

Network Systems
Science & Advanced
Computing
Biocomplexity Institute
& Initiative
University of Virginia

Estimation of COVID-19 Impact in Virginia

October 5th, 2022

(data current to Sept 30th – October 4th)

Biocomplexity Institute Technical report: TR BI-2022-1770



BIOCOMPLEXITY INSTITUTE

biocomplexity.virginia.edu

About Us

- Biocomplexity Institute at the University of Virginia
 - Using big data and simulations to understand massively interactive systems and solve societal problems
- Over 20 years of crafting and analyzing infectious disease models
 - Pandemic response for Influenza, Ebola, Zika, and others



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Overview

- **Goal:** Understand impact of COVID-19 mitigations in Virginia
- **Approach:**
 - Calibrate explanatory mechanistic model to observed cases
 - Project based on scenarios for next 4 months
 - Consider a range of possible mitigation effects in "what-if" scenarios
- **Outcomes:**
 - Ill, Confirmed, Hospitalized, ICU, Ventilated, Death
 - Geographic spread over time, case counts, healthcare burdens

Key Takeaways

Projecting future cases precisely is impossible and unnecessary.

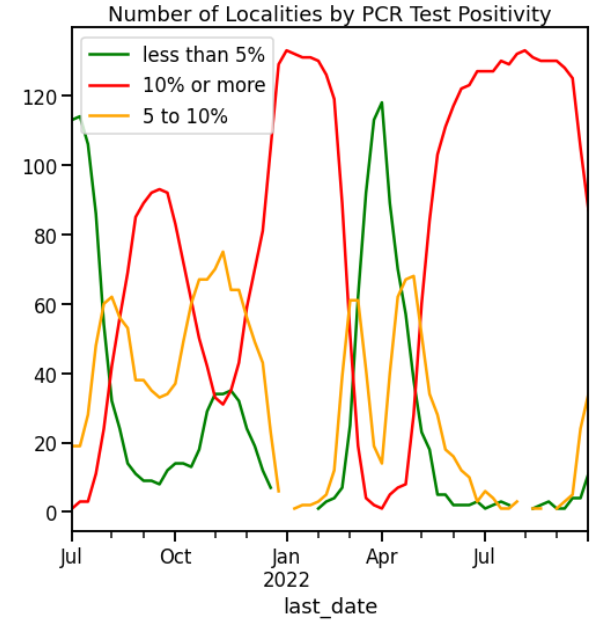
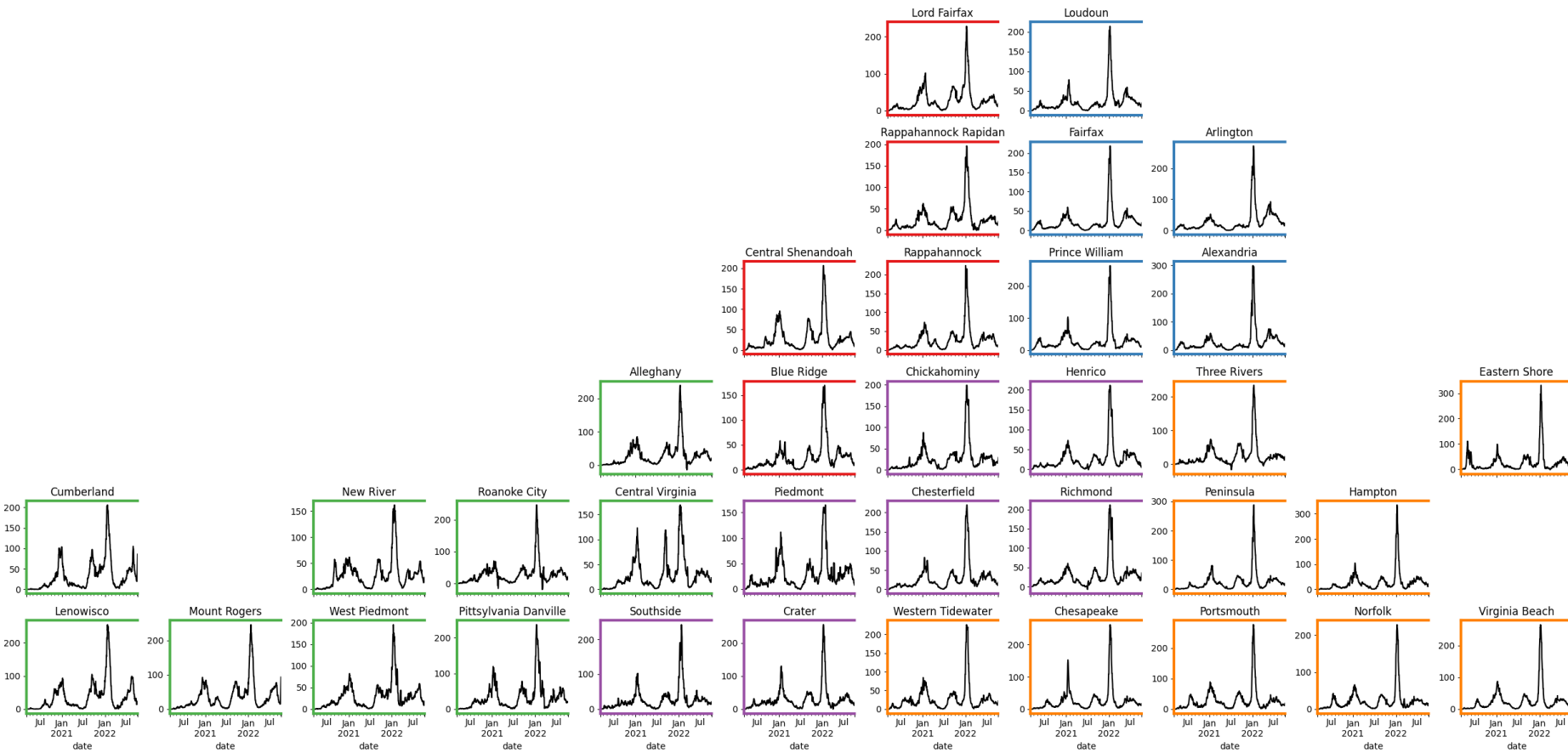
Even without perfect projections, we can confidently draw conclusions:

- **Case rates continue their decline, hospitalizations continue steady decline**
- VA weekly case rate continues decline to 99/100K from 109/100K
 - US weekly case rate is down considerably as well at 89/100K from 105/100K
 - VA hospital occupancy (rolling 7 day mean of 545 down from 599 a week ago) has continued to decline
- Projections anticipate continued declines in cases as well as hospitalizations
- Potential for rebounds due to seasonal forces and/or novel sub-variants in the Fall remains
- Model updates:
 - Maintained Booster Scenarios by slowing down the rate of vaccination adopting the rate of 3rd dose rollout
 - Current monitoring still not finding a definite candidate for Variant X, though BQ.1.1 or BA.2.75.2 remain likely, 50% prevalence pushed back to Nov 15th

The situation continues to change. Models continue to be updated regularly.

Situation Assessment

Case Rates (per 100k) and Test Positivity

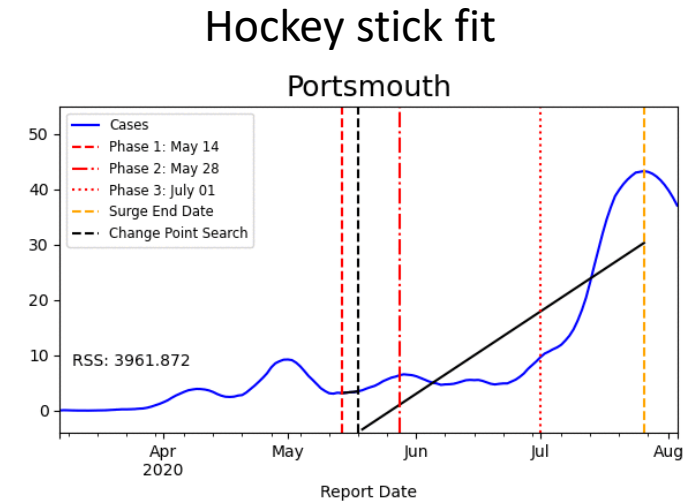


County level RT-PCR test positivity
Green: <5.0% (or <20 tests in past 14 days)
Orange: 5.0%-10.0% (or <500 tests and <2000 tests/100k and >10% positivity over 14 days)
Red: >10.0% (and not "Green" or "Yellow")

District Trajectories

Goal: Define epochs of a Health District's COVID-19 incidence to characterize the current trajectory

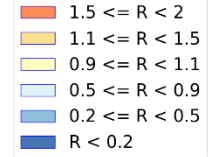
Method: Find recent peak and use hockey stick fit to find inflection point afterwards, then use this period's slope to define the trajectory



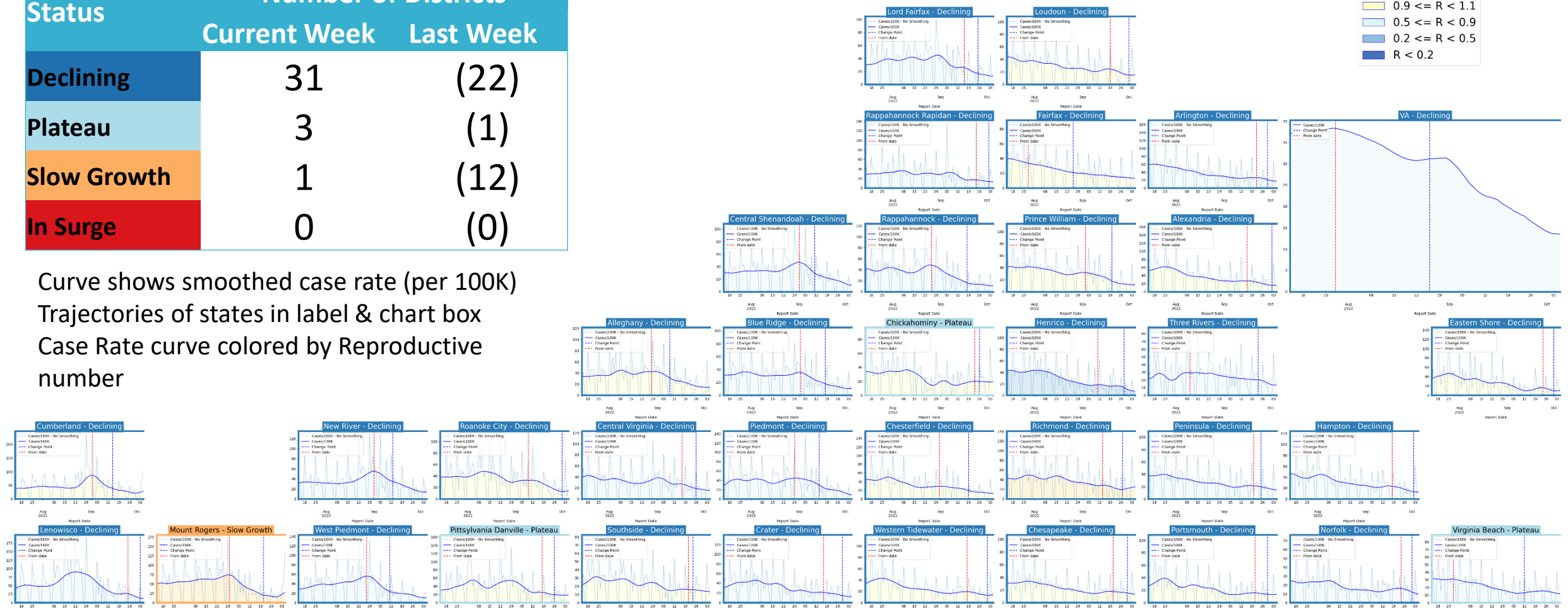
| Trajectory | Description | Weekly Case Rate Slope (per 100k) | Weekly Hosp Rate Slope (per 100k) |
|--------------------|---|-----------------------------------|-----------------------------------|
| Declining | Sustained decreases following a recent peak | slope < -0.88/day | slope < -0.07/day |
| Plateau | Steady level with minimal trend up or down | -0.88/day < slope < 0.42/day | -0.07/day < slope < 0.07/day |
| Slow Growth | Sustained growth not rapid enough to be considered a Surge | 0.42/day < slope < 2.45/day | 0.07/day < slope < 0.21/day |
| In Surge | Currently experiencing sustained rapid and significant growth | 2.45/day < slope | 0.21/day < slope |

District Case Trajectories – last 10 weeks

| Status | Number of Districts | |
|-------------|---------------------|-----------|
| | Current Week | Last Week |
| Declining | 31 | (22) |
| Plateau | 3 | (1) |
| Slow Growth | 1 | (12) |
| In Surge | 0 | (0) |



Curve shows smoothed case rate (per 100K)
 Trajectories of states in label & chart box
 Case Rate curve colored by Reproductive number

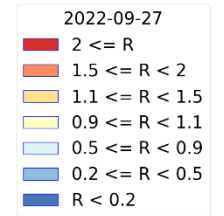


District Hospital Trajectories – last 10 weeks

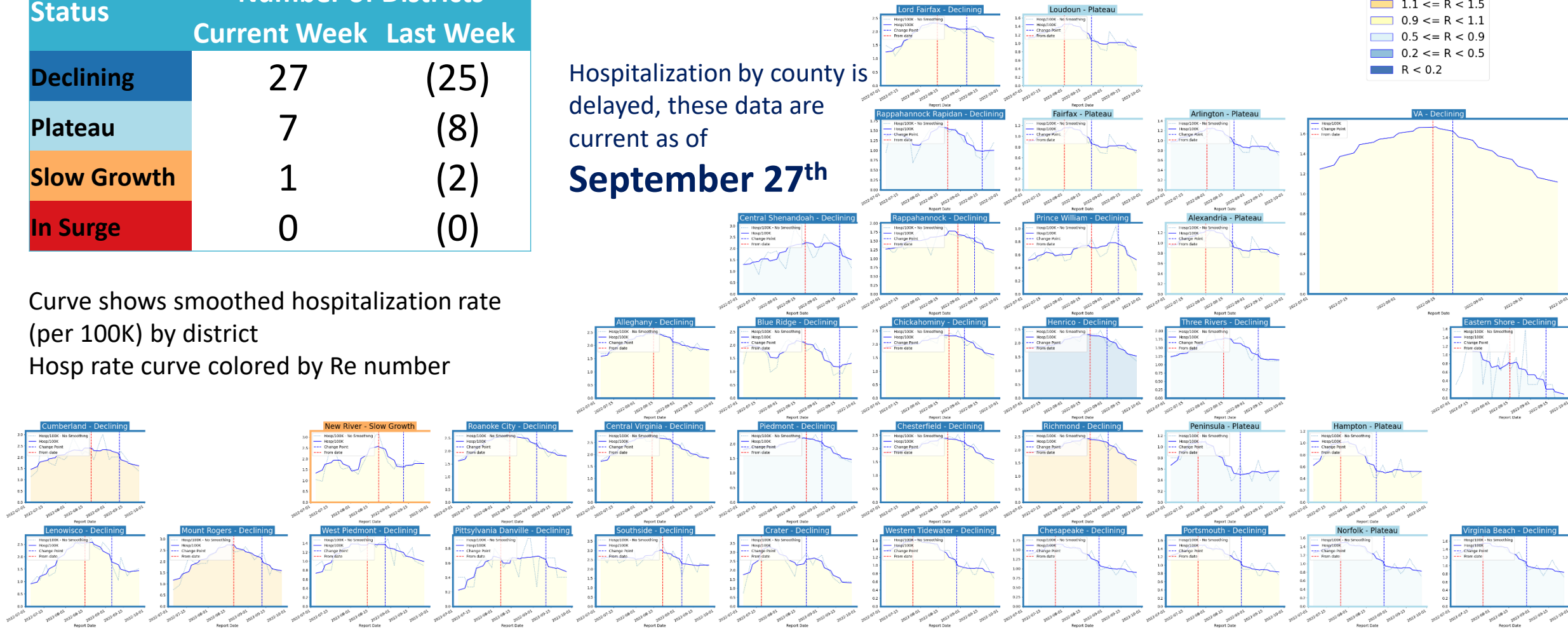
| Status | Number of Districts | |
|-------------|---------------------|-----------|
| | Current Week | Last Week |
| Declining | 27 | (25) |
| Plateau | 7 | (8) |
| Slow Growth | 1 | (2) |
| In Surge | 0 | (0) |

Hospitalization by county is delayed, these data are current as of

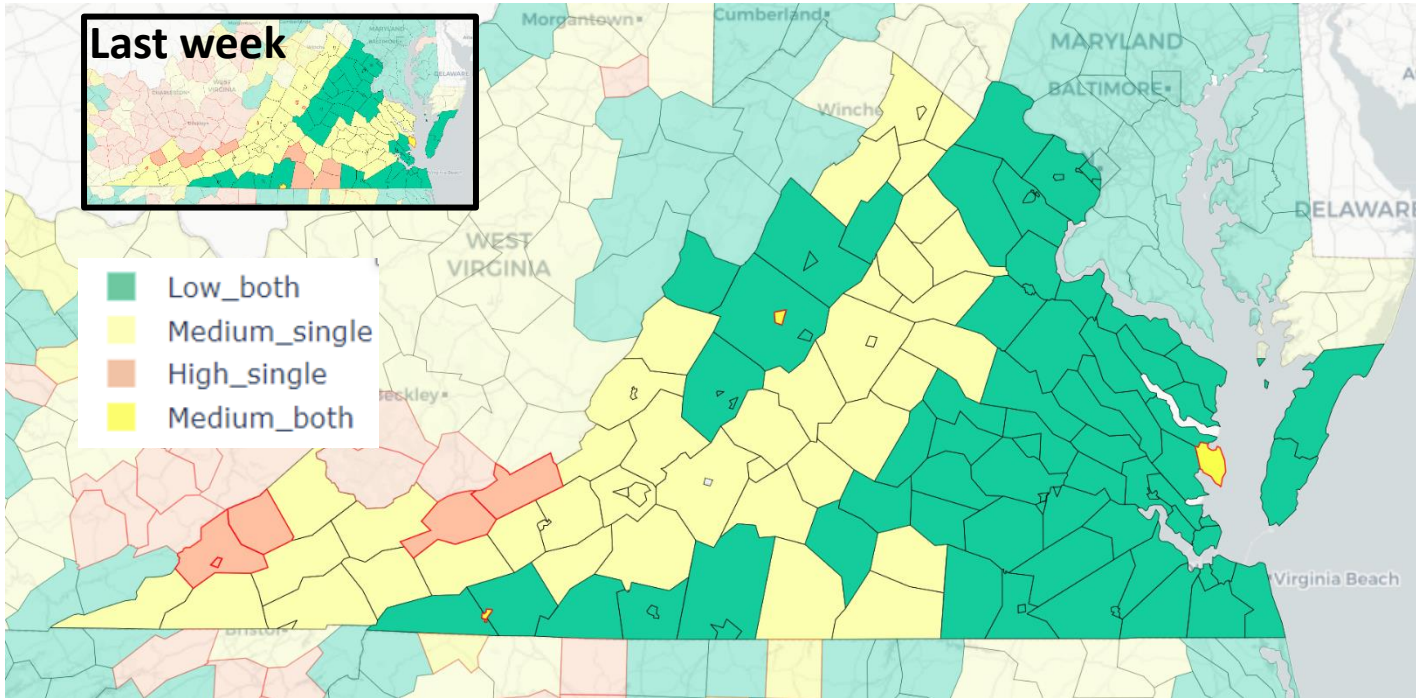
September 27th



Curve shows smoothed hospitalization rate (per 100K) by district
Hosp rate curve colored by Re number



