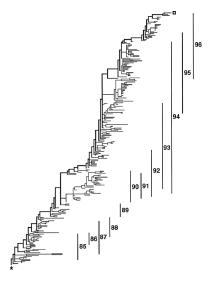
Influenza Incidence Patterns (lab confirmed)



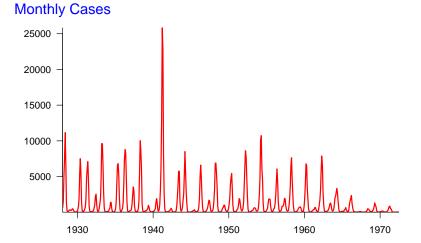
Earn, Dushoff & Levin 2002, Trends in Ecology and Evolution 17, 334-340

Influenza Evolution

Molecular phylogenetic reconstruction of influenza A/H3N2 evolution, 1985–1996 (Fitch *et al.* 1997)

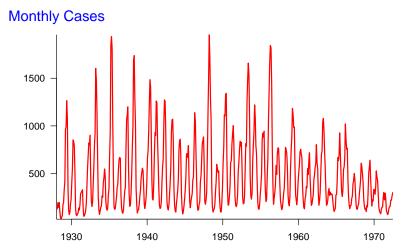


Measles in New York City, 1928–1972



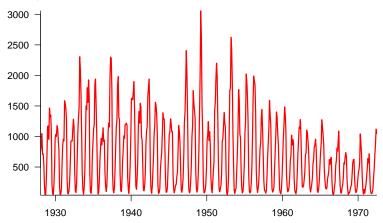
Instructor: David Earn Mathematics 4MB3/6MB3 Mathematical Biology

Mumps in New York City, 1928–1972

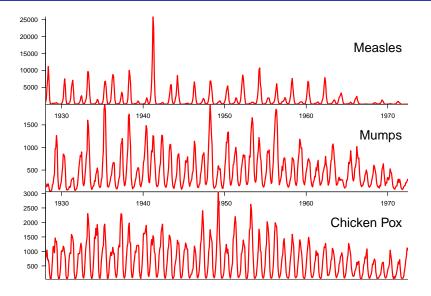


Chicken Pox in New York City, 1928–1972

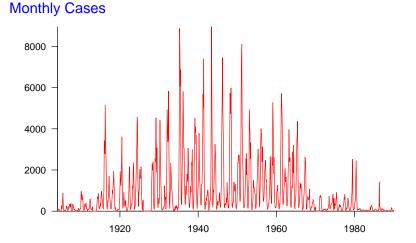
Monthly Cases



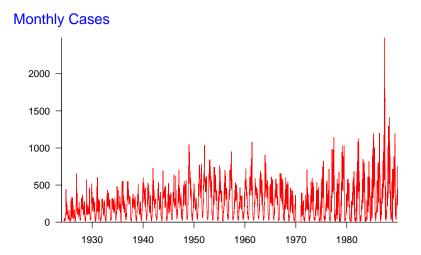
Childhood diseases in New York City, 1928–1972



Measles in Ontario, 1904–1989

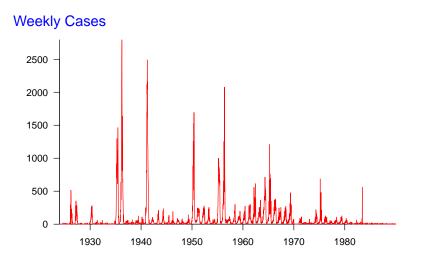


Chicken Pox in Ontario, 1924–1989



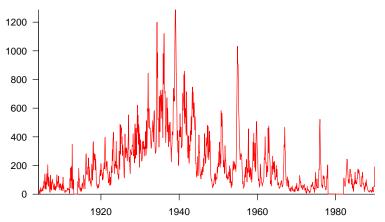
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Rubella in Ontario, 1924–1989

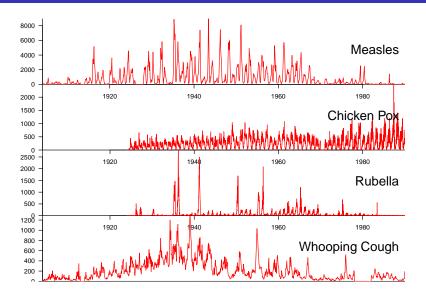


Whooping Cough in Ontario, 1904–1989





Childhood diseases in Ontario, 1904–1989



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Mathematics and Statistics

$$\int_{M} d\omega = \int_{\partial M} \omega$$

Mathematics 4MB3/6MB3 Mathematical Biology

Instructor: David Earn

Lecture 10 Epidemic Data III Monday 29 Jan 2018 Comment from TA on Assignment 1:

"For a few of the groups, I would recommend that they look over the work that their group members have done. Question 2c and 2d in particular were closely related and there were a few obvious cases where the students had not communicated with each other "

Assignment 2:

Due Monday 5 February 2018 in class (and by e-mail) at 11:30am.

