

# Covid-19 Infection Analysis

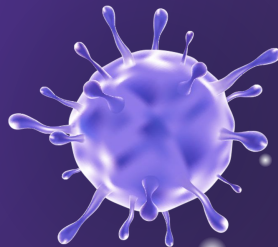


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Program: Data Analytics

Group Name: Endless Knot

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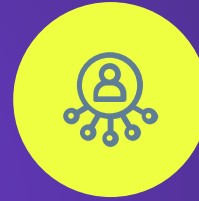


# INTRODUCTION



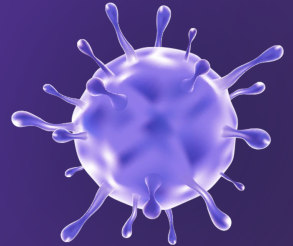
## TOPIC

What is the likelihood of being infected by Covid-19? How is infection affected by factors such as vaccination rates, gender, and ethnicity?



## PURPOSE

It is important to analyze future trends following the Covid-19 pandemic to understand the prevalence of infection within the American population.






## Questions Answered

- Are certain populations more likely to be infected than others?
- How do these factors affect the other?
- What other factors should be considered in identifying risks of infection?



## Data Sources

We gathered data from reliable organizations such as New York Times and the Center for Disease Control (CDC) which provide csv files on their findings. Our data identifies:

- vaccination rates
  - gender ratios
  - ethnicity statistics
- 



## Entity Relationship Diagram



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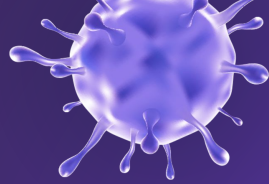
## Entity Relationship Diagram

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- ## Entity Relationship Diagram

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# Data Analysis

- Ordinary Least Squares (OLS)
- Linear regression
- SVM support vector machine
- Autoregressive Integrated Moving Average (ARIMA) for Time series model

coefficient of determination: 0.5703788862651746

## OLS Regression Results

```
=====
Dep. Variable:          Date      R-squared:                0.607
Model:                  OLS       Adj. R-squared:           0.599
Method:                 Least Squares
Date:                  Sat, 02 Apr 2022
Time:                  00:12:10   Prob (F-statistic):       0.00
No. Observations:      5320      Log-Likelihood:          -48837.
Df Residuals:          5212      AIC:                    9.789e+04
Df Model:              107       BIC:                    9.860e+04
Covariance Type:       nonrobust
=====
```

## SARIMAX Results

```
=====
Dep. Variable:          recovered   No. Observations:      5320
Model:                  ARIMA(5, 1, 0)   Log Likelihood        -55067.188
Date:                  Fri, 01 Apr 2022   AIC                   110146.376
Time:                  23:52:26          BIC                   110185.850
Sample:                0              HQIC                  110160.168
Covariance Type:       opg
=====
              coef    std err          z      P>|z|    [0.025    0.975]
-----
ar.L1         -0.8148      0.007    -124.881    0.000     -0.828     -0.802
ar.L2         -0.7459      0.008    -97.523    0.000     -0.761     -0.731
ar.L3         -0.5328      0.007    -76.355    0.000     -0.546     -0.519
ar.L4         -0.3472      0.006    -58.862    0.000     -0.359     -0.336
ar.L5         -0.2130      0.004    -52.946    0.000     -0.221     -0.205
sigma2        5.937e+07  1.79e-10   3.31e+17    0.000   5.94e+07   5.94e+07
=====
Ljung-Box (L1) (Q):                2.19   Jarque-Bera (JB):        5558677.83
Prob(Q):                          0.14   Prob(JB):                0.00
Heteroskedasticity (H):            0.12   Skew:                    6.56
Prob(H) (two-sided):              0.00   Kurtosis:               160.83
=====
```

