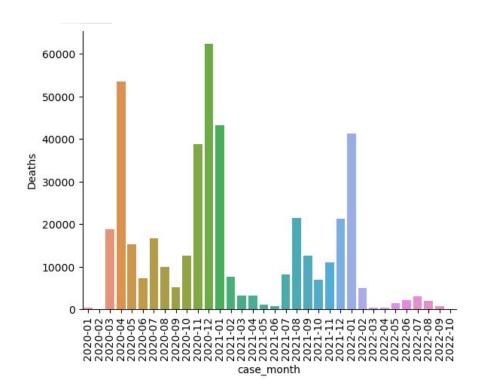
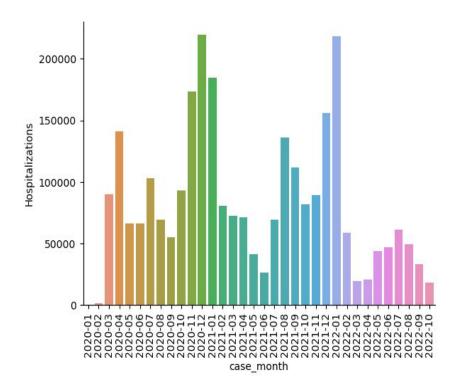
Effects of Geography & Demography on COVID-19 Hospitalizations and Deaths (Part 2)

- Daniel Johnson
- Brian Li
- Braden Lockwood
- Evan Powell
- Jowaki Merani
- Mason Hawks

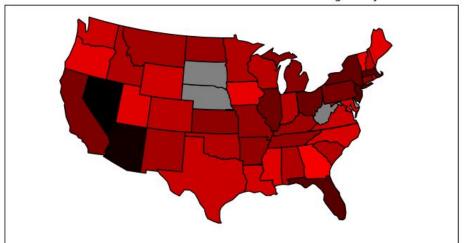
Statistics & EDA Results

- → Target (hosp/death) visualizations over different features (age, month, etc)
- → State ranking T-tests for hosps/deaths per capita
- → Visualizations for illness severity by age and race
- → Feature ranking based on P-values and critical regions