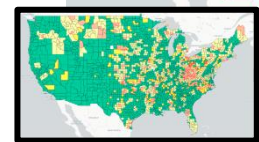
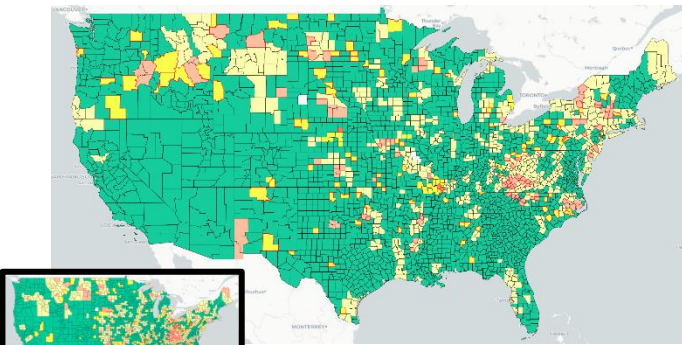
CDC's COVID-19 Community Levels



Red outline indicates county had 200 or more cases per 100k in last week

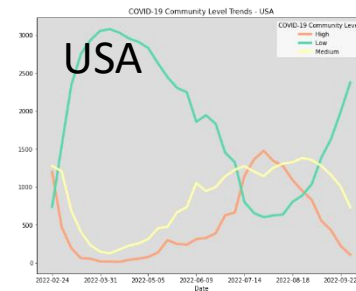
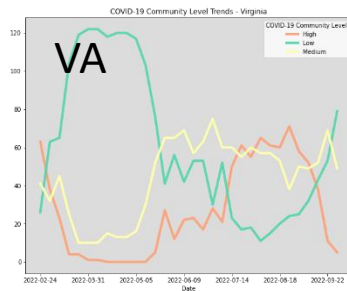
Pale color indicates either beds or occupancy set the level for this county

Dark color indicates both beds and occupancy set the level for this county



Last week

7-Oct-22

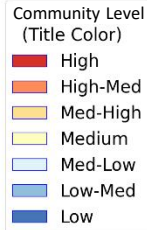


| COVID-19 Community Levels - Use the Highest Level that Applies to Your Community | | | | |
|--|---|--------|------------|--------|
| New COVID-19 Cases Per 100,000 people in the past 7 days | Indicators | Low | Medium | High |
| Fewer than 200 | New COVID-19 admissions per 100,000 population (7-day total) | <10.0 | 10.0-19.9 | ≥20.0 |
| | Percent of staffed inpatient beds occupied by COVID-19 patients (7-day average) | <10.0% | 10.0-14.9% | ≥15.0% |
| 200 or more | New COVID-19 admissions per 100,000 population (7-day total) | NA | <10.0 | ≥10.0 |
| | Percent of staffed inpatient beds occupied by COVID-19 patients (7-day average) | NA | <10.0% | ≥10.0% |

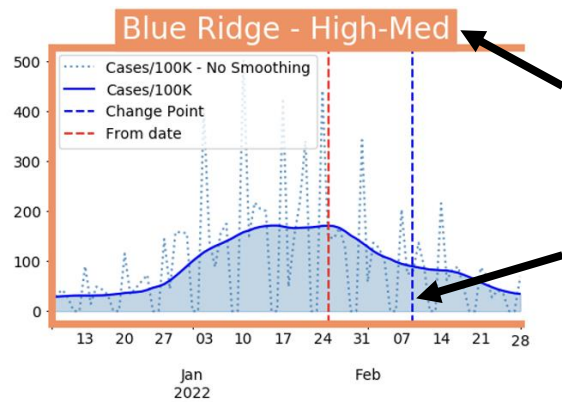
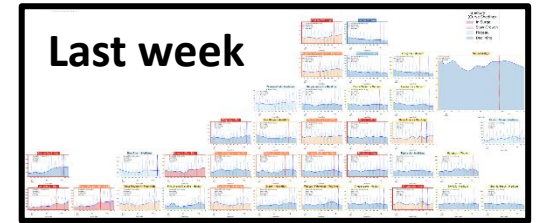
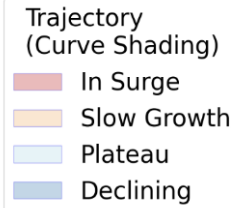
The COVID-19 community level is determined by the higher of the new admissions and inpatient beds metrics, based on the current level of new cases per 100,000 population in the past 7 days



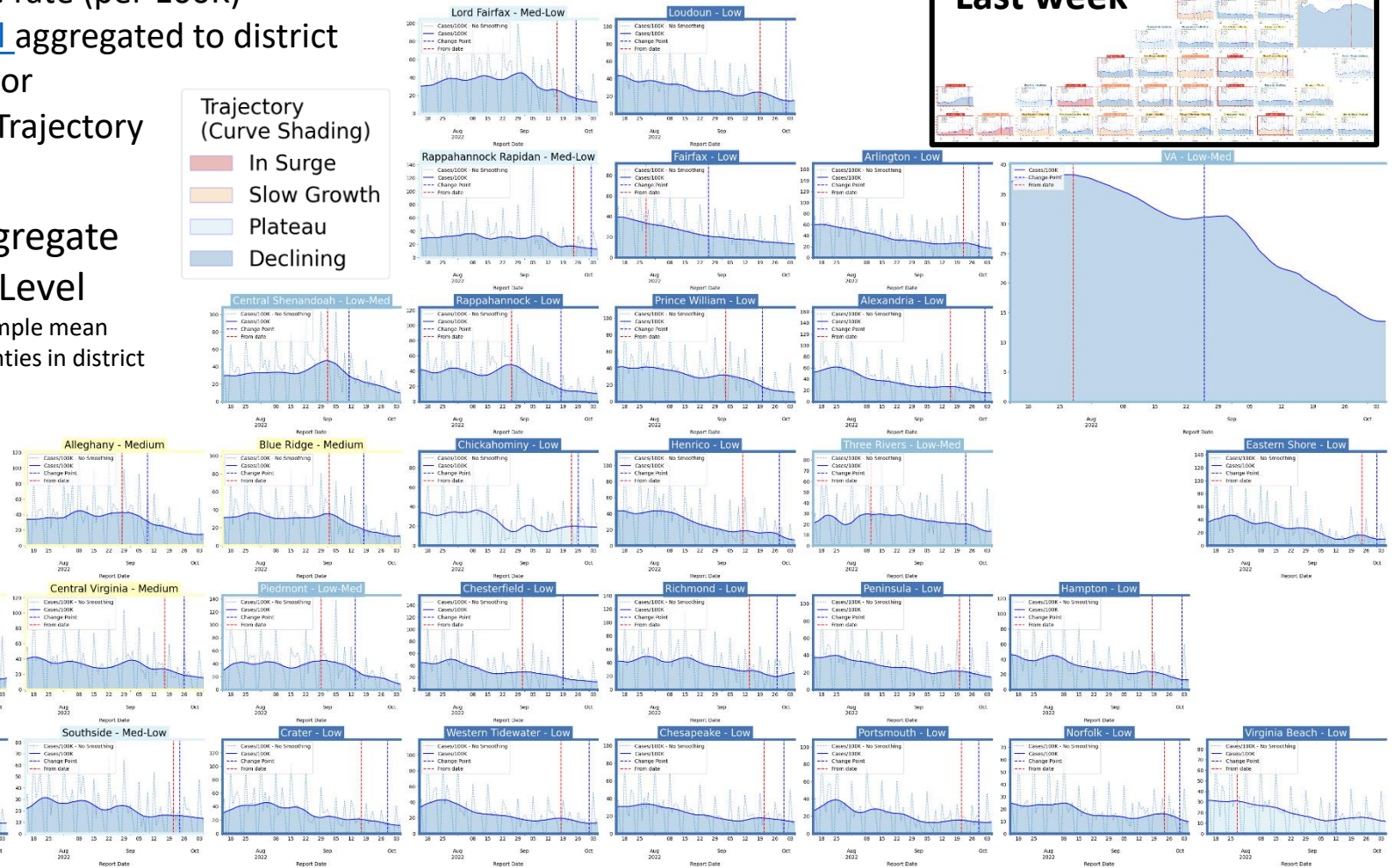
District Trajectories with Community Levels



Curve shows smoothed case rate (per 100K)
 CDC's new Community Level aggregated to district level in label & chart box color
 Case Rate curve colored by Trajectory



District's Aggregate Community Level
 Aggregate level a simple mean of all levels for counties in district
 Case rate Trajectory



Estimating Daily Reproductive Number – Redistributed gap

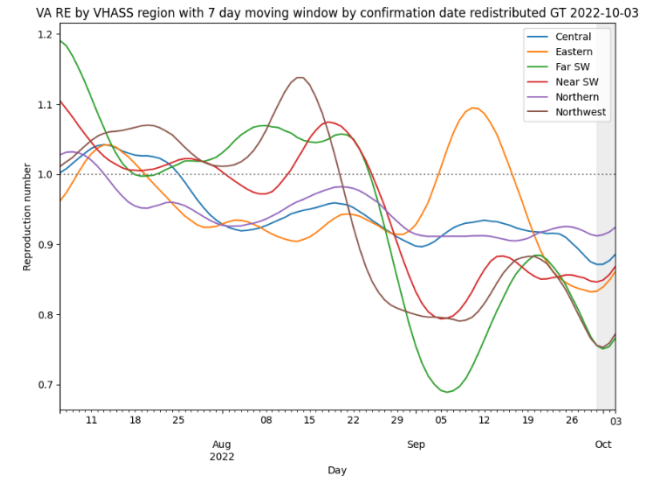
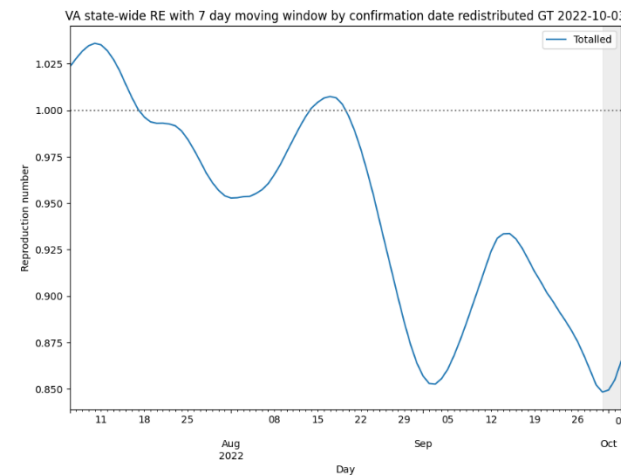
October 3rd Estimates

| Region | Date Confirmed R_e | Date Confirmed Diff Last Week |
|------------|-------------------------|----------------------------------|
| State-wide | 0.865 | 0.006 |
| Central | 0.886 | 0.014 |
| Eastern | 0.861 | -0.027 |
| Far SW | 0.766 | -0.121 |
| Near SW | 0.868 | 0.093 |
| Northern | 0.924 | 0.035 |
| Northwest | 0.772 | -0.025 |

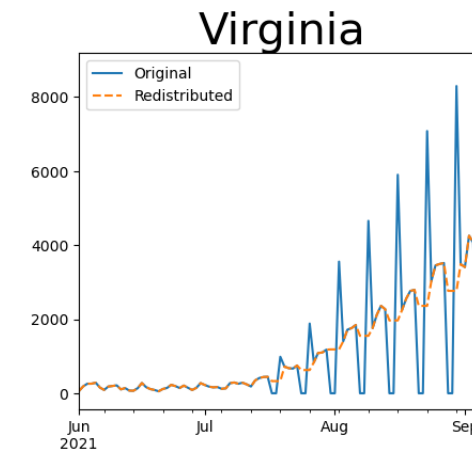
Methodology

- Wallinga-Teunis method (EpiEstim¹) for cases by confirmation date
- Serial interval: updated to discrete distribution from observations (mean=4.3, Flaxman et al, Nature 2020)
- Using Confirmation date since due to increasingly unstable estimates from onset date due to backfill

1. Anne Cori, Neil M. Ferguson, Christophe Fraser, Simon Cauchemez. A New Framework and Software to Estimate Time-Varying Reproduction Numbers During Epidemics. American Journal of Epidemiology, Volume 178, Issue 9, 1 November 2013, Pages 1505–1512, <https://doi.org/10.1093/aje/kwt133>



Skipping Weekend Reports & holidays biases estimates
Redistributed “big” report day to fill in gaps, and then estimate R from
“smoothed” time series

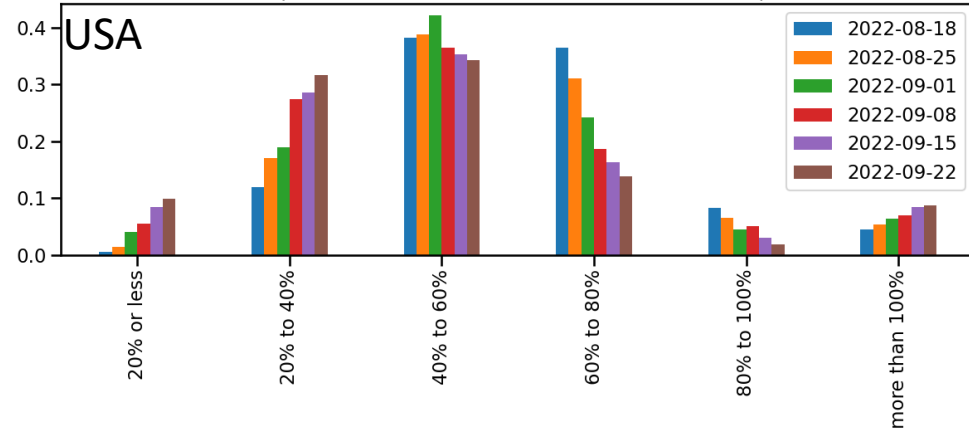


Wastewater Monitoring

Wastewater provides a coarse early warning of COVID-19 levels in communities

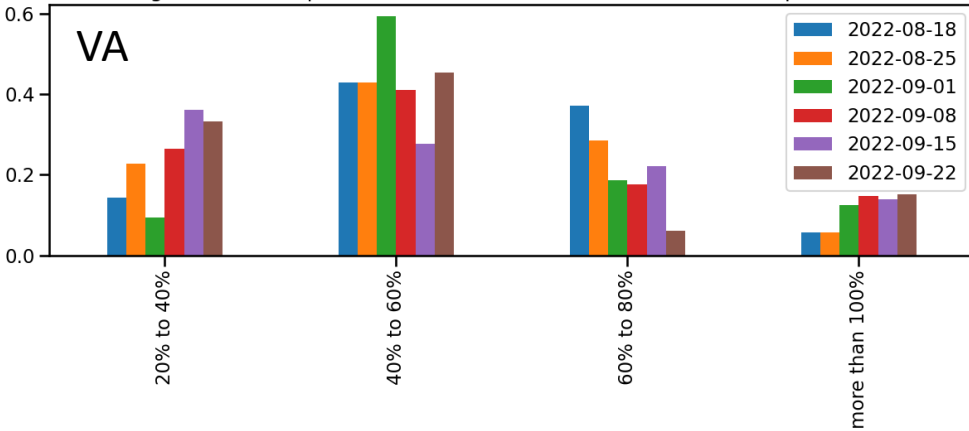
- Overall in the US, there is an increase in sites with increased levels of virus compared to 15 days ago
- Current virus levels are at or exceeding max of previous historical levels, has slowed, though more sites are entering upper quintiles