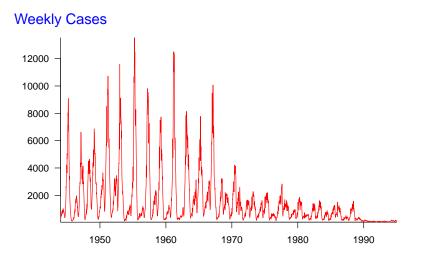
Midterm test: We agreed on:

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- *Time:* 7:00pm to 9:00pm
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Recurrent epidemics of childhood infections

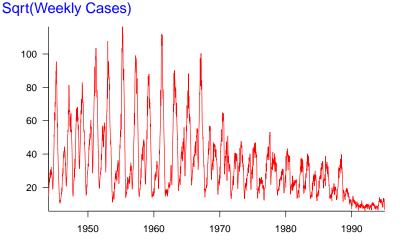
- Childhood diseases in New York City, 1928–1972
- Childhood diseases in Ontario, 1904–1989

Measles incidence in England and Wales, 1944–1995



Instructor: David Earn

Measles incidence in England and Wales, 1944–1995



Why study measles epidemics?

- ~ 90,000 children died from measles in 2016.
- A major cause of vaccine-preventable deaths.
- Potential impact in developed countries during vaccine scares (e.g., MMR scare in UK in 1990s).
- Understand past patterns
- Predict future patterns
- Manipulate future patterns
- Develop vaccination strategy that can...



Other reasons to model infectious disease epidemics

Mathematical models make hypotheses and inferences precise

- Give better advice to policymakers
- Make better predictions
- Host-pathogen dynamics are important aspects of ecosystem dynamics
 - Infectious disease models more likely to be successful than predator-prey models
- Excellent data for human infectious diseases
 - Models can be tested!

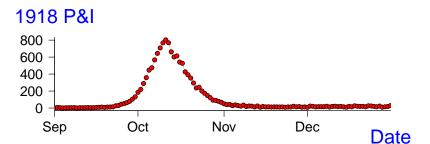
Modelling population dynamics childhood infections

- The basic SIR model cannot explain recurrent epidemics.
- What should we do?...
 - **1** Get depressed, drop the course.
 - 2 Keep developing models until we can explain recurrent epidemics.
- First, let's talk about tools that allow us to make our questions about time series data more precise.

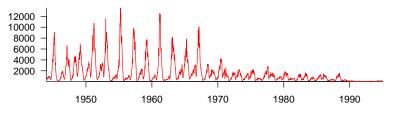
Epidemic Data Analysis

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Time Plots of Temporal Epidemic Patterns

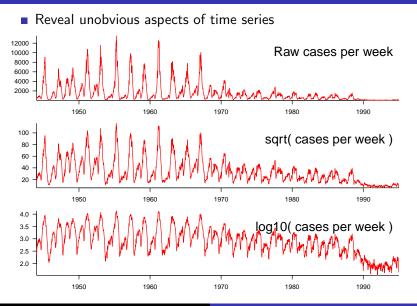


Weekly Measles in England and Wales



50/87

Time Plots of Transformed Data



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- Reveal trends clouded by noise or seasonality
- Moving Average:

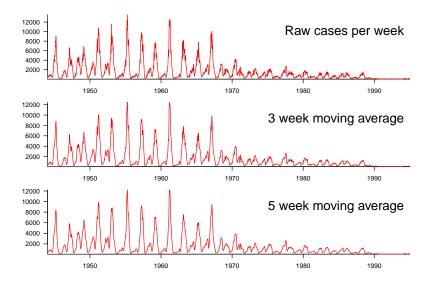
$$x_t \to \frac{1}{2a+1} \sum_{i=-a}^{a} x_{t+i}$$

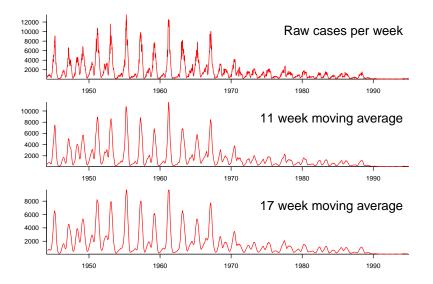
Replace original data points x_t with averages of nearby points.