### Warnings:

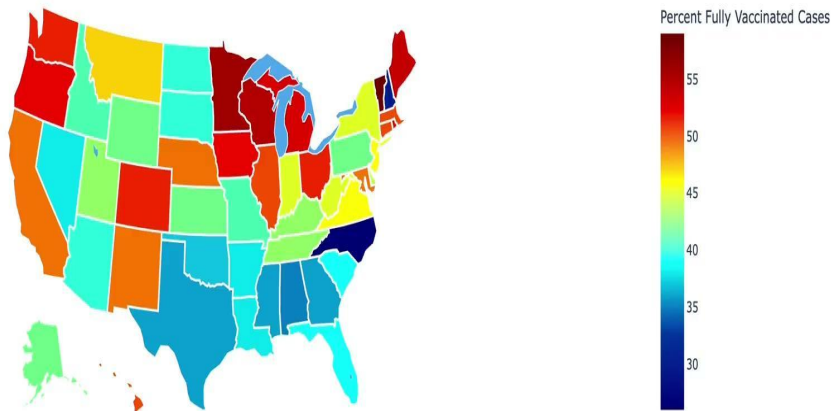
- [1] Covariance matrix calculated using the outer product of gradients (complex-step).
- [2] Covariance matrix is singular or near-singular, with condition number 5.21e+32. Standard errors may be unstable.





# Interactive Element

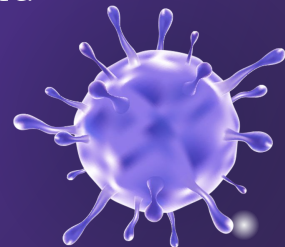
US Vaccination Rates



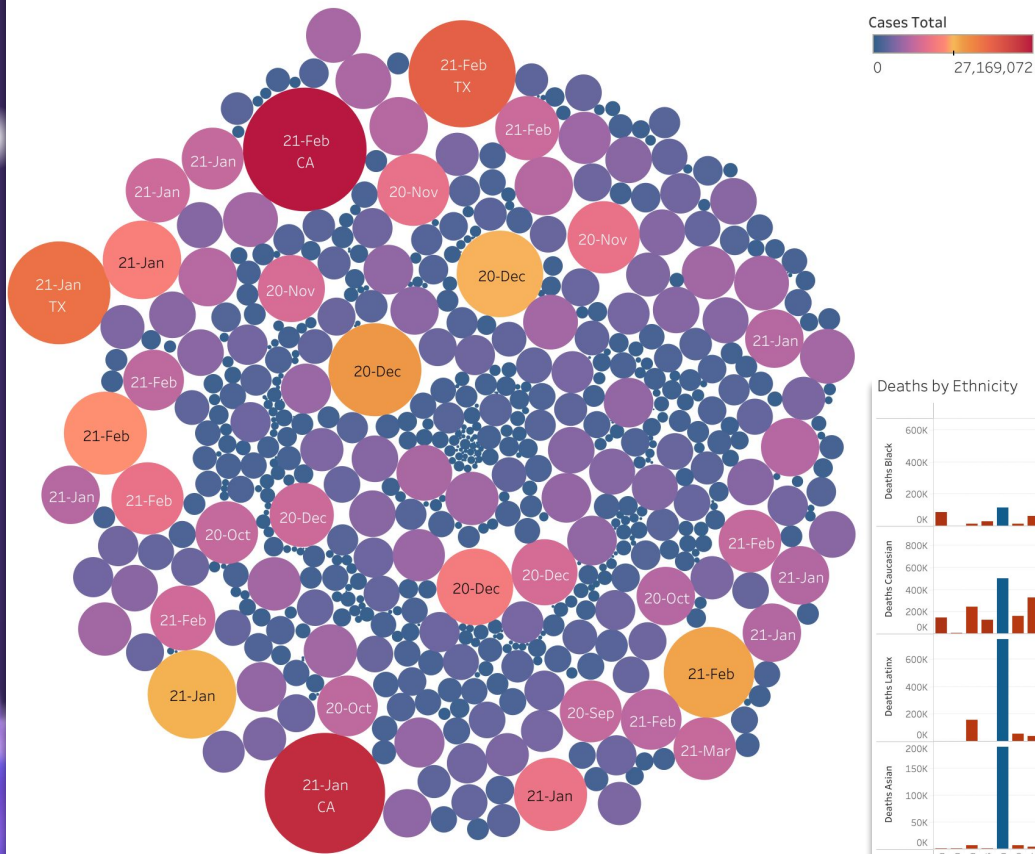
*Vaccination Rates by State*

Plotly

- Map with options to toggle different factors of each state such as:
  - Vaccination rates
  - Infections
  - Gender data



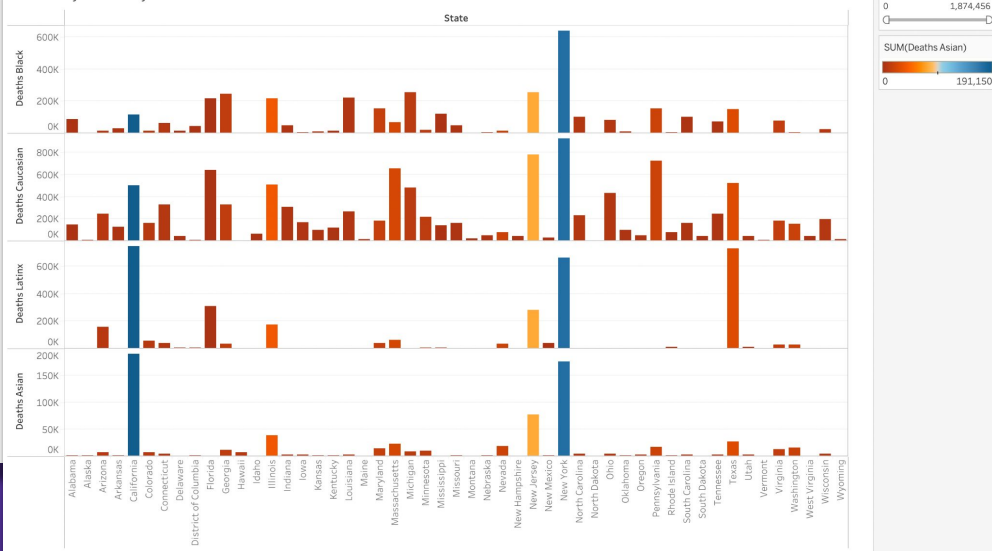
## All Cases Bubble



# Visualization

- Tableau
- Correlation graphs

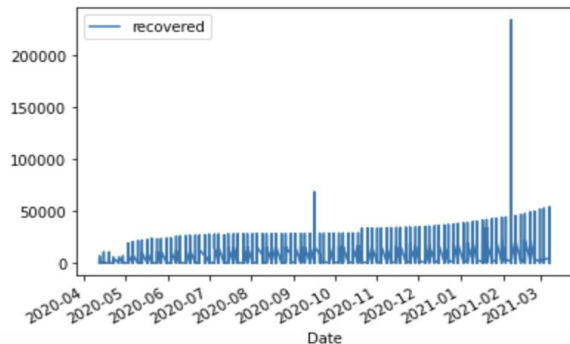
## Deaths by Ethnicity



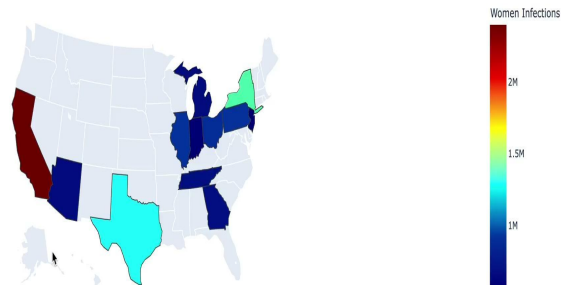
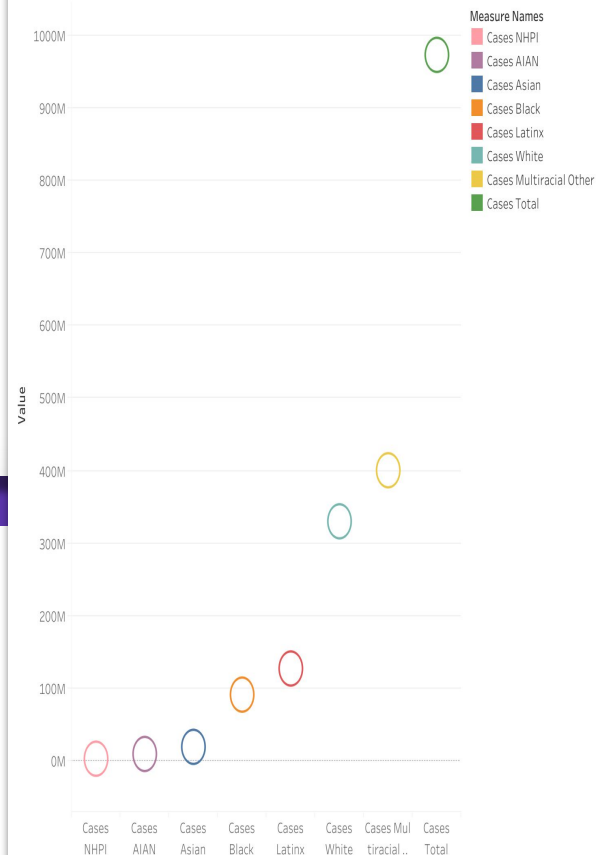
# Dashboard

recovered

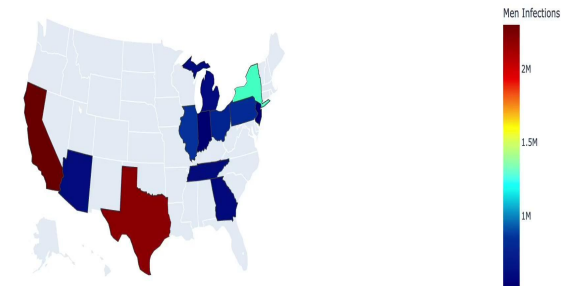
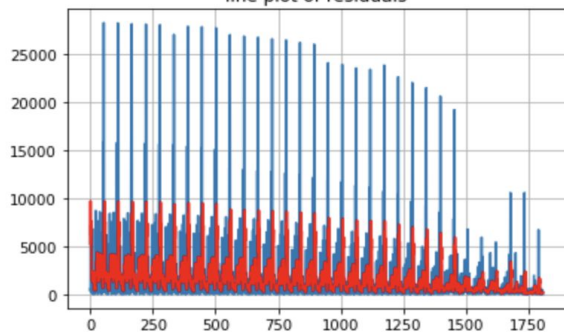
Date	recovered
2021-03-07	305.0
2021-03-07	10148.0
2021-03-07	5319.0
2021-03-07	1233.0
2021-03-07	16328.0



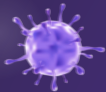
All Covid Cases Graph



line plot of residuals







# Results of Analysis

- No significant differences between gender infection rates
- More vaccinations → lower cases and death rates
- People who were vaccinated and boosted had lower amounts of cases and fatalities
- On the time series analysis, the line plot is created, showing the expected values (blue) compared to the rolling forecast predictions (red). Values show some trends and are in the correct scale.





## What We Would Do Differently

- Before deciding on a topic, look into data resources
- Compare more factors from different data sets
  - Identify more primary keys to connect data
- Added other factors such as economic impact

## Recommendations for Future Analysis

- Statistical analysis can be used with bigger data sets
- Explore other factors that may have a correlation with the data we already have
- Expand to global research
  - countries that are not as ethnically homogenous
- Project future peak infections



# Gender

<https://www.genderscilab.org/gender-and-sex-in-covid19/#CaseDeathRatebySex>

# Ethnicity

<https://covidtracking.com/race/dashboard>



# Vaccination

[d.cdc.gov/covid-data-tracker/#vaccinations\\_vacc-total-admin-rate-total](https://d.cdc.gov/covid-data-tracker/#vaccinations_vacc-total-admin-rate-total)

Check out our project at:  
[https://github.com/antirose/  
CovidInfectionAnalysis](https://github.com/antirose/CovidInfectionAnalysis)

