Analyzing Foodborne Disease Outbreaks Over Time

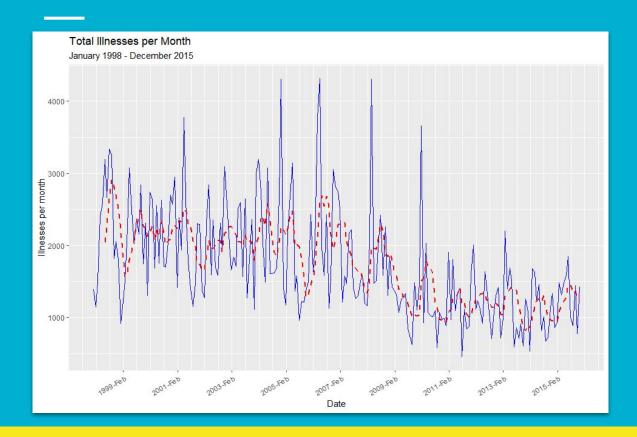
A look into illnesses and more

Group 12

- Trina Shores (group leader)
- Steven Macapagal
- Journey Martinez
- Yuan Yao
- Heather Nagy
- Kenneth Porter

Summary of prior findings

A reminder of what our data look like



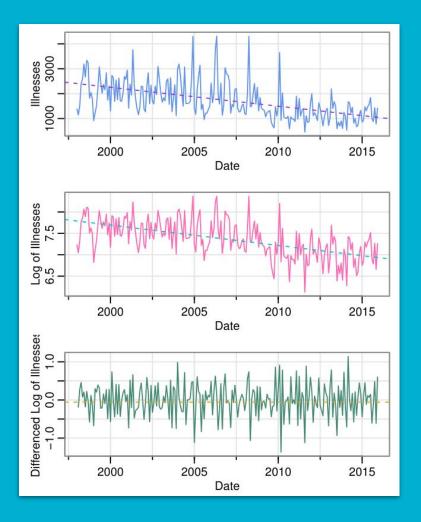
Solid blue line represents the original time series

Dotted red line represents a filtered time series over a 6-month period

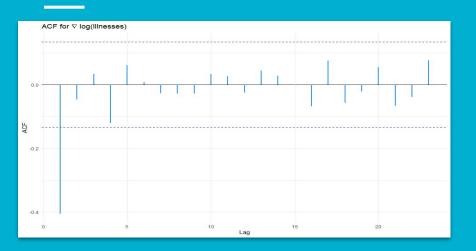
Stationarity

Variability is much greater from 1998 to 2010 and decreases from 2011 to 2015. Illnesses also seem to be trending downward and might be seasonal.

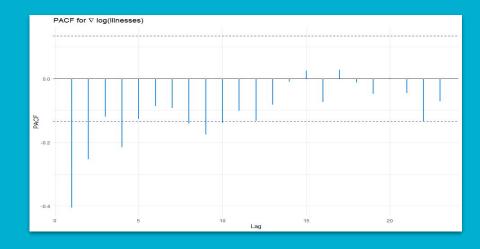
Log transforming and differencing achieved a more constant variance and got rid of the trend, giving us stationary data to work with.



ACF and PACF



The partial autocorrelogram appears to tail off over time. The first few lags are significant, and the magnitude of PACF decreases over time. Only the first lag appears to be significant, while the ACF is not significantly different from zero afterwards.



Parameter Estimation of MA(1)

- MLE estimation (unconditional least squares) with the sarima function in the astsa library
- Using p = 0, d= 1 (using log data) and q = 1 yielded best results
 - Values converged = > reasonable model
 - Conditional SS = -1.010483
 - Unconditional SS = -1.017639
 - MA(1) term significant

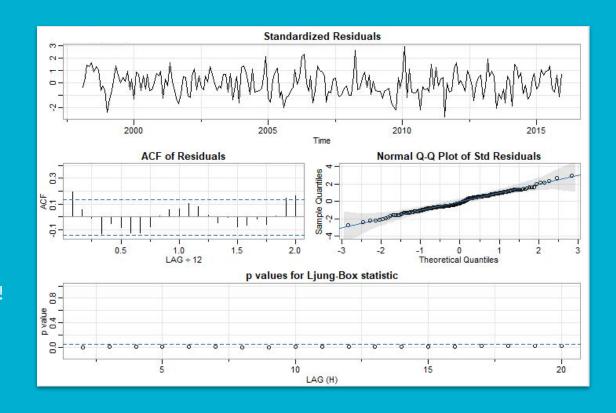
$$\hat{\theta} = -.9103, \hat{\sigma}_w^2 = 0.1296$$

 $x_t = \text{difference in log(Illnesses)}$ for time t

$$x_t = \omega_t - .9103_{(0.02)}\omega_{t-1}$$

Estimation Output of MA(1)

- Scattered, Normally distributed residuals
- ACF shows no departure from model assumptions
- p-values for Q tests are all less than 0.05
 - Residuals are correlated!