US Historic percentile of current detected levels over the past weeks



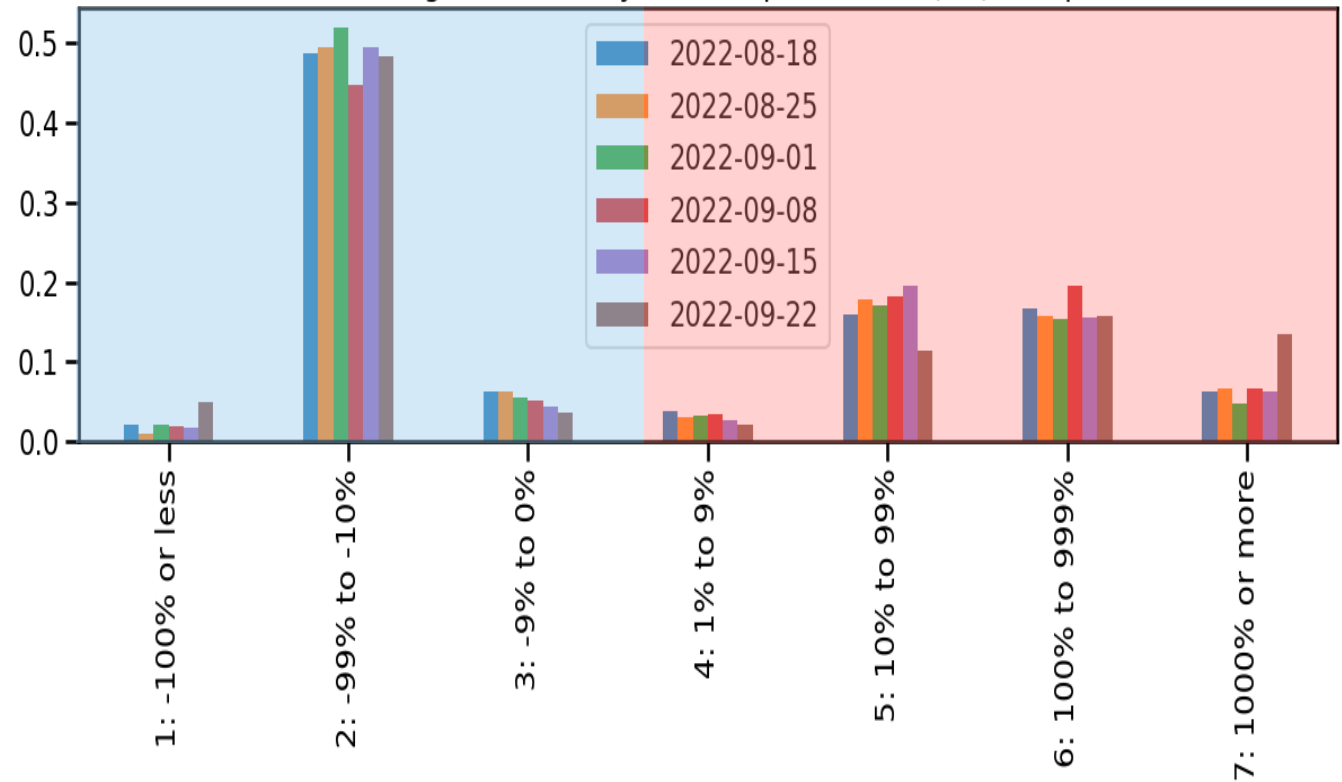
category

Virginia - Historic percentile of current detected levels over the past weeks



category

Percent Change over 15 days for the past weeks (US) - Proportions



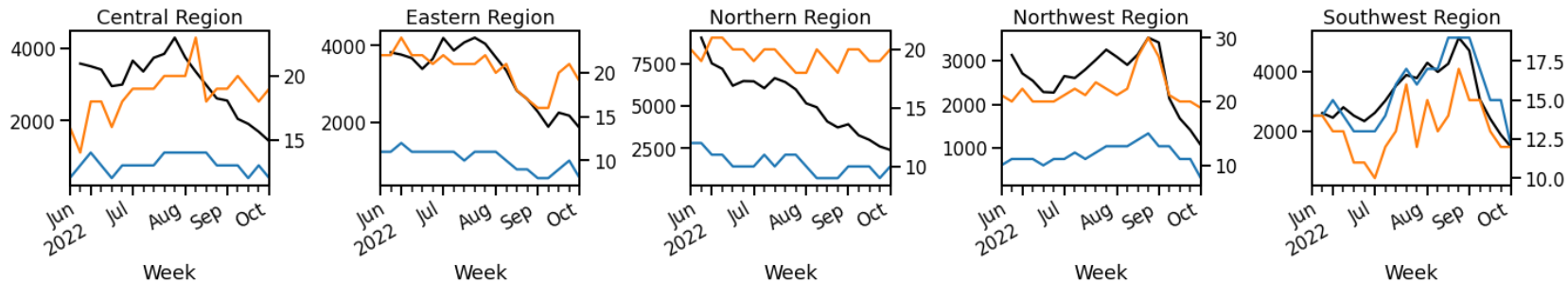
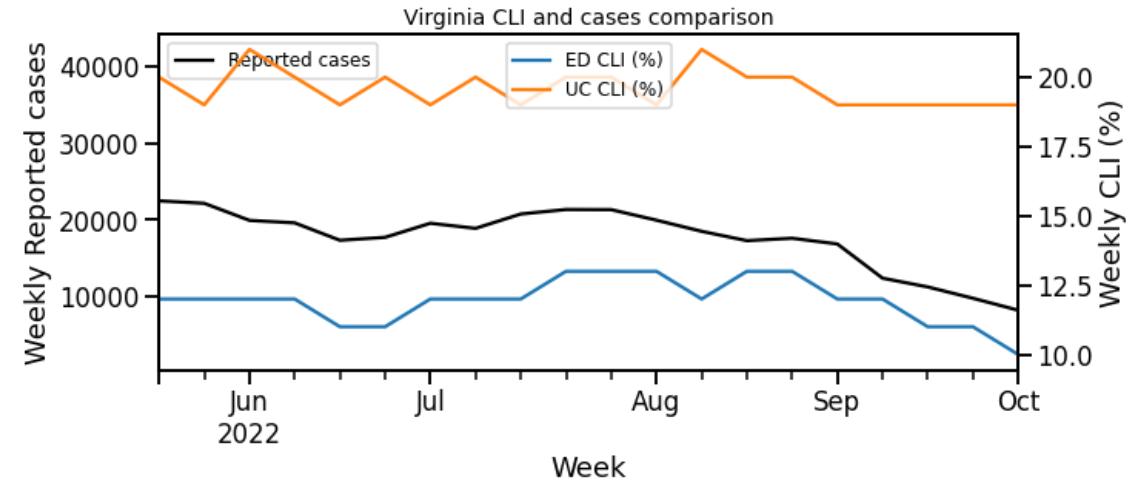
category

Data Source: [CDC Data Tracker](https://data.cdc.gov/)

COVID-like Illness Activity

COVID-like Illness (CLI) gives a measure of COVID transmission in the community

- Emergency Dept (ED) based CLI is more correlated with case reporting
- Urgent Care (UC) is a leading indicator but prone to some false positives
- **Current trends in UC CLI have been in a plateau since mid-May state-wide, with some regional fluctuations**

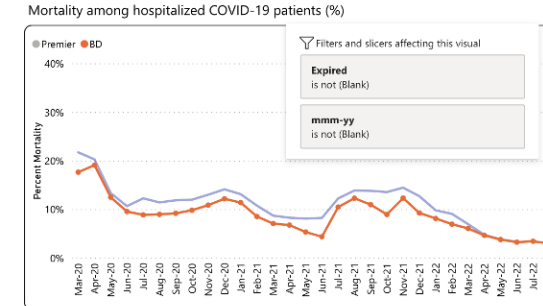
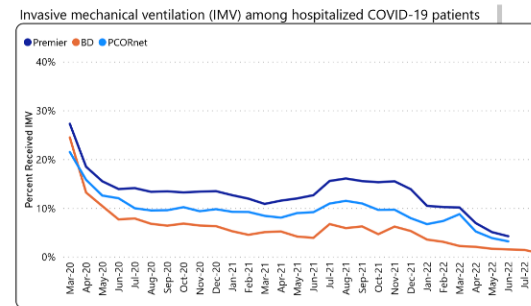
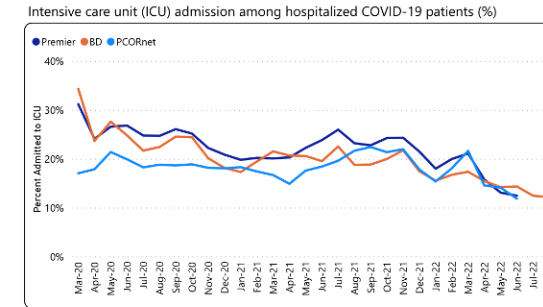
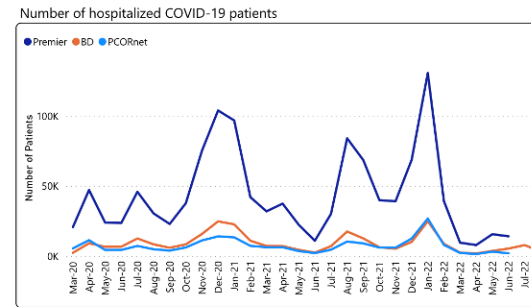


Hospitalizations and Severe Outcomes

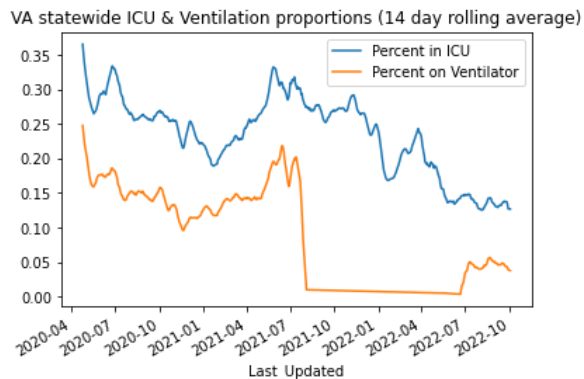
Data Source: [CDC Data Tracker](#)

Proportion of most severe outcomes decreasing among those who are hospitalized

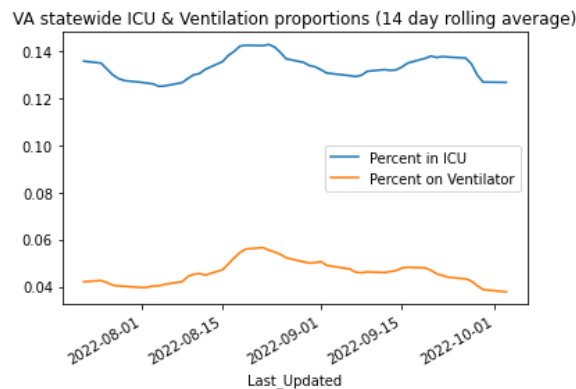
- ICU has declined from ~20% of hospitalized to nearly 10% since initial Omicron wave
- Also seen across all age-groups
- Similar levels of decline seen in VA
- Regionally more variation



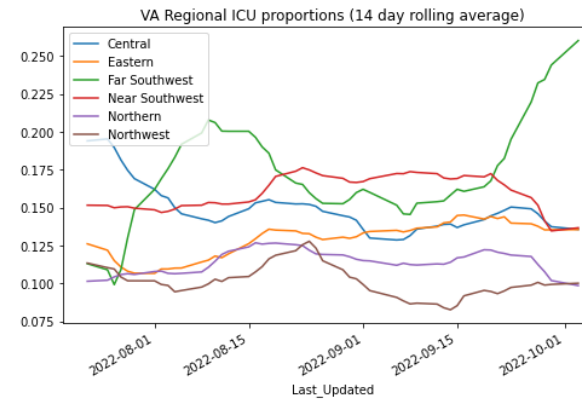
Virginia wide – full pandemic



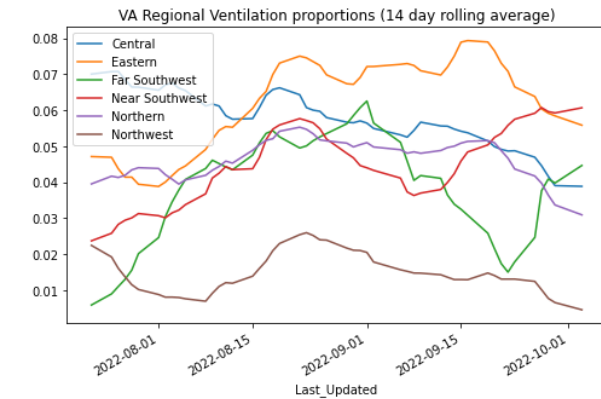
Virginia wide – recent



Virginia Regional ICU percent



Virginia Regional Ventilation %



SARS-CoV2 Variants of Concern

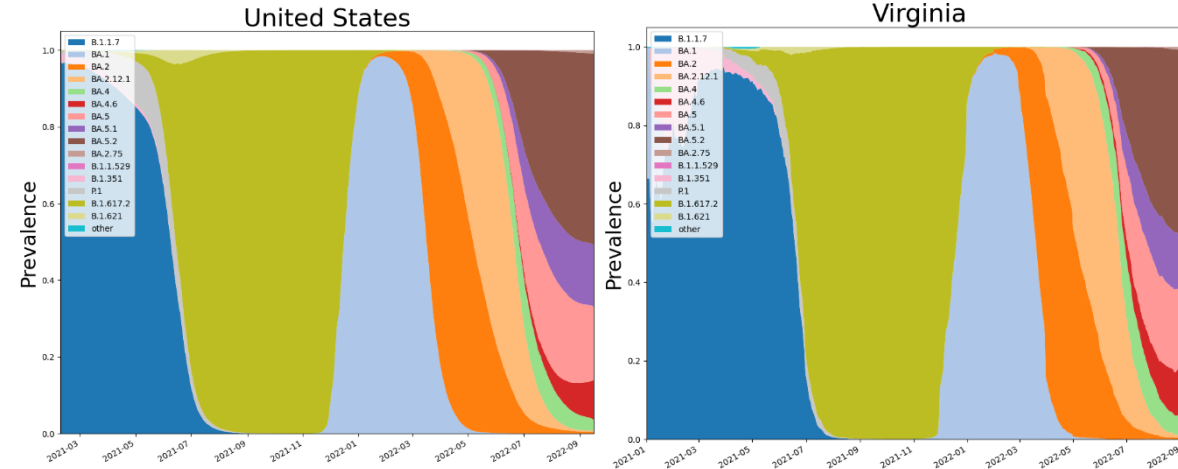


Emerging new variants will alter the future trajectories of pandemic and have implications for future control

- Emerging variants can: Increase transmissibility, increase severity (more hospitalizations and/or deaths), and limit immunity provided by prior infection and vaccinations

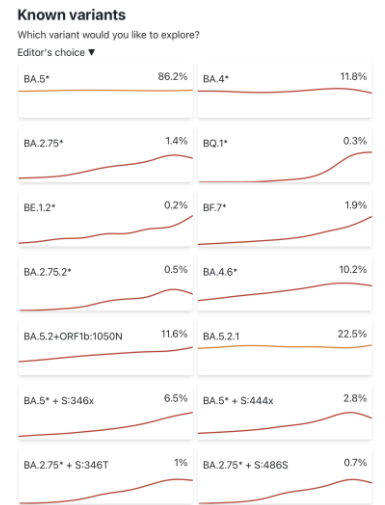
Omicron Updates

- BA.4.6 remains slow growing to about 16% from 15% last week
- BA.5 continues to slowly yield share to other variants, now down to 78% from 80% last week
- BF.7 showing strong growth up to 4.4 from 1.2% last week
- BQ.1.1 recently seeing growth in England and other countries that mimics past variants of concern that have gone on to dominate
- BA.2.72.2 also shows signs as being a potential candidate as a future variant of concern, has recently [been shown to have significant immune escape](#)



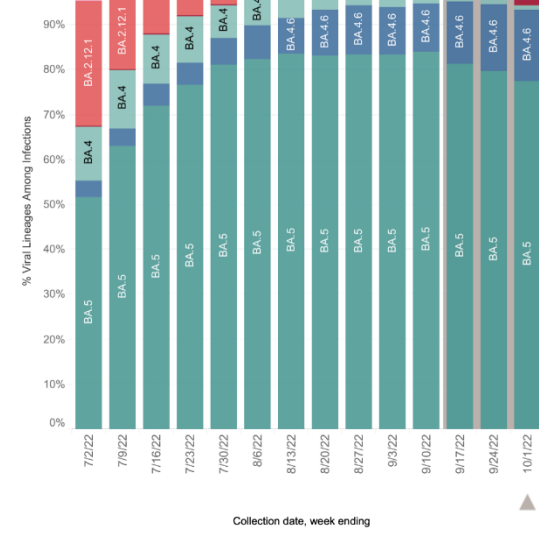
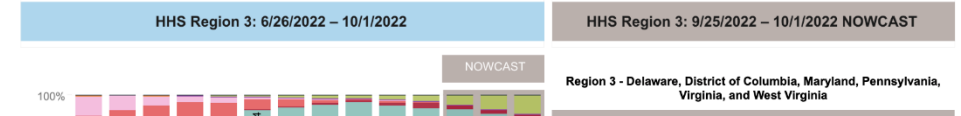
covSPECTRUM

Enabled by data from GISAID



COV-spectrum

“Editor’s choice”
Variants to watch



Region 3 - Delaware, District of Columbia, Maryland, Pennsylvania, Virginia, and West Virginia

| WHO label | Lineage # | US Class | %Total | 95%PI |
|-----------|-----------|----------|--------|------------|
| Omicron | BA.5 | VOC | 77.5% | 74.8-79.9% |
| | BA.4.6 | VOC | 15.8% | 13.4-18.6% |
| | BF.7 | VOC | 4.4% | 2.7-6.9% |
| | BA.2.75 | VOC | 1.3% | 0.8-2.1% |
| | BA.4 | VOC | 1.0% | 0.9-1.2% |
| | BA.2.12.1 | VOC | 0.0% | 0.0-0.0% |
| | BA.2 | VOC | 0.0% | 0.0-0.0% |
| | B.1.1.529 | VOC | 0.0% | 0.0-0.0% |
| | BA.1.1 | VOC | 0.0% | 0.0-0.0% |
| Delta | B.1.617.2 | VBM | 0.0% | 0.0-0.0% |
| Other | Other* | | 0.0% | 0.0-0.0% |