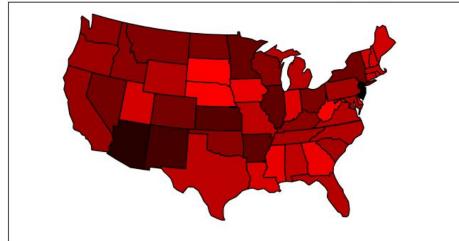




Deaths Per 100,000 Max=Black=324.85 Min=Red=4.95 Missing=Gray



Hospitalizations Per 100,000 Max=Black=2173.65 Min=Red=0.09 Missing=Gray



```
top_25_pop_deathspc = [deaths_pc[state] for state in top_25_population]
                      bottom_25_pop_deathspc = [deaths_pc[state] for state in bottom_25_population]
                      pop_pval = ttest_ind(top_25_pop_deathspc, bottom_25_pop_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 Population Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                   Python
                  P-Value for Means of Top 25 vs Bottom 25 Population Deaths Per Capita: 0.048
< 0.05
                      top_10_pop_deathspc = [deaths_pc[state] for state in top_10_population]
                      bottom_40_pop_deathspc = [deaths_pc[state] for state in bottom_40_population]
                      pop_pval = ttest_ind(top_10_pop_deathspc, bottom_40_pop_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 Population Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                   Python
                  P-Value for Means of Top 10 vs Bottom 40 Population Deaths Per Capita: 0.089
> 0.05
                      top_25_pop_hospspc = [hosps_pc[state] for state in top_25_population]
                      bottom_25_pop_hospspc = [hosps_pc[state] for state in bottom_25_population]
                      pop_pval = ttest_ind(top_25_pop_hospspc, bottom_25_pop_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 Population Hosps Per Capita: {pop_pval/2:.3f}")
                                                                                                                                    Python
> 0.05
                   P-Value for Means of Top 25 vs Bottom 25 Population Hosps Per Capita: 0.230
                      top_10_pop_hospspc = [hosps_pc[state] for state in top_10_population]
                      bottom_40_pop_hospspc = [hosps_pc[state] for state in bottom_40_population]
                      pop_pval = ttest_ind(top_10_pop_hospspc, bottom_40_pop_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 Population Hosps Per Capita: {pop_pval/2:.3f}")
                                                                                                                                    Python
> 0.05
                   P-Value for Means of Top 10 vs Bottom 40 Population Hosps Per Capita: 0.427
```

```
top_25_education_deathspc = [deaths_pc[state] for state in top_25_education]
                      bottom_25_education_deathspc = [deaths_pc[state] for state in bottom_25_education]
                      pop_pval = ttest_ind(top_25_education_deathspc, bottom_25_education_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 Education Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                    Python
> 0.05
                  P-Value for Means of Top 25 vs Bottom 25 Education Deaths Per Capita: 0.383
                      top_10_education_deathspc = [deaths_pc[state] for state in top_10_education]
                      bottom_40_education_deathspc = [deaths_pc[state] for state in bottom_40_education]
                      pop_pval = ttest_ind(top_10_education_deathspc, bottom_40_education_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 Education Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                    Python
                  P-Value for Means of Top 10 vs Bottom 40 Education Deaths Per Capita: 0.400
> 0.05
                      top_25_education_hospspc = [hosps_pc[state] for state in top_25_education]
                      bottom_25_education_hospspc = [hosps_pc[state] for state in bottom_25_education]
                      pop_pval = ttest_ind(top_25_education_hospspc, bottom_25_education_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 Education Hosps Per Capita: {pop_pval/2:.3f}")
> 0.05
               \cdots P-Value for Means of Top 25 vs Bottom 25 Education Hosps Per Capita: 0.167
                      top_10_education_hospspc = [hosps_pc[state] for state in top_10_education]
                      bottom_40_education_hospspc = [hosps_pc[state] for state in bottom_40_education]
                      pop_pval = ttest_ind(top_10_education_hospspc, bottom_40_education_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 Education Hosps Per Capita: {pop_pval/2:.3f}")
               「487 ✓ 0.1s
                                                                                                                                    Python
                  P-Value for Means of Top 10 vs Bottom 40 Education Hosps Per Capita: 0.206
> 0.05
```

```
top_25_GDP_deathspc = [deaths_pc[state] for state in top_25_GDP]
                      bottom_25_GDP_deathspc = [deaths_pc[state] for state in bottom_25_GDP]
                      pop_pval = ttest_ind(top_25_GDP_deathspc, bottom_25_GDP_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 GDP Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                   Python
                  P-Value for Means of Top 25 vs Bottom 25 GDP Deaths Per Capita: 0.039
< 0.05
                      top_10_GDP_deathspc = [deaths_pc[state] for state in top_10_GDP]
                      bottom_40_GDP_deathspc = [deaths_pc[state] for state in bottom_40_GDP]
                      pop_pval = ttest_ind(top_10_GDP_deathspc, bottom_40_GDP_deathspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 GDP Deaths Per Capita: {pop_pval/2:.3f}")
                                                                                                                                   Python
> 0.05
                  P-Value for Means of Top 10 vs Bottom 40 GDP Deaths Per Capita: 0.057
                      top_25_GDP_hospspc = [hosps_pc[state] for state in top_25_GDP]
                      bottom_25_GDP_hospspc = [hosps_pc[state] for state in bottom_25_GDP]
                      pop_pval = ttest_ind(top_25_GDP_hospspc, bottom_25_GDP_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 25 vs Bottom 25 GDP Hosps Per Capita: {pop_pval/2:.3f}")
               7497 ✓ 0.1s
                                                                                                                                    Python
> 0.05
                  P-Value for Means of Top 25 vs Bottom 25 GDP Hosps Per Capita: 0.160
                      top_10_GDP_hospspc = [hosps_pc[state] for state in top_10_GDP]
                      bottom_40_GDP_hospspc = [hosps_pc[state] for state in bottom_40_GDP]
                      pop_pval = ttest_ind(top_10_GDP_hospspc, bottom_40_GDP_hospspc, equal_var=False).pvalue
                      print(f"P-Value for Means of Top 10 vs Bottom 40 GDP Hosps Per Capita: {pop_pval/2:.3f}")
               7507 ✓ 0.1s
                                                                                                                                    Python
> 0.05
                  P-Value for Means of Top 10 vs Bottom 40 GDP Hosps Per Capita: 0.182
```

```
blue_states_deathspc = [deaths_pc[state] for state in blue_states]
    red_states_deathspc = [deaths_pc[state] for state in red_states]
    pop_pval = ttest_ind(blue_states_deathspc, red_states_deathspc, equal_var=False).pvalue
    print(f"P-Value for Means of Red State vs Blue State Deaths Per Capita: {pop_pval/2:.3f}")

V 0.1s

P-Value for Means of Red State vs Blue State Deaths Per Capita: 0.116
```