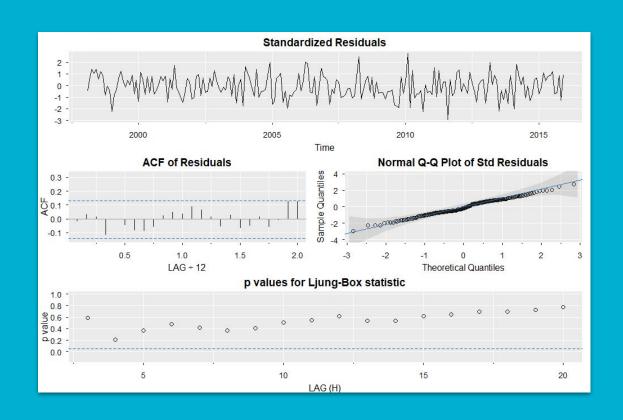
Parameter estimation of ARIMA(1,1,1)

- MLE estimation (unconditional least squares) with the sarima function in the astsa library
- Using p = 1, d= 1 (using log data) and q = 1 yielded best results
 - Values converged = > reasonable model
 - Conditional SS = -1.014162
 - Unconditional SS = -1.039521
 - AR(1) and MA(1) terms significant
 - Smaller AIC and BIC than AR(1) and MA(1) models
 - $\hat{\phi} = 0.2193, \hat{\theta} = -0.9351, \hat{\sigma}_w^2 = 0.1241$

```
x_t = 	ext{difference in log(Illnesses)} 	ext{ for time } t x_t = 0.2193_{(0.07)}x_{t-1} - .9351_{(0.02)}\omega_{t-1} + \omega_t
```

Estimation Output of ARIMA(1,1,1)

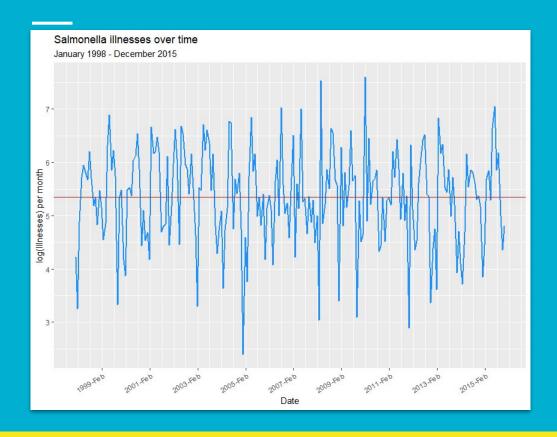
- Scattered, Normally distributed residuals
- ACF shows no departure from model assumptions
- p-values for Q tests all greater than 0.05
 - Residuals are uncorrelated



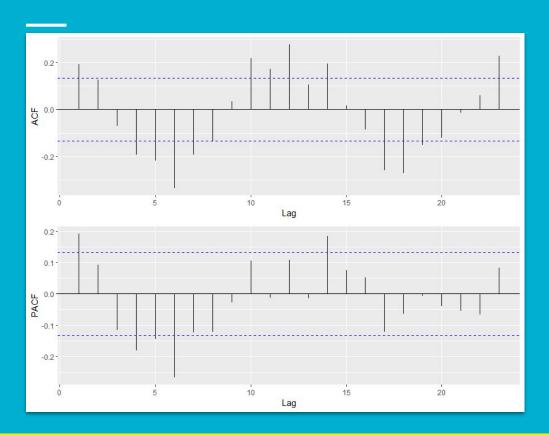
Our plan

- 1. Compare subsets of the data to see if there is an underlying pattern to foodborne disease outbreaks.
- Compare models of foodborne illness to models of hospitalizations related to foodborne illness.
- Compare different models to our previously established ARIMA(1, 1, 1) model on fit characteristics and prediction.

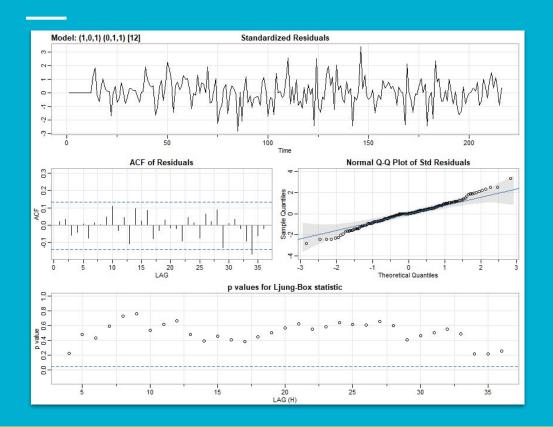
Analyzing source of outbreaks



- Plot of log(illnesses) for salmonella cases appears to be stationary
- Seems to have a seasonal component



- Seasonal differencing needed: ACF is large and significant for every 12th lag, seems to tail off slowly
- ACF and PACF have significant 1st lags



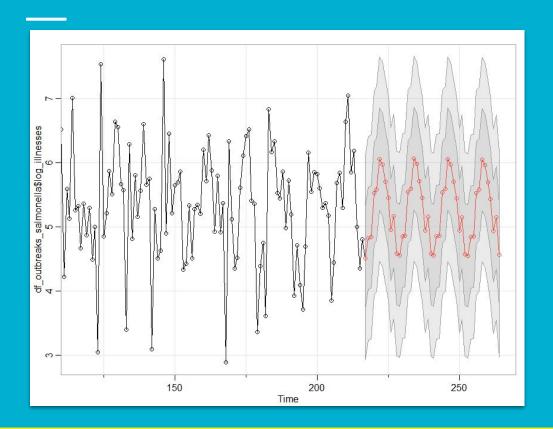
SARIMA(1, 0, 1) x (0, 1, 1)₁₂

- Residuals appear to be normally distributed white noise
- P-values for Ljung-Box statistic appear to be nonsignificant