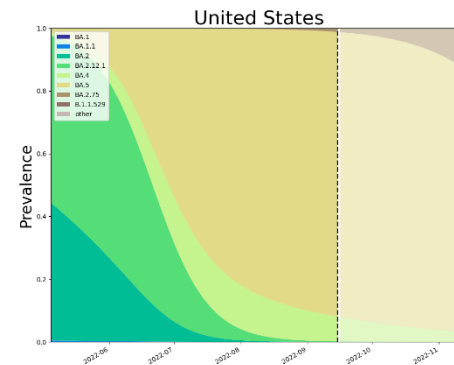
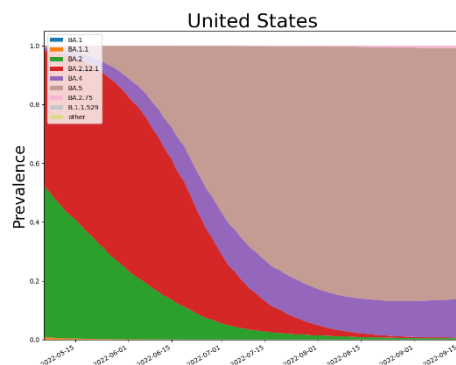
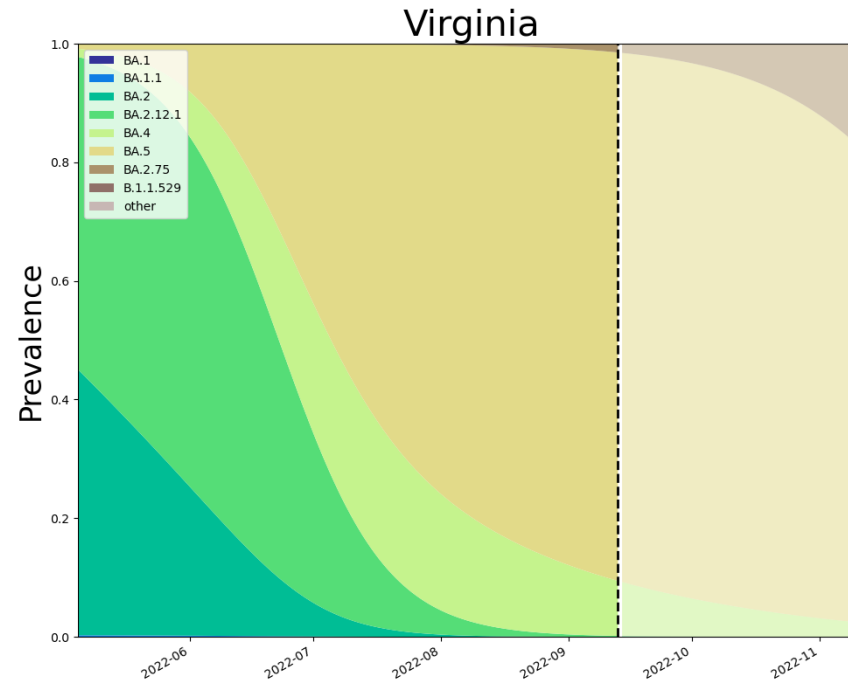
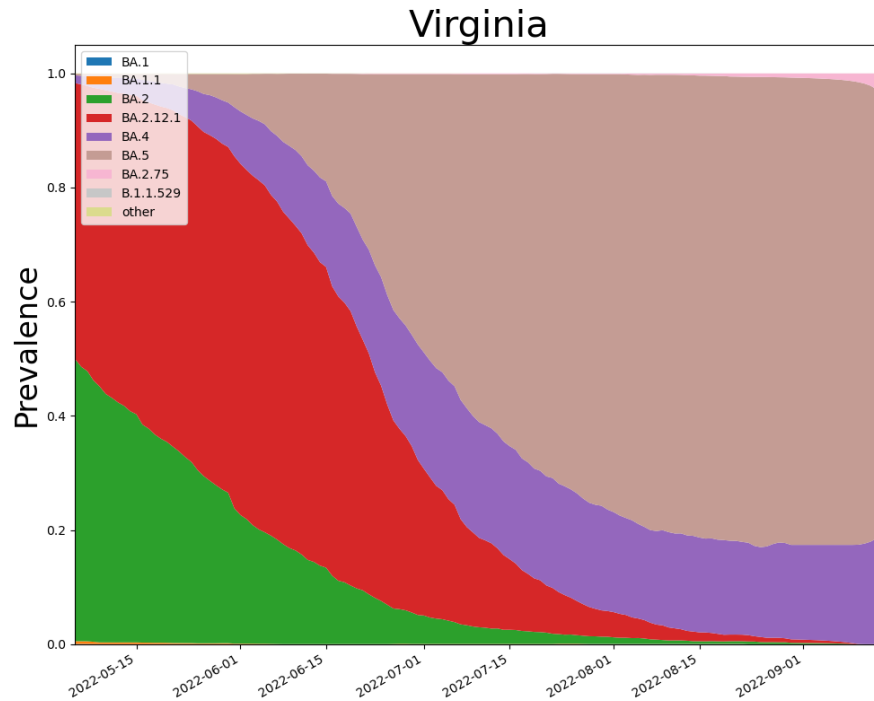
* Enumerated lineages are US VOC and lineages circulating above 1% nationally in at least one week period. "Other" represents the aggregation of lineages which are circulating <1% nationally during all weeks displayed.
 ** These data include Nowcast estimates, which are modeled projections that may differ from weighted estimates generated at later dates
 # AY.1-AY.133 and their sublineages are aggregated with B.1.617.2. BA.1, BA.3 and their sublineages (except BA.1.1 and its sublineages) are aggregated with B.1.1.529. Except BA.2.12.1, BA.2.75 and their sublineages, BA.2 sublineages are aggregated with BA.2. Except BA.4.6, sublineages of BA.4 are aggregated to BA.4. Except BF.7, sublineages of BA.5 are aggregated to BA.5. Sublineages of BA.1.1 and BA.2.75 are aggregated to the parental BA.1.1 and BA.2.75 respectively. Previously, BA.2.75 was aggregated with BA.2, and BF.7 was aggregated with BA.5. Lineages BA.4.6, BF.7, and many BA.2.75 contain the spike substitution R346T.



SARS-CoV2 Omicron and Sub-Variants

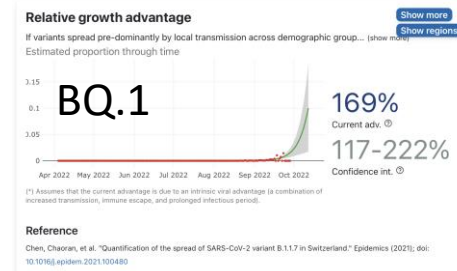
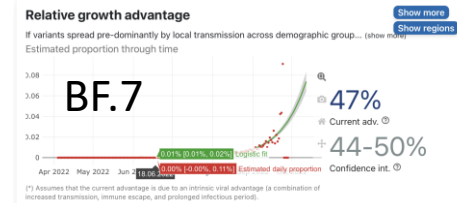
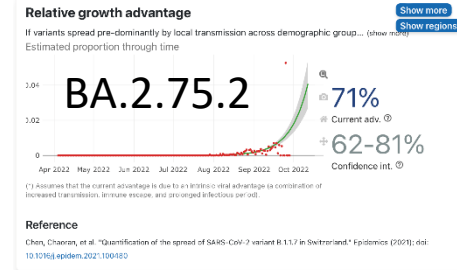
As detected in whole Genomes in public repositories

VoC Polynomial Fit Projections



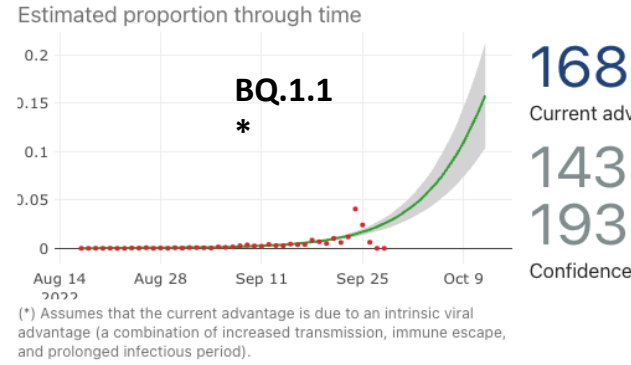
Note: Data lags force projections to start in past. Everything from dotted line forward is a projection.

Relative Growth Advantages

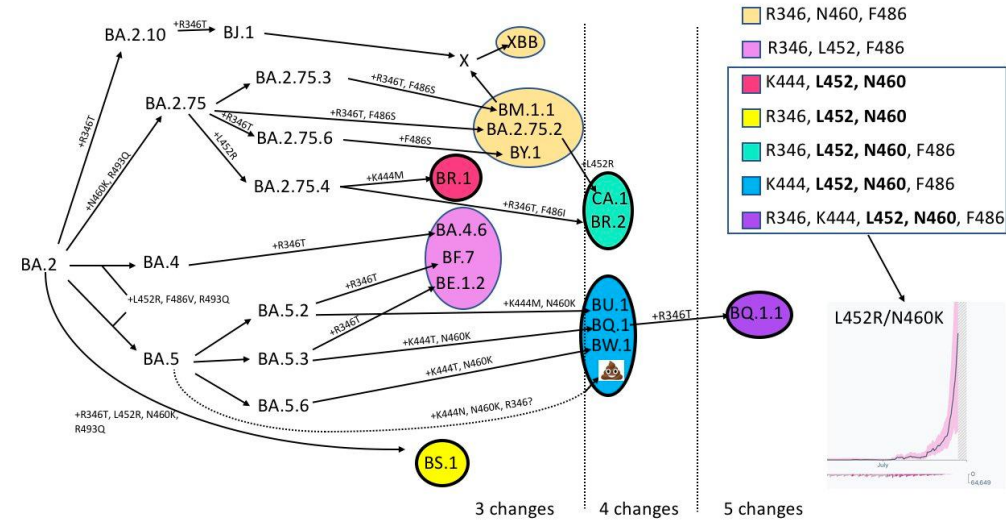


Pandemic Pubs

1. BQ.1.1, a descendant of BA.5, shows strong growth advantage in recent sequence surveillance.
 Current Bivalent booster doses target BA.5 making it a well-timed and a worthwhile intervention.

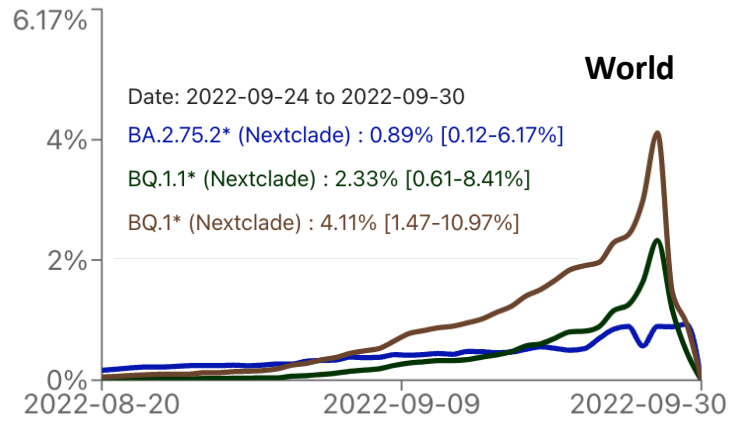
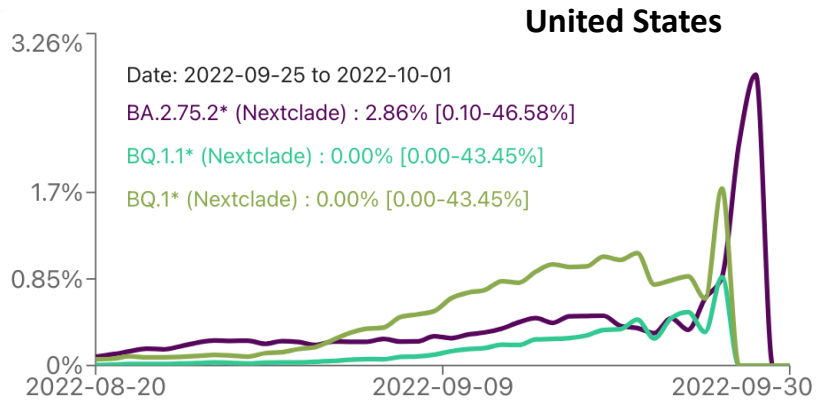


Omicron lineages with changes at 346, 444, 452, 460, and 486



https://twitter.com/dfocosi/status/1574319177602670592?s=12&t=myre59DuQ1H31TI_-yd7DQ

World



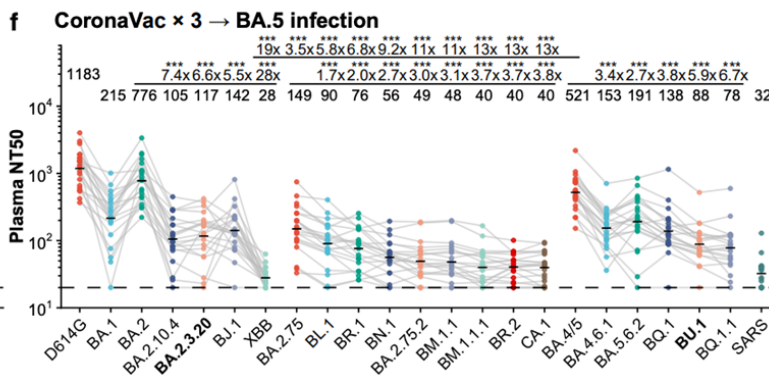
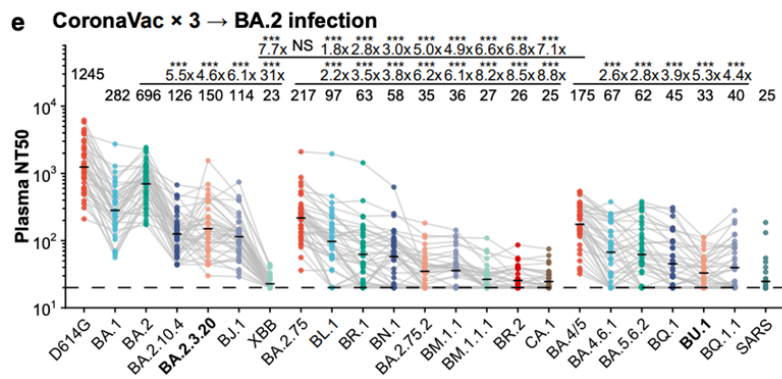
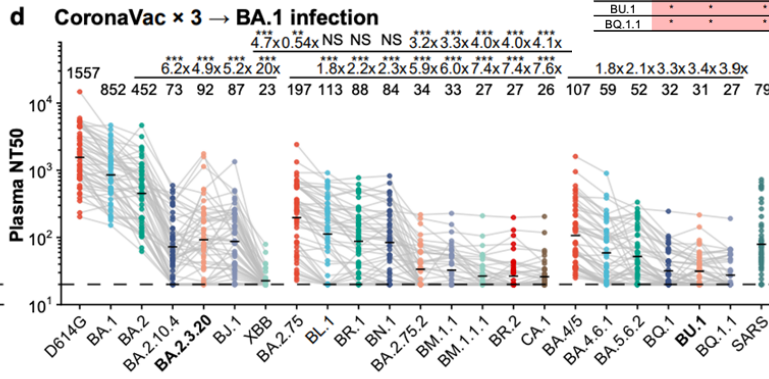
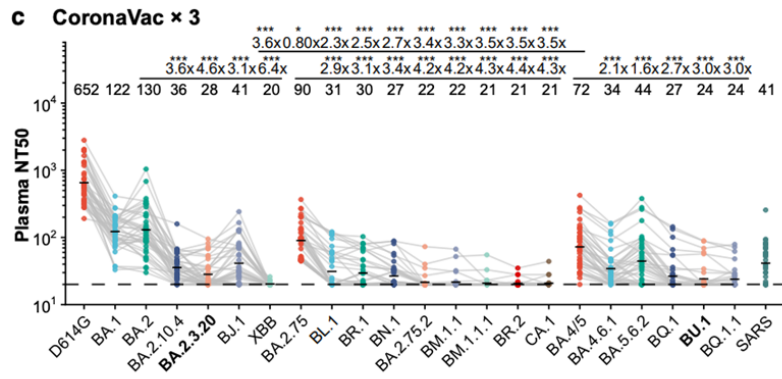
| description | number _sequen ces | submitted _past_10_ days | relative _growth _advant age | relative_growt h_advantage_low | relative_growt h_advantage _high |
|---|--------------------|--------------------------|------------------------------|--------------------------------|----------------------------------|
| BA.4.6, BF.7, any BA.4/5+S:346T/S:44 etc. | 67699 | 11538 | 31.64% | 31.39% | 31.89% |
| BA.2.75.2, BQ.1, BM.1.1, BY.1, BA.2.3.20, ... | 3853 | 1465 | 73.37% | 70.96% | 75.78% |
| BQ.1.1, BN.1, BM.1.1.1, CA.1, ... | 714 | 345 | 101.44% | 93.21% | 109.68% |
| XBB | 66 | 40 | 157.47% | 111.26% | 203.67% |

Estimates from surveillance using mutational profiles approximating BQ.1, BQ.1.1, and BA.2.75.2
https://cov-spectrum.org/explore/World/AllSamples/from=2022-08-17&to=2022-10-04/variants?nextcladePangoLineage=BA.2.75.2*&nextcladePangoLineage1=bq.1.1*&nextcladePangoLineage2=bq.1&analysisMode=CompareEquals&

<https://cov-spectrum.org/collections/54>

Pandemic Pubs

2. Both BQ.1.1 and XBB show substantial immune escape and significant negation of many pharmaceutical interventions. XBB is significantly more immune evasive than BA.2.75.2 and BQ.1.1 against plasma from all breakthrough infections, comparable to or even exceeding the level of escape displayed by SARS-CoV-1 against SARS-CoV-2 convalescent plasma.



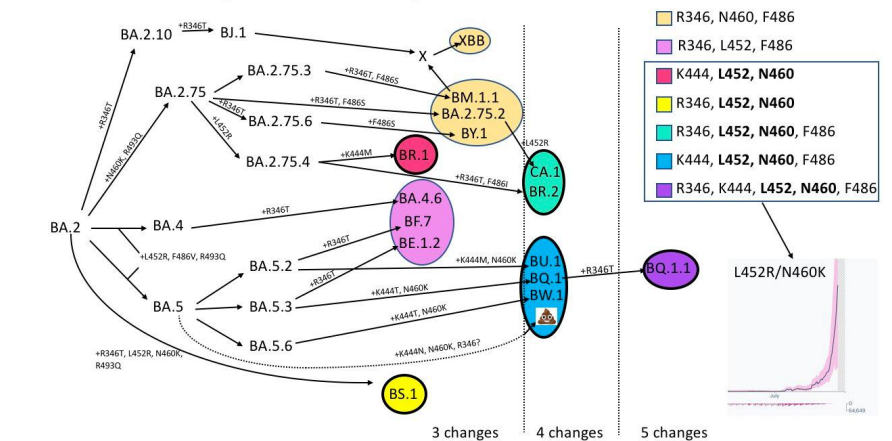
a

| Pango lineages | REGN 10933 | REGN 10987 | REGN10933 +10987 | COV2-2196 | COV2-2130 | COV2-2196+2130 | BR1-196 | BR1-198 | BR1-196+198 | S309 | DXP-604 | LY-CoV 1404 | SA58 | SA55 | SA55+SA58 | Additional RBD mutations | |
|----------------|------------|------------|------------------|-----------|-----------|----------------|---------|---------|-------------|------|---------|-------------|------|------|-----------|--------------------------|---|
| BA.2 | * | 590 | 821 | 4312 | 2130 | 6.3 | 8.2 | 8530 | 8990 | 8610 | 852 | 219 | 0.9 | 5.1 | 7.2 | 7.8 | |
| BA.2.3.20 | 121 | * | 199 | 15 | * | 26 | 14 | * | 24 | * | 897 | 181 | 9.7 | 20 | 4.6 | 7.8 | K444R+N450D+L452M+N460K+R493Q |
| BA.2.10.4 | * | * | * | * | 289 | 501 | 2109 | 7990 | 3984 | 706 | 6348 | 1.3 | 4.3 | 4.9 | 5.0 | | K444R+N450D+L452M+N460K+R493Q+S494P |
| BJ.1 | * | * | * | 3076 | * | 5985 | 7609 | * | * | 709 | 166 | * | 8163 | 3.7 | 8.6 | | D399H+R346T+L388I+V445P+G446S+N460K+F486V |
| XBB | * | * | * | * | * | * | * | * | * | 963 | * | * | 8805 | 5.3 | 9.8 | | D399H+R346T+L388I+V445P+G446S+N460K+F486S+F490S+R493Q |
| BA.2.75 | 278 | * | 410 | 119 | 352 | 121 | 1730 | 6622 | 3861 | 672 | 5920 | 2.2 | 246 | 4.3 | 9.6 | | |
| BL.1 | 260 | * | 511 | 93 | * | 174 | 1251 | * | 3075 | 508 | 7193 | 2.8 | 7975 | 6.3 | 10 | | R346T |
| BR.1 | 319 | * | 679 | 117 | * | 170 | 1992 | * | 3160 | 564 | 6689 | * | 1616 | 5.9 | 9.7 | | L452R+K444M |
| BN.1 | 390 | * | 701 | 59 | 303 | 109 | 4101 | * | 8444 | 6979 | 8901 | 1.7 | 4960 | 5.7 | 9.4 | | K356T+F490S |
| BN.1 | 344 | * | 599 | 70 | * | 166 | 3683 | * | 7791 | * | 6012 | 3.3 | 8295 | 4.9 | 9.0 | | R346T+K356T+F490S |
| BA.2.75.2 | * | * | * | * | * | * | * | * | * | 852 | * | 3.0 | 8922 | 5.9 | 9.7 | | R346T+F486S |
| BM.1.1 | * | * | * | * | * | * | * | * | * | 879 | * | 2.3 | 8823 | 5.2 | 8.9 | | R346T+F486S |
| BM.1.1.1 | * | * | * | * | * | * | * | * | * | 956 | * | 1.9 | 8082 | 4.8 | 10.5 | | R346T+F486S+F490S |
| BR.2 | * | * | * | * | * | * | * | * | * | 921 | * | 2.6 | 7263 | 4.7 | 10.5 | | R346T+L452R+F486I |
| CA.1 | * | * | * | * | * | * | * | * | * | 897 | * | 3.2 | 6927 | 6.0 | 11.5 | | R346T+L452R+F486S |
| BA.4/5 | 520 | 709 | * | 23 | 40 | 7124 | * | * | 1055 | 6264 | 0.8 | 3.9 | 5.0 | 4.5 | | | |
| BA.4.6.1 | 2338 | 5402 | * | * | * | 4763 | * | 7809 | 4456 | 4634 | 1.2 | 50 | 4.8 | 9.9 | | R346T | |
| BA.5.6.2 | * | * | * | * | * | 4636 | * | 7883 | 1408 | 5892 | 1662 | 58 | 5.1 | 8.9 | | K444T | |
| BQ.1 | * | * | * | * | * | * | * | * | 1709 | * | 1905 | 44 | 6.6 | 9.2 | | K444T+N460K | |
| BU.1 | * | * | * | * | * | * | * | * | 1082 | * | 26 | 5.6 | 5.3 | 10.3 | | K444M+N460K | |
| BQ.1.1 | * | * | * | * | * | * | * | * | 5581 | * | * | 900 | 5.9 | 10.5 | | R346T+K444T+N460K | |

b

IC50 of hACE2 (μg/mL)

Omicron lineages with changes at 346, 444, 452, 460, and 486



Researchers in Beijing characterize multiple concerning immune escape variants using plasma from breakthrough infections in those vaccinated with three doses of CoronaVac. Similar to BQ.1.1, XBB also escapes Evusheld and Bebtelovimab. BU.1, BR.2, BM.1.1.1, CA.1, and XBB all displayed moderate hACE2 binding capability relative to previously widely circulating variants of BA.2 and BA.5.