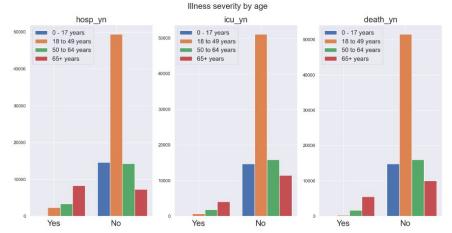
> 0.05

> 0.05

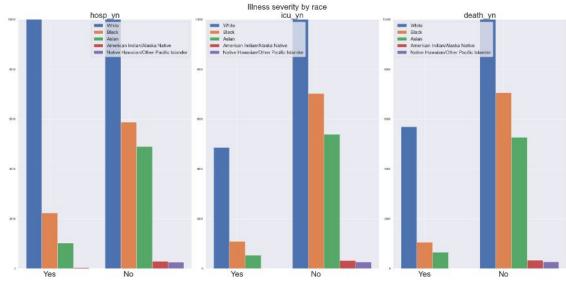
```
blue_states_hospspc = [hosps_pc[state] for state in blue_states]
red_states_hospspc = [hosps_pc[state] for state in red_states]
pop_pval = ttest_ind(blue_states_hospspc, red_states_hospspc, equal_var=False).pvalue
print(f"P-Value for Means of Red State vs Blue State Hosps Per Capita: {pop_pval/2:.3f}")

[51] 
V 0.1s

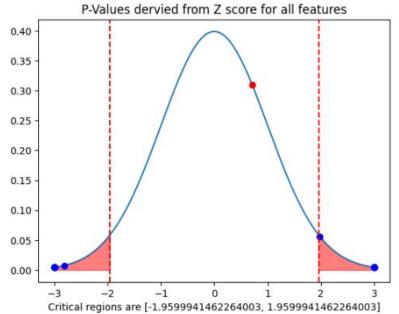
P-Value for Means of Red State vs Blue State Hosps Per Capita: 0.123
```



As you can see there is major imbalances in our dataset. For example there are vastly more many datapoint for people ages 18-48 then other ages.



Here we can see most features are statistically significant with one being not statistically significant. This poses a challenge in feature selection since the statistical test shows that almost all features are significant.