SARIMA(1, 0, 1) x (0, 1, 1)₁₂

Estimated coefficients:

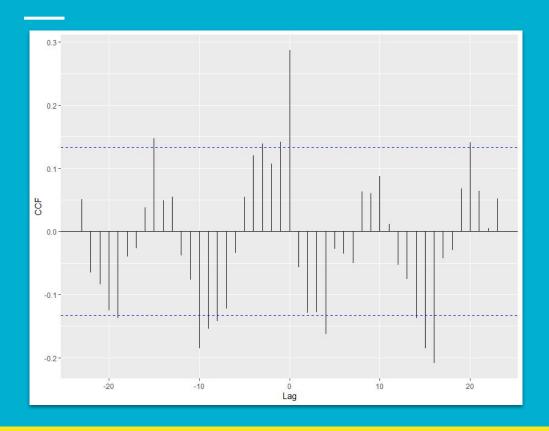
$$Ø_1 = 0.76$$
 $θ_1 = -0.84$
 $Θ_1 = -1$
 $σ^2 = 0.59$, df = 200



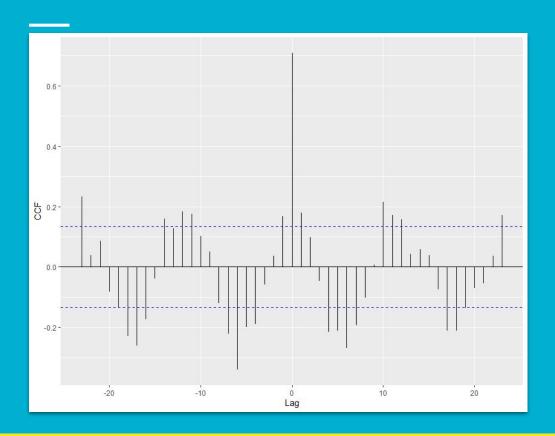
Seasonal forecasts of log(salmonella)

Relationships between illnesses and hospitalizations

CCF of overall illnesses and hospitalizations



- Strongest cross-correlation between illnesses and hospitalizations in the same period
- Appears to have some seasonality



CCF of log(illnesses) and log(hospitalizations)

- Seasonal patterns of illness still apparent
- Strongest
 cross-correlations in same
 period (hospitalizations
 follow a diagnosed illness
 closely)

Comparisons to baseline ARIMA(1, 1, 1) model

Steps

Modeling process:

- 1. Model formulation
- 2. Model estimation
- 3. Model diagnostics
- 4. Model selection

We will also compare their forecasts to the actual data released for 2016 through 2019.

Models:

- 1. ARIMA(1, 1, 1) [baseline]
- 2. ARIMA(1, 1, 0) [Li et al. (2021)]
- 3. ARIMA(1, 1, 1) + GARCH(1, 0)
- 4. ARIMA(1, 1, 1) \times (1, 0, 1)₁₂
- 5. ARIMA(1, 1, 1) \times (0, 1, 1)₁₂
- 6. Prophet (examined later)

Comparing models suggested in prior literature

Li, et al. (2021) suggested an ARIMA(1, 1, 0) model to describe the incidence of foodborne illnesses over time.

- Performance and prediction compared to ARIMA(1, 1, 1) model we had previously found to fit our data

Li, S., Peng, Z., Zhou, Y., & Zhang, J. (2021). Time series analysis of foodborne diseases during 2012-2018 in Shenzhen, China. *Journal of Consumer Protection and Food Safety*, 17(2), 83-91.

Journal of Consumer Protection and Food Safety (2022) 17:83–91 https://doi.org/10.1007/s00003-021-01346-w

Journal of Consumer Protection and Food Salety

RESEARCH ARTICLE



Time series analysis of foodborne diseases during 2012–2018 in Shenzhen. China

Siguo Li¹0 · Zhao Peng² · Yan Zhou¹ · Jinzhou Zhang¹

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Abstract

The present study aimed to use the autoregressive integrated moving merage (ARIMA) model to forecast foodborne disease incidence in Sherwhen ciny and help guide efforts to prevent foodborne disease. The data of foodborne disease surveillance network of the disease surveillance network. The indirect data from January 2012 to December 2012 was used for the model-constructing, while the data from January 2018 to December 2018 was used for the model-constructing, while the data from January 2018 to December 2018 was used for sense insender presenting error (MARF) was used of to sense the performance of the model. The monthly foodborne disease incidence from January 2012 to December 3017 in Shenzhes was between 975 and 32.863 with an incidence rate absolute presenting error (MARF) was used to assess the performance of the model. The monthly foodborne disease incidence from January 2012 to December 2017 in Shenzhes was between 975 and 32.863 with an incidence rate scale incidence incidence series, yelding a MAPE of 5.9%. The mathematical formula of the ARIMA (1,10) model was considered advantage of the ARIMA (1,10) model. The model can be considered adequate for predicting future foodborne disease incidence in Shenzhes and one and in the decisions making processes.