<https://www.biorxiv.org/content/10.1101/2022.09.15.507787v3>

https://twitter.com/yunlong_cao/status/1577343549120872448

<https://twitter.com/dfocosi/status/1574319177602670592?s=12&t=myre59DuQ1H31TI-yd7DQ>

Pandemic Pubs (last week)

1. Representative survey estimates 7.3% of the US adult population (~18.5 million) reported having long COVID by July
2. A propensity matched cohort from the UK found that 2+ doses of vaccine decreased adjusted long COVID risk by 41% (95% CI 31%-50%)
3. BA.2.75.2 neutralised, on average, at titers ~6.5-times lower than BA.5, making BA.2.75.2 the most neutralisation-resistant variant evaluated to date
4. Retrospective cohort study of 6,245,282 older adults (age ≥65 years) who had medical encounters between 2/2020–5/2021, indicates that people with COVID-19 were at significantly increased risk for a new diagnosis of Alzheimer's disease within 360 days after the initial COVID-19 diagnosis

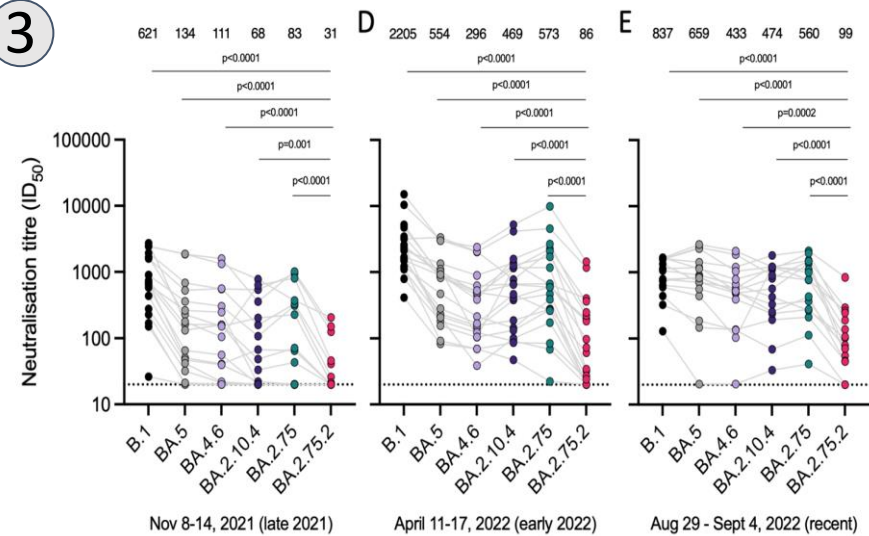
1

3

4

Prevalence of Long COVID Among All Respondent (with or without COVID)

| | Total | Long COVID | Crude Prevalence of Long COVID % (95% CI) | Age and sex direct-standardized prevalence of long COVID* % (95% CI) | Crude prevalence ratio (PR) PR (95% CI) | Adjusted prevalence ratio (aPR)** aPR (95% CI) | Estimated Number with Long COVID |
|---------------|----------------|----------------|---|--|---|--|----------------------------------|
| | Weighted N (%) | Weighted N (%) | | | | | |
| Total | 3,042 (100.0) | 222 (100.0) | 7.3 (6.1, 8.5) | | | | 18,533,864 |
| Age | | | | | | | |
| 18-24 | 365 (12.0) | 17 (7.8) | 4.8 (0.9, 8.6) | 4.4 (1.8 - 10.0) | 0.45 (0.27, 0.76) | 0.50 (0.30, 0.84) | 1,445,641 |
| 25-34 | 547 (18.0) | 58 (26.1) | 10.6 (6.6, 14.6) | 10.0 (6.8 - 14.6) | Ref | Ref | 4,837,339 |
| 35-44 | 495 (16.3) | 43 (19.1) | 8.6 (5.3, 11.9) | 9.0 (6.1 - 13.1) | 0.81 (0.56, 1.18) | 0.87 (0.60, 1.26) | 3,539,968 |
| 45-54 | 498 (16.4) | 37 (16.5) | 7.4 (4.6, 10.2) | 7.4 (5.1 - 10.7) | 0.70 (0.47, 1.03) | 0.72 (0.49, 1.07) | 3,058,088 |
| 55-64 | 508 (16.7) | 41 (18.2) | 8.0 (5.5, 10.5) | 8.3 (6.0 - 11.2) | 0.75 (0.51, 1.10) | 0.80 (0.55, 1.18) | 3,373,163 |
| 65+ | 629 (20.7) | 27 (12.2) | 4.3 (3.0, 5.6) | 4.2 (3.0 - 5.8) | 0.41 (0.26, 0.63) | 0.43 (0.27, 0.66) | 2,261,131 |
| Gender | | | | | | | |
| Male | 1,443 (47.4) | 72 (32.4) | 5.0 (3.6, 6.3) | 5.0 (3.8 - 6.5) | Ref | Ref | 6,004,972 |
| Female | 1,516 (49.8) | 144 (64.8) | 9.5 (7.5, 11.5) | 9.4 (7.7 - 11.6) | 1.90 (1.45, 2.50) | 1.84 (1.40, 2.42) | 12,009,944 |



In 18 random blood donor samples in Stockholm, sampled recently BA.2.75.2 was neutralised, on average, five-fold less potently than BA.5. These are recent samples in a city that has good vaccine coverage and likely relatively high prior infection rates. Researchers also report the sensitivity of emerging omicron sublineages BA.2.75.2, BA.4.6, and BA.2.10.4 to neutralisation by a panel of clinically relevant and pre-clinical monoclonal antibodies, as well as by serum from blood donated in Stockholm, Sweden.

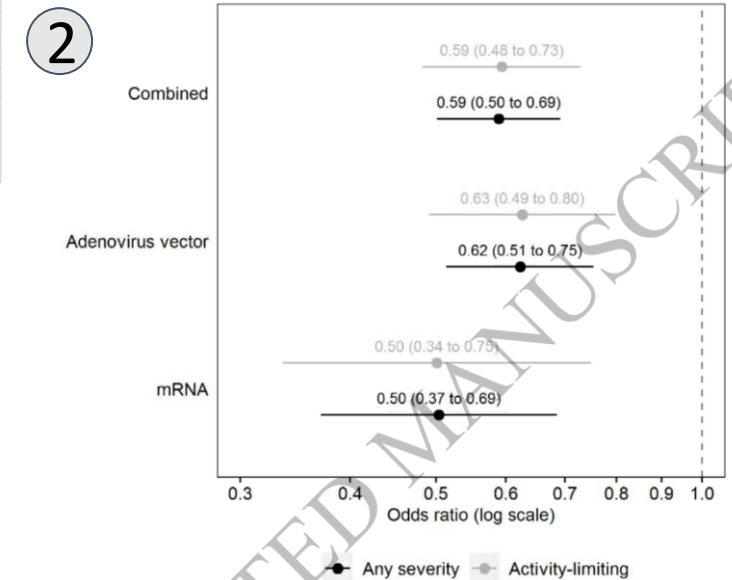
<https://www.biorxiv.org/content/10.1101/2022.09.16.508299v2>

<https://twitter.com/benjmurrell/status/1570862185819303937?s=12&t=sl45IDMQ7GGmC2KakXPNz>

NY researchers conducted a population-representative survey, June 30-July 2, 2022, of a random sample of 3,042 United States adults. Using questions developed by the United Kingdom's Office of National Statistics, we estimated the prevalence by sociodemographics, adjusting for gender and age. An estimated 7.3% (95% CI: 6.1-8.5%) of all respondents reported long COVID, approximately 18,533,864 adults. One-quarter (25.3% [18.2-32.4%]) of respondents with long COVID reported their day-to-day activities were impacted 'a lot' and 28.9% had SARS-CoV-2 infection >12 months ago.

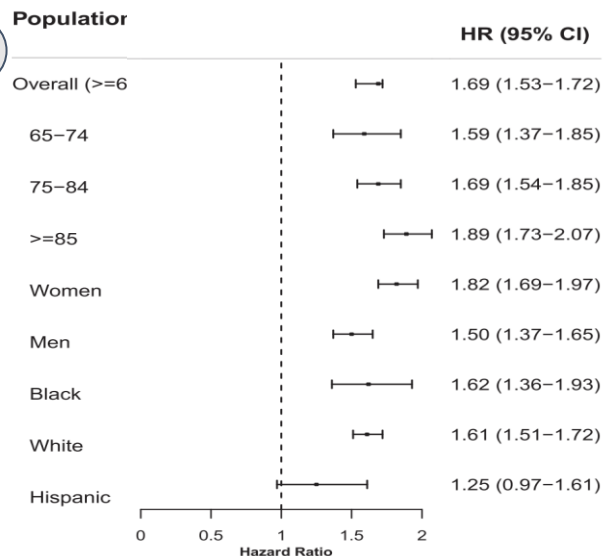
<https://www.medrxiv.org/content/10.1101/2022.09.12.22279862v1>

2



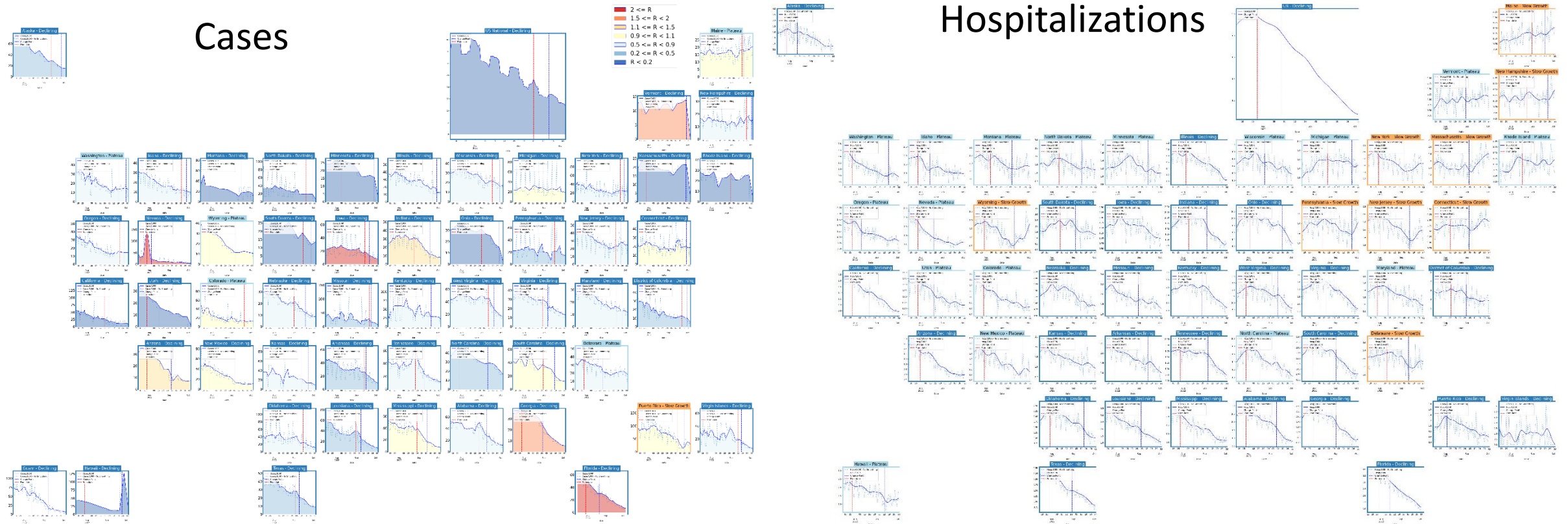
Oxford researchers investigated Long Covid incidence by vaccination status in a random sample of UK adults from April 2020 to November 2021. Persistent symptoms were reported by 9.5% of 3,090 breakthrough SARS-CoV-2 infections and 14.6% of unvaccinated controls (adjusted odds ratio 0.59, 95% CI: 0.50-0.69), emphasising the need for public health initiatives to increase population-level vaccine uptake. Matched study participants who were double-vaccinated at time of infection to control participants who were unvaccinated when infected and remained so at their first follow-up visit ≥12 weeks later. Most double-vaccinated participants (3,057, 98.9%) were infected after 17 May 2021, when the Delta variant dominated in the UK, while nearly all unvaccinated participants (3,082, 14 99.7%) were infected before this date.

<https://academic.oup.com/ofid/advance-article/doi/10.1093/ofid/ofac464/6696170?login=false>
<https://twitter.com/DFisman/status/1570901408211402752>

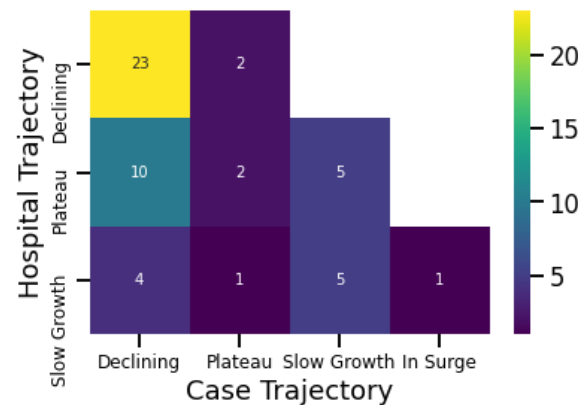


<https://content.iospress.com/articles/journal-of-alzheimers-disease/jad220717>

United States Case & Hospitalizations



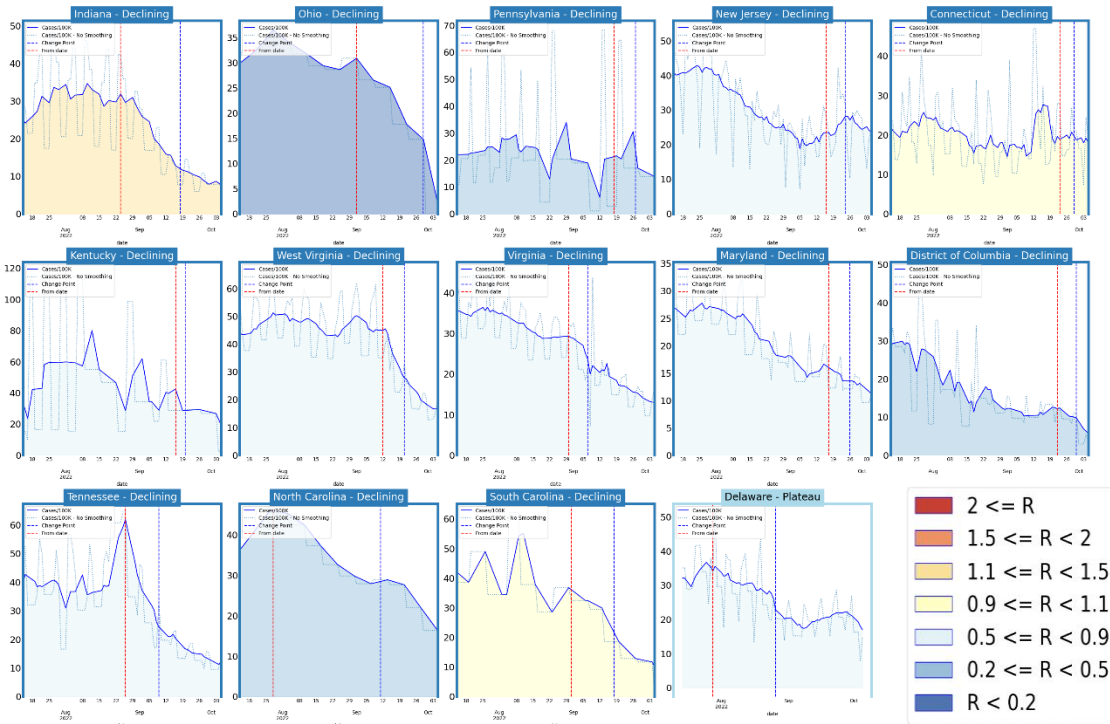
| Status | Number of States | |
|--------------------|------------------|-----------|
| | Current Week | Last Week |
| Declining | 48 | (37) |
| Plateau | 5 | (5) |
| Slow Growth | 1 | (11) |
| In Surge | 0 | (1) |