state_fips_code

res_state

res_county

age_group

race

ethnicity process

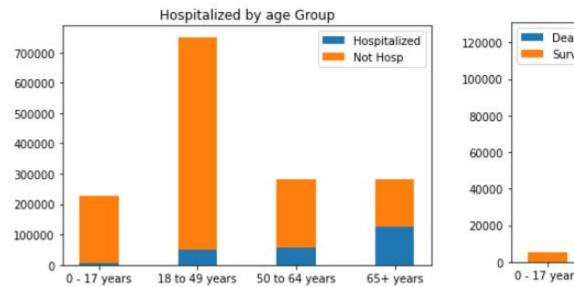
exposure_yn current_status

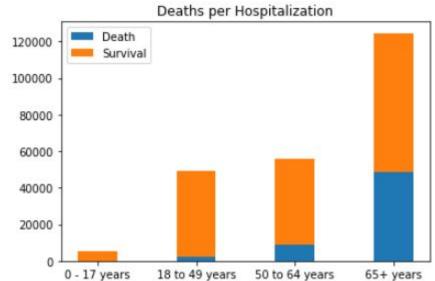
symptom_status

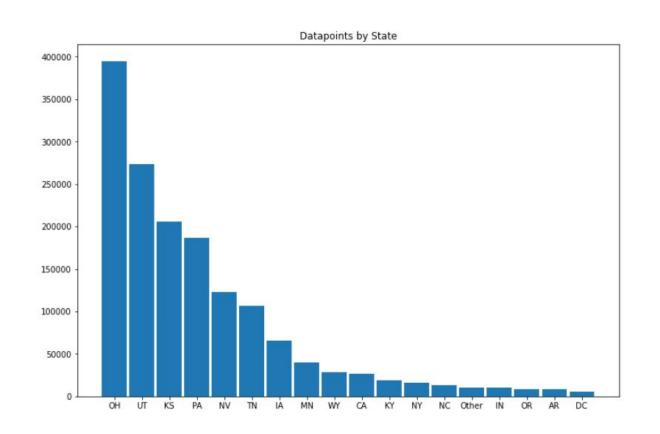
county_fips_code case positive specimen

case_onset_interval







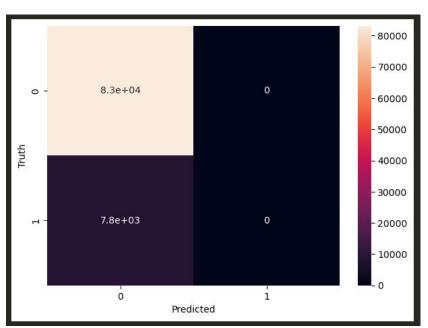


Classification

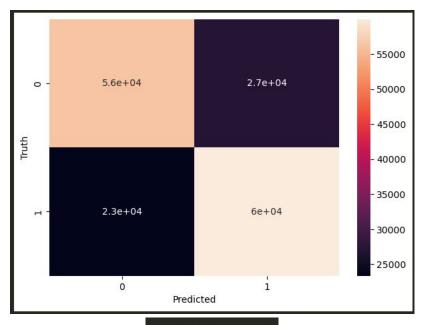
- Tried to classify hospitalization, death
- Models used: KNN, Decision Trees, NBC
- Techniques used: bagging, SMOTE, feature selection

Confusion matrix - KNN (Hospitalization)





After SMOTE



0.9138499406619489

0.6975989812039334

Scores - KNN (hospitalization)

Before SMOTE

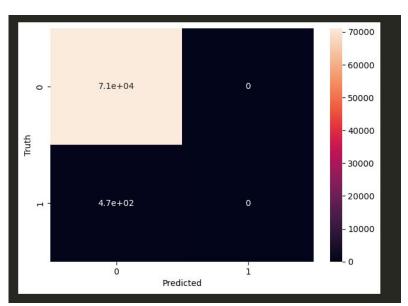
| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|------------|
| 9 | 0.91 | 1.00 | 0.95 | 83164 |
| 1 | 0.00 | 0.00 | 0.00 | 7840 |
| | | | | V.2/2/2/20 |
| accuracy | | | 0.91 | 91004 |
| macro avg | 0.46 | 0.50 | 0.48 | 91004 |
| weighted avg | 0.84 | 0.91 | 0.87 | 91004 |

After SMOTE

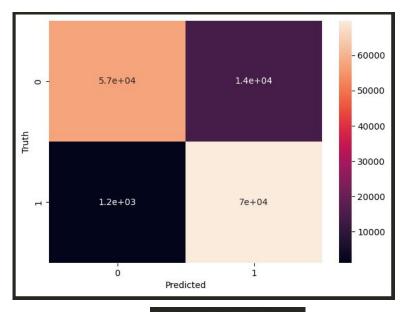
| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.71 | 0.68 | 0.69 | 83146 |
| 1 | 0.69 | 0.72 | 0.70 | 83325 |
| accuracy | | | 0.70 | 166471 |
| macro avg | 0.70 | 0.70 | 0.70 | 166471 |
| weighted avg | 0.70 | 0.70 | 0.70 | 166471 |

Confusion matrix - KNN (Death)

Before SMOTE



After SMOTE



0.9934316260219411

0.8942884079587988

Scores - KNN (Deaths)

Before SMOTE

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 9 | 0.99 | 1.00 | 1.00 | 71085 |
| 1 | 0.00 | 0.00 | 0.00 | 470 |
| accuracy | | | 0.99 | 71555 |
| macro avg | 0.50 | 0.50 | 0.50 | 71555 |
| weighted avg | 0.99 | 0.99 | 0.99 | 71555 |

After SMOTE

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.98 | 0.81 | 0.88 | 71199 |
| 1 | 0.83 | 0.98 | 0.90 | 70933 |
| accuracy | | | 0.89 | 142132 |
| macro avg | 0.91 | 0.89 | 0.89 | 142132 |
| weighted avg | 0.91 | 0.89 | 0.89 | 142132 |

KNN Summary

- Model performed marginally better with more than 20 neighbors.
- No significant improvement after 100 neighbors. **0.6161553664001538**

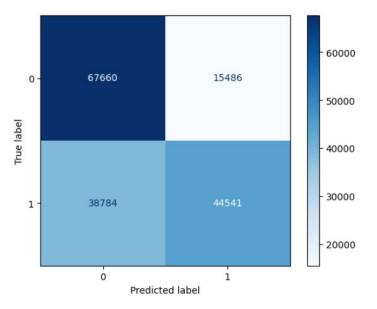
0.6975989812039334

- Training time was long.
- Predicting time was actually around the same.
- Bagging no interesting results to show.