Keywords Foodborne disease · ARIMA · Time series analysis · Forecasting foodborne diseases · Foodborne disease incidence

1 Introduction

Foodborne disease is one of the most important public health issues: in both developed and developing countries (Saular 2017). Generally, foodborne disease results from the consumption of food contaminated with pathogens such as bacteria, vinues, pranties or with polsonous chemicals or bio-toxine (Binsis 2017). Although the disease is usually mild and self-limiting, due to the high number of individuals affected each year, foodborne disease searts a substantial societies coming branch on the polsonic and healthcare.

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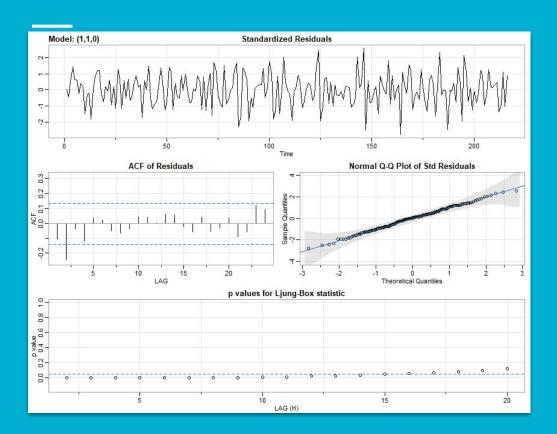
- Shenzhen Center for Disease Control and Prevention, Longyuan Road 8, Shenzhen 518055, China
- Department of Nutrition and Food Hygiene, Hubei Key Laboratory of Food Nutrition and Sakhy, Tongji Medical College, Huzzhong University of Science and Technology, Hanekone Road 13, Wuhan 430030, China

system (Wu et al. 2018). The World Health Organization (WHO) estimated that 31 foodborne hazards caused 600 million foodborne disease case and 420,000 deaths world-wide in 2010 WHO 2017). China faces various and unprised in 2015 who world-wide in 2010 WHO 2017. China faces various and unprised and 94,117 m cases of bacterial foodborne diseases occur every year, or which 3.357 m are hospitalized and 8530 die with a mortality rate of 9.1/100,000 (Mass et al. 2011). Furthermore, a minority of patients with foodborne diseases seek formal medical care, and informative tests are backen of 5.00 feet of 1.00 feet of

China is currently developing and implementing a footborne disease survillance system arross the country However, the foodborne disease surveillance system remains in the early stage of development in a stepwise finalian the system has various limitations (Lin et al. 2018). In Stemelne, a city of Guangdone Province, a citywa the vivillance network on infectious distribes patients was estabilated in 2019. Sameles were collected at intreats from both ideed in 2019. Sameles were collected at interest from both the contract of the c



ARIMA(1, 1, 0) [Li et al., 2021]



- Diagnostics show residuals are not white noise, still have some autocorrelation
- Estimated coefficients:

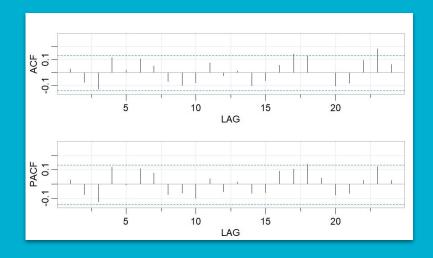
$$\varnothing_1 = -0.41 (0.06)$$

 $\sigma^2 = 0.16, df = 214$

- AIC = 1.02 BIC = 1.05

ARIMA(1,1,1) + GARCH(1,0)

- Some small dependence left in ARIMA(1,1,1) squared residuals
 - Average of residuals = 0.012
- ARIMA(1,1,1) + GARCH(1,0)
 on transformed data does not
 have significant alpha term
- GARCH(1,0) not needed



```
Error Analysis:
         3.65873
                     0.46265
         0.50357
                     0.06260
                     0.02054
         0.14864
                     0.08138
alpha1
         0.01944
                                         0.8112
        10,00000
                     3.93673
                                 2.540
                                         0.0111 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Log Likelihood:
              normalized: -0.4526084
```

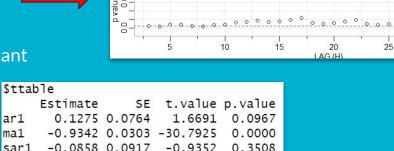
Seasonality

- There appears to be a seasonal component
 - ACF cuts off at lag 12, PACF tails off
 - Try adding SMA1 term
- ARIMA(1,1,1)x(0,1,1)₁₂ is possible
 - However, AR1 is not significant
- ARIMA(0,1,1)x(0,1,1) did not have white noise residuals
- $ARIMA(1,1,1)x(1,1,1)_{12}$
 - AR1 and SAR1 terms not significant