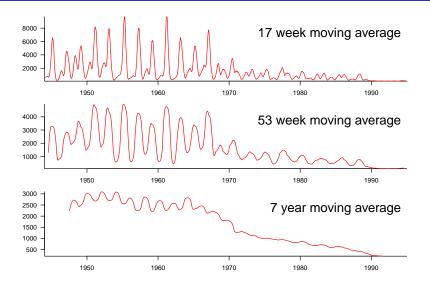
Linear filter:

$$x_t o \sum_{i=-\infty}^{\infty} \lambda_i x_{t+i}$$

- Generalization of moving average.
- Weights λ_i can be nonlinear functions of *i*.







Correlation

- Recurrent epidemics with number of cases in the past and the future.
- Given *N* pairs of observations of different quantities, {(*x_i*, *y_i*) : *i* = 1,..., *N*}, the *correlation coefficient* is defined to be

$$r = \frac{\sum_{i=1}^{N} (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^{N} (x_i - \bar{x})^2 \sum_{i=1}^{N} (y_i - \bar{y})^2}}$$

where \bar{x} and \bar{y} are the means of $\{x_i\}$ and $\{y_i\}$, respectively.

Correlation

Properties of the correlation coefficient:

- $-1 \le r \le 1$ (Proof? Cauchy-Schwarz inequality)
- r = 1 \iff all points lie on a line with positive slope ("complete positive correlation")
- $r = -1 \iff$ all points lie on a line with negative slope ("complete negative correlation")
- $r \simeq 0 \implies$ "uncorrelated"
- Interpretation: r² is the proportion of the variance in y explained by a linear function of x.

Derivations and discussions:

- MathWorld on r^2 , Wikipedia on r^2
- Wikipedia on general coefficient of determination

- Given a single sequence of observations {*x_t* : *t* = 1,...,*N*}, we can compute the correlation of each observation with the observation *k* time steps in the future.
- Thus, we consider the pairs of observations
 {(x_t, x_{k+t}) : t = 1,..., N k} and define the *autocorrelation coefficient at lag k* to be

$$r_{k} = \frac{\sum_{t=1}^{N-k} (x_{t} - \bar{x}_{1,N-k}) (x_{k+t} - \bar{x}_{k+1,N})}{\sqrt{\sum_{t=1}^{N-k} (x_{t} - \bar{x}_{1,N-k})^{2} \sum_{t=1}^{N-k} (x_{k+t} - \bar{x}_{k+1,N})^{2}}}$$

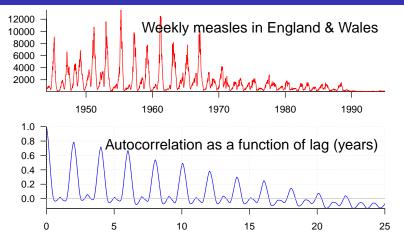
where $\bar{x}_{1,N-k}$ and $\bar{x}_{k+1,N}$ are the means of first and last N-k observations, respectively.

If number of observations N is large and lag $k \ll N$ then

$$r_k \simeq rac{\sum_{t=1}^{N-k} (x_t - ar{x})(x_{k+t} - ar{x})}{\sum_{t=1}^{N} (x_t - ar{x})^2}$$

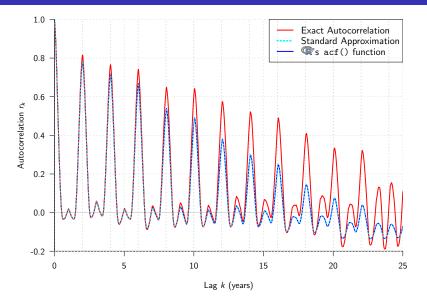
- Approximation of r_k is worse for larger lags k
- Plot of autocorrelation r_k as a function of lag k is called the correlogram.

Correlogram



Peaks in correlogram ⇒ periodicities in original time series.
 Correlograms of temporal segments are often informative.

Correlogram: exact vs. approximate r_k



Spectral Density

- Can we compute the dominant periods in the time series? (Rather than estimating them by eye from the correlogram.)
- Express the time series as a Fourier series:

$$x_t = a_0 + \left(\sum_{p=1}^{(N/2)-1} \left(a_p \cos \omega_p t + b_p \sin \omega_p t\right)\right) + a_{N/2} \cos \pi t \,,$$

where $\omega_p = 2\pi p/N$.

Compute the *Fourier coefficients* {a_p}, {b_p} by taking inner products with cos ω_pt and sin ω_pt.

Spectral Density

Fourier coefficients of x_t are:

$$a_0 = \bar{x} = \frac{1}{N} \sum_t x_t ,$$

$$a_p = \frac{2}{N} \sum_t x_t \cos \omega_p t , \qquad b_p = \frac{2}{N} \sum_t x_t \sin \omega_p t ,$$

$$a_{N/2} = \frac{1}{N} \sum_t (-1)^t x_t ,$$

where sum is over observation times.

• Estimated power spectral density (PSD) at frequency ω_p is*:

$$I(\omega_p) = rac{N}{4\pi} \left(a_p^2 + b_p^2
ight)$$

*The normalization by $N/4\pi$ is the convention chosen by Chatfield (2004, "Analysis of Time Series: An Introduction"). Other normalization conventions are also in common use.



Mathematics and Statistics

$$\int_{M} d\omega = \int_{\partial M} \omega$$

Mathematics 4MB3/6MB3 Mathematical Biology

Instructor: David Earn

Lecture 11 Epidemic Data Tools Wednesday 31 Jan 2018

Announcements

Assignment 2:

Due Monday 5 February 2018 in class (and by e-mail) at 11:30am.

- Midterm test: We agreed on:
 - Date: Thursday 8 March 2018
 - Time: 7:00pm to 9:00pm
 - Location: TBA

Please consider...

5 minute Student Respiratory Illness Survey:

https://surveys.mcmaster.ca/limesurvey2/index.php/893454

Please complete this anonymous survey to help us monitor the patterns of respiratory illness, over-the-counter drug use, and social contact within the McMaster community. There are no risks to filling out this survey, and your participation is voluntary. You do not need to answer any questions that make you uncomfortable, and all information provided will be kept strictly confidential. Thanks for participating.

-Dr. Marek Smieja (Infectious Diseases)

Last time...

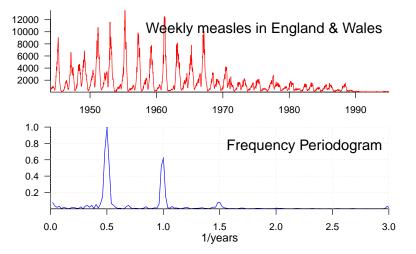
- Statistical description of time series: time plot, moving average
- Correlation coefficient: properties
- Autocorrelation
- Correlogram
- Exact vs. approximate autocorrelation
- Power spectral density (PSD)

- There are many different ways to express the power spectral density (aka *power spectrum*).
- Most common/useful equivalence is that the power spectrum is the discrete Fourier transform of the correlogram:

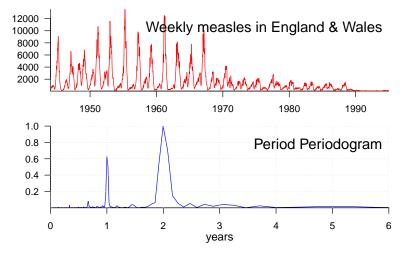
$$I(\omega_p) = \frac{1}{\pi} \left(r_0 + 2 \sum_{k=1}^{N-1} r_k \cos \omega_p k \right)$$

Plot of estimated power spectrum as a function of frequency ω_p is called the *frequency periodogram* or just the *periodogram*.

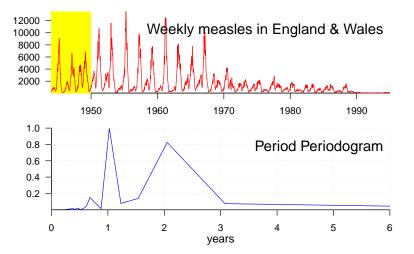
Spectral Density



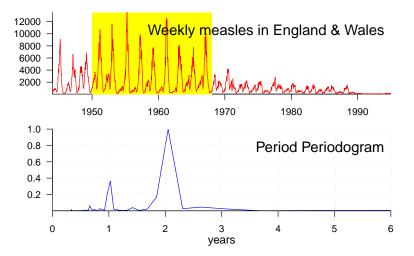
Spectral Density



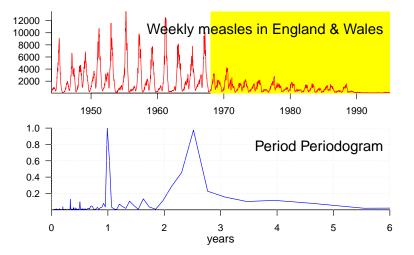
Spectral Density of Temporal Segments



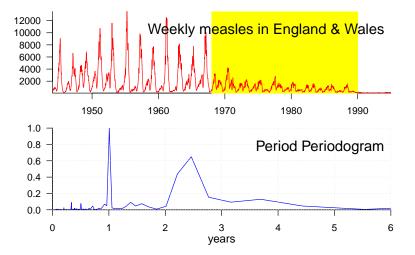
Spectral Density of Temporal Segments



Spectral Density of Temporal Segments



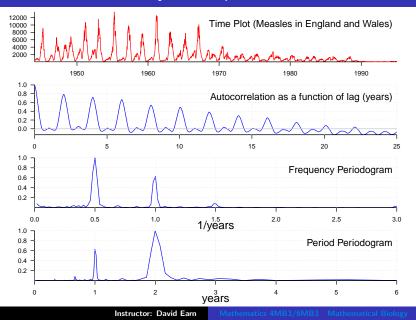
Spectral Density of Temporal Segments



Spectral Density Properties

- Periodogram is discrete Fourier transform of correlogram
- Same information in correlogram and periodogram
- Periodogram usually easier to interpret
- In (R), calculate power spectrum with spectrum()
- The power spectrum $I(\omega_p)$ partitions the variance in the time series with respect to frequency ω_p .
 - Parseval's theorem implies $\frac{1}{N} \sum_{t} (x_t \bar{x})^2 = \frac{1}{2\pi N} \sum_{p>0} I(\omega_p)$. But $\frac{1}{N} \sum_{t} (x_t - \bar{x})^2 = \text{Var}\{x_t\}$, hence $I(\omega_p)/(2\pi N)$ is the proportion of the variance in the time series associated with period $2\pi/\omega_p$. [For details, see Chatfield (2004).]

Basic Time Series Analysis of Epidemic Data



Spectral Density of Temporal Segments

- Pre-war measles
- Post-war pre-vaccination measles
- Vacinnation era measles
- Vacinnation era measles until 1990

Time series analysis functions

R has built-in tools for time series analysis:

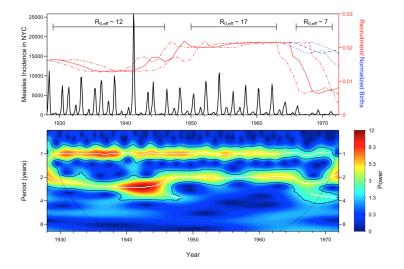
- Time plot: plot() etc.
- Linear filter (e.g., moving average): filter()
- Correlogram (auto-correlation function): acf()
- Periodogram (power spectrum): spectrum()

You will use all of these functions in Assignment 4.

More sophisticated spectral method

- Traditional power spectrum measures frequency content of entire time series.
- Wavelet decomposition is local in time.
 - Reveals changes in the spectrum over time without having to identify distinct temporal segments yourself.
 - Nice intro to wavelet analysis of time series: Torrence and Compo (1998) "A Practical Guide to Wavelet Analysis" Bulletin of the American Meteorological Society 79, 61 - 78
 - $\blacksquare \exists \mathbf{Q} \mathsf{P} \mathsf{packages}$ for wavelet analysis of time series (*e.g.*, WaveletComp, wavelets), and at least one book on wavelet methods in @

Wavelet Spectrum of Monthly Measles in New York City



Krylova & Earn 2013, J. R. Soc. Interface 10, 20130098

Wavelet Spectrum of Weekly Measles in New York City

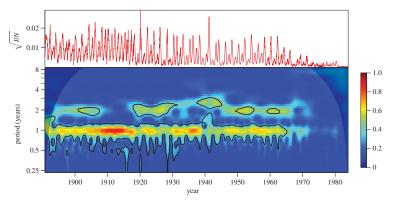


Figure 5. Observed measles dynamics in NYC from 1891 to 1984. (a) Square root of measles case reports, normalized by total concurrent population. (b) Colour depth plot of a continuous wavelet transform of the square root of normalized observed NYC measles cases (colour warmth scales with spectral power and 95% significance contours are shown in black). Shaded regions in the upper left and right indicate the cone of influence.