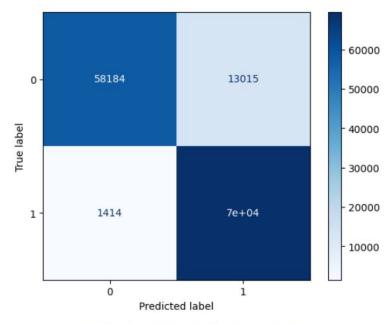


Conclusion: bad data

Decision Trees: Hosp. and Death

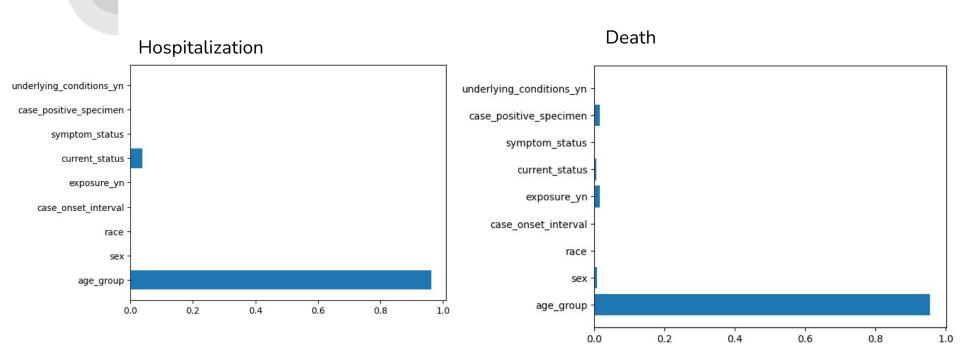


| | | precision | recall | f1-score | support |
|----------|------|-----------|--------|----------|---------|
| | 0 | 0.64 | 0.81 | 0.71 | 83146 |
| | 1 | 0.74 | 0.53 | 0.62 | 83325 |
| accui | racy | | | 0.67 | 166471 |
| macro | avg | 0.69 | 0.67 | 0.67 | 166471 |
| weighted | avg | 0.69 | 0.67 | 0.67 | 166471 |

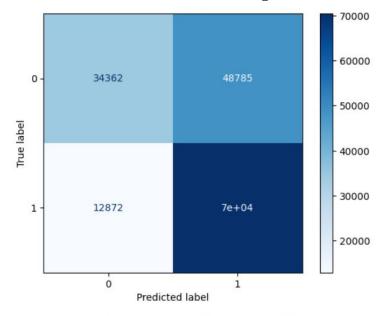


| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.98 | 0.82 | 0.89 | 71199 |
| 1 | 0.84 | 0.98 | 0.91 | 70933 |
| accuracy | | | 0.90 | 142132 |
| macro avg | 0.91 | 0.90 | 0.90 | 142132 |
| weighted avg | 0.91 | 0.90 | 0.90 | 142132 |

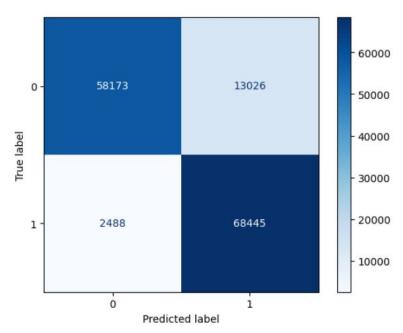
Tree Feature Importance



NBC: Hosp. and Death



| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.73 | 0.41 | 0.53 | 83147 |
| 1 | 0.59 | 0.85 | 0.70 | 83324 |
| accuracy | | | 0.63 | 166471 |
| macro avg | 0.66 | 0.63 | 0.61 | 166471 |
| weighted avg | 0.66 | 0.63 | 0.61 | 166471 |



| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.96 | 0.82 | 0.88 | 71199 |
| 1 | 0.84 | 0.96 | 0.90 | 70933 |
| accuracy | | | 0.89 | 142132 |
| macro avg | 0.90 | 0.89 | 0.89 | 142132 |
| weighted avg | 0.90 | 0.89 | 0.89 | 142132 |

Classification: Some Conclusions and Concerns

- Data points did not take into account time or geography
- No incorporation of vaccine impacts
- Hosp Vs. Death