ar1

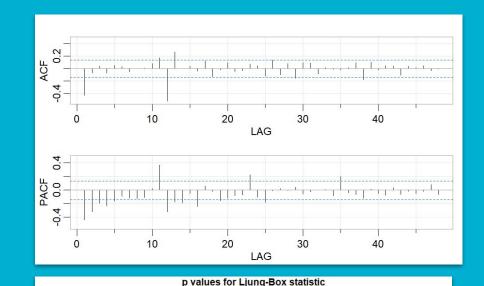
ma1

sar1



0.0000

-9.2709



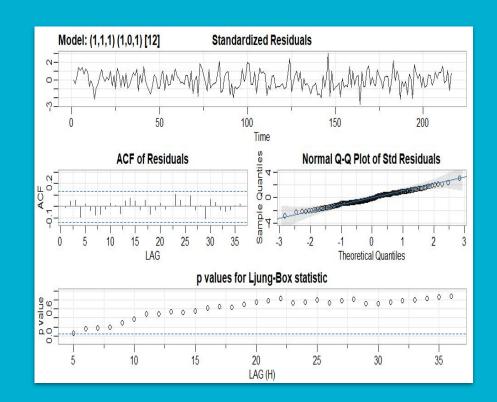
35

```
$ttable
                  SE t.value p.value
     Estimate
ar1
                                0.0000
ma1
      -0.9093 0.0857 -10.6136
```

Selecting seasonal component

- ARIMA(1,1,1)x(1,0,1)₁₂ has significant estimates
- Slightly smaller AIC compared to ARIMA(1,1,1)
- Residuals appear to be white noise
- AIC = 0.769, BIC = 0.847
 - AIC = 0.787, BIC = 0.834 for baseline ARIMA

24	Estimate	SE	t.value	p.value
ar1	0.1692	0.0729	2.3201	0.0213
ma1	-0.9403	0.0226	-41.5258	0.0000
sar1	0.9509	0.0912	10.4234	0.0000
sma1	-0.8796	0.1443	-6.0948	0.0000



$ARIMA(1,1,1) \times (1,0,1)_{12}$

- Less dependence in squared residuals compared to ARIMA(1,1,1)
- Mean(residuals) = 0.006
 - Closer to zero than mean of ARIMA(1,1,1) residuals
- Less biased forecast



```
Estimate
                  SE t.value p.value
       0.1692 0.0729
                       2.3201
                               0.0213
ar1
ma1
      -0.9403 0.0226 -41.5258
                               0.0000
sar1
       0.9509 0.0912
                      10.4234
                               0.0000
      -0.8796 0.1443
                      -6.0948
                               0.0000
sma1
```

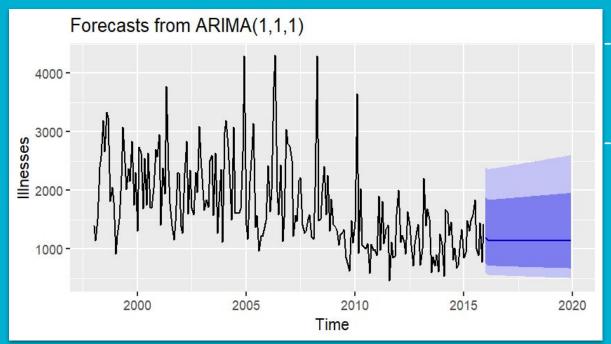
$$\begin{aligned} x_t &= (1+\phi)x_{t-1} - \phi x_{t-2} + \ \Phi x_{t-12} - \Phi(\phi+1)x_{t-13} + \Phi \phi x_{t-14} + w_t + \theta w_{t-1} + \Theta w_{t-12} + \Theta \theta w_{t-13} \\ x_t &= 1.1692x_{t-1} - 0.1692x_{t-2} + \ 0.9509x_{t-12} - 1.1118x_{t-13} + 0.1608x_{t-14} + w_t - 0.9403w_{t-1} - 0.8796w_{t-12} + 0.8271w_{t-13} \end{aligned}$$

Models/Forecasts comparison

Model Comparisons

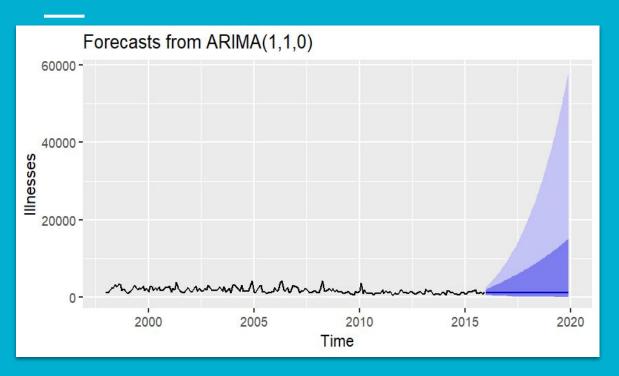
Model	AIC	BIC
ARIMA(1, 1, 1)	0.787	0.834
ARIMA(1, 1, 0)	1.020	1.051
ARIMA(1,1,1)x(1,0,1) ₁₂	0.769	0.847
ARIMA(1,1,1)x(0,1,1) ₁₂	0.881	0.946
ARIMA(1,1,1)-GARCH(1,0)	0.926	0.989

Forecast for ARIMA(1, 1, 1)