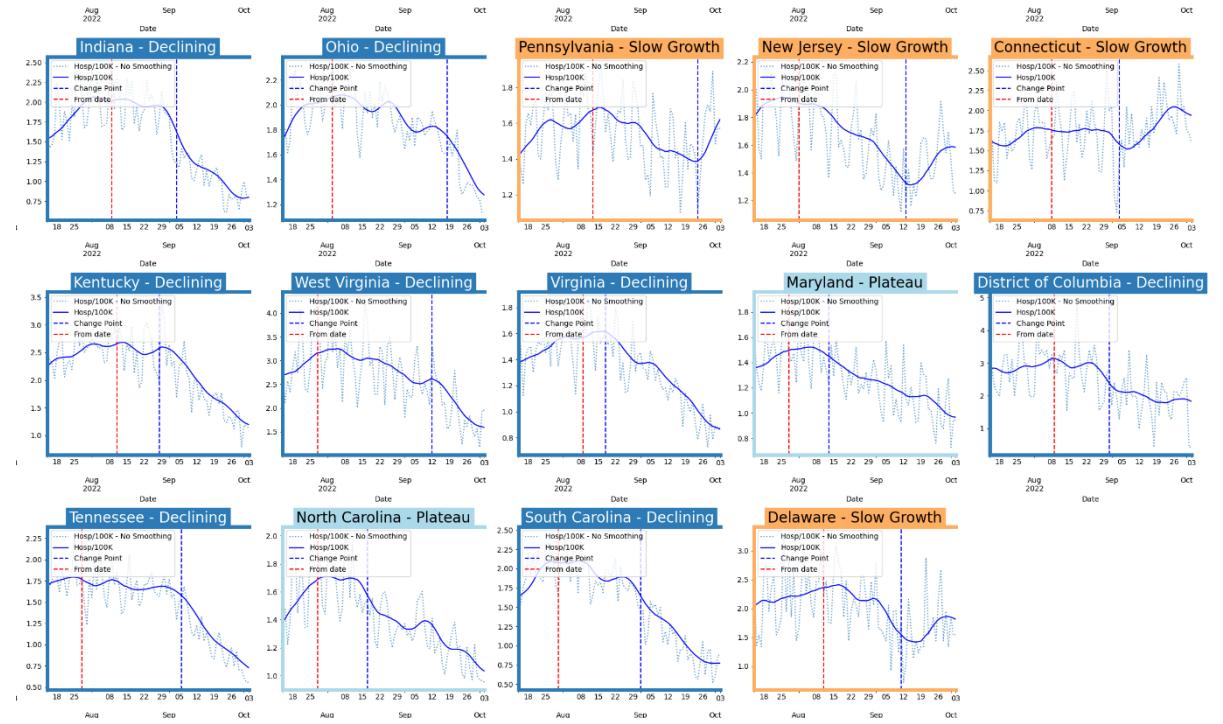
| Status | Number of States | |
|--------------------|------------------|-----------|
| | Current Week | Last Week |
| Declining | 27 | (26) |
| Plateau | 17 | (21) |
| Slow Growth | 9 | (6) |
| In Surge | 0 | (1) |

Virginia and Her Neighbors

Cases



Hospitalizations

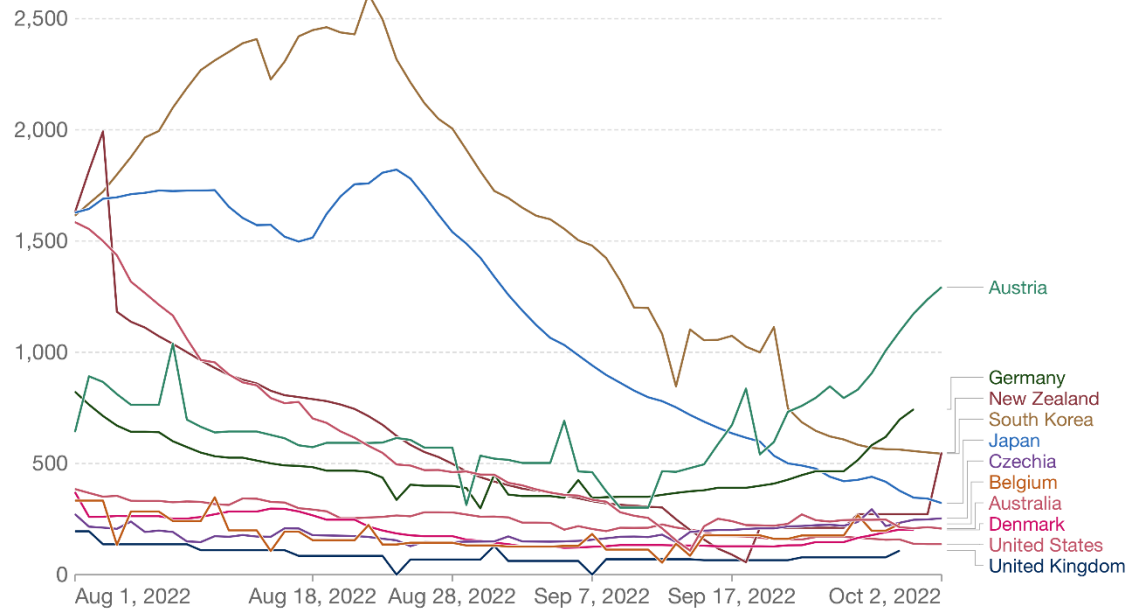


Around the World – Various trajectories

Confirmed cases

Daily new confirmed COVID-19 cases per million people

7-day rolling average. Due to limited testing, the number of confirmed cases is lower than the true number of infections.



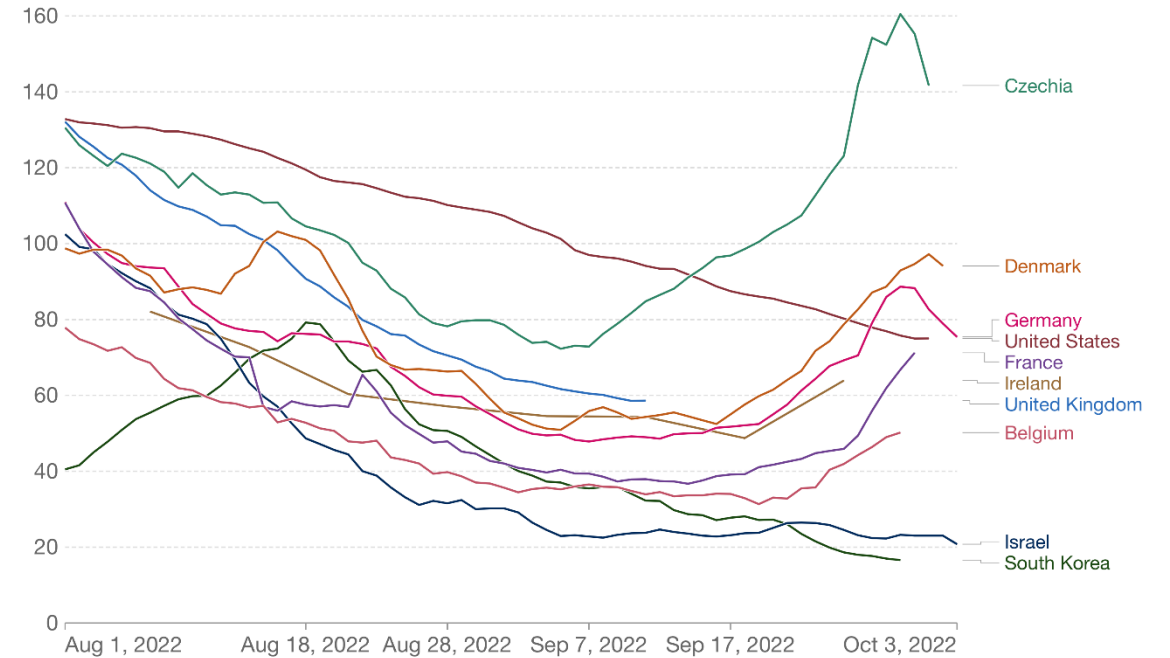
Source: Johns Hopkins University CSSE COVID-19 Data

CC BY

Hospitalizations

Weekly new hospital admissions for COVID-19 per million people

Weekly admissions refer to the cumulative number of new admissions over the previous week.



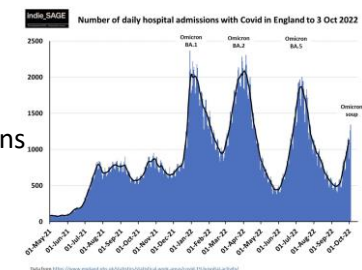
Source: Official data collated by Our World in Data

CC BY

[Our World in Data](https://ourworldindata.org/)

UNIVERSITY of VIRGINIA

[National Health England](https://www.nhs.uk/)
Rapid rise in hospitalizations



Zip code level weekly Case Rate (per 100K)

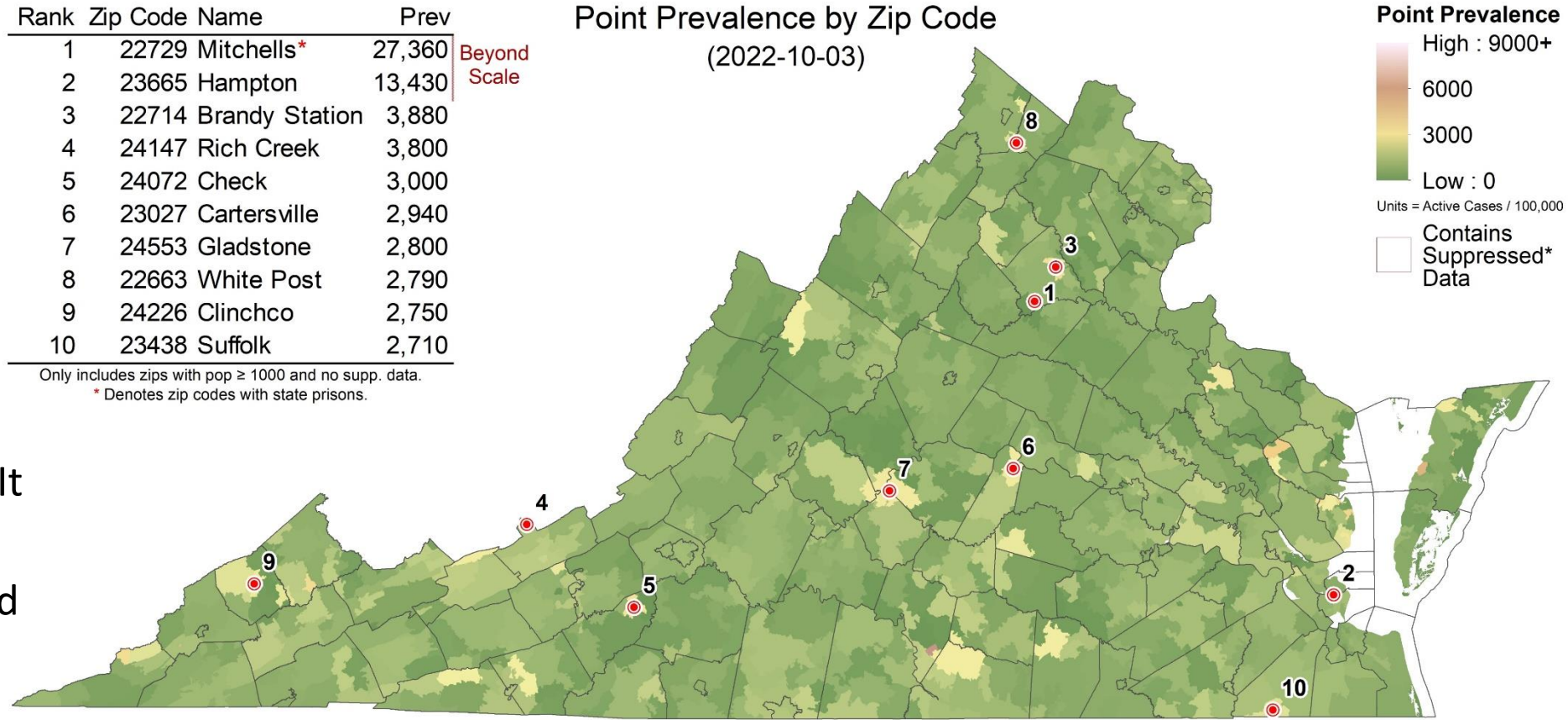
Case Rates in the last week by zip code

- Statewide rates have fallen significantly. We will adjust the color ramp if trends continue.
- High prevalence areas are spread randomly across Virginia (no spatial autocorrelation).
- Mitchells, VA has by far the highest prevalence. It is home to a prison.
- Some counts are low and suppressed to protect anonymity. Those are shown with a dark red outline.

| Rank | Zip Code | Name | Prev |
|------|----------|----------------|--------|
| 1 | 22729 | Mitchells* | 27,360 |
| 2 | 23665 | Hampton | 13,430 |
| 3 | 22714 | Brandy Station | 3,880 |
| 4 | 24147 | Rich Creek | 3,800 |
| 5 | 24072 | Check | 3,000 |
| 6 | 23027 | Cartersville | 2,940 |
| 7 | 24553 | Gladstone | 2,800 |
| 8 | 22663 | White Post | 2,790 |
| 9 | 24226 | Clinchco | 2,750 |
| 10 | 23438 | Suffolk | 2,710 |

Only includes zips with pop ≥ 1000 and no supp. data.
 * Denotes zip codes with state prisons.

Point Prevalence by Zip Code (2022-10-03)



Based on Spatial Empirical Bayes smoothed point prevalence, with an 8:1 ascertainment ratio, for week ending 2022-10-03.

Risk of Exposure by Group Size and HCW prevalence

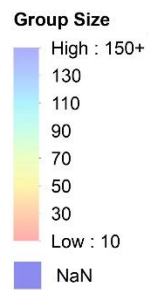
Case Prevalence in the last week by zip code used to calculate risk of encountering someone infected in a gathering of randomly selected people

- **Group Size:** Assumes **8 undetected infections** per confirmed case (ascertainment rate from recent seroprevalence survey) and shows minimum size of a group with a 50% chance an individual is infected by zip code (e.g., in a group 2 in Mitchells, there is a 50% chance someone will be infected).
- **HCW ratio:** Case rate among health care workers (HCW) in the last week using patient facing health care workers as the denominator / general population's case prevalence. Note Williamsburg City.

| Rank | Zip Code | Name | Size |
|------|----------|----------------|------|
| 1 | 22729 | Mitchells* | 2 |
| 2 | 23665 | Hampton | 5 |
| 3 | 22714 | Brandy Station | 18 |
| 4 | 24147 | Rich Creek | 18 |
| 5 | 24072 | Check | 23 |
| 6 | 23027 | Cartersville | 23 |
| 7 | 24553 | Gladstone | 24 |
| 8 | 22663 | White Post | 24 |
| 9 | 24226 | Clinchco | 25 |
| 10 | 23438 | Suffolk | 25 |

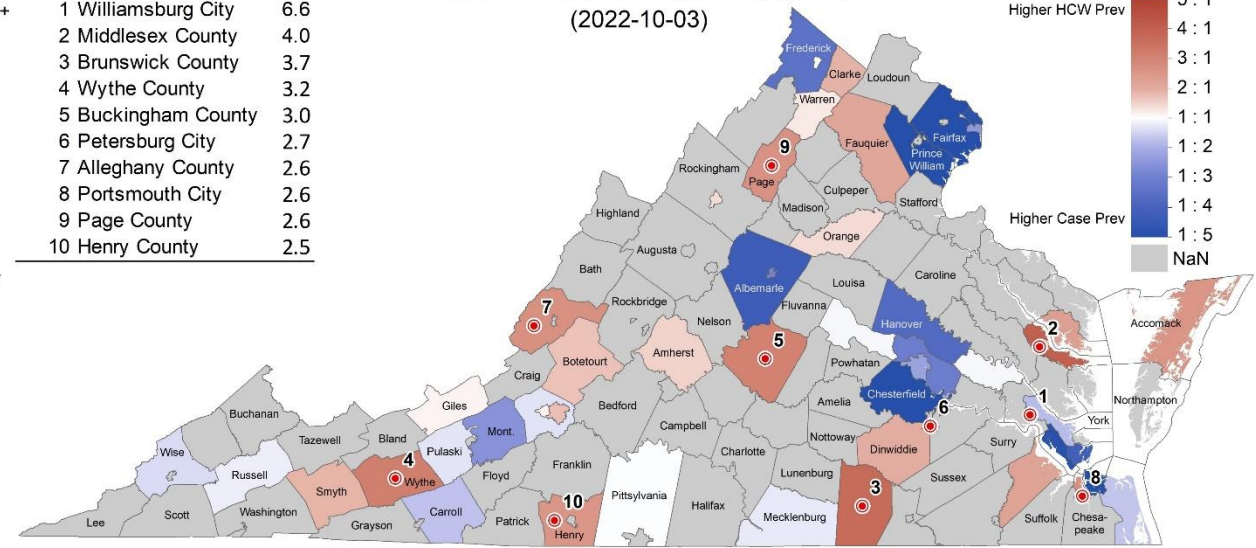
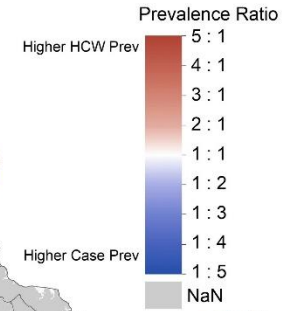
Only includes zips with pop ≥ 1000 and no supp. data.
* Denotes zip codes with state prisons.

Group Size Needed for 50% Likelihood of ≥1 Infected



| Rank | Name | Ratio |
|------|-------------------|-------|
| 1 | Williamsburg City | 6.6 |
| 2 | Middlesex County | 4.0 |
| 3 | Brunswick County | 3.7 |
| 4 | Wythe County | 3.2 |
| 5 | Buckingham County | 3.0 |
| 6 | Petersburg City | 2.7 |
| 7 | Alleghany County | 2.6 |
| 8 | Portsmouth City | 2.6 |
| 9 | Page County | 2.6 |
| 10 | Henry County | 2.5 |

HCW Prevalence / Case Prevalence (2022-10-03)



Note: This assumes that the ascertainment rate of healthcare workers is double that of the public.

Based on Spatial Empirical Bayes smoothed point prevalence, with an 8:1 ascertainment ratio, for week ending 2022-10-03.

Current Hot-Spots

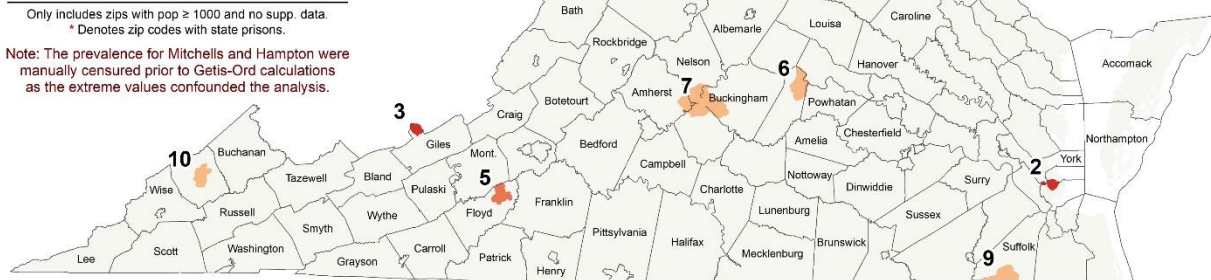
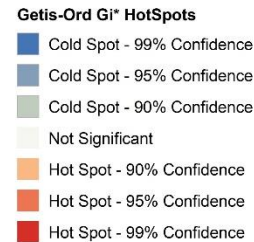
Case rates that are significantly different from neighboring areas or model projections

- **Spatial:** Getis-Ord G_i^* based hot spots compare clusters of zip codes with weekly case prevalence higher than nearby zip codes to identify larger areas with statistically significant deviations
- **Temporal:** The weekly case rate (per 100K) projected last week compared to observed by county, which highlights temporal fluctuations that differ from the model's projections. **Note new color ramp (down 75%).**
- Spatial hot spots are appearing sporadically across the Commonwealth. Models slightly underpredicted the Far Southwest, and slightly overpredicted Central Virginia, Northern Virginia, and the Tidewater.