Hempel & Earn 2015, J. R. Soc. Interface 12, 20150024

Statistical Modelling of Time Series

Instructor: David Earn Mathematics 4MB3/6MB3 Mathematical Biology

Statistical Modelling of Time Series

- Imagine time series $\{X_t\}$ is generated by random processes.
- Simplest case: X_t (number of cases at time t) is simply a random variable with a known distribution,

$$X_t = \mu + Z_t \tag{(*)}$$

where $\mu = \text{time}$ average number of cases and $\{Z_t\} = \text{sequence of random variables with zero mean.}$

- Might be a reasonable model for importation of new, infectious individuals into a focal community.
- Bad model for epidemics: ignores transmission from one individual to another.
 - There must be a correlation between the number of individuals in the focal community who are infected now and the number who will be infected in the near future.

Statistical Modelling of Time Series: AR and MA

- So, imagine that that successive data points in {X_t} are correlated.
- For example, perhaps the data are generated by an autoregressive (AR) process:

 $X_{t}-\mu = \alpha_{1}(X_{t-1}-\mu) + \alpha_{2}(X_{t-2}-\mu) + \cdots + \alpha_{p}(X_{t-p}-\mu) + Z_{t},$

where the α_i are constants that determine the degree of correlation along the time series.

 Alternatively, the data might be generated by a moving average (MA) process:

$$X_t - \mu = \beta_0 Z_t + \beta_1 Z_{t-1} + \dots + \beta_q Z_{t-q},$$

where the β_i are constants that define a weighted average.

Statistical Modelling of Time Series: ARMA

More generally, the data might be generated by an autoregressive moving average "ARMA(p, q)" process:

$$X_{t} - \mu = \alpha_{1}(X_{t-1} - \mu) + \alpha_{2}(X_{t-2} - \mu) + \dots + \alpha_{p}(X_{t-p} - \mu) + \beta_{0}Z_{t} + \beta_{1}Z_{t-1} + \dots + \beta_{q}Z_{t-q}.$$

Statistical Modelling of Time Series: ARIMA

Finally, an autoregressive integrated moving average "ARIMA(p, d, q)" model includes weighted differences of the time series:

$$X_{t} - \mu = \alpha_{1}(X_{t-1} - \mu) + \alpha_{2}(X_{t-2} - \mu) + \dots + \alpha_{p}(X_{t-p} - \mu) + \gamma_{1}(X_{t-1} - X_{t-2}) + \gamma_{2}(X_{t-2} - X_{t-3}) + \dots + \beta_{0}Z_{t} + \beta_{1}Z_{t-1} + \dots + \beta_{q}Z_{t-q}.$$

- The "I" in ARIMA refers to the original time series X_t, which is an "integrated" version of the differenced time series.
- Technically, an ARIMA model is just an ARMA model with differently labelled coefficients, but explicit differences are often helpful conceptually (*e.g.*, they can "stationarize" a time series).

What kind of process generated our data?

- How can we tell if our data were generated by such a process? Can we identify an AR(p), MA(q) or ARMA(p,q) process?
- Compare time plots of these processes with time plot of our data? (Comparison by eye often challenging/unreliable.)
- Compare autocorrelation functions (correlograms) of these processes with correlogram of our data? (Better.)
- Compare power spectra (periodograms) of these processes with periodogram of our data? (Even better.)
- Compare wavelet spectra of these processes with wavelet spectrum of our data? (Better yet.)

Statistical Modelling of Time Series: ARMA fitting

- Looking at the power spectra of ARMA models would be instructive.
- But is there a better approach to discovering if an ARMA model could explain our data?
- Find the best fit ARMA parameters by minimizing the residual sum of squares. e.g., for an AR model, minimize:

$$S = \sum_{t=p+1}^{N} \left[(x_t - \mu) - \alpha_1 (x_{t-1} - \mu) - \cdots - \alpha_p (x_{t-p} - \mu) \right]^2.$$

- More generally, we can find the best fit parameters of an ARIMA(p, d, q) model
 - Non-trivial, but there are standard methods
 - Compare models with Akaike Information Criterion (AIC)
- See Earn (2009) review article for more discussion of this.