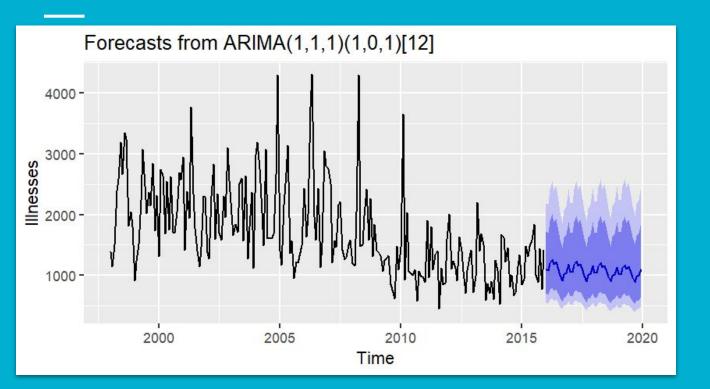
- The model mean's forecast appears to match mean of data
- But it doesn't appear to capture seasonal volatility well

Forecast for ARIMA(1, 1, 0)



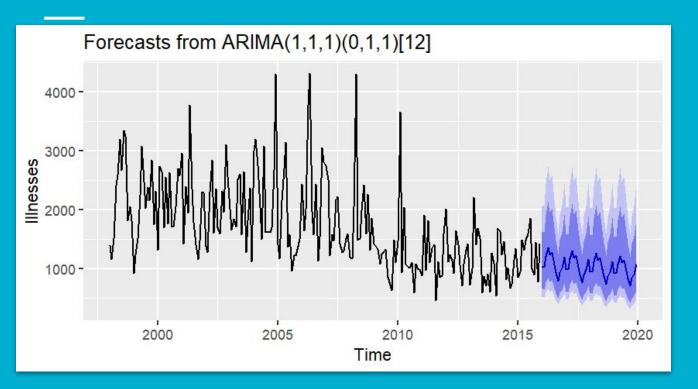
- Clearly not an ideal forecast due to the high upper CI bound
- However, the mean forecast appears to be accurate

Forecast for ARIMA(1,1,1)x(1,0,1)12



Good
 performance
 for forecasting
 compared to
 actuals

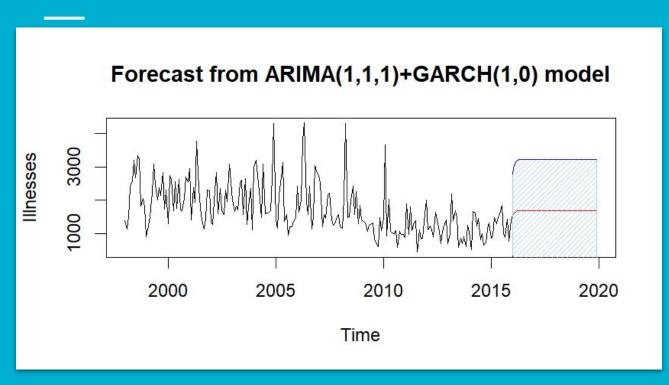
Forecast for ARIMA(1,1,1)x(0,1,1)12



- This model

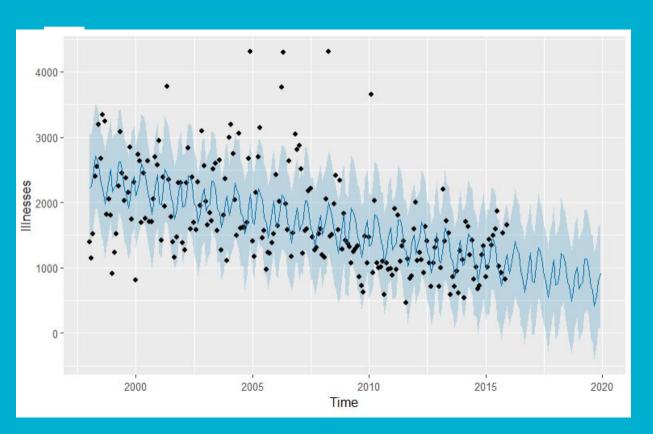
 appears to
 capture more
 of volatility
- This model did not have a significant AR1 term

Forecast for ARIMA(1,1,1) + GARCH(1,0)



- Like the non seasonal ARIMA models but the lower CI limit is 0
- Similar pattern for forecasting using log data vs untransformed data

Forecast for Prophet



- The model forecast appears to do a good job of capturing the seasonality of the historical data
- The forecast also appears to have a slight negative trend

Forecasting Comparisons

Model	RMSE
ARIMA(1, 1, 1)	489.42
ARIMA(1, 1, 0)	474.94
ARIMA(1,1,1)x(1,0,1) ₁₂	488.63
ARIMA(1,1,1)x(0,1,1) ₁₂	487.91
ARIMA(1,1,1)-GARCH(1,0)	552.43
Prophet	638

Data from 2016 and beyond

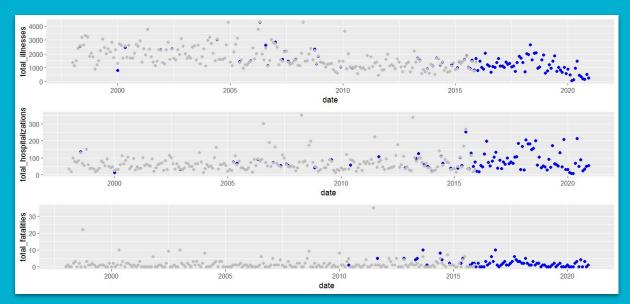
Data1: Foodborne illness data set (1998 - 2015):