Spatial Hotspots

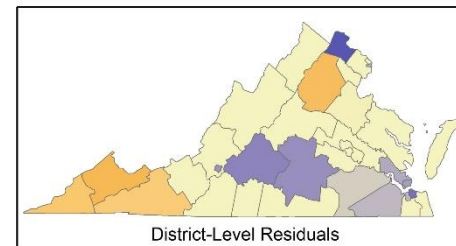
| Spot | Zip Code | Name | Conf. |
|------|----------|----------------|-------|
| 1 | 22729 | Mitchells | 99% |
| 2 | 23665 | Hampton | 99% |
| 3 | 24147 | Rich Creek | 99% |
| 4 | 22714 | Brandy Station | 99% |
| 5 | 24072 | Check | 95% |
| 6 | 23027 | Cartersville | 90% |
| 7 | 24553 | Gladstone | 90% |
| 8 | 22663 | White Post | 90% |
| 9 | 23438 | Suffolk | 90% |
| 10 | 24226 | Clinchco | 90% |

Point Prevalence Hot Spots by Zip Code (2022-10-03)

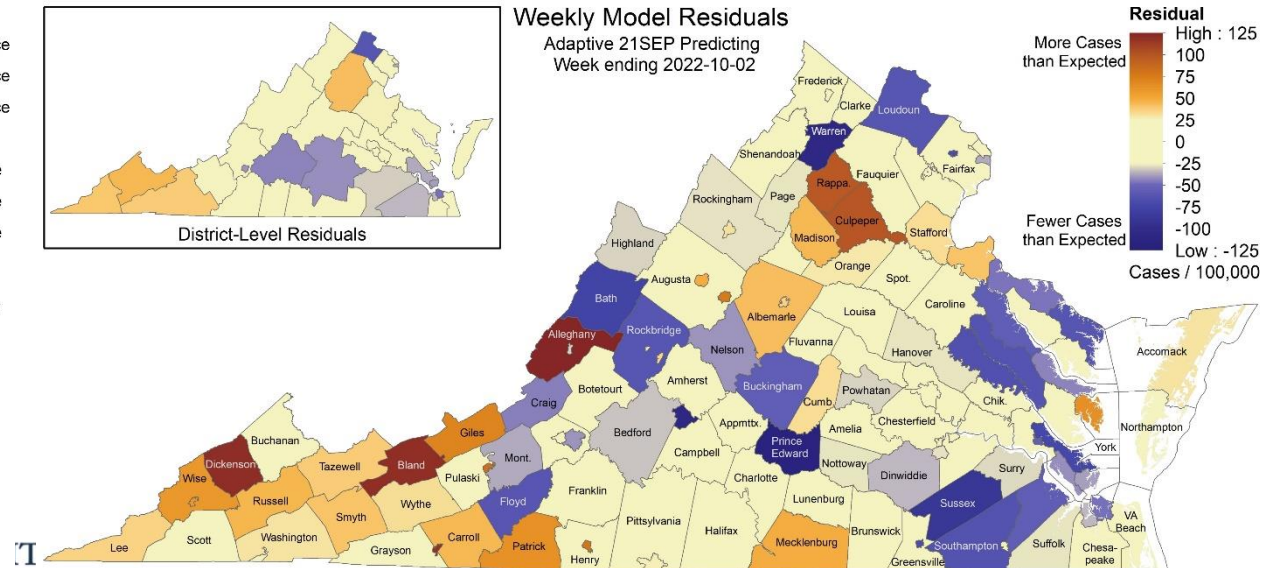
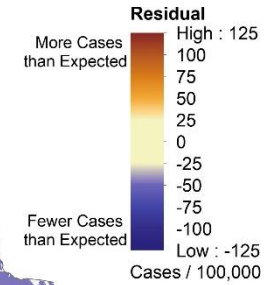


Based on Global Empirical Bayes smoothed point prevalence for week ending 2022-10-03.

Clustered Temporal Hotspots



Weekly Model Residuals
Adaptive 21SEP Predicting
Week ending 2022-10-02

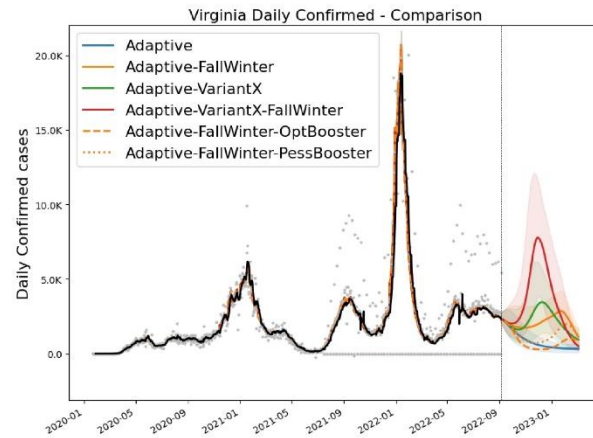


Health District Level Moran's I = 0.054294, Z-Score = 1.394733, P-Value = 0.163096
No Residual Autocorrelation Detected

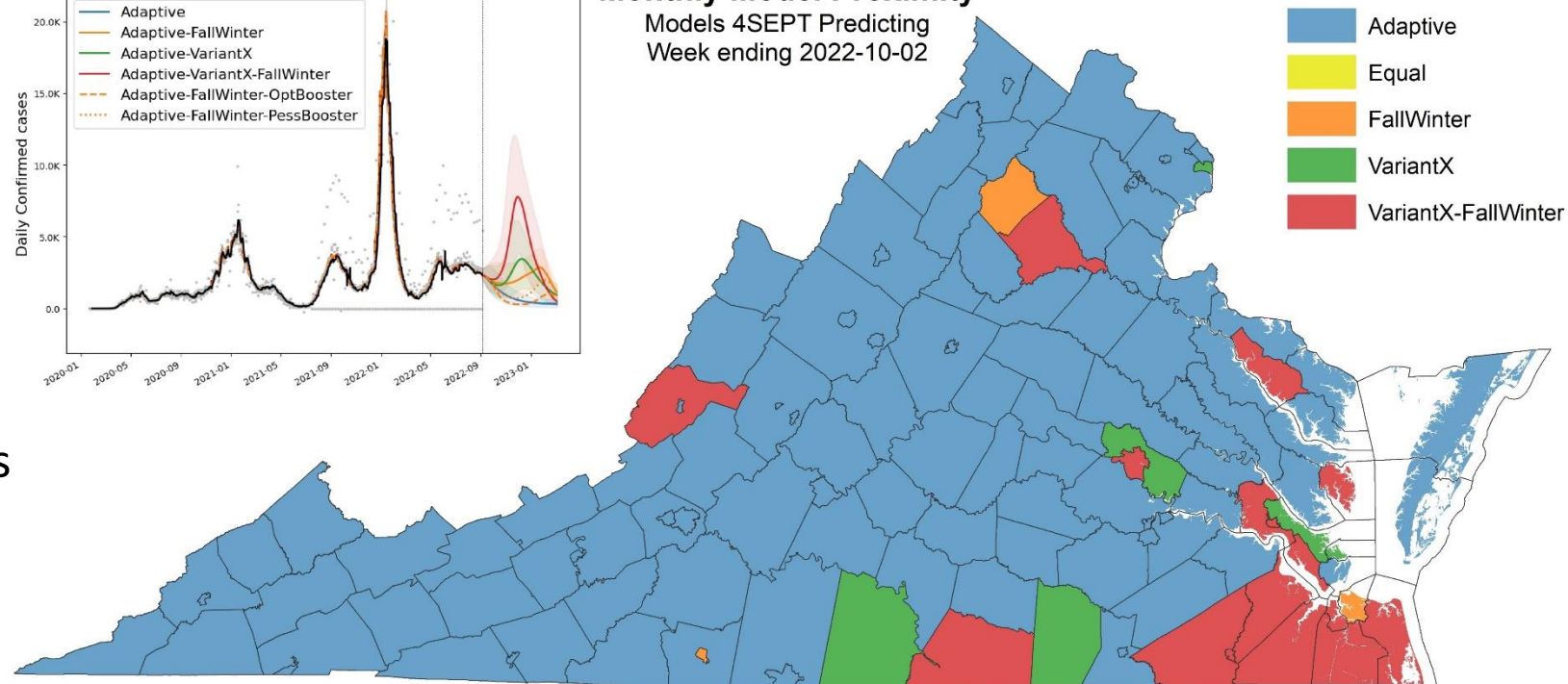
Scenario Trajectory Tracking

Which scenario from a month ago did the projection for each county track closest?

- One month out separates the projections more and reveals larger overall patterns.
- Booster scenarios not included as vaccination efforts have only recently begun.
- Overall state level models were on track, largely tracking the Adaptive scenario over Variant X. This is expected, as no novel variants supplanted BA.4/5 in the time since the modeling run.
- Currently only a handful of counties, mostly in the Southeast, are tracking FallWinter scenarios more closely than the Adaptive scenario.



Monthly Model Proximity
Models 4SEPT Predicting
Week ending 2022-10-02

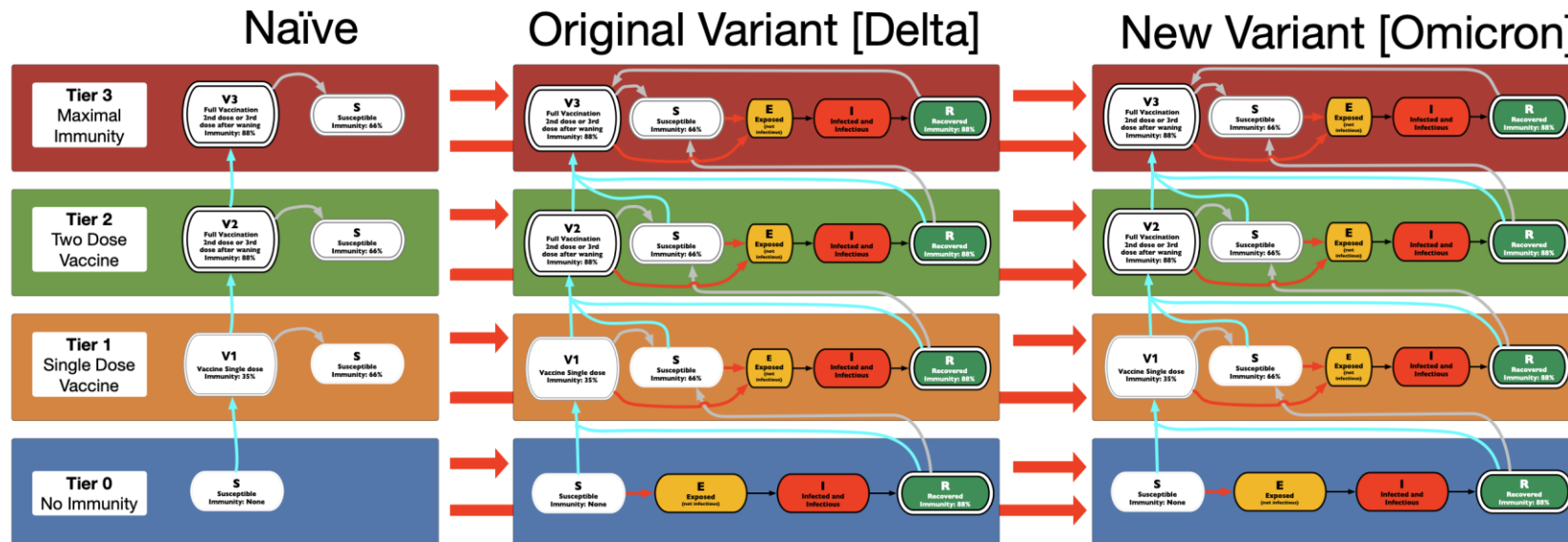


Model Update – Adaptive Fitting

Model Structure Extended for more sub-variants

Omicron sub-variants escape immunity induced by previous sub-variants

- Multiple strain support allows representation of differential protection based on immunological history (BA.1, BA.2, BA.2.12.1, BA.4/5, and future variants (VariantX))
- Each sub-variant has differing levels of immune escape to previous sub-variants, the prevalences are based on observations for fitting purposes, and projections use estimated future prevalences
- Adaptive fitting approach continues to use simulation to generate the full distribution of immune states across the population



Adaptive Fitting Approach

Each county fit precisely, with recent trends used for future projection

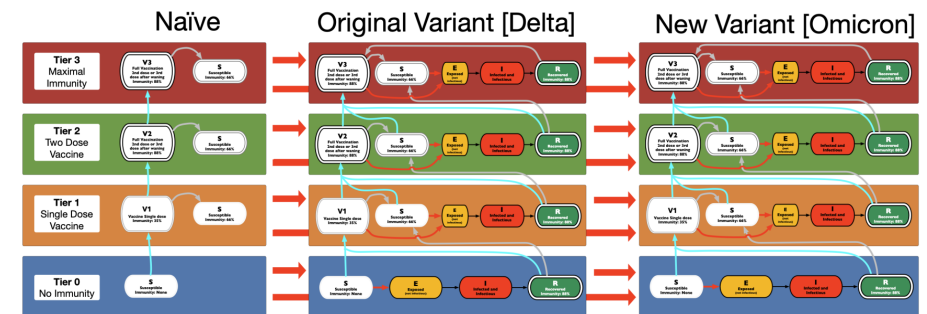
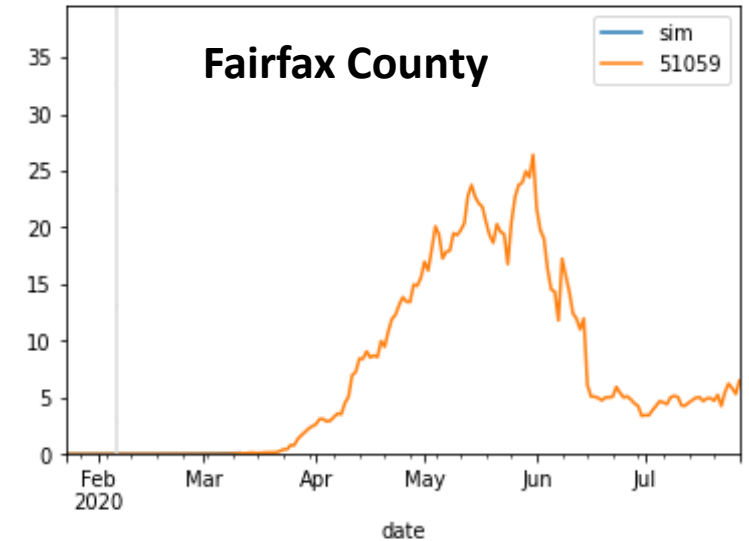
- Allows history to be precisely captured, and used to guide bounds on projections

Model: An alternative use of the same meta-population model, PatchSim with multiple tiers of immunity

- Allows for future “what-if” Scenarios to be layered on top of calibrated model
- Allows for waning of immunity and for partial immunity against different outcomes (eg lower protection for infection than death)

External Seeding: Steady low-level importation

- Widespread pandemic eliminates sensitivity to initial conditions, we use steady 1 case per 10M population per day external seeding



Using Ensemble Model to Guide Projections

Ensemble methodology that combines the Adaptive with machine learning and statistical models such as:

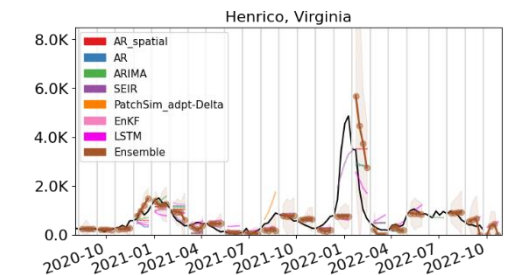
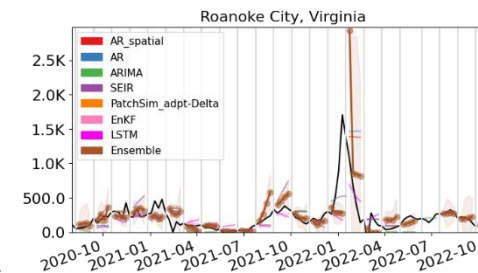
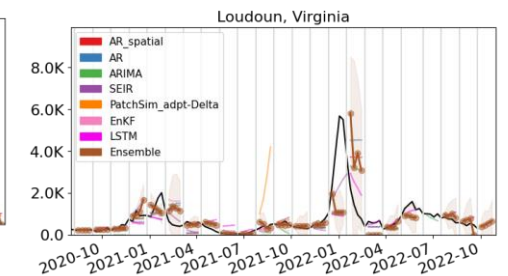
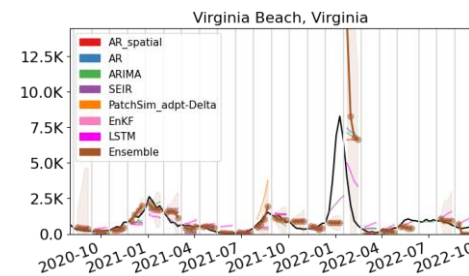
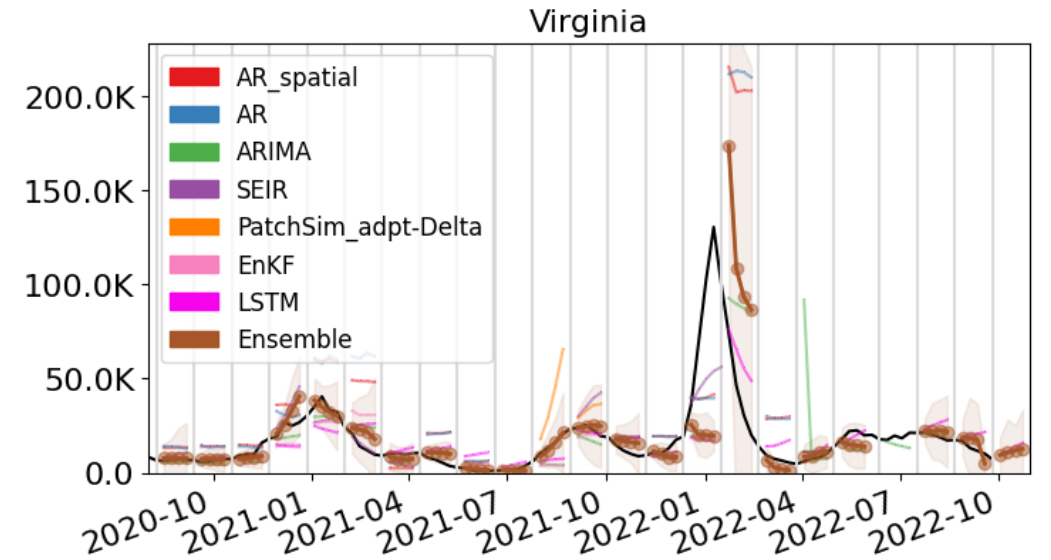
- Autoregressive (AR, ARIMA)
- Neural networks (LSTM)
- Kalman filtering (EnKF)

Weekly forecasts done at county level.

Models chosen because of their track record in disease forecasting and to increase diversity and robustness.

Ensemble forecast provides additional 'surveillance' for making scenario-based projections.

Also submitted to CDC Forecast Hub.



Calibration Approach

- **Data:**
 - County level case counts by date of onset (from VDH)
 - Confirmed cases for model fitting
- **Calibration:** fit model to observed data and ensemble's forecast
 - Tune transmissibility across ranges of:
 - Duration of incubation (5-9 days), infectiousness (3-7 days)
 - Undocumented case rate (1x to 7x) guided by seroprevalence studies
 - Detection delay: exposure to confirmation (4-12 days)
 - Approach captures uncertainty, but allows model to precisely track the full trajectory of the outbreak
- **Project:** future cases and outcomes generated using the collection of fit models run into the future
 - **Mean trend from last 7 days of observed cases and first week of ensemble's forecast used**
 - Outliers removed based on variances in the previous 3 weeks
 - 2 week interpolation to smooth transitions in rapidly changing trajectories
- **Outcomes:** Data driven by shift and ratio that has least error in last month of observations
 - Hospitalizations: 3 days from confirmation, 6.8% of cases hospitalized
 - Deaths: 11 days from confirmation, 1.45% of cases die



COVID-19 in Virginia: Summary

Dashboard Updated: 10/5/2022
Data entered by 5:00 PM the prior day.



| Cases, Hospitalizations and Deaths | | | | | |
|------------------------------------|-----------------------|------------------------------------|-----------------------|------------------------|-----------------------|
| Total Cases* | | Total Hospital Admissions** | | Total Deaths | |
| 2,093,393 | | 56,344 | | 21,919 | |
| (New Cases: 1,138) [^] | | | | | |
| Confirmed [†] | Probable [†] | Confirmed [†] | Probable [†] | Confirmed [†] | Probable [†] |
| 1,482,278 | 611,115 | 52,871 | 3,473 | 18,208 | 3,711 |

* Includes both people with a positive test (Confirmed), and symptomatic with a known exposure to COVID-19 (Probable).

** Hospitalization of a case is captured at the time VDH performs case investigation. This underrepresents the total number of hospitalizations in Virginia.

[^]New cases represent the number of confirmed and probable cases reported to VDH in the past 24 hours.

[†] VDH adopted the updated CDC COVID-19 confirmed and probable surveillance case definitions on August 27, 2020. Found here: <https://www.cdc.gov/nndss/conditions/coronavirus-disease-2019-covid-19/case-definition/2020/08/05/>

Source: Cases - Virginia Electronic Disease Surveillance System (VEDSS); data entered by 5:00 PM the prior day.

| Outbreaks | |
|-------------------------|----------------------------------|
| Total Outbreaks* | Outbreak Associated Cases |
| 9,912 | 161,992 |

* At least two (2) lab confirmed cases are required to classify an outbreak.

| Testing (PCR Only) | |
|-------------------------------------|---|
| Testing Encounters PCR Only* | Current 7-Day Positivity Rate PCR Only** |
| 15,035,997 | 10.5% |

* PCR refers to "Reverse transcriptase polymerase chain reaction laboratory testing."

** Lab reports may not have been received yet. Percent positivity is not calculated for days with incomplete data.

| Multisystem Inflammatory Syndrome in Children | |
|---|---------------------|
| Total Cases* | Total Deaths |
| 181 | 1 |

*Cases defined by CDC HAN case definition: <https://emergency.cdc.gov/han/2020/han00432.asp>

Accessed 9:00am October 5, 2022

<https://www.vdh.virginia.gov/coronavirus/>



Scenarios – Transmission Conditions

- Variety of factors continue to drive transmission rates
 - Seasonal impact of weather patterns, travel and gatherings, fatigue and premature relaxation of infection control practices
- **Waning Immunity:** Omicron waning with a mean of 4 months
- **Projection Condition Ingredients:**
 - **Adaptive:** Controls remain as currently experienced into the future with NO influence from other conditions (eg seasonal, variants, etc.)
 - **Seasonal (Fall-Winter boosting):** Controls remain the same, however, seasonal forcing similar to past Fall-Winter waves is added from Sept-Feb
 - **Vaccine Booster Campaign (Booster):** Reformulated booster available this fall provides improved immunity against Omicron sub-variants
 - **New Variants (VariantX):** As of yet unidentified novel sub-variant with similar immune escape but no transmission advantage emerges 4 months after the last significant sub-variant and grows at a similar rate

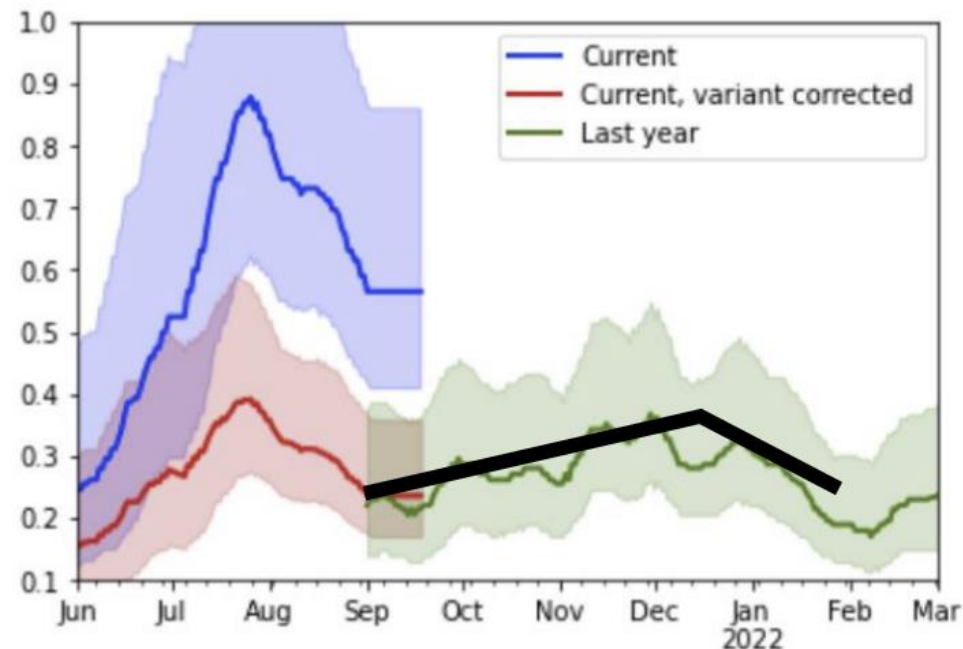
Scenarios – FallWinter

September – February saw strong waves of transmission for both years

- Based on analyses of the past 2 seasons we generate a “coarse baseline transmission boost”
 - In 2021 the distribution of fitted model transmissibility was nearly identical between these periods when corrected for Delta’s increased transmissibility
- **FallWinter** captures these “transmission drivers” from the past and uses them as if they were to occur again this season

Fitting:

Black line represents the coarsely fitted base transmissibility



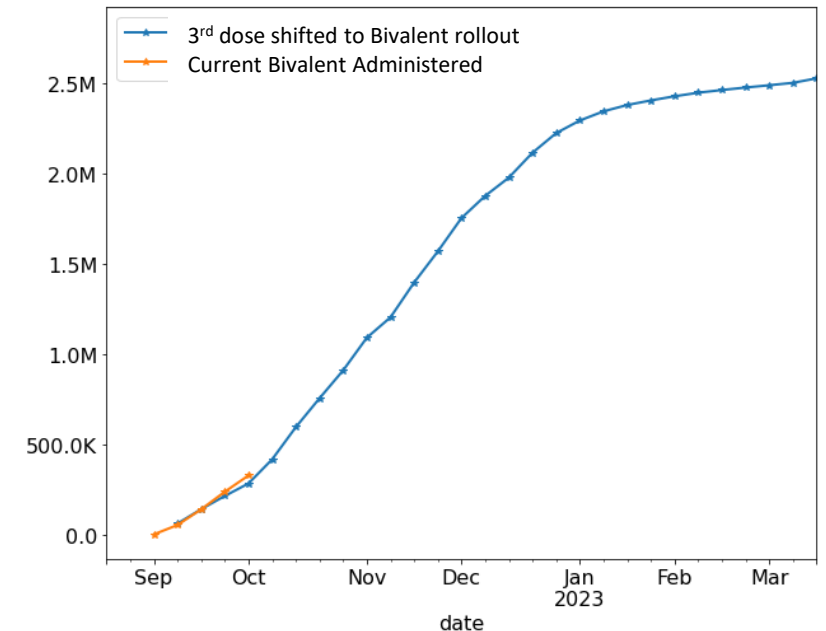
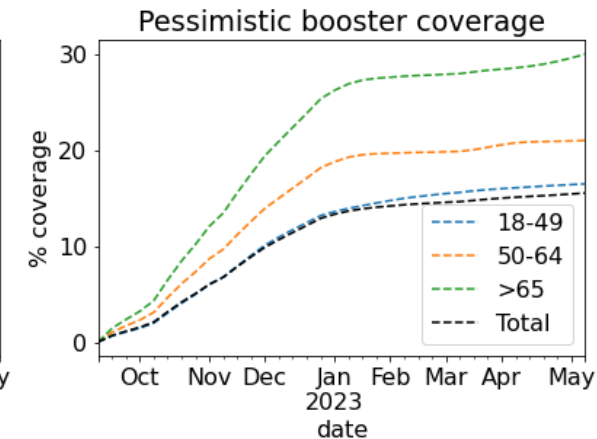
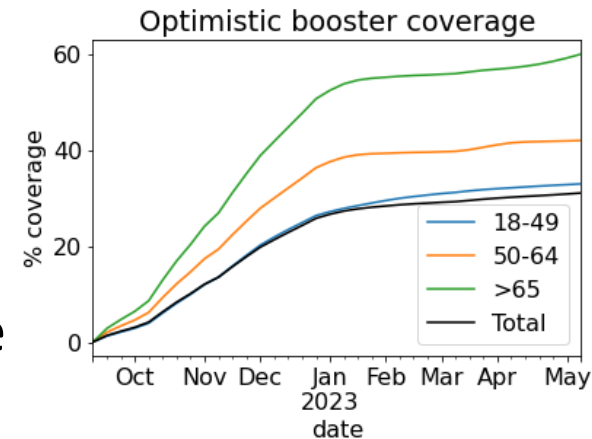
Scenarios – Optimistic vs. Pessimistic Booster Coverage

Reformulated Boosters available now

- Assuming Vax efficacy for BA.4/5 and previous variants is 80% against symptomatic illness
- Campaign starts in mid-September following the ongoing rollout
- Pace of vaccination follows that of 3rd booster (adjusted after initial uptake did not keep pace with seasonal influenza vax uptake)
- Variant X has same immune escape to these vaccines as against BA.5 (40%)

Optimistic coverage: 33% of pop (same as 3rd dose)

Pessimistic coverage: 16% of pop (half of 3rd dose)

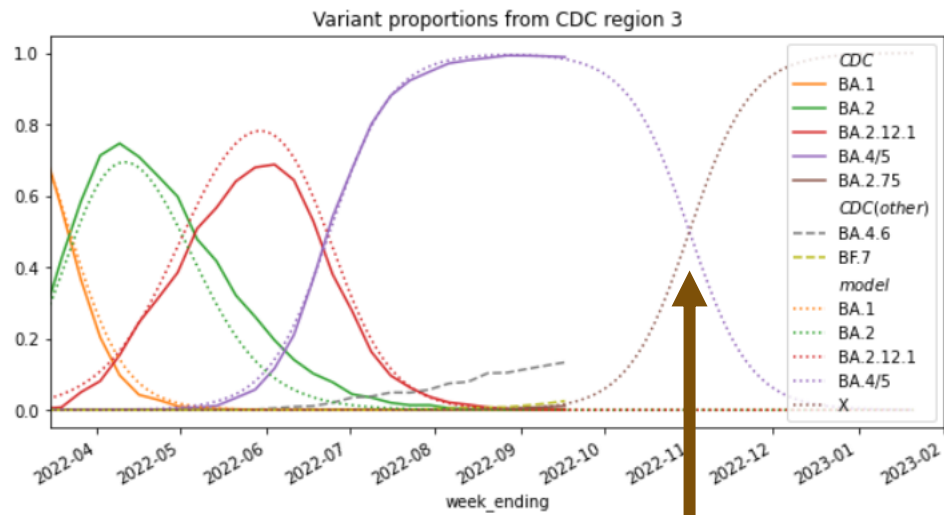


Scenarios – Variant X

Omicron sub-variants seem to be emerging and then dominating with some regularity

- ECDC currently monitoring BA.2 + L452X and BA.2.75 as VOI and XAK as well as other BA.4 and BA.5 strains as Variants under Monitoring
- BA.2.75.2 remains main sub-variant characterized enough to suggest future dominance, but has yet to demonstrate rapid growth in US
- Hypothetical future sub-variant, **VariantX**, may continue the pattern. Assumes similar growth and level of immune escape against previous sub-variants as BA.4/5 (same transmissibility and 40% immune escape against BA.4/5, higher for other sub-variants).

Sub-Variants with Fitted Prevalences and Hypothetical Future waves



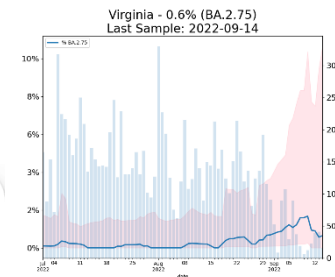
7-Oct-22

Variant X reaches 50% on Nov 15th

Variants of Interest

| WHO label | Lineage + additional mutations | Country first detected (community) | Spike mutations of interest | Year and month first detected | Impact on transmissibility | Impact on immunity | Impact on severity | Transmission in EU/EEA |
|-----------|--------------------------------|------------------------------------|-----------------------------|-------------------------------|----------------------------|--------------------|--------------------|------------------------|
| Omicron | BA.2 + L452X (x) | n/a | L452X | n/a | No evidence | Increased (7) | No evidence | Detected (a) |
| Omicron | BA.2.75 | India | (y) | May 2022 | No evidence | Increased (9, 10) | No evidence | Detected (a) |

BA.2.75 detected in VA (very limited samples)



Variants under Monitoring

| WHO label | Lineage + additional mutations | Country first detected (community) | Spike mutations of interest | Year and month first detected | Impact on transmissibility | Impact on immunity | Impact on severity | Transmission in EU/EEA |
|-----------|--------------------------------|------------------------------------|-----------------------------|-------------------------------|----------------------------|--------------------|--------------------|------------------------|
| Omicron | XAK | Germany | (y) | June 2022 | No evidence | No evidence | No evidence | Detected (a) |
| Omicron | BA.4 + R346X (x) | n/a | R346X | n/a | No evidence | No evidence | No evidence | Detected (a) |
| Omicron | BA.5 + R346X (x) | n/a | R346X | n/a | No evidence | No evidence | No evidence | Detected (a) |

Projection Scenarios – Combined Conditions

| Name | Txm | Variant | Description |
|---------------------------------|------------|---------|---|
| Adaptive | C | SQ | Likely trajectory based on conditions remaining similar to the current experience, includes immune escape due to Omicron |
| Adaptive-FallWinter | FallWinter | SQ | Like Adaptive, with seasonal forcing of FallWinter added on |
| Adaptive-FallWinter-OptBooster | FallWinter | SQ | Like Adaptive-Fall Winter but with Optimistic Booster (90% of seasonal influenza coverage) |
| Adaptive-FallWinter-PessBooster | FallWinter | SQ | Like Adaptive-Fall Winter but with Pessimistic Booster (45% of seasonal influenza coverage) |
| Adaptive-VariantX | C | X | Like Adaptive, with emergence of a speculative unknown variant 4 months after BA.4/5 with similar level of immune escape and equal transmissibility |
| Adaptive-VariantX-FallWinter | FallWinter | X | Like Adaptive-VariantX but with the seasonal force of FallWinter added on |

Transmission:

C = Current levels persist into the future

FallWinter = Transmission rates learned from Sept through February of past seasons are estimated and added as a seasonal boosting to baseline transmission rates

Variant:

SQ = Status quo of current transmission driver from BA.5 remains the same (eg already significantly past dominance, thus no significant major driving of transmission anticipated)

X = Speculative novel sub-variant scenario, emerges 4 months after current variant, similar levels of escape

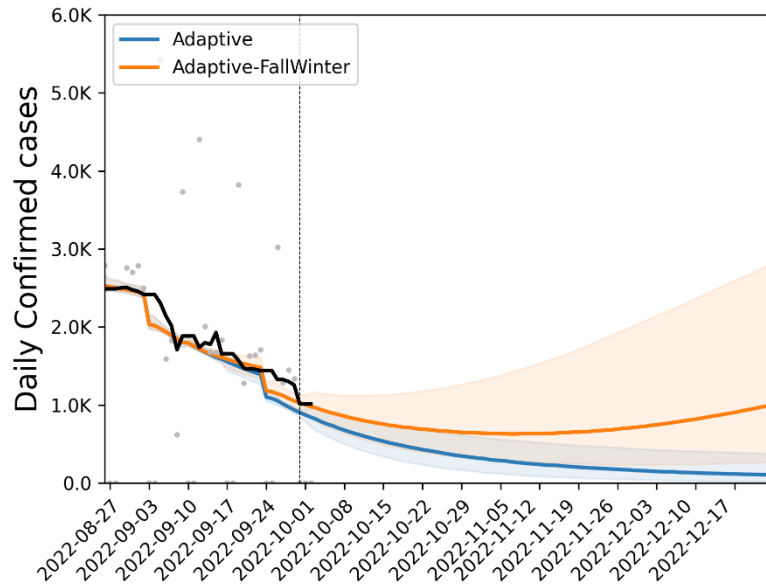
Model Results

Previous projections comparison - Cases

- Previous projections continue to track observed cases
- Projection from 2 weeks ