

8 Epidemic Data

9 Epidemic Data II

10 Epidemic Data III

11 Epidemic Data Tools

12 Epidemic Data Tools II



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:**

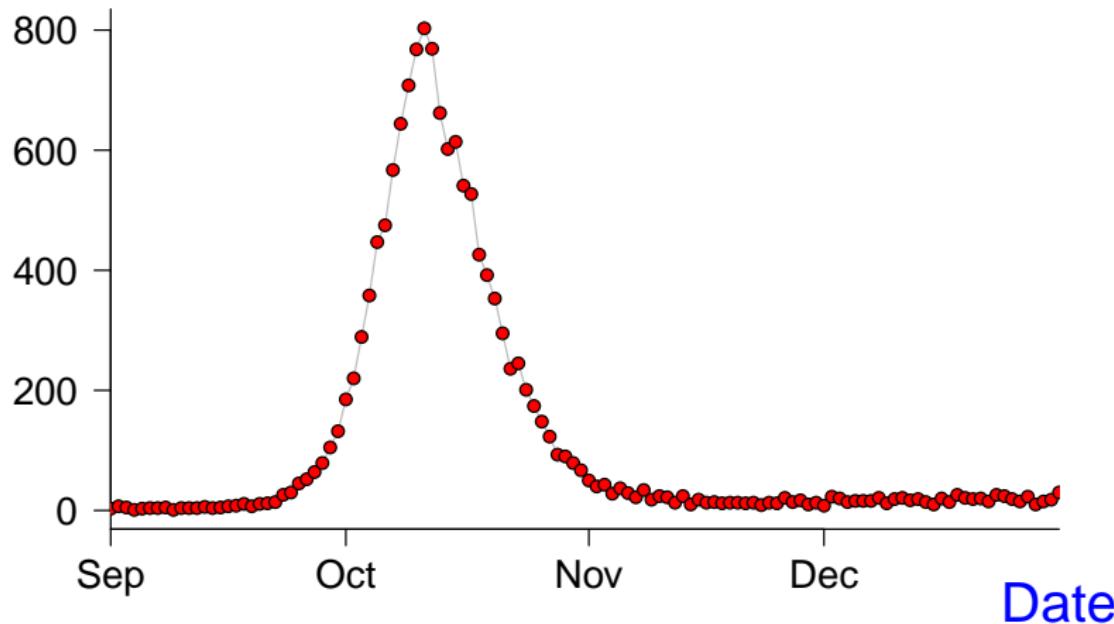
Due Monday 5 February 2018 in class (and by e-mail) at 11:30am.

- **Midterm test:**

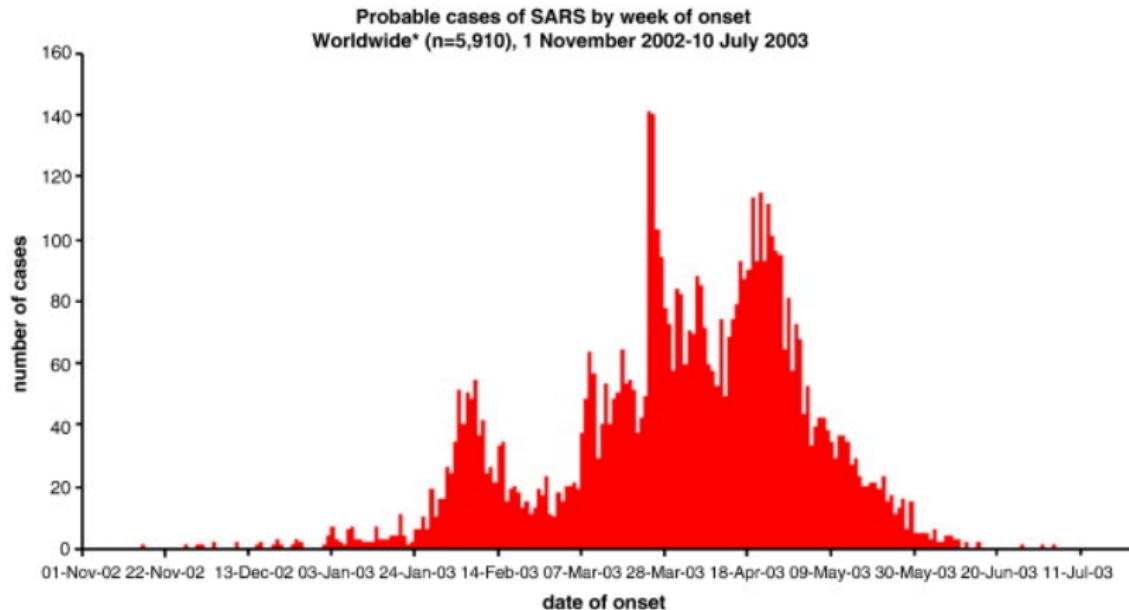
- *Date:* week of 5–9 March? or 12–16 March?
- *Time:* TBA
- *Location:* TBA

P&I Mortality, Philadelphia, 1918

P&I Deaths

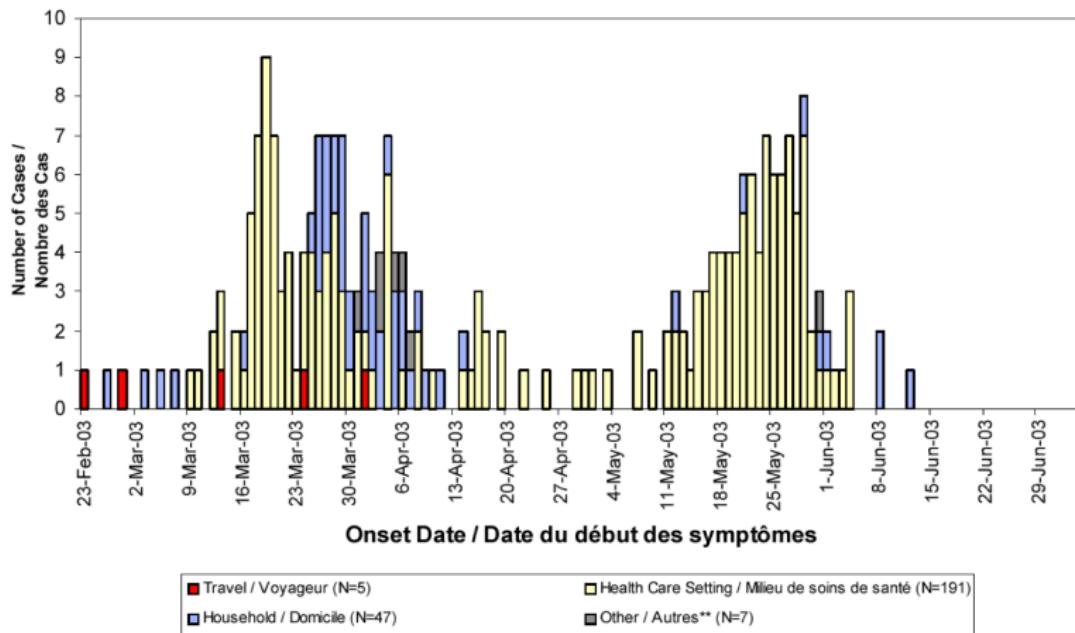


SARS in 2003 (Worldwide)



*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.

SARS in 2003 (Toronto)

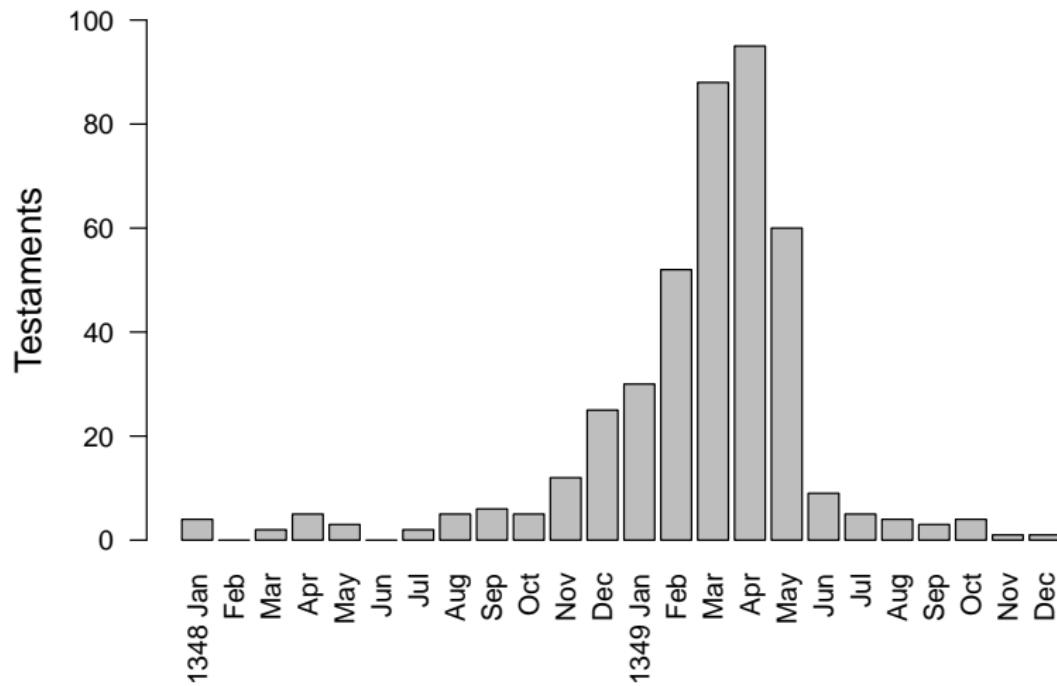


$N = 249$ (of 250 reported)

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



London Bill of Mortality, 26 Sept to 3 Oct 1665

The Diseases and Casualties this Week,



Bortive	6
Aged	50
Ague	1
Apoxyxie	2
Chilblains	42
Chirfomes	11
Cold	1
Consumption	99
Convallion	63
Cough	1
Dropyle	22
Drown'd at St. Martin in the Fields	1
Feaver	268
Fiftula	2
Flor and Small-pox	4
Flux	1
Found dead in the Fields at St. Mary Iffington	1
Males	687
Christned Females	78
In all	146
Decreased in the Burials this Week	1837
Parishes clear of the Plague	7
Males	3212
Baried Females	3248
In all	6460
Plague— ⁵⁵³³	

London 45		From the 26 of September to the 3 of October,		1665	
Bar. Reg.	Bar. Reg.	Bar. Reg.	Bar. Reg.	Bar. Reg.	Bar. Reg.
St. Leon Woodforne	16 15	St. George Bosophilene	1 1	St. Martin Ludgate	12 10
Albionnes Baring	40 34	St. George in St. Paul's	16 25	St. Martin Orgar	3 3
Albionnes Crosse	41 41	St. James Duke place	27 23	St. Martin Outwich	6 5
Albionnes Hoyleane	7 17	St. James Garlickhitche	16 13	St. Martin Vintry	44 44
Albionnes Little	3 17	St. John Bowie	11 10	St. Michael Fridayes	4 4
Albionnes Newgate	3 17	St. John Evangelist	12 9	St. Michael Oldhorne	4 4
Albionnes the Wal	33 28	St. Katharine Coleman	20 12	St. Michael Buffawen	10 7
S'w'lings	13 5	St. Katharine Creechard	14 29	St. Michael Cornhill	4 3
S' Andrew Hubbard	16 44	St. Lawrence Jewry	5 5	St. Michael Queenhithe	15 12
S' Andrew Undershaft	20 44	St. Margarete Pannay	14 10	St. Michael Querne	25 23
S' Andrew Werches	18 22	St. Mary Aldermary	16 33	St. Michael Royal	4 3
S' Ann Blackfriars	57 30	St. Mary Pavill	5 4	St. Michael Woudborne	10 9
S' Antholme Parb	7 4	St. Margaret Lothbury	7 6	St. Michael Brendford	6 3
S' Antholme Parb	12 12	St. Margarete Mofes	1 1	St. Nicholas Acon	4 2
S' Bermondsey Exchange	12 7	St. Margarete Newgate	13 3	St. Nicholas Colehale	2 2
S' Bermonsey Fyneke	12 7	St. Mary Abbat	12 3	St. Nicholas Olaves	9 8
S' Bermer Gracechurch	4 2	St. Mary Abchurch	12 1	St. Olave Hartfurd	1 1
S' Bermer Paulwharf	15 7	St. Mary Aldermanbury	14 4	St. Olave Jewry	13 14
S' Bermer Shereches	2 11	St. Mary Aldemary	4 1	St. Olave Newfowre	7 4
S' Bouth Billinggate	8 8	St. Mary le Bow	1 1	St. Pancras Soperiane	1 1
S' Chichester	44 59	St. Mary le Bow	6 1	St. Peter Cheape	1 1
S' Chichester	12 12	St. Mary Magdalene	10 6	St. Peter Cornhill	6 6
S' Cirencester Bathsheba	7 2	St. Mary Magdalene	4 4	St. Peter Paulwharf	10 10
S' Dionis Backbor	9 2	St. Mary Mauncen	4 4	St. Peter Poer	
S' Danfys East	18 24	St. Mary Somerses	4 4	St. Steven Colemanfie	33 28
S' Edmund Lambreth	3 1	St. Mary Savynge	5 5	St. Steven Walbrook	3 2
S' Eichelbrough	7 4	St. Mary Woolchurche	7 7	St. Thomas Aquin	6 5
S' Faith	8 6	St. Mary Woolnoth	7 5	St. Thomas Asyng	10 9
S' Fister	8 6	St. Martin Iwengengiane	2 2	Trinity Parf	
S' Gabriel Fincharch	3 3				
described in the 16 Parishes without the Wall— ⁴⁵			39 Buried	1149 Plague	948
S' Andrew Holborn	173 151	S' Bowbly Aldgate	371 338	Sextuors Southwark	364 352
S' Bartholomew Great	157 151	S' Bowbly Billingsgate	333 121	S' Sepulchre Parf	117 95
S' Bartholomew Leie	7 75	S' Dunstans Well	59 59	S' Thomas Southwark	40 20
S' Bridge	92 67	S' George Southwark	140 153	Trinity Minotors	24 20
S' Bridewell Prencipall	33 23	S' Gilre Crispologe	196 200	At the Pelehouse	8 8
S' Bouth Alderger	71 64	S' Olave Southwark	378 208		
described in the 16 Parishes within the Wall— ⁴⁵			2258 Buried	2258 Plague	1912
S' Cate in the Brads	95 78	Lambeth Parf	49 39	S' Mary Allington	35 31
S' Chichester	14 12	S' Leonard Shoreditch	95 91	S' Mary Whitemarsh	38 301
S' James Clerkenwe	48 48	S' Margarete Remyndene	128 106	Rocholme Parf	21 18
S' Kath. near the Tower	153 9	S' Mary Newington	81 81	Soppy Parf	67 631
described in the 12 Parishes in Middlesex and Sury			40 Buried	1623 Plague	1469
S' Clement Danes	13 8 10	S' Martin in the fields	109 143	S' Margarete Welfmire	309 297
S' Paul Covenard	35 84	S' Mary Savoy	119 16	Veretey at the Petham	4
described in the 5 Parishes in the City and Liberties of Welfmire			690 Buried	590 Plague	590

M

Mortality Bills are typically handwritten

London 29: From the 4 th of July to the 11 th of the same 1665																	
Buried			Plag.			Buried			Plag.			Buried			Plag.		
St Alban Woodstreet	2		St Clement Eastchapp-			St Margaret Newfisht			St Michael Crookedla.			St Michael Queenhithe	4		3		
Alhallows Bark-	2		St Dionys Backchurch	1		St Margret Patrons			St Michael Quern	7		St Michael Royal					
Alhallows Breadfreet			St Dunstans East	2		St Mary Abchurch			St Michael Woodstreet			St Mildred Breadfreet					
Alhallows Great	1		St Edmund Lumbardst.	2		St Mary Aldermanbury			St Mildred Poultrey			St Nicholas Acons					
Alhallows Honilane			St Ethelborough	2		St Mary Alde mary			St Nicholas Coleababy			St Nicholas Olaves					
Alhal ows Less	1		St Faiths	1		St Mary le Bow			St Olave Hartfreet			St Olave Jewry					
Alhallows Lombardstr.			St Gabriel Fenchurch			St Mary Bothaw			St Olave Silverstreet			St Pancras Soperlane	+		1		
Alhallows Staining	1		St George Botolphane			St Mary Colechurch			St Peter Cheap			St Peter Cornhill					
Alhallows the Wall	4	3	St Gregories by St. Paul			St Mary Hill			St Peter Paulisharf			St Peter Paullwharf					
St Alphege			St Hellen	2		St Mary Mag. Milkstr.			St Peter Poor			St Steven Colemanstr.	2		1		
St Andrew Hubbard			St James Dukes place	1		St Mary Mag. Oldfish			St Steven Walbrook			St Steven Walbrook					
St Andrew Underhafe	3		St James Garlickhithe			St Mary Mounthaw			St Swithin			St Thomas Apostle	1		1		
St Andrew Wardrobe			St John Baptis			St Mary Summercer	2		Trinity Parish			Trinity Parish					
St Anne Alderfae	1		St John Evangelist			St Mary Straining			St Vedast alias Fosters								
St Anne Blackfriars	7	6	St John Zichary			St Mary Woolchurch											
St Anthonies Parish			St Katharine Coleman	1		St Mary Woolnoth											
St Aufins Parish			St Katharine Creechur.			St Martins Iremonger											
St Barthol Exchange	1		St Lawrence Jewry			St Martins Ludgate	2										
St Benner Fynck			St Lawrence Pountney			St Martins Orgars											
St Benner Gracechurch	2		St Leonard Eastcheap			St Martins Outwich	1										
St Benner Paulisharf	7		St Leonard Fosterlane			St Martins Vintrey	1										
St Benner Sherehog			St Magnus Parish	1		St Matthew Frydaystr.											
St Boroloph Billinggate			St Margaret Lothbury			St Michael Balliflaw	5	4									
Christ Church			St Margaret Moles			St Michael Cornhill											
Se Christophs	2	3															
Christened in the 27 Parishes within the walls												86	Plague		28		
St Andrew Holborn	66	40	St Boroloph Aldergate	11	7	St George Southwark	13		St Sepulchres Parish	117	81						
St Bartholomew Great	4	4	St Boroloph Aldgate	27	4	St Giles Cripplegate	103	47	St Thomas Southwark	7	5						
St Bartholomew Less			St Boroloph Bishopgate	37	20	St Olave Southwark	20		Trinity Minories								
St Bridge	24	14	St Dunstan West	19	9	St Saviour Southwark	21	1	At the Pesthouse	6	6						
Bridewell Preinct																	
Christened in the 15 Parishes without the walls												473	Plague		273		
Christ Church			St Kath.near the Tower	7	1	St Mary Islington	3	2	St Paul Shadwell								
St John at Hackney	1		Lambeth Parish	7	1	St Mary Newington	3	2	Rotherhithe Parish	7	3						
St Giles in the Fields	268	213	St Leonard Shoreditch	21	13	St Mary Whitechapel	16	3	Stepney Parish	47	1						
St James Clerkenwel	5		St Magdalens Bermond.	14													
Buried in the 27 Parishes within the walls												455	Plague		286		