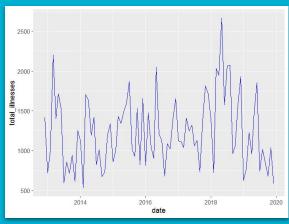
https://www.kaggle.com/datasets/cdc/foodborne-diseases

Data2: National Outbreak Reporting System(1998 - 2020)

https://wwwn.cdc.gov/norsdashboard

Validate 2016 and beyond data





2013 - 2020 illness from data 2 Choose 2016 - 2019(48 month) For forecast comparing

Real data and forecasting of ARIMA(1,1,1)

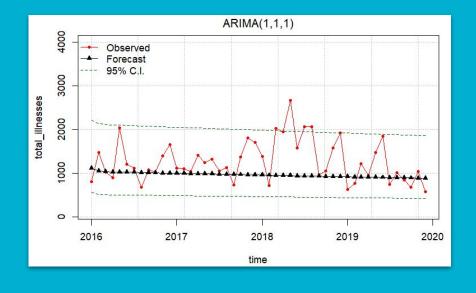
Model output from data before 2016

Forecast for data between 2016 - 2019

Data 1

Data 2

\$ttable
Estimate SE t.value p.value
ar1 0.1998 0.0714 2.7971 0.0056
ma1 -0.9337 0.0216 -43.2306 0.0000



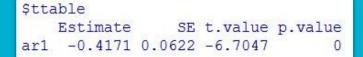
Real data and forecasting of ARIMA(1,1,0)

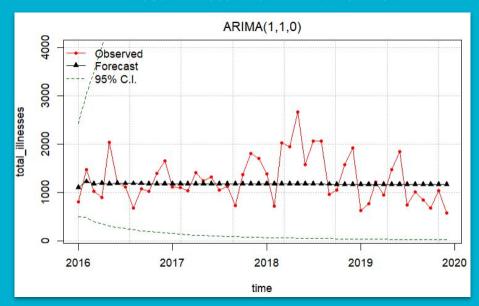
Model output from data before 2016

Data 1

```
$ttable
Estimate SE t.value p.value
ar1 -0.4078 0.0624 -6.535 0
```

Data 2





Real data and forecasting of SARIMA(1,1,1)x(1,0,1)

0.0426

0.0000

0.0000

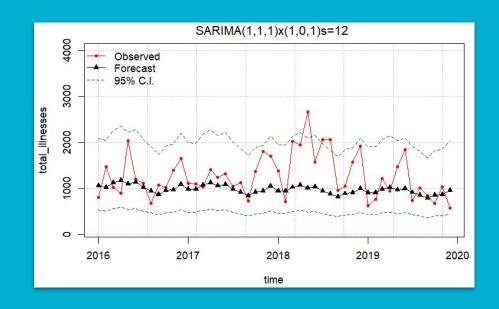
0.0000

Model output from data before 2016

\$ttable Estimate t.value p.value Data 1 0.1692 0.0729 2.3201 0.0213 -0.9403 0.0226 -41.5258 0.0000 ma1 0.9509 0.0912 10.4234 0.0000 sar1 sma1 -0.8796 0.1443 -6.09480.0000

Estimate t.value p.value Data 2 0.1501 0.0736 2.0395 ar1 -0.9375 0.0237 -39.5128 ma1 0.9474 0.0951 9.9615 sar1 -0.8755 0.1472 -5.9484sma1

Sttable



Real data and forecasting of SARIMA(1,1,1)x(0,1,1)₁₂

Model output from data before 2016

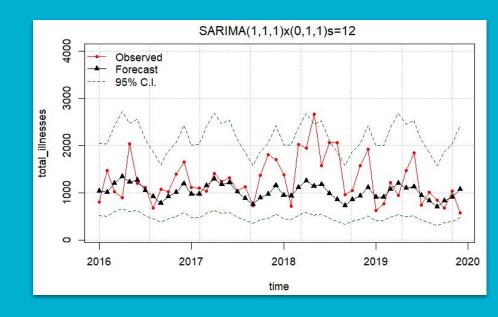
Data 1

\$ttal	\$ttable						
A. A. C. A.	Estimate	SE	t.value	p.value			
ar1	0.1371	0.0752	1.8244	0.0696			
ma1	-0.9393	0.0290	-32.3892	0.0000			
sma1	-0.9093	0.0857	-10.6136	0.0000			

Data 2

044-1-1-1

				ore	Şttar
ue	p.valu	t.value	SE	Estimate	
65	0.126	1.5346	0.0759	0.1164	ar1
00	0.000	-31.1513	0.0300	-0.9336	ma1
00	0.000	-11.0127	0.0820	-0.9034	sma1
	0.00	-31.1513	0.0300	-0.9336	ma1



Real data and forecasting of ARIMA(1,1,1)-GARCH(1,0)