But handwriting is usually very clear

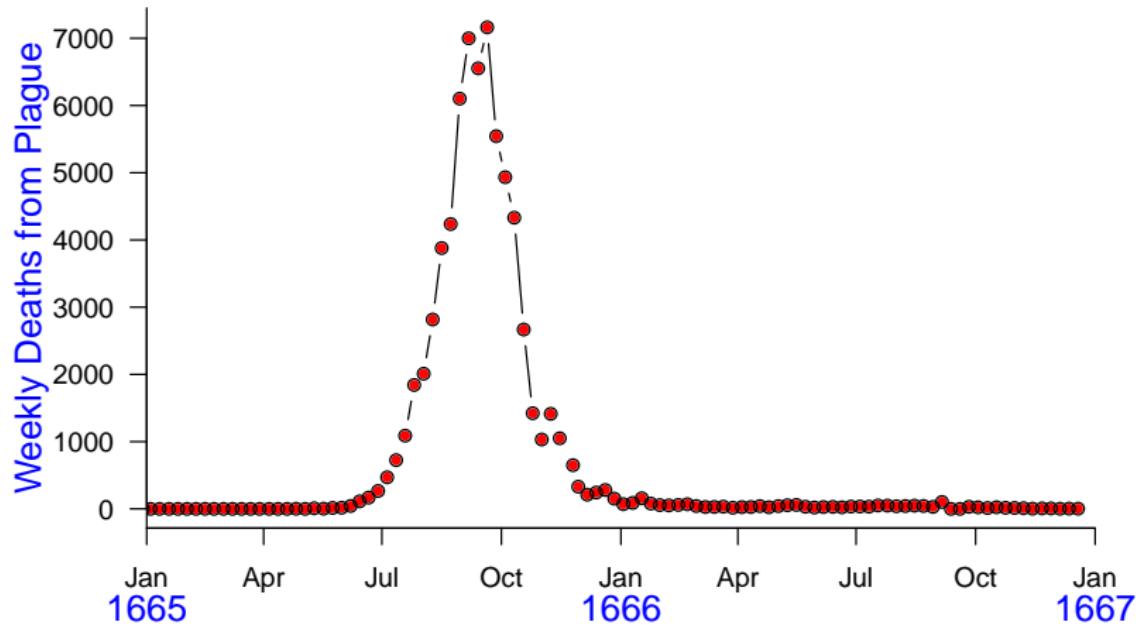
A historical ledger page from London, dated 29th [month]. The page is divided into columns for 'Buried.' and 'Plag.'. The data is organized by parish.

	Buried.	Plag.
St Alban Woodstreet	2	1
Alhallows Bark-	2	
Alhallows Breadstreet	1	
Alhallows Great	1	

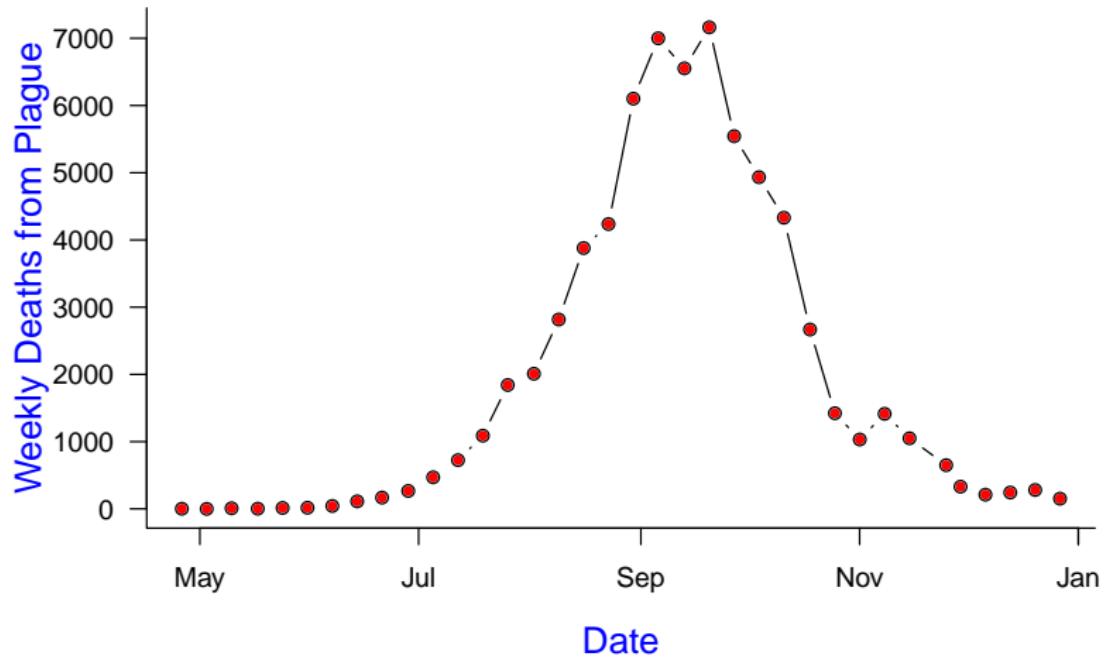
But handwriting is usually very clear

St Christopher's —————— Christened in 97 the Parishes :		
St Andrew Holborn ——————	66	40
St Bartholomew Great	+	+
St Bartholomew Less ——————		
St Bridget ——————	24	14
Bridewell Precept ——————	1	1
Christened in the 16 Parishes :		

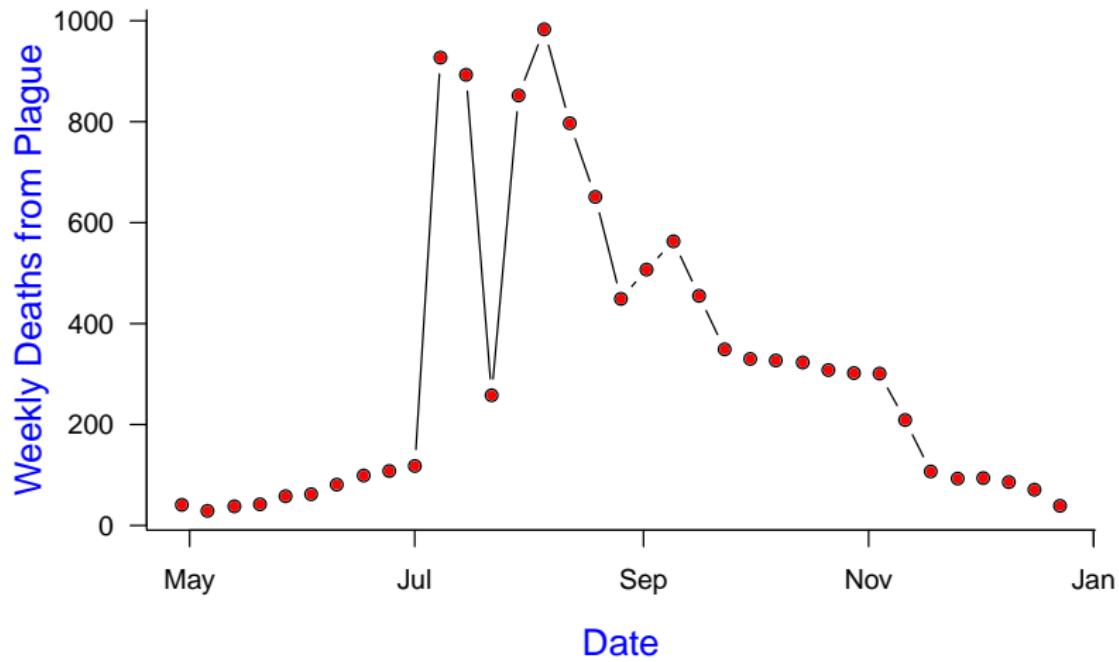
The Great Plague of London, 1665



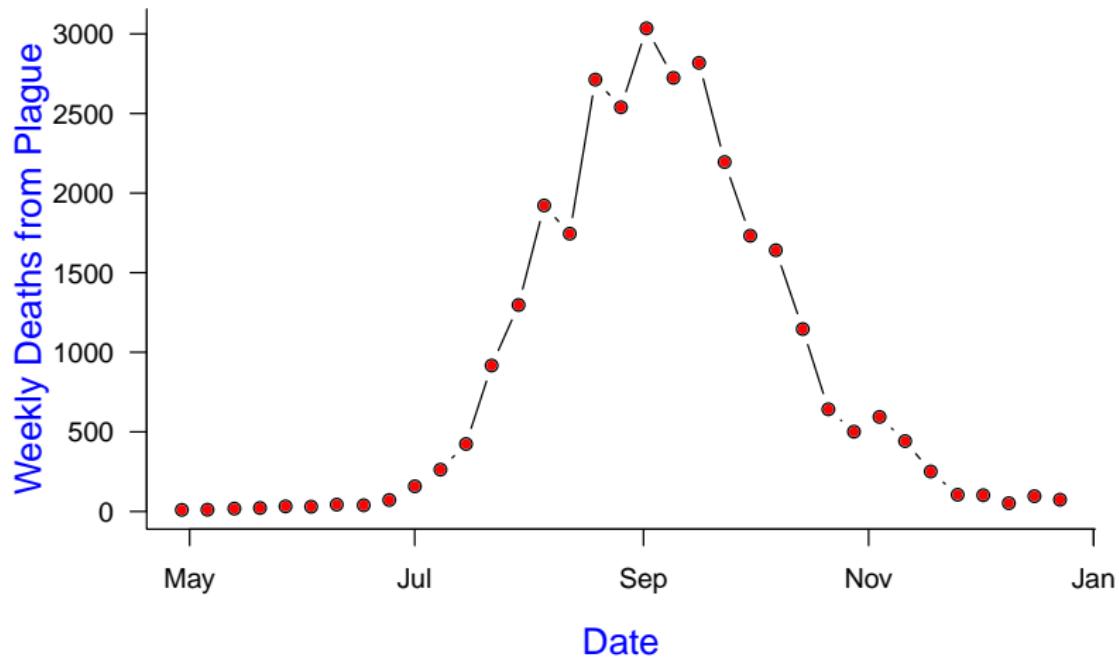
The Great Plague of London, 1665



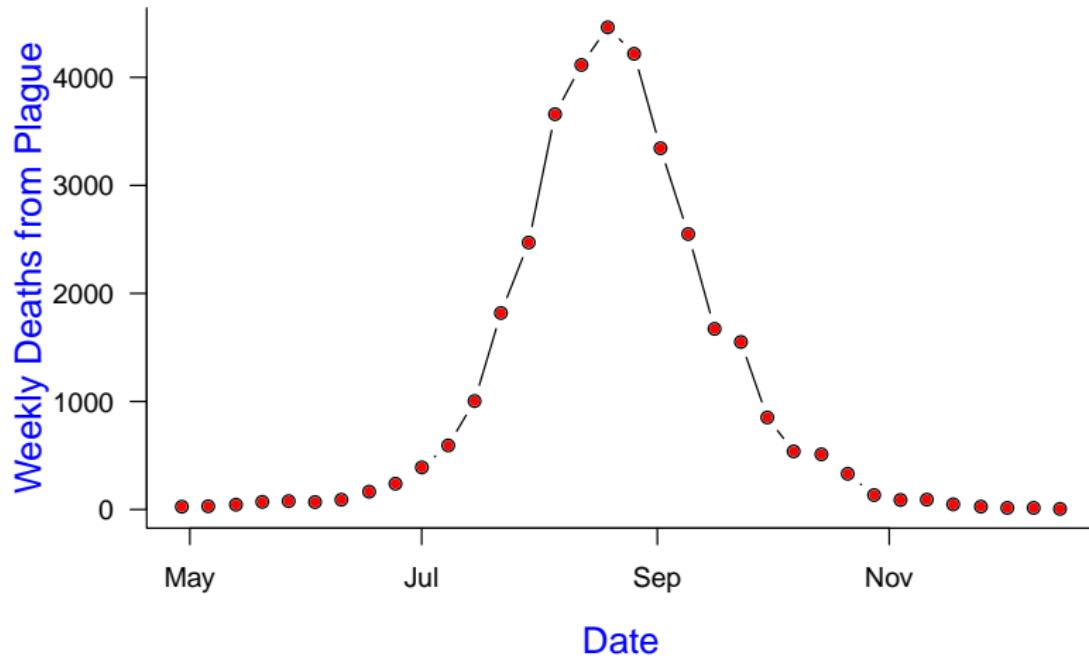
London Plague of 1593



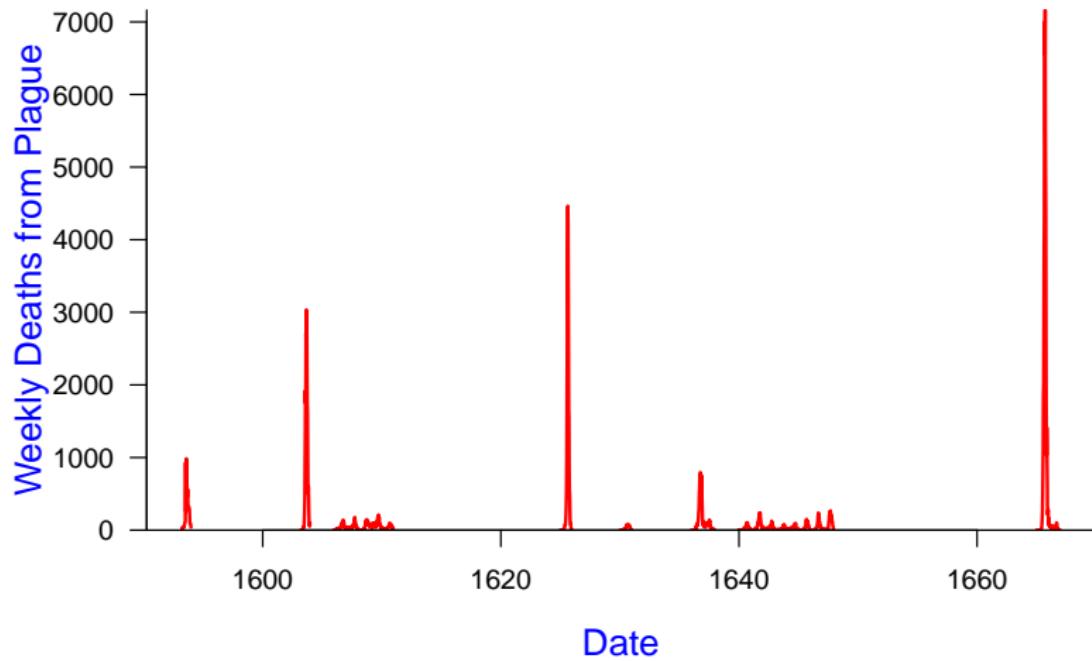
London Plague of 1603



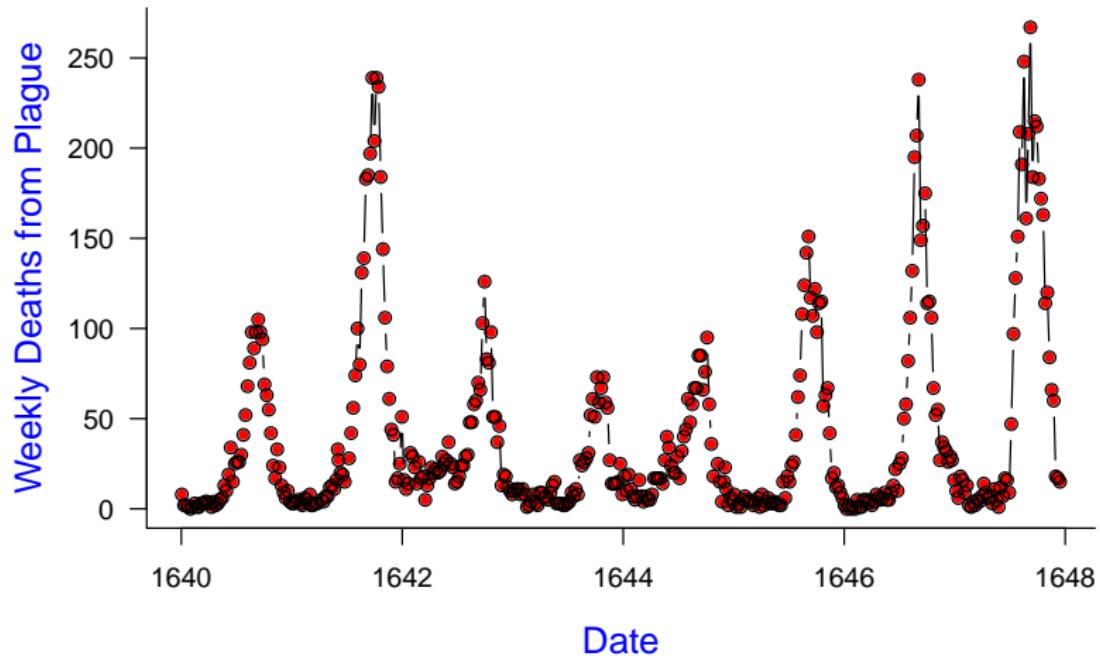
London Plague of 1625



Weekly Deaths from Plague in London, 1592–1666



Weekly Plague in London, 1640–1648



Some Plague Facts

- Plague epidemics recorded from Roman times to early 1900s.
- $\gtrsim 1/3$ Europe's population died in "Black Death" of 1348
 - ~ 300 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](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:

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

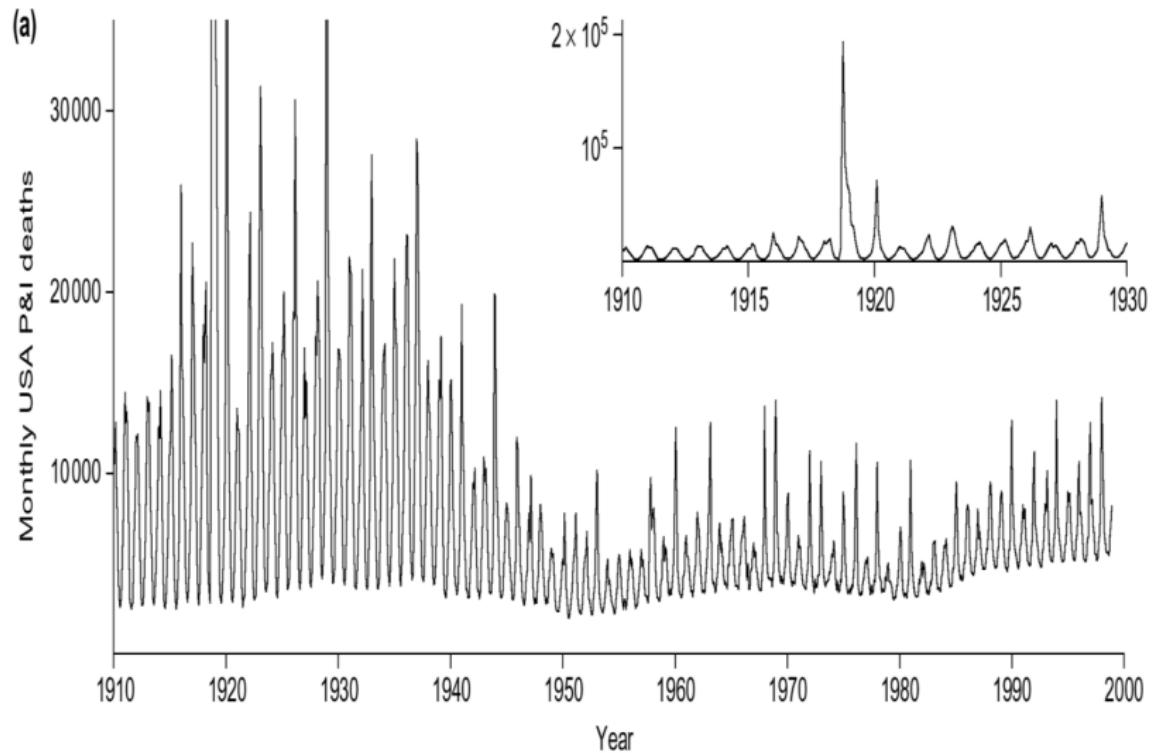
■ Midterm test: We agreed on:

- *Date:* Thursday 8 March 2018
- *Time:* 7:00pm to 9:00pm
- *Location:* TBA

Visualization of entire course of the Great Plague

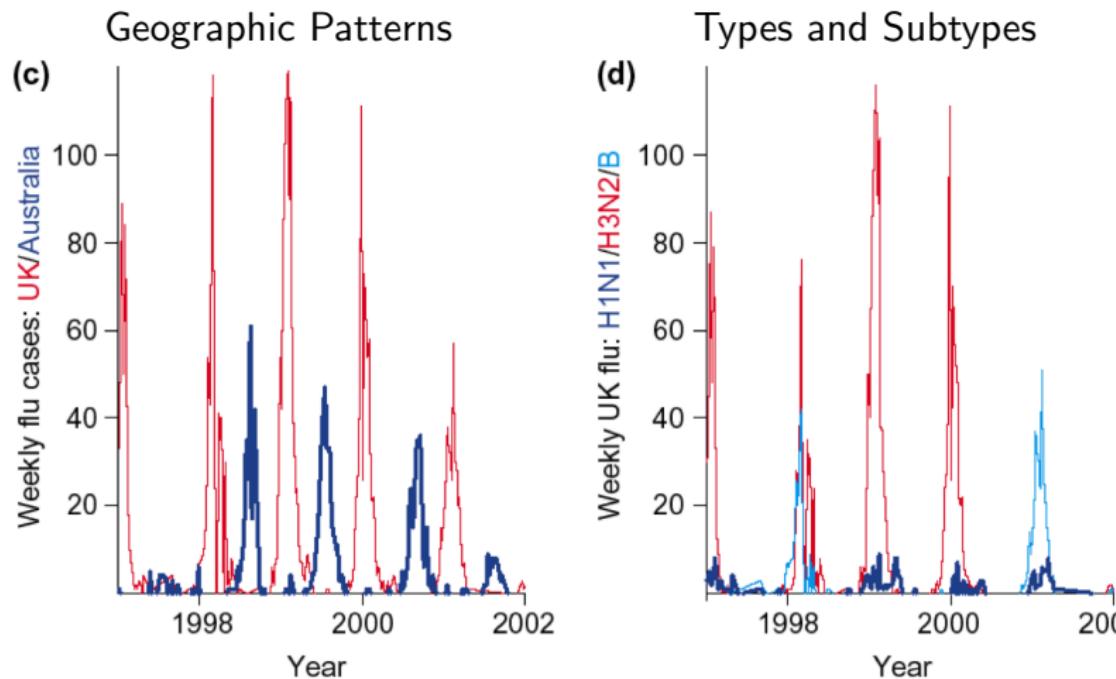
- What happened after initial spatial spread?
- Visualize full spatial epidemic structure
- Show magnitude of epidemic in each parish with cylinder.
- **Epidemic Visualization** (EpiVis) software by Junling Ma.

P&I mortality in U.S.A., 1910–1998



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

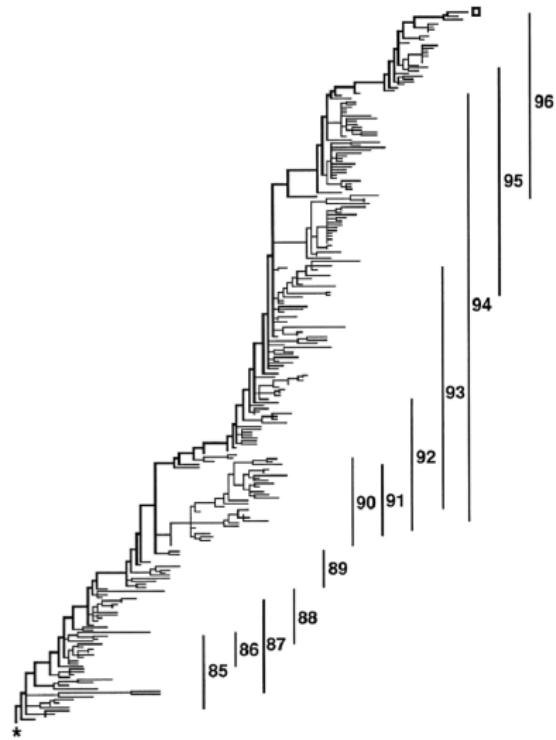
Influenza Incidence Patterns (lab confirmed)



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

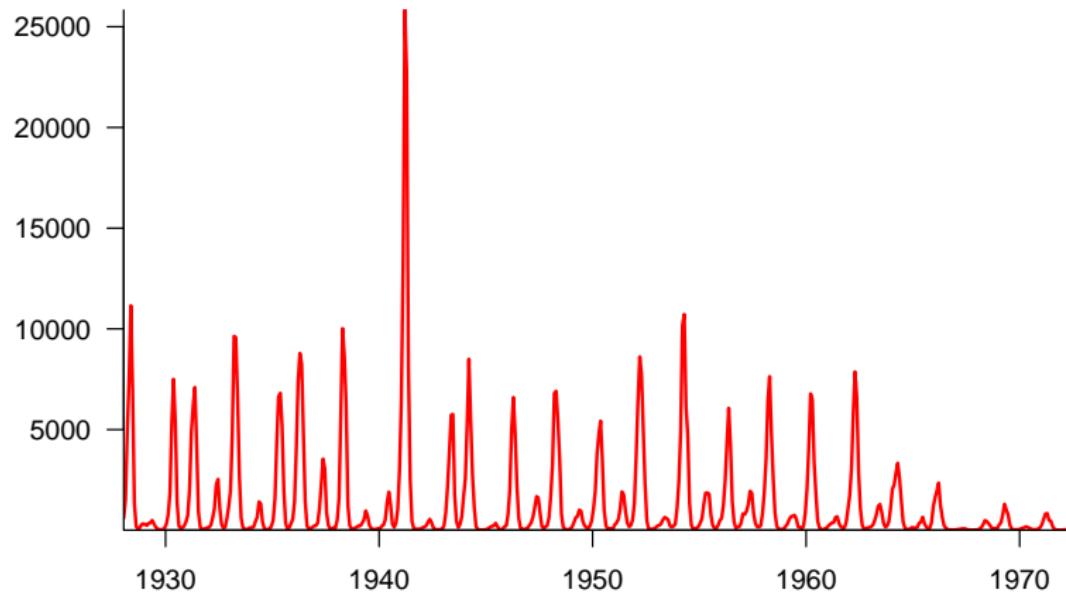
Influenza Evolution

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



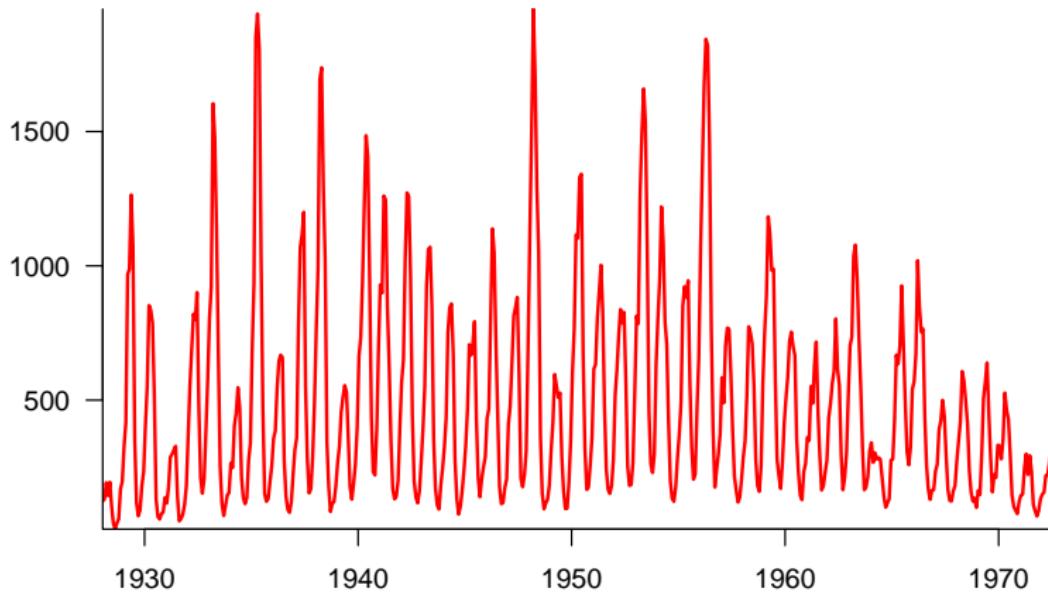
Measles in New York City, 1928–1972

Monthly Cases



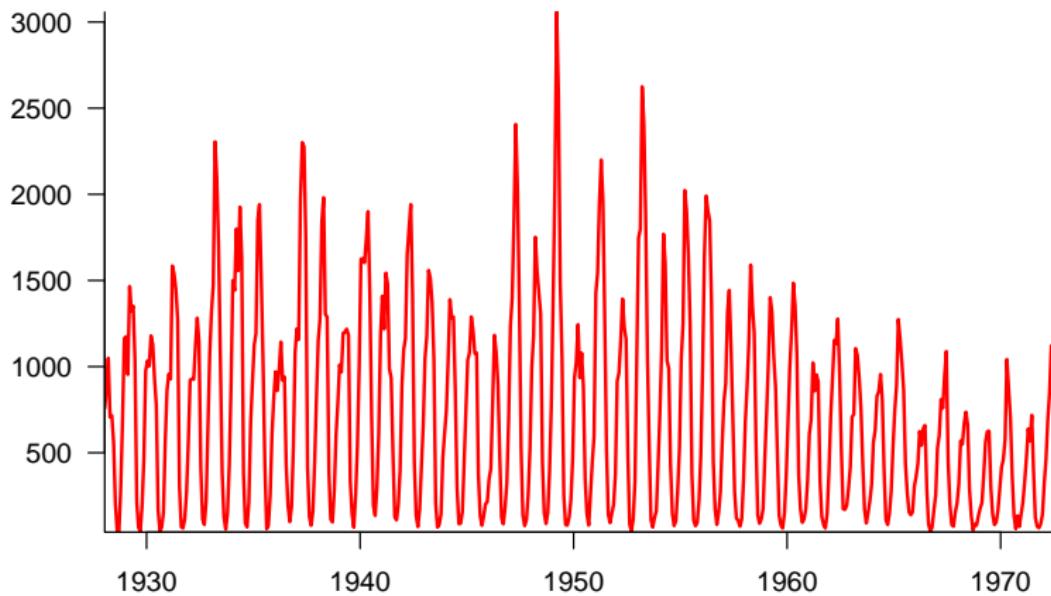
Mumps in New York City, 1928–1972

Monthly Cases

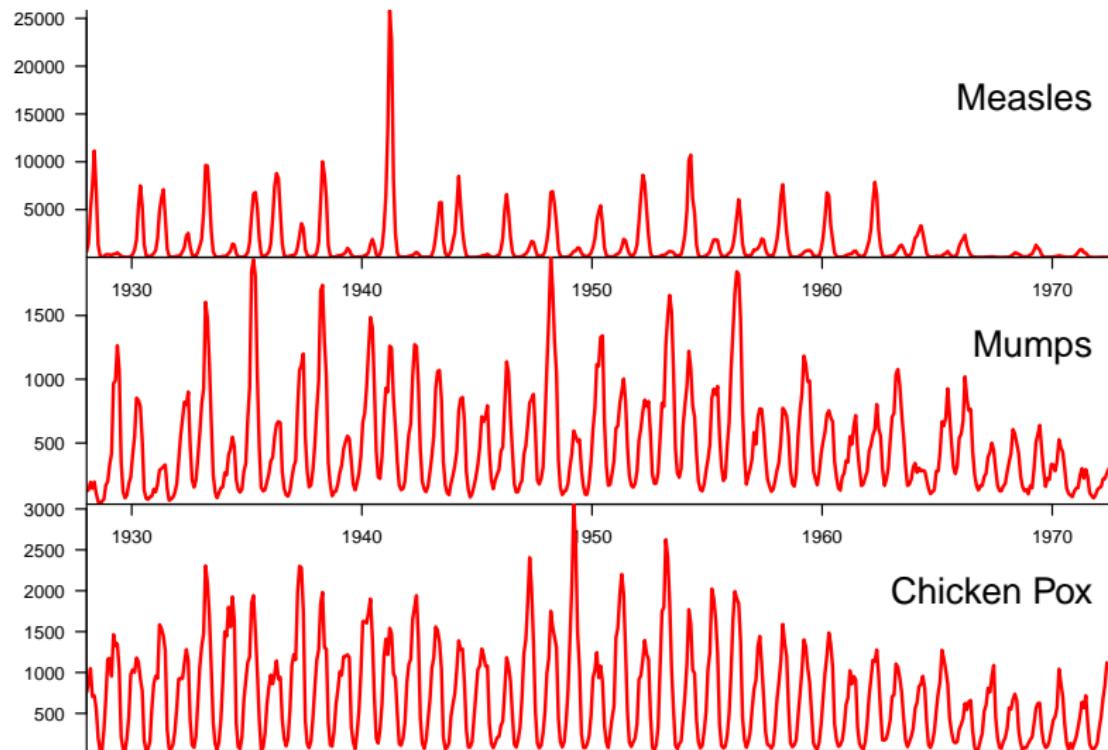


Chicken Pox in New York City, 1928–1972

Monthly Cases

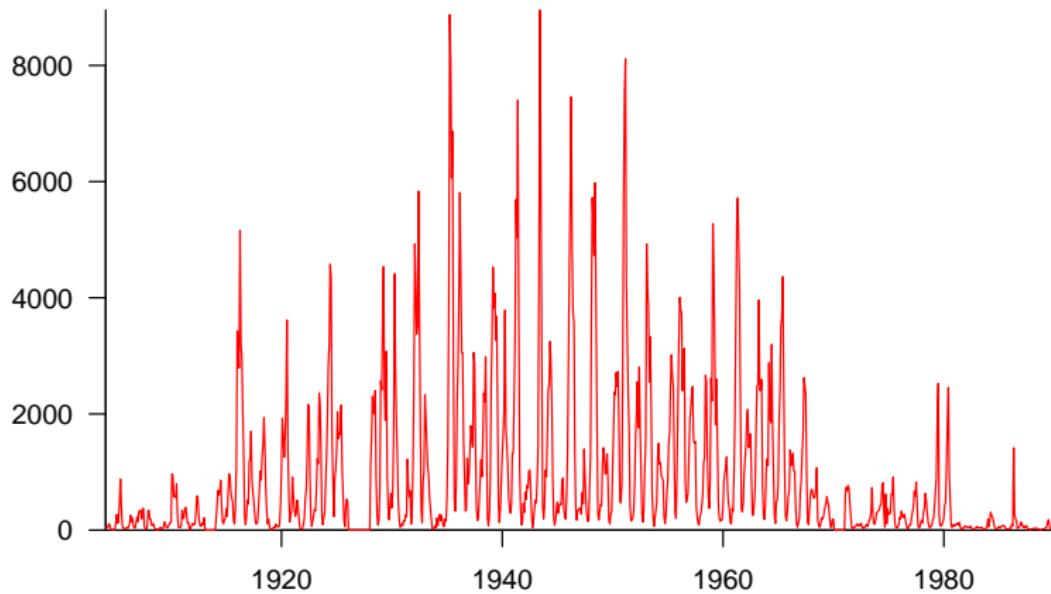


Childhood diseases in New York City, 1928–1972



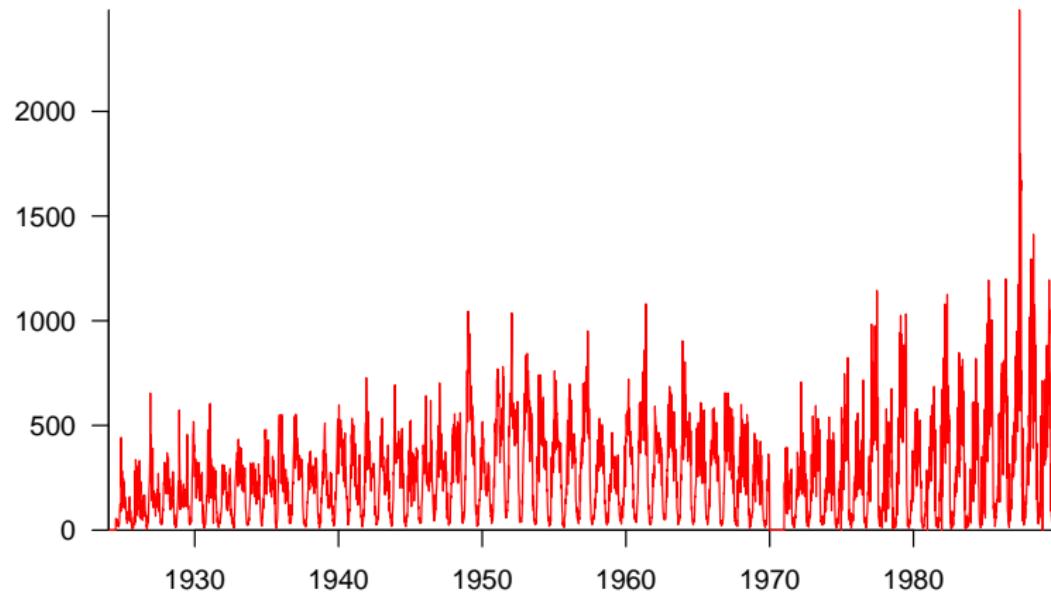
Measles in Ontario, 1904–1989

Monthly Cases



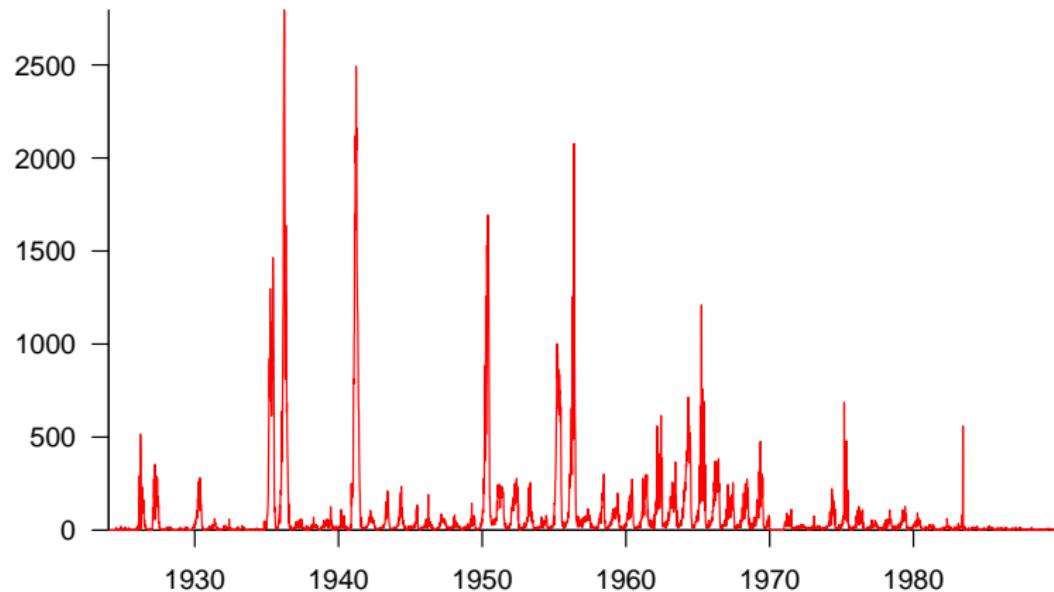
Chicken Pox in Ontario, 1924–1989

Monthly Cases



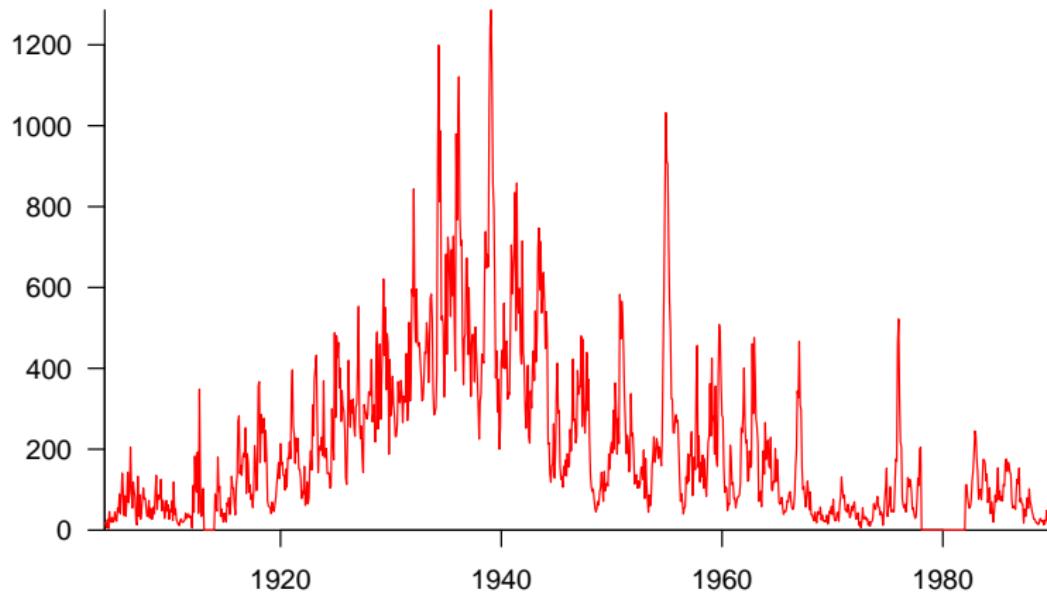
Rubella in Ontario, 1924–1989

Weekly Cases

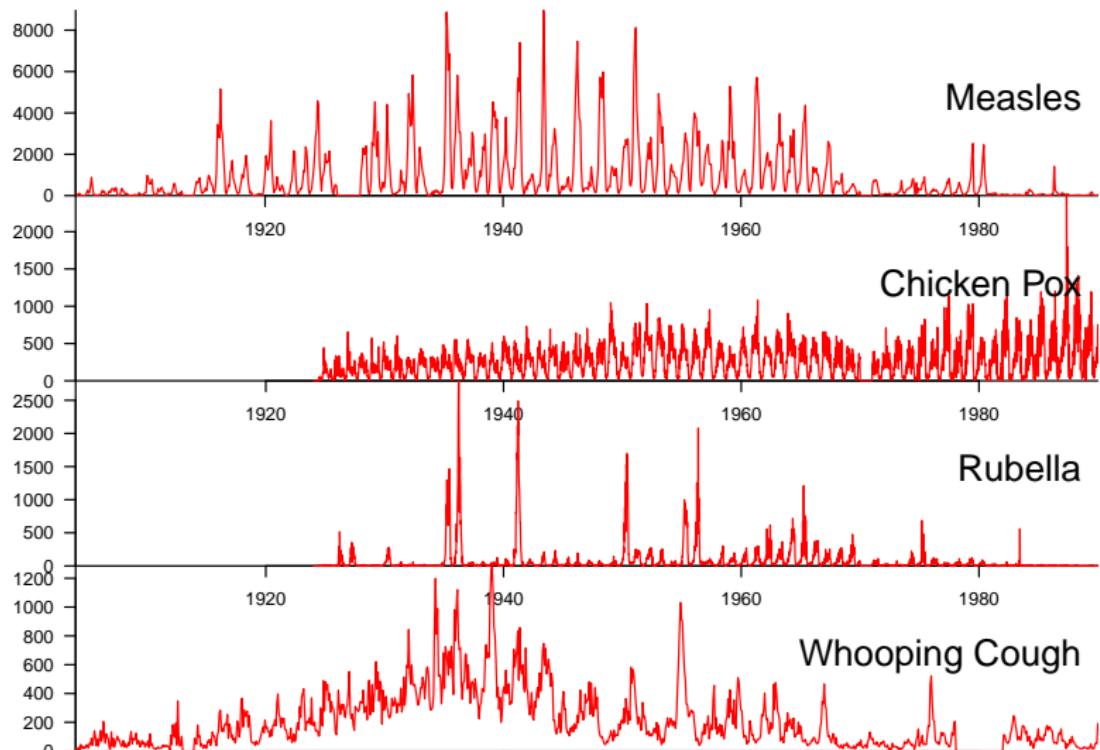


Whooping Cough in Ontario, 1904–1989

Monthly Cases



Childhood diseases in Ontario, 1904–1989



Ontario Disease Notification Data

Province of Ontario

YEAR: 1939 COUNTY..... MUNICIPALITY.....

Month	Week End.	COUNTY.....												MUNICIPALITY.....													
		CSM		C.P.		DIP.		DYS. A/B		EN. LETH.		ERYS.		G.C.		FLU.		INF. JAUN.		G.M.		MEAS.		MUMPS		PARA. TYPH.	
		C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D	C	D
Jan.	7 1			452	1	3	0	1	0			5	1	101	0	8	1	17	0	17	0	670	1	56	0	2	0
	14 2	2	1	490	0	8	0					5	0	82	0	21	1	18	0	18	0	850	0	92	0	1	0
	21 3	2	1	511	0	9	3			0	1	5	0	89	0	16	2	26	0	22	0	932	0	98	0		
	28 4	1	0	384	0	2	0					2	0	73	0	164	0	10	0	28	0	933	1	24	0		
	Total	5	2	1931	1	27	3	1	0	0	1	17	0	218	4	71	0	65	2	3385	2	270	0	3	0		
Feb.	4 5			355	0	7	1	1	0			3	0	83	0	57	1	24	0	25	0	1335	1	110	0	2	0
	11 6	2	1	363	0	1	0	1	0			7	0	82	0	27	1	41	1	29	0	1033	0	91	0	1	0
	18 7	2	1	354	1	2	0					4	1	68	0	103	1	35	0	44	0	1161	0	59	0		
	25 8	1	1	308	0	2	0					9	0	560	1	77	0	19	0	28	0	999	0	73	0		
	Total	5	3	1980	1	27	3	1	0			23	1	23	1	3	1	12	0	158	1	3381	0	3	0		
Mar.	4 9	1	1	271	0	7	1	3	1			7	0	93	0	114	19	21	0	40	0	131	2	109	0	1	0
	11 10			239	0	7	0	2	0			8	1	61	0	137	18	31	0	32	0	845	0	91	0	2	0
	18 11			166	0							6	0	66	0	1322	6	5	0	59	0	969	2	69	0	1	0
	25 12	1	2	236	0	1	0	1	0			7	0	63	0	806	16	9	0	20	0	879	0	120	0	case	PAH
	Total	8	3	118	0	15	1	6	1			28	1	283	0	613	4	66	0	151	0	353	1	389	0	34	0
Apr.	1 13	2	0	139	0	3	0	1	0			8	0	95	0	667	6	1	0	24	0	950	0	89	0	3	0
	8 14	2	0	162	0	1	0	1	0			5	0	67	0	731	22			14	0	790	0	65	0	1	0
	15 15	2	0	108	0	1	0			0	1	11	0	41	0	529	16	2	0	16	0	745	0	56	0		
	22 16	1	1	134	0	2	0	1	0	1	1	6	0	64	0	245	8	2	0	26	0	845	0	54	0		
	29 17	5	1	167	0	4	0	2	0	2	1	3	0	55	0	124	9	2	1	13	0	746	1	120	0		
	Total	12	2	110	0	10	0	3	0			33	0	312	0	216	1	1	0	24	0	450	0	334	0	47	0

Dominion Bureau of Statistics Disease Notification Data

VITAL STATISTICS BRANCH - COMMUNICABLE DISEASE SECTION

Cases of ~~Influenza~~ Reported by Provincial Health Departments, Year 1924

WEEK ENDING	P.E.I.	N.S.	N.B.	QUE.	ONT.	MAN.	SASK.	ALTA.	B.C.	CANADA
	W19-20	W19-20								
Jan 5		11					1			12
2	12	29					18			47
3	19	37					32			69
4	26	75 152		68	181	36	13 64	97	4 88 602	
5 FEB 2	12	1					53			66
6	9	5					40			45
7	16	31					14			45
8	23	- 2 50	1 2	267	202	48	4 111	116	1 7 797	
9 MAR 1		2					21			23
10	1						9			9
11	15	3					11			14
12	22	60					34			94
13	29	2 61		144	140	52	15 90	15	7 17 515	
14 APR 5		9					11			20
15	12	1					12			13
16	19	26	1				8			35
17	26	14 50	3 4	42	140	39	16 47	67	5 33 394	
18 MAY 3		26					2			28



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

Announcements

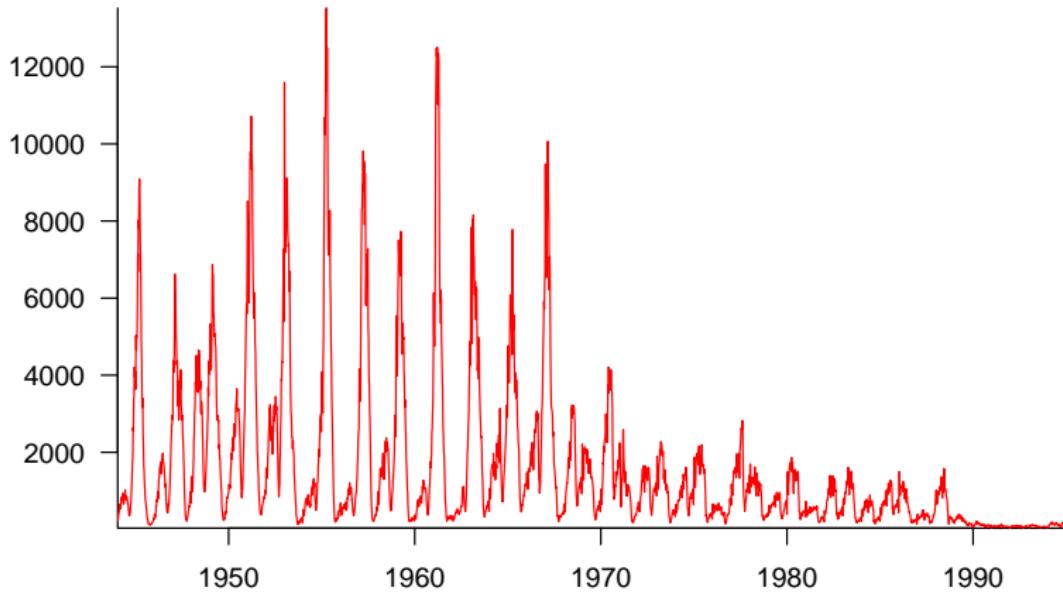
- Comment from TA on Assignment 1:
“For a few of the groups, I would recommend that they look over the work that their group members have done. Question 2c and 2d in particular were closely related and there were a few obvious cases where the students had not communicated with each other.”
- **Assignment 2:**
Due Monday 5 February 2018 in class (and by e-mail) at 11:30am.
- **Midterm test:** We agreed on:
 - *Date:* Thursday 8 March 2018
 - *Time:* 7:00pm to 9:00pm
 - *Location:* TBA

Recurrent epidemics of childhood infections

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

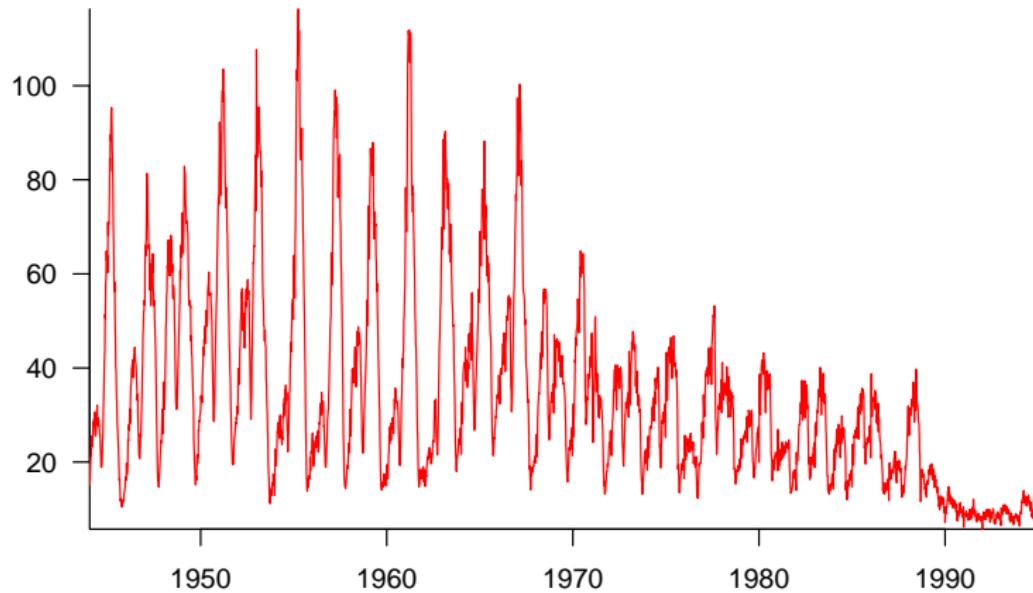
Measles incidence in England and Wales, 1944–1995

Weekly Cases



Measles incidence in England and Wales, 1944–1995

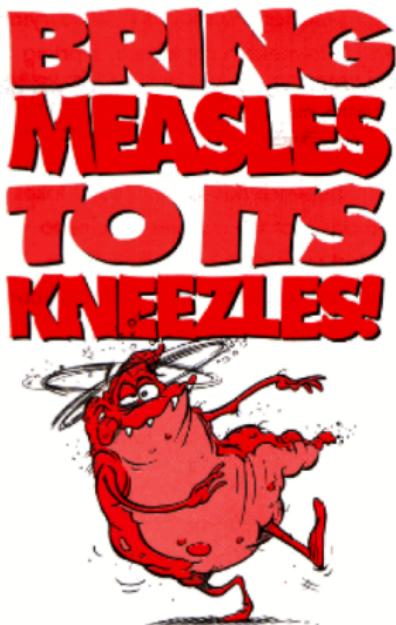
Sqrt(Weekly Cases)



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!

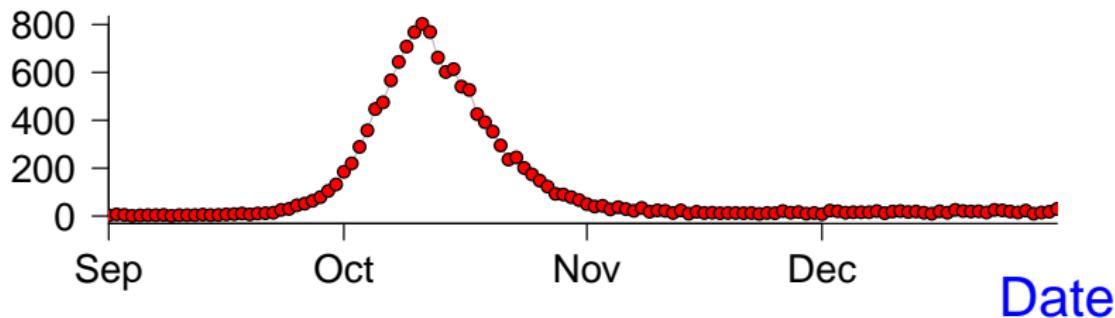
Modelling population dynamics childhood infections

- 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.

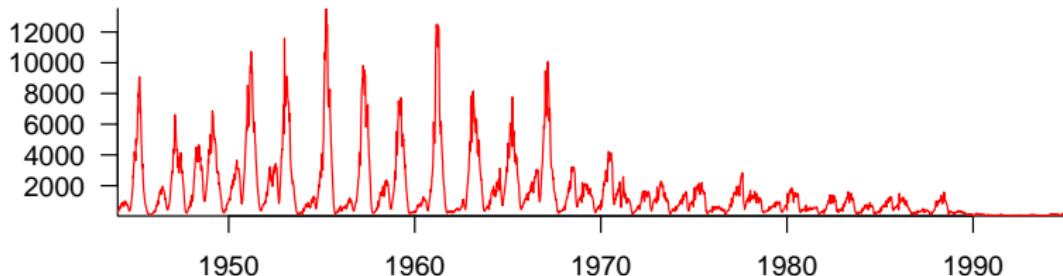
Epidemic Data Analysis

Time Plots of Temporal Epidemic Patterns

1918 P&I

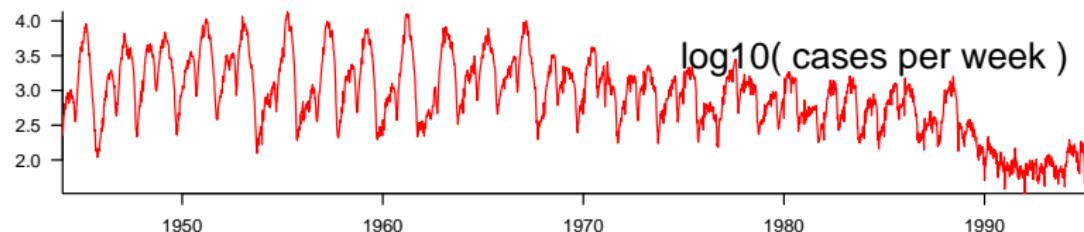
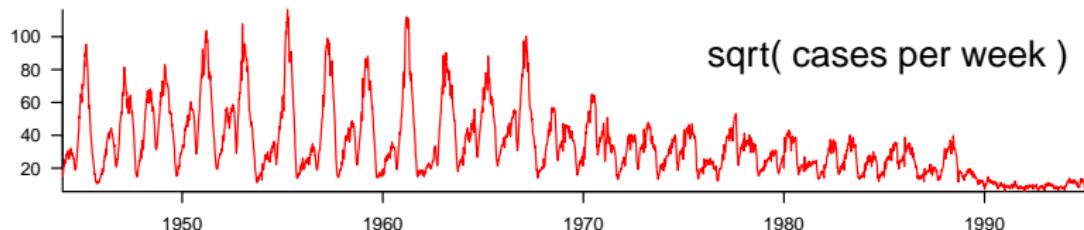
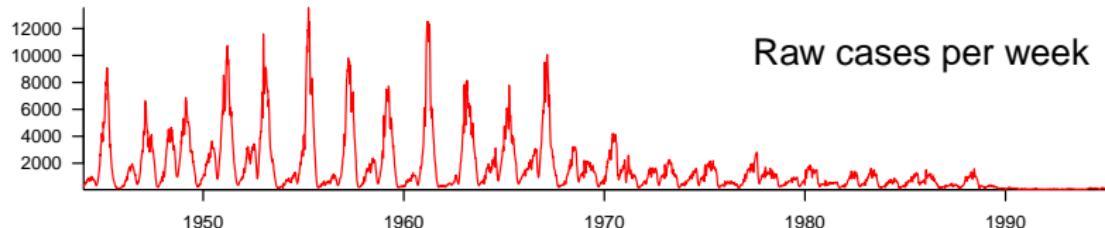


Weekly Measles in England and Wales



Time Plots of Transformed Data

- Reveal unobvious aspects of time series



Times Plots of Smoothed Data

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

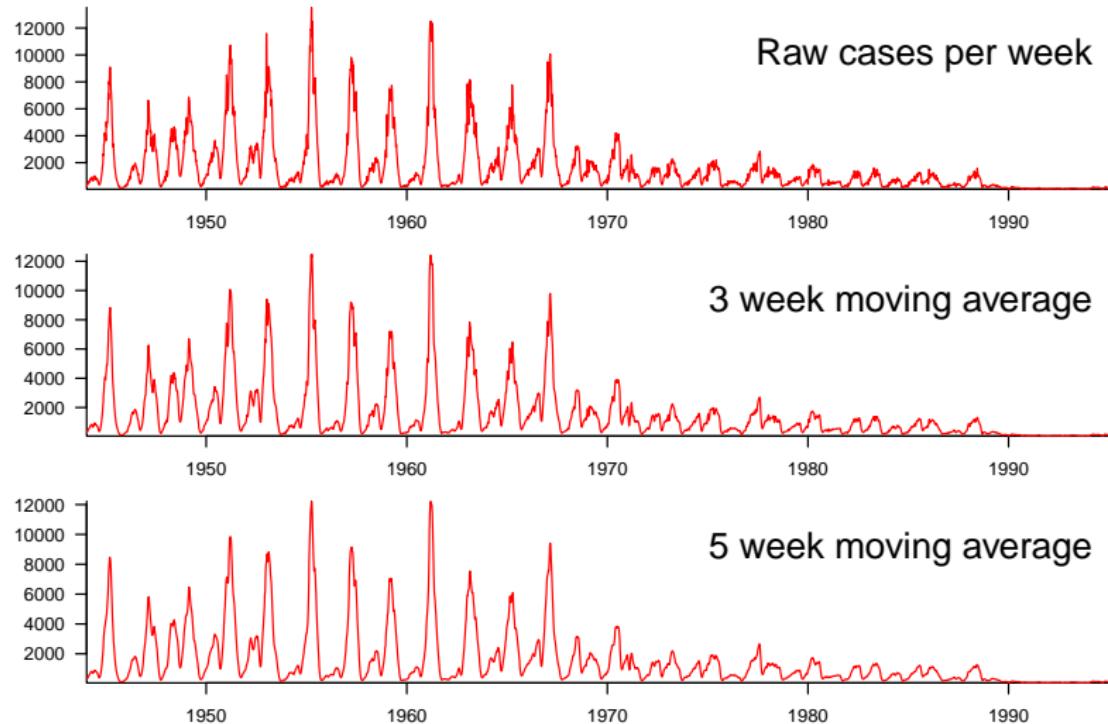
$$x_t \rightarrow \frac{1}{2a+1} \sum_{i=-a}^a x_{t+i}$$

- Replace original data points x_t with averages of nearby points.
- *Linear filter:*

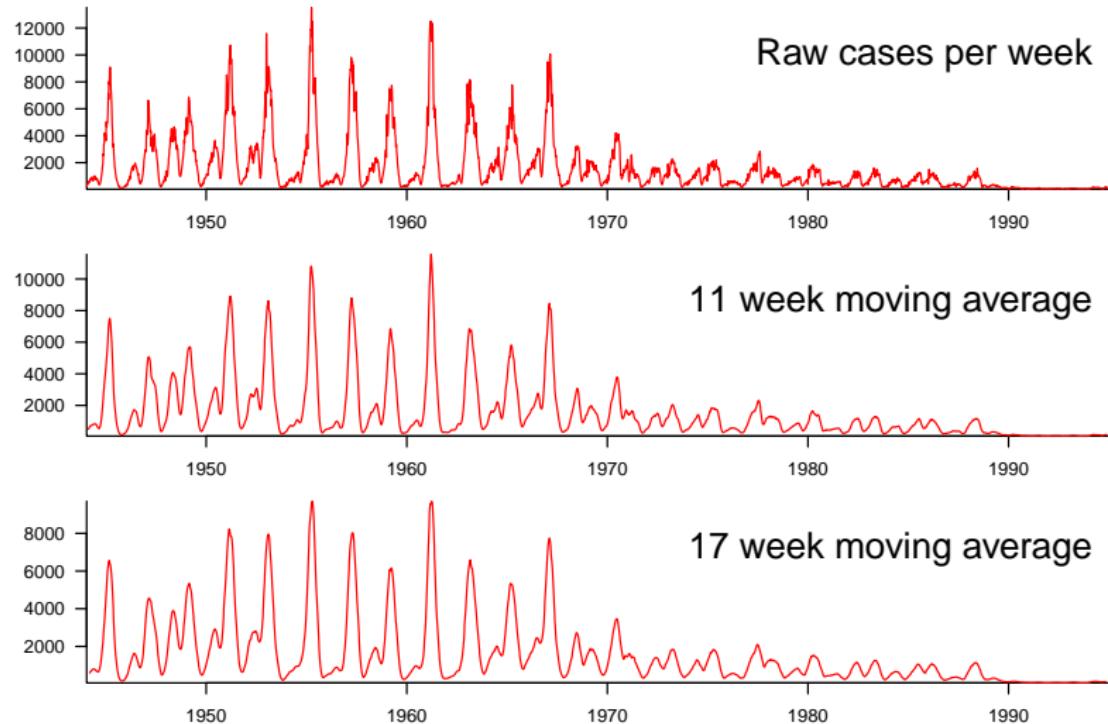
$$x_t \rightarrow \sum_{i=-\infty}^{\infty} \lambda_i x_{t+i}$$

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

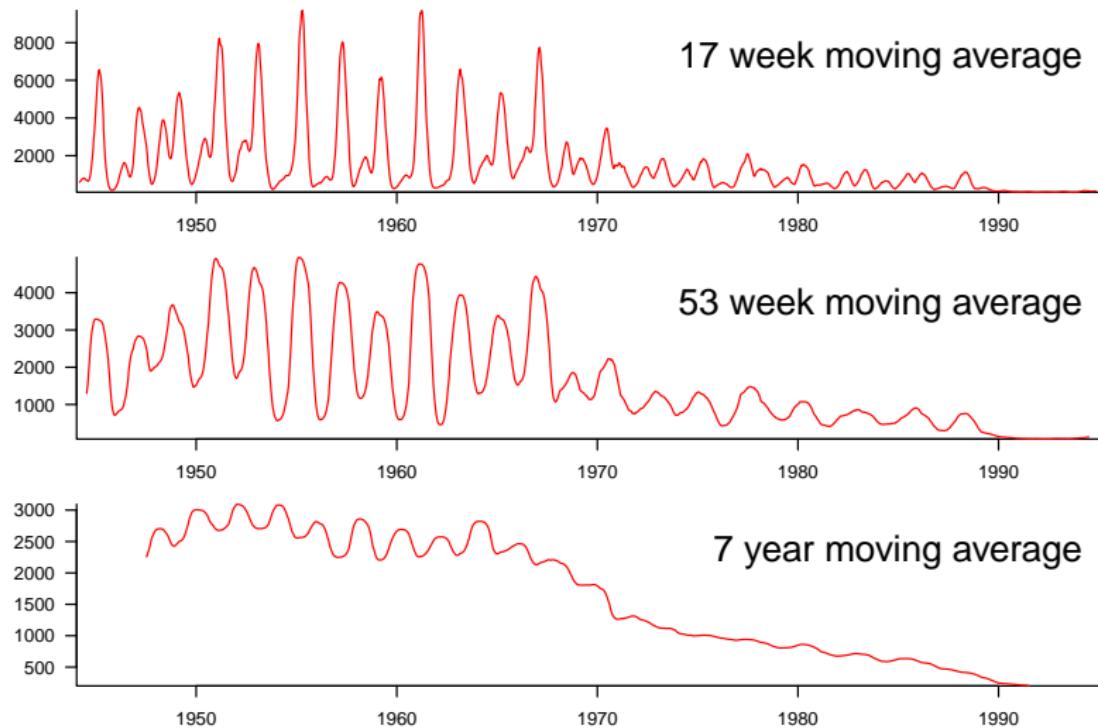
Times Plots of Smoothed Data



Times Plots of Smoothed Data



Times Plots of Smoothed Data



Correlation

- Recurrent epidemics \implies number of cases now is correlated with number of cases in the past and the future.
- Given N pairs of observations of different quantities, $\{(x_i, y_i) : i = 1, \dots, N\}$, 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:

- $-1 \leq r \leq 1$ (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:* r^2 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](#)

Autocorrelation

- Given a single sequence of observations $\{x_t : t = 1, \dots, N\}$, we can compute the correlation of each observation with the observation k time steps in the future.
- Thus, we consider the pairs of observations $\{(x_t, x_{k+t}) : t = 1, \dots, N - k\}$ 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.

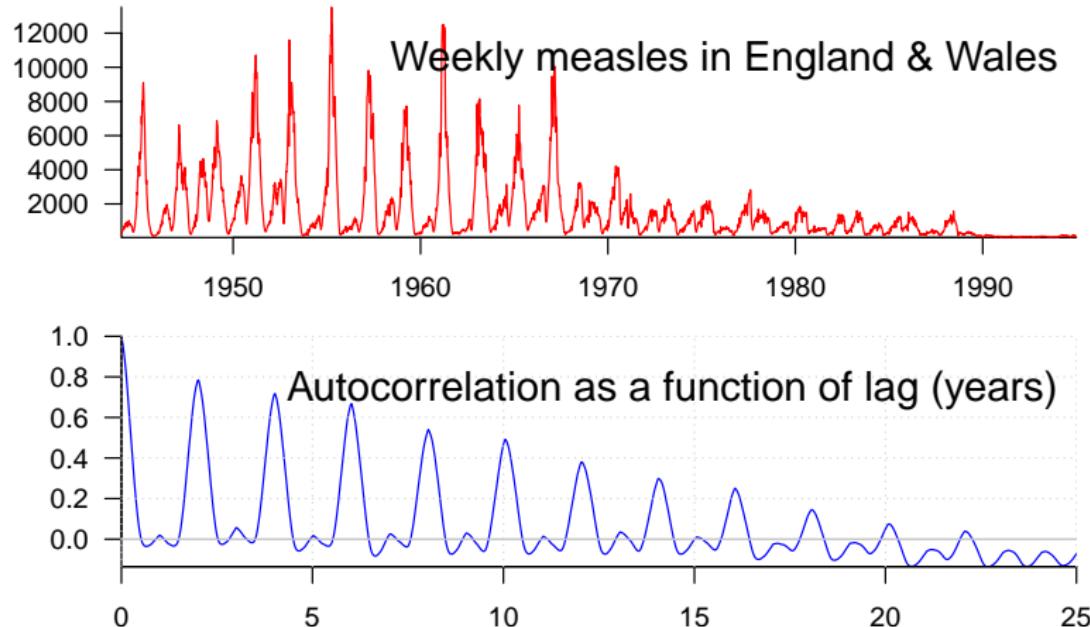
Autocorrelation

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

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

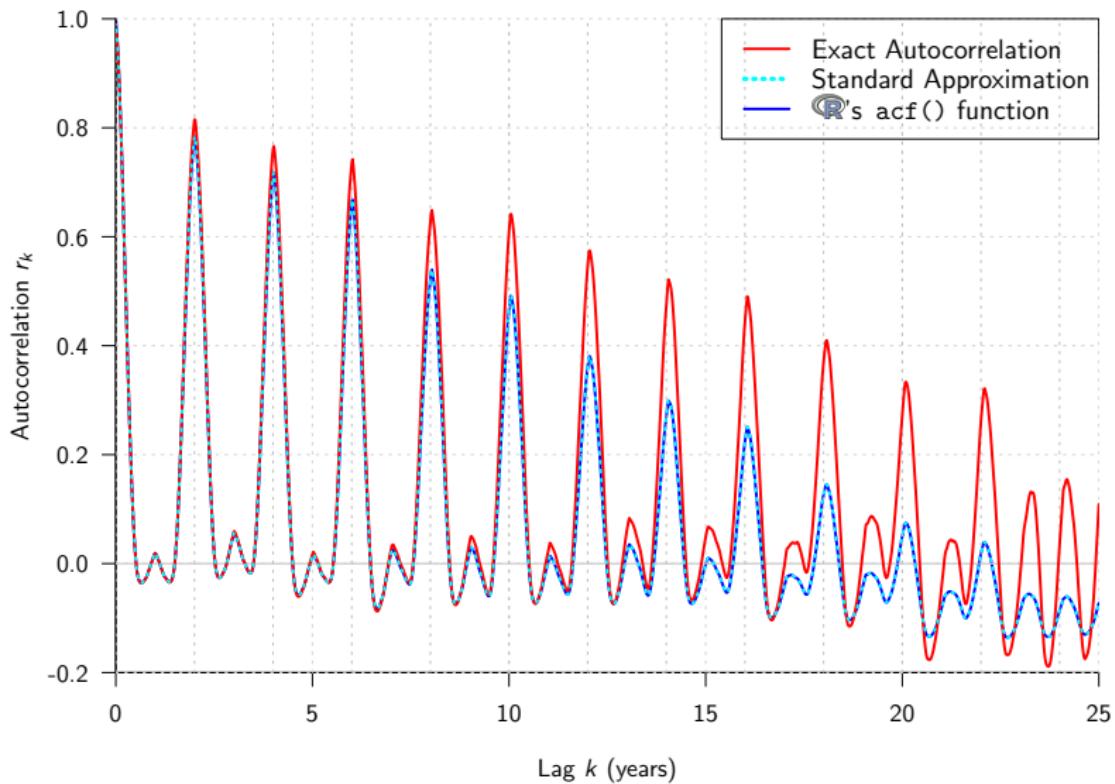
- Approximation of r_k is worse for larger lags k
- Plot of autocorrelation r_k as a function of lag k is called the *correlogram*.

Correlogram



- Peaks in correlogram \implies 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](#):

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

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

- Compute the [Fourier coefficients](#) $\{a_p\}$, $\{b_p\}$ by taking inner products with $\cos \omega_p t$ and $\sin \omega_p t$.

Spectral Density

- Fourier coefficients of x_t are:

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

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

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

where sum is over observation times.

- Estimated power spectral density (PSD) at frequency ω_p is^{*}:

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

*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)

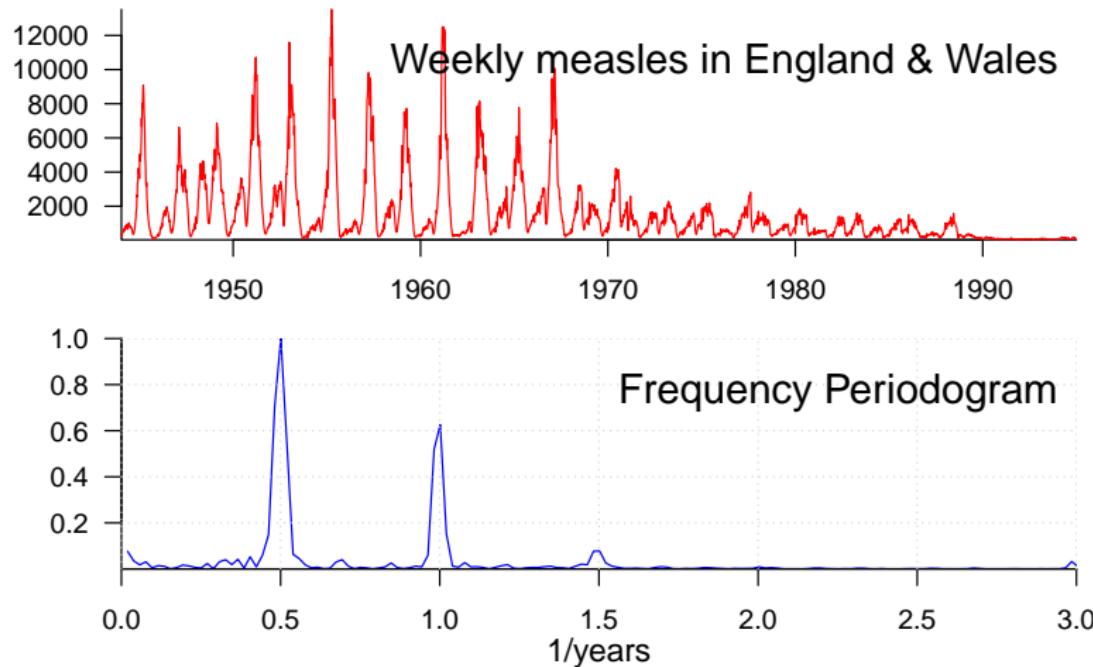
Spectral Density

- 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:

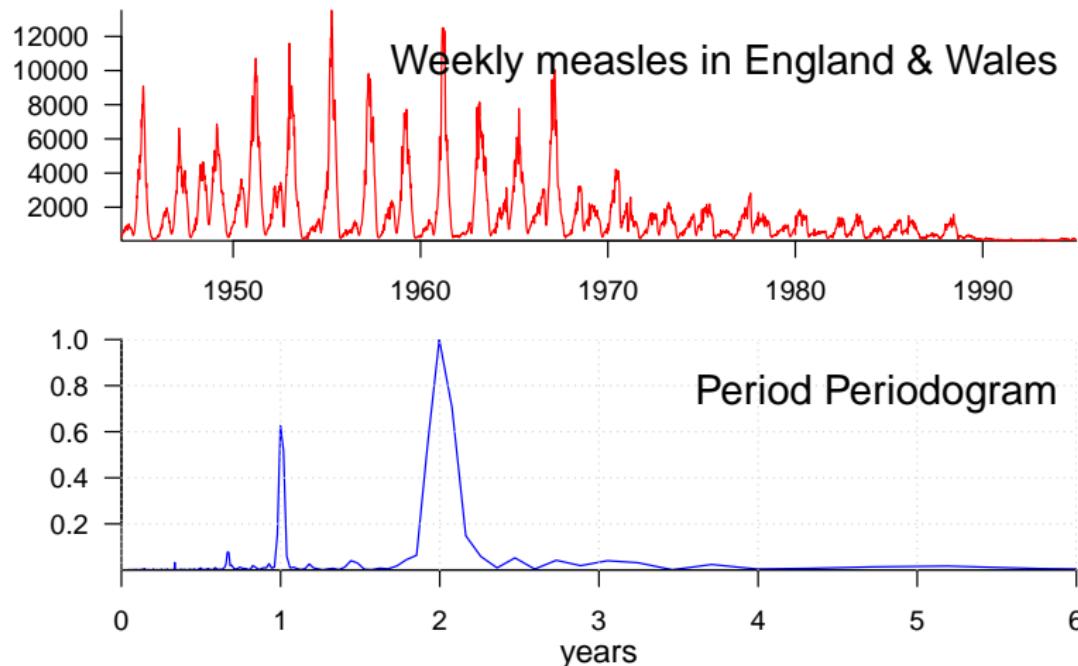
$$I(\omega_p) = \frac{1}{\pi} \left(r_0 + 2 \sum_{k=1}^{N-1} r_k \cos \omega_p k \right)$$

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

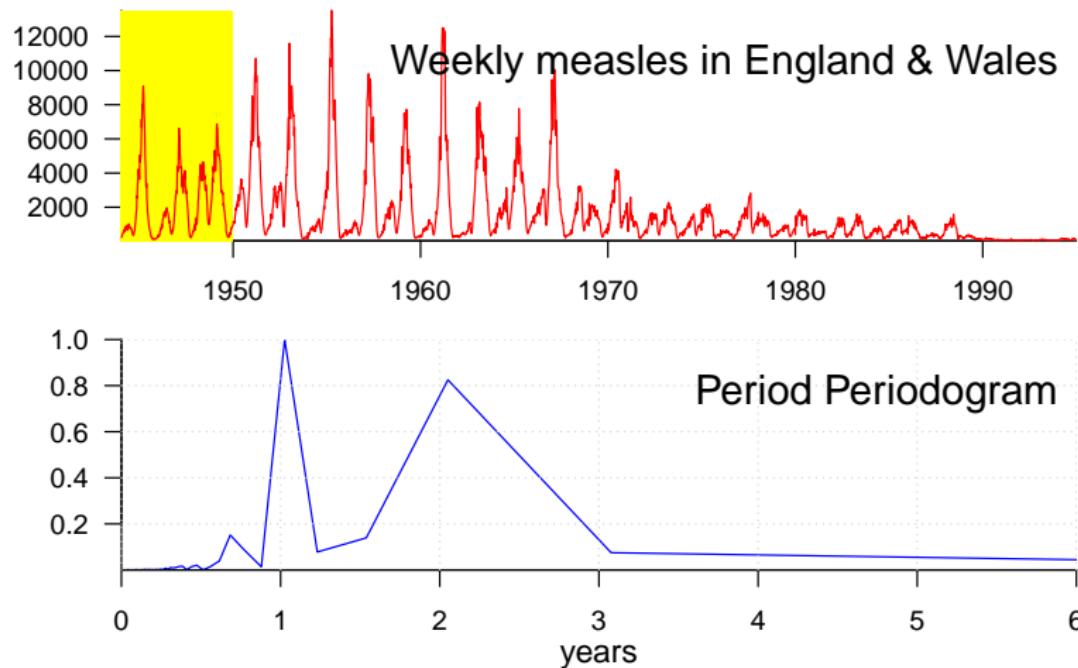
Spectral Density



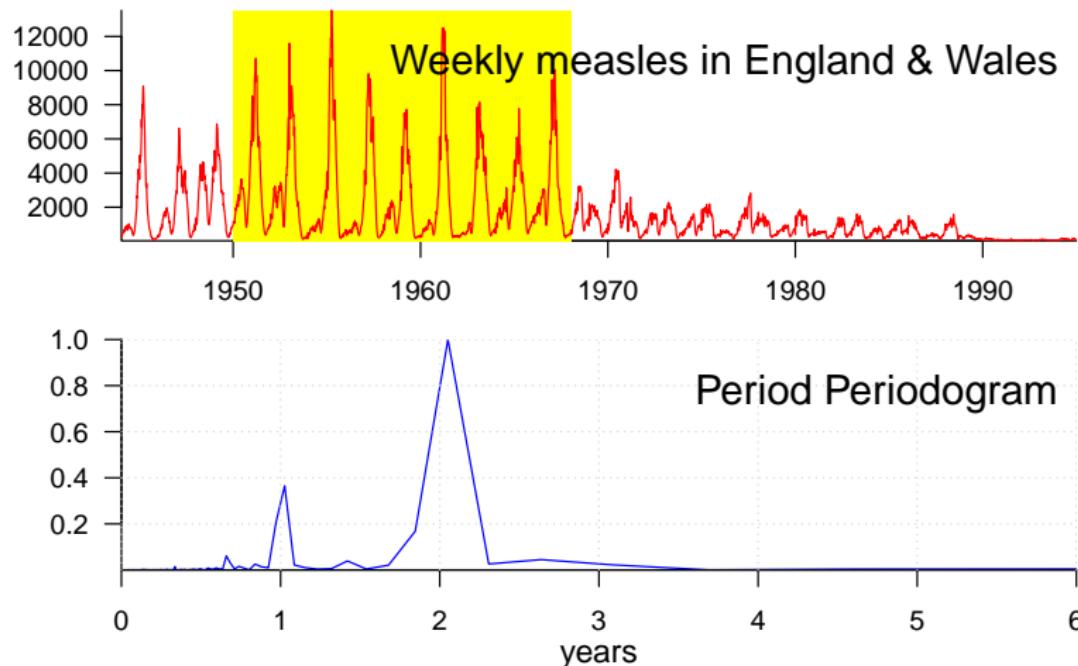
Spectral Density



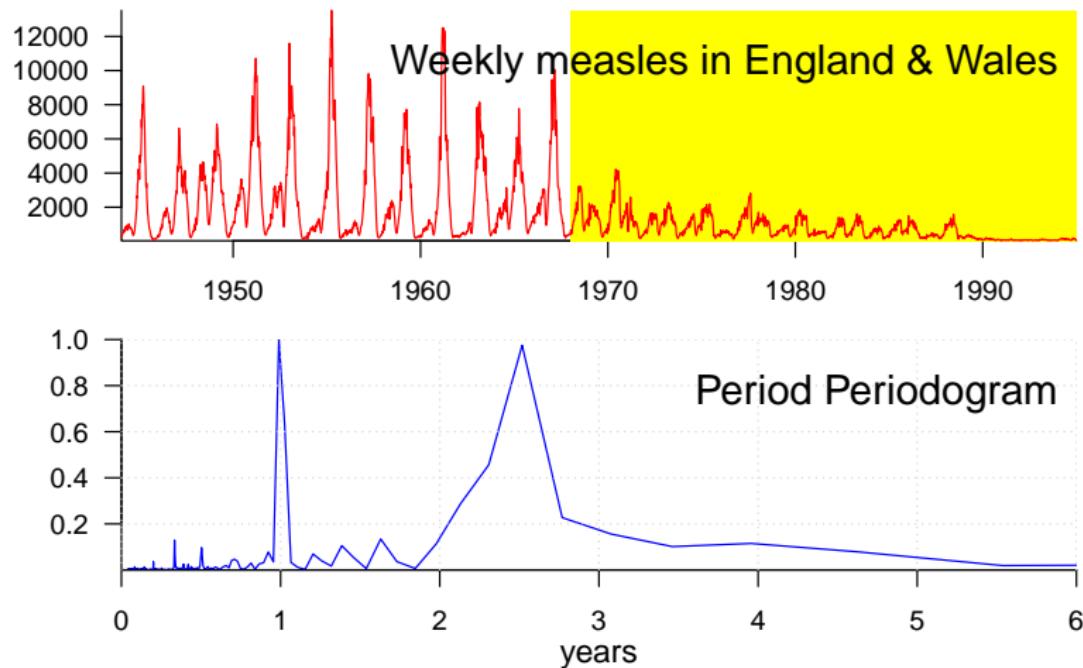
Spectral Density of Temporal Segments



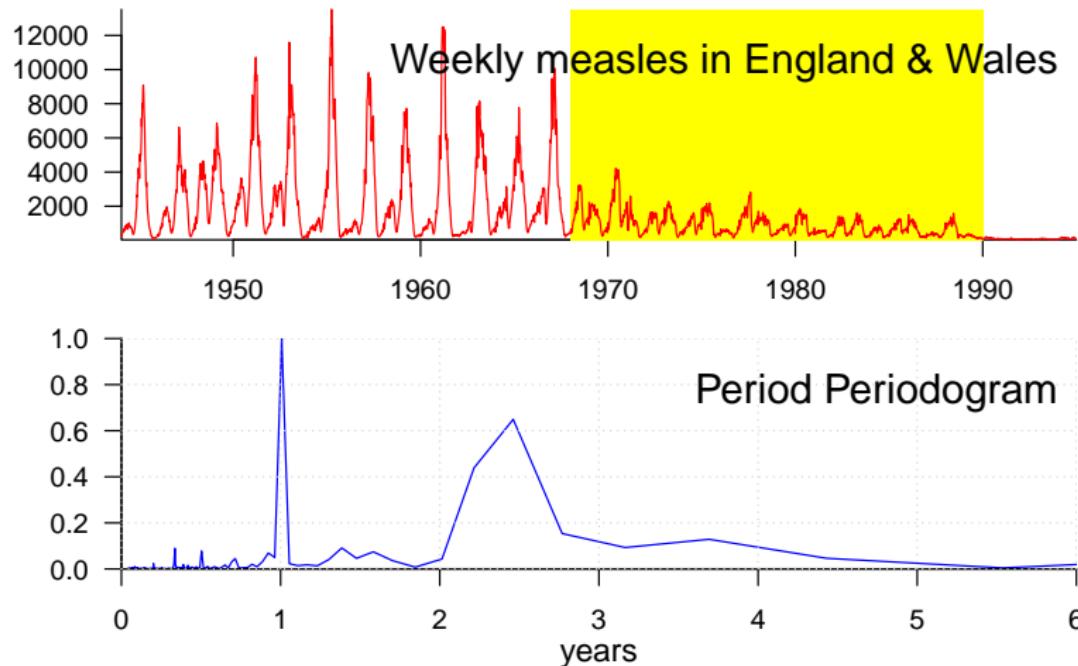
Spectral Density of Temporal Segments



Spectral Density of Temporal Segments



Spectral Density of Temporal Segments

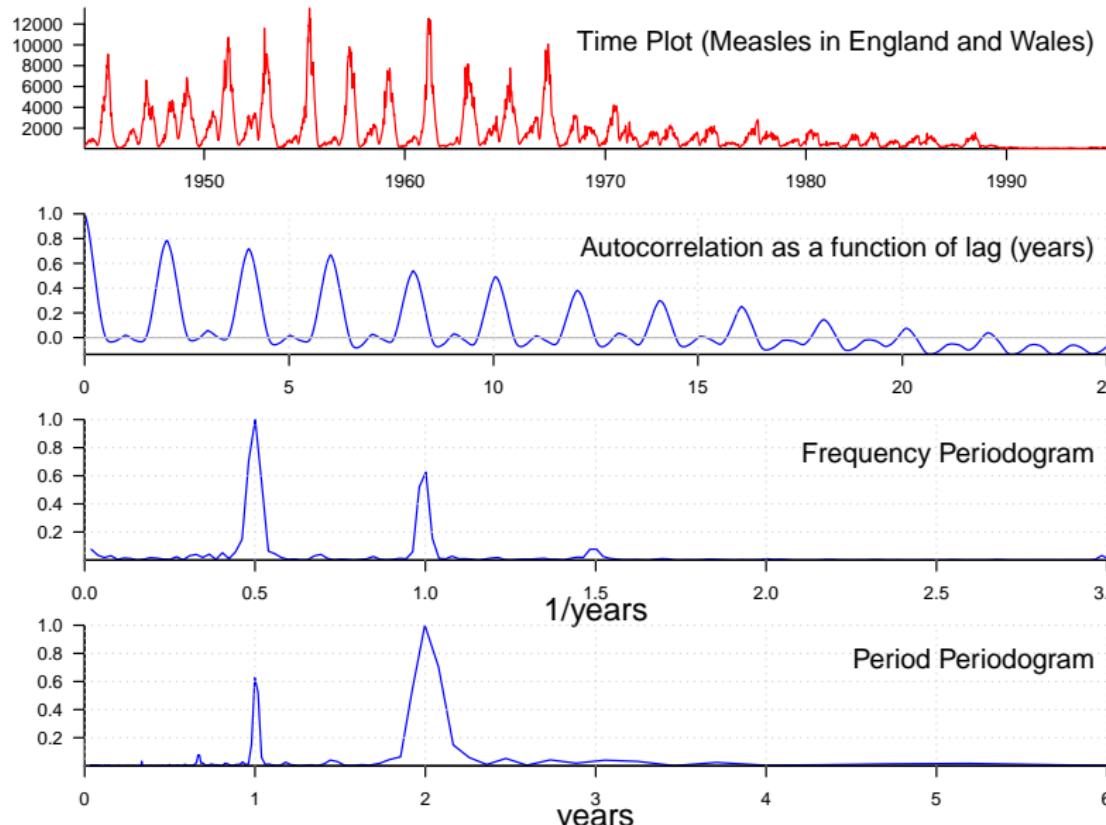


Spectral Density Properties

- Periodogram is discrete Fourier transform of correlogram
- Same information in correlogram and periodogram
- Periodogram usually easier to interpret
- In , 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
- Vaccination era measles
- Vaccination era measles until 1990

Time series analysis functions



has built-in tools for time series analysis:

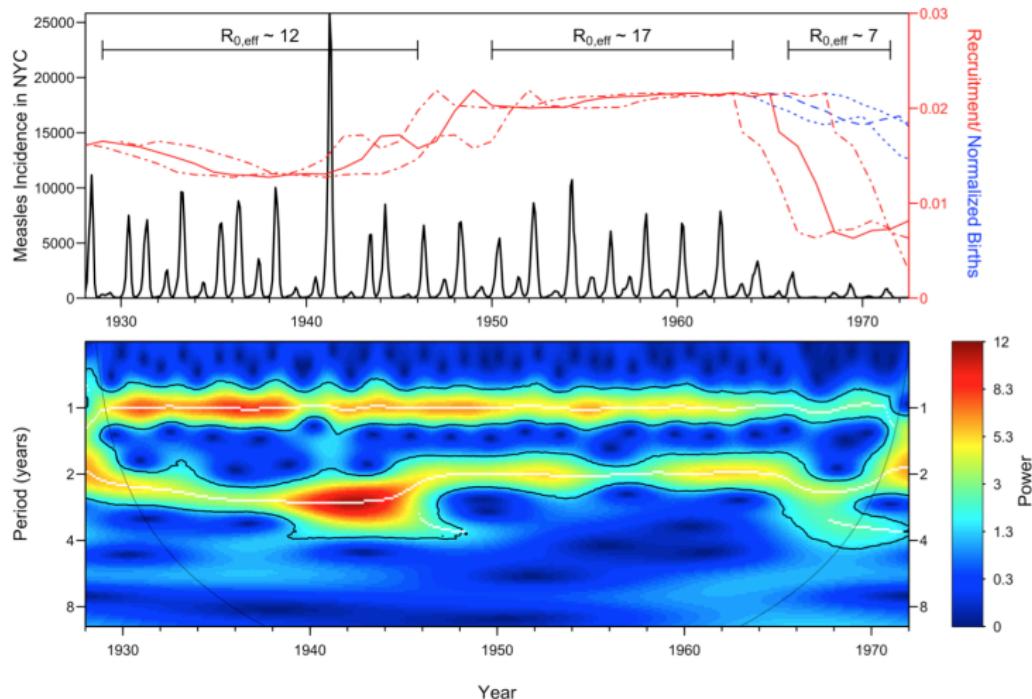
- Time plot: `plot()` etc.
- Linear filter (e.g., moving average): `filter()`
- Correlogram (auto-correlation function): `acf()`
- Periodogram (power spectrum): `spectrum()`

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
 - $\exists \text{ } \text{R}$ 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

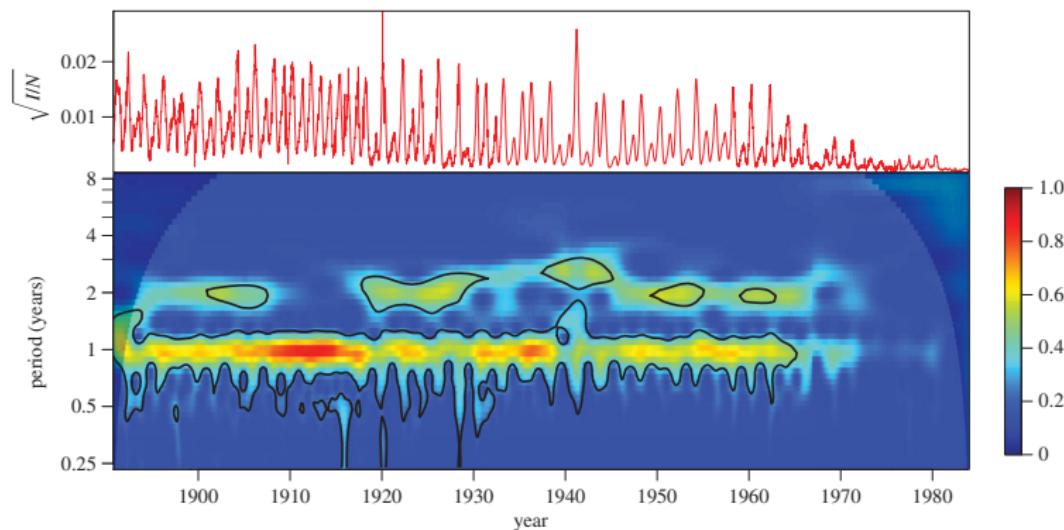


Figure 5. Observed measles dynamics in NYC from 1891 to 1984. (a) Square root of measles case reports, normalized by total concurrent population. (b) Colour depth plot of a continuous wavelet transform of the square root of normalized observed NYC measles cases (colour warmth scales with spectral power 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

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,

$$X_t = \mu + Z_t \quad (*)$$

where μ = time average number of cases
and $\{Z_t\}$ = 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 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*:

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

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:*

$$\begin{aligned} X_t - \mu = & \alpha_1(X_{t-1} - \mu) + \alpha_2(X_{t-2} - \mu) + \cdots + \alpha_p(X_{t-p} - \mu) \\ & + \beta_0 Z_t + \beta_1 Z_{t-1} + \cdots + \beta_q Z_{t-q}. \end{aligned}$$

Statistical Modelling of Time Series: ARIMA

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

$$\begin{aligned} X_t - \mu = & \alpha_1(X_{t-1} - \mu) + \alpha_2(X_{t-2} - \mu) + \cdots + \alpha_p(X_{t-p} - \mu) \\ & + \gamma_1(X_{t-1} - X_{t-2}) + \gamma_2(X_{t-2} - X_{t-3}) + \cdots \\ & + \beta_0 Z_t + \beta_1 Z_{t-1} + \cdots + \beta_q Z_{t-q}. \end{aligned}$$

- 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:

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

- 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)**, which penalizes models that have more parameters
 - See [Earn \(2009\)](#) review article for more discussion of this.



Mathematics
and Statistics

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

Mathematics 4MB3/6MB3 Mathematical Biology

Instructor: David Earn

Lecture 12
Epidemic Data Tools II
Friday 2 Feb 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:* BSB-B154

Time series tools discussed so far...

- Statistical description of time series:
time plot, moving average, correlation coefficient,
autocorrelation, correlogram, power spectral density (PSD),
periodogram, wavelet spectrum
- Time series models:
AR, MA, ARMA, ARIMA

Statistical Modelling of Time Series

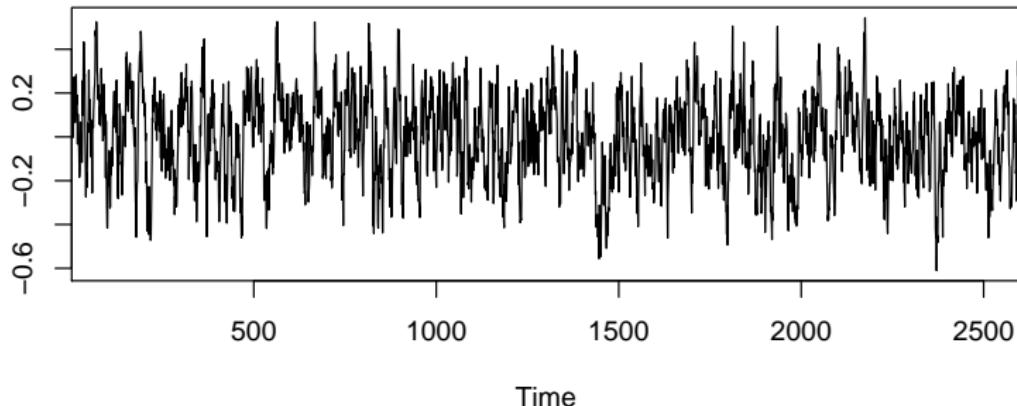
How to do it in ...

- Simulate any ARIMA(p, d, q) model with `arima.sim()`
- Fit an AR model to a time series with `ar()`
- Fit an ARIMA model to a time series with `arima()`
- Alternatively, there are specialized time series modelling packages.

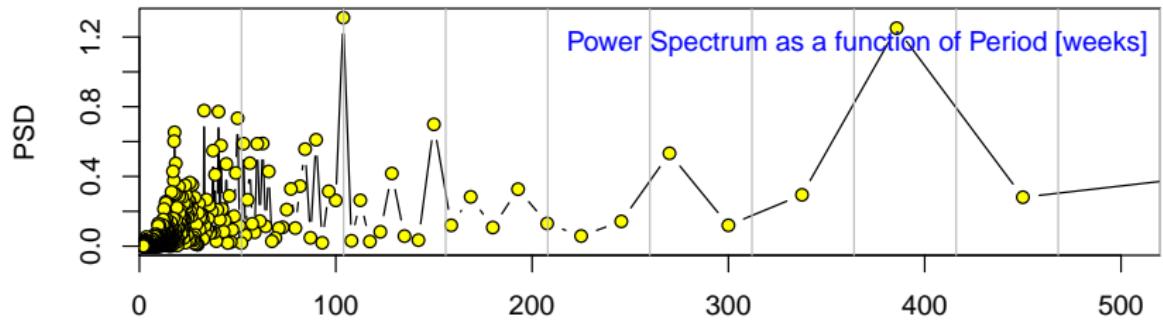
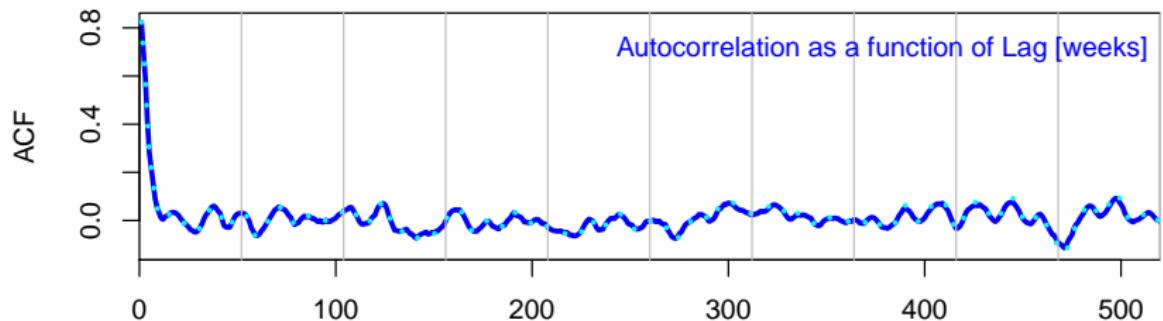
ARMA Example (50 years of weekly data)

```
my.model <- list(ar=c(1,-0.5,0.5,-0.25),ma=c(-0.25,0.5))
my.sim <- arima.sim(n=52*50,model=my.model,sd=0.1)
plot(my.sim,main="ARMA Example",ylab="",xaxs="i")
```

ARMA Example



ARMA Example (ACF and PSD up to 10 year lag)



Statistical Modelling of Time Series: Forecasting

- Once we have a fitted model, we can then use it to *forecast* future observations
- *Validate* this procedure by using part of the data to fit the model and then forecast the remainder of the data (*cf. cross-validation*)
- How successful is this likely to be for an infectious disease time series?
 - Conceivably good for **chicken pox in NYC**.
 - Not for **measles**... at least not for the main patterns...

Statistical Modelling of Time Series: Limitations

- Best to remove mean, trend and seasonality before fitting an ARMA model
 - But this means we will remove the aspects of the data about which we care most!
- The fitted parameters of an ARMA model have no obvious biological meaning
 - The model completely ignores any understanding we have of infectious disease transmission
- Statistical models use the time series itself to parameterize an ARMA (or more general) process
 - It would be better to have a model that we can parameterize from independently collected data and then see if that model can explain the observed time series

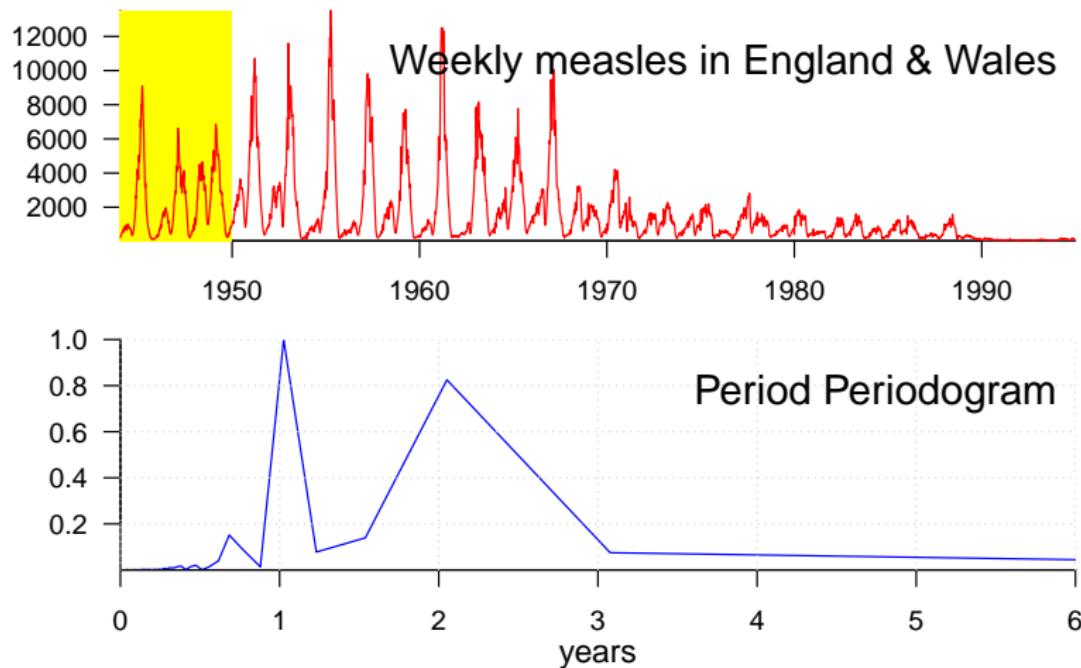
Mechanistic Mathematical Modelling

- SIR and all that...
- Takes into account transmission process...

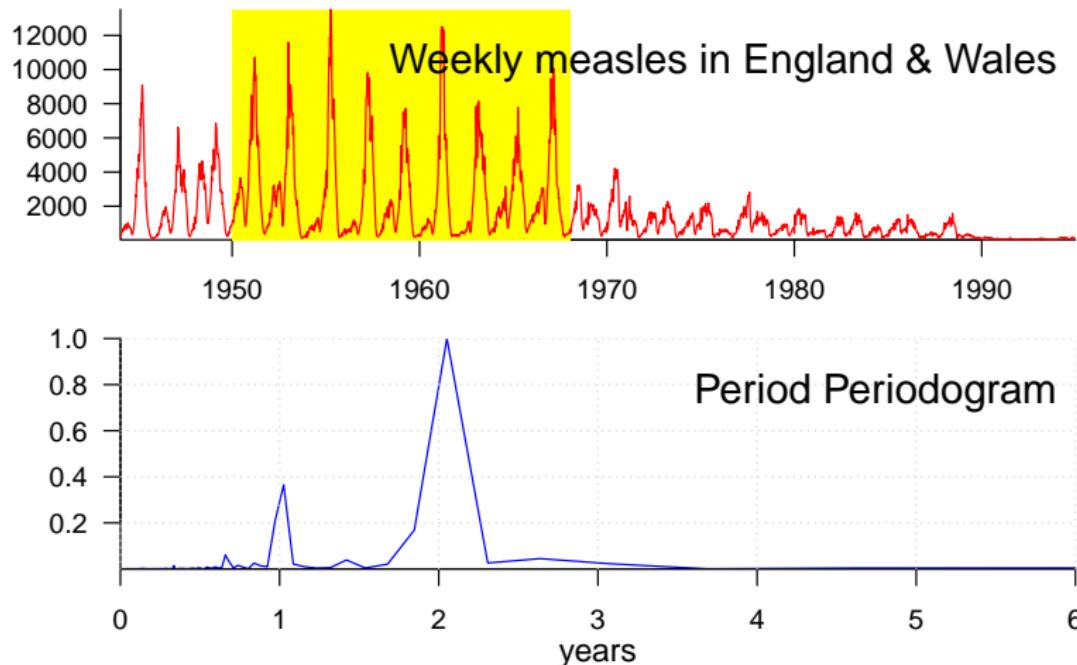
- So why did we just spend time talking about statistical modelling?
 - Important to be familiar with time series models that are in common use.
 - Helps us appreciate the value of mechanistic modelling.
 - Some processes that affect disease dynamics might be better modelled as ARMA or similar processes.
 - Weather (e.g., perhaps model $\beta = \beta(t)$ as an ARMA process)
 - Immigration
 - Ruling out an ARMA model (or at least one with a modest number of parameters) is a step towards finding a good model.

Mechanistic Modelling of Recurrent Epidemics

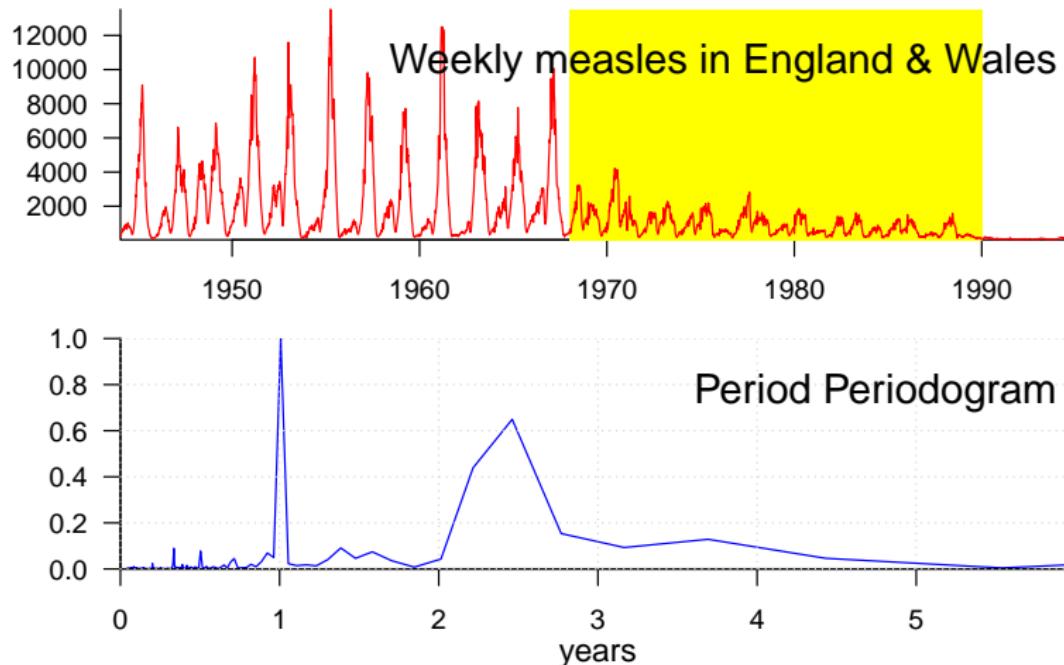
What causes changes in frequency content over time?



What causes changes in frequency content over time?

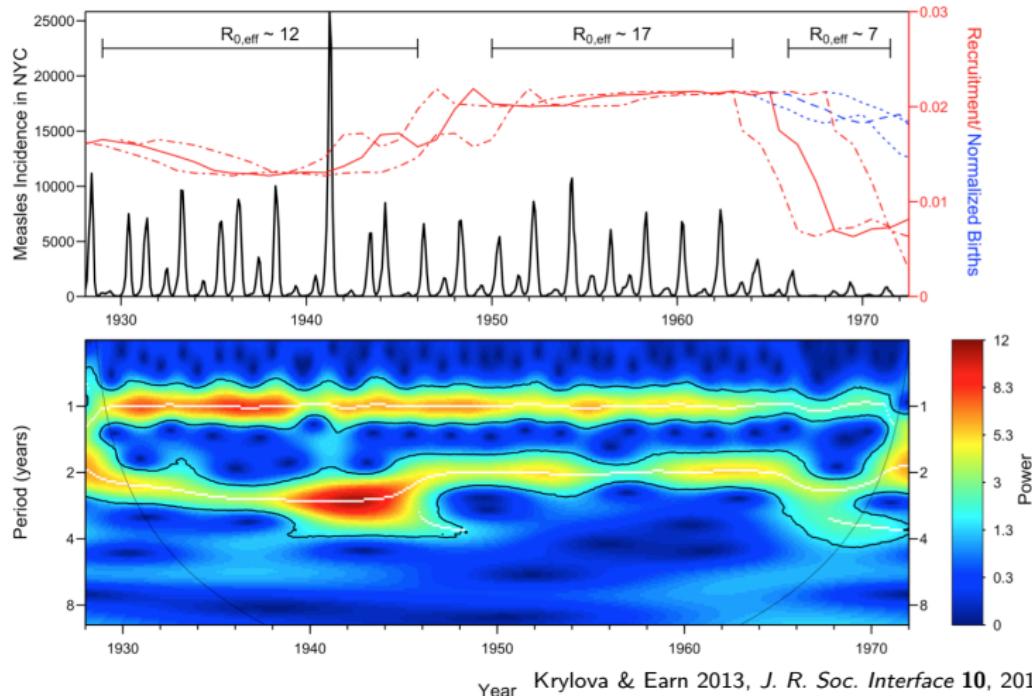


What causes changes in frequency content over time?



What causes changes in frequency content over time?

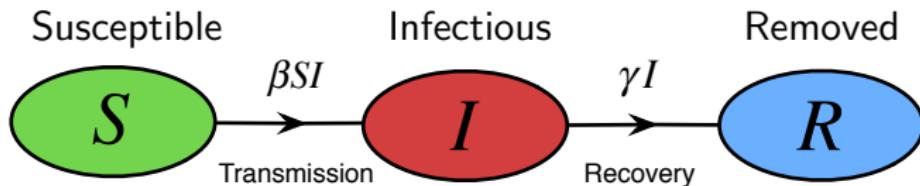
Measles in New York City



Mechanistic Epidemic Modelling: Principles

- Consider the biological mechanisms involved in disease transmission and spread
- Model mechanisms and infer their effects
- Start as simple as possible!
- Rule out simple models by comparing results with observed time series of incidence or mortality
- Add complexity one step at a time, so key mechanisms can be identified
- Ideally converge on simplest possible model that can explain observed patterns

The SIR model: Flow Chart and Parameters



$$\frac{dS}{dt} = -\beta SI$$

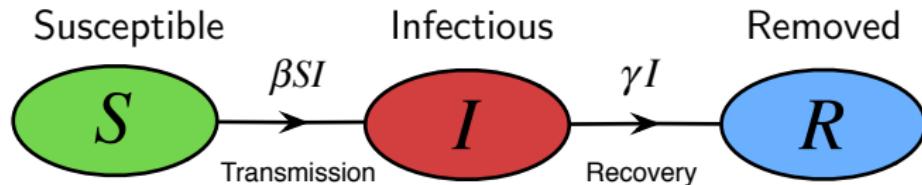
$$\frac{dI}{dt} = \beta SI - \gamma I$$

$$\frac{dR}{dt} = \gamma I$$

■ Parameters:

- Transmission rate β
- Recovery rate γ
(or Removal rate)

The SIR model: Derived Parameters



$$\frac{dS}{dt} = -\beta SI$$

$$\frac{dI}{dt} = \beta SI - \gamma I$$

$$\frac{dR}{dt} = \gamma I$$

■ Derived Parameters:

- Initial growth rate $\beta - \gamma$
- Mean infectious period $\frac{1}{\gamma}$
- Basic Reproduction Number

$$\mathcal{R}_0 = \frac{\beta}{\gamma}$$

Basic SIR Model: Important Results

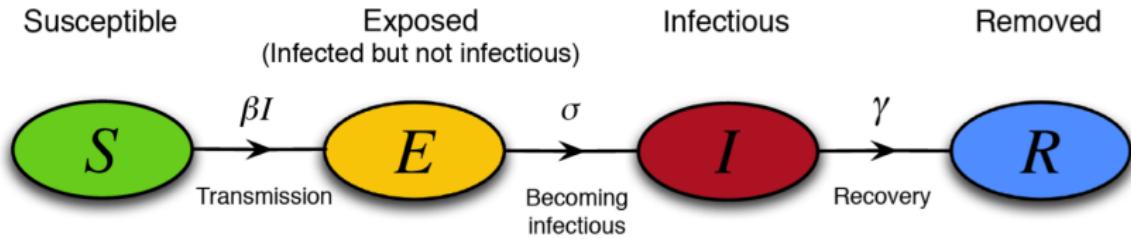
- Epidemic occurs if and only if $\mathcal{R}_0 > 1$
- Exact solution for phase portrait
- Single epidemic, then disease disappears
- Exact formula for final size as a function of \mathcal{R}_0

- Cannot explain diseases that persist
- Cannot explain recurrent cycles of epidemics

What are we missing?



SEIR Model: flow chart

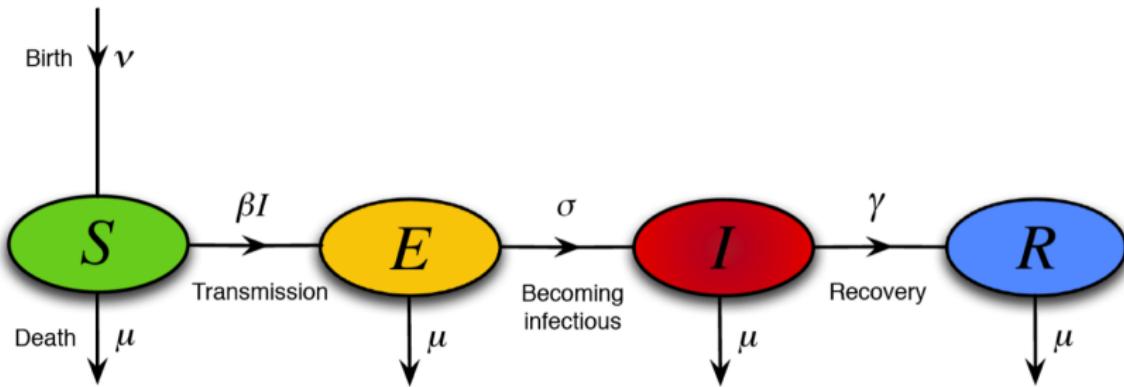


- Introduces only one new parameter (σ)
- Mean latent period ($1/\sigma$) can often be estimated
- But... effect of inclusion of exposed class usually small

What are we **still** missing?



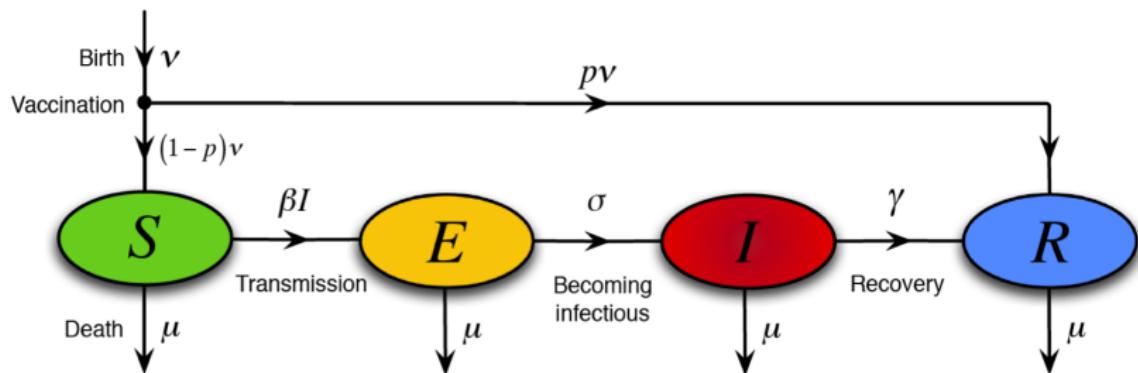
SEIR Model with vital dynamics: flow chart



New Parameters:

- Birth rate (ν for natality)
- Death rate (μ for mortality)
- Mean latent period ($1/\sigma$)

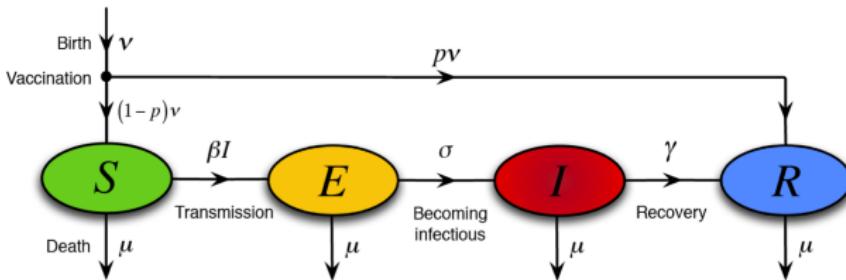
SEIR with vital dynamics and vaccination: flow chart



New Parameters:

- Birth rate (ν for natality)
- Death rate (μ for mortality)
- Mean latent period ($1/\sigma$)
- Proportion vaccinated (p)

SEIR with vital dynamics and vaccination: Equations



$$\frac{dS}{dt} = \nu(1 - p) - \beta SI - \mu S$$

$$\frac{dE}{dt} = \beta SI - \sigma E - \mu E$$

$$\frac{dI}{dt} = \sigma E - \gamma I - \mu I$$

$$\frac{dR}{dt} = \nu p + \gamma I - \mu R$$

- Birth rate (ν for natality)
- Death rate (μ for mortality)
- Proportion vaccinated (p)
- Transmission rate (β)
- Mean latent period ($1/\sigma$)
- Mean infectious period ($1/\gamma$)

SEIR with vital dynamics and vaccination: Analysis

- \mathcal{R}_0 ?
 - Biological derivation: (assuming $\nu = \mu$ and $p = 0$)
$$\mathcal{R}_0 = \beta \times \frac{\sigma}{\sigma+\mu} \times \frac{1}{\gamma+\mu} \quad \simeq \frac{\beta}{\gamma} \quad \because \frac{1}{\mu} \gg \max\left(\frac{1}{\sigma}, \frac{1}{\gamma}\right)$$
 - Mathematical derivation:
 $\mathcal{R}_0 = 1$ is stability boundary
- Final size ? Not well defined (because of continuous source of new susceptibles).
- Equilibria ?