Model output from data before 2016

Error Analysis: Estimate Std. Error t value Pr(>|t|) 3.65004 0.47009 0.50416 0.06355 0.14052 0.01639 < 2e-16 *** omega 0.836

0.06644

0.207

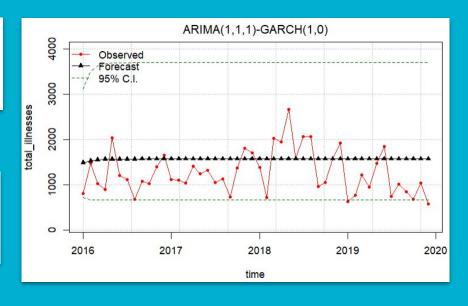
Data 2

Data 1

alpha1

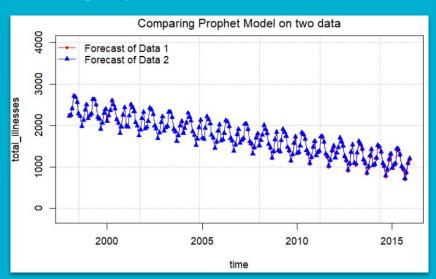
0.01375

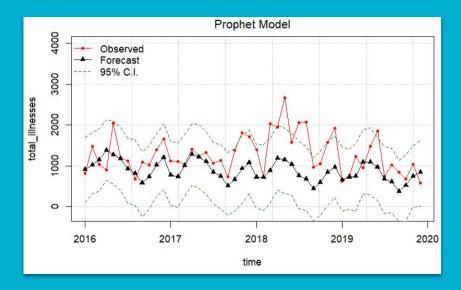
Error A	Analysis:				
	Estimate	Std. Error	t value	Pr(> t)	
mu	3.74032	0.48335	7.738	9.99e-15	***
ar1	0.49186	0.06530	7.532	5.00e-14	***
omega	0.14107	0.01671	8.444	< 2e-16	***
alpha1	0.03958	0.07106	0.557	0.578	



Real data and forecasting of Prophet Model

Fitting output from data before 2016





Conclusions

Key takeaways

- 1. Model formulation and estimation. We constructed six models (two ARIMA, two SARIMA, ARIMA-GARCH, Prophet) and compared their fit and forecasts.
- 2. Model selection. We chose the ARIMA(1, 1, 1) \times (1, 0, 1)₁₂ model, based on these criteria:
 - The seasonal ARIMA model has lower AIC and BIC values.
- The forecast captures the seasonality of foodborne disease outbreaks better than non-seasonal models.
- 3. Validation. The RMSE of the forecast is the lowest of all models that include seasonality.

Learnings

Key Concepts:

- Importance of the data
- Time changes everything
- Impact of the Data

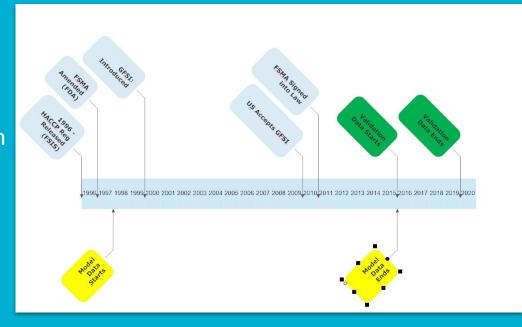
Importance of Data

- Data Cleaning
 - Source of the information
 - Understanding the basics of the project
 - Consumer habits
 - 2022 Sal 572 (92 / 2)Backyard Poultry



Time Changes Everything

- Regulatory Changes
 - Time Delays in Implementation
- Improvements in Data Collection
- Improvements in Testing



"Salmonella contamination on broiler chickens (carcasses) decreased by 56 percent from 1995, before the HACCP final rule was announced, to 2000. The number of foodborne illness cases attributed to Salmonella on broilers was 190,000 lower in 2000 than in 1995." — Williams and Ebel study, 2012

Outbreaks

- 1999 Hot Dogs LM 100
- 2006 Spinach EC 205
- 2006 Taco Bell EC 71
- 2009 -Peanut Butter SAL 714
- 2011 Canteloupes LM 147
- 2011 Ground Turkey SAL 136
- 2013 Chicken Sal 634
- 2015 Chipotle EC 55
- 2015 Mexi Cucumbers SAL -907

- 2016 Flour EC 63
- 2017 Leafy Greens EC 25
- 2018 Romaine EC 272
- 2019 Ground Beef EC 209
- 2019 Flour EC 167



Impacts of the Data

- Determines success of the industry
- Basis for Regulatory Goals and Changes
 - Healthy People Goals CDC
 - USDA / FDA Strategy
- Regulatory Changes -
 - Sal Adulterant Kiev / Cordon Blue
 - Flour Mills
 - Exclusion of Supply





USDA Declares Salmonella an Adulterant in Breaded Stuffed Raw Chicken Products

"Today's announcement is an important moment in U.S. food safety because we are declaring Salmonella an adulterant in a raw poultry product," said Sandra Eskin, USDA Deputy Under Secretary for Food Safety. "This is just the beginning of our efforts to improve public health."

Questions, comments, suggestions?

Thank you for your engagement and feedback!

Surprisingly good of Prophet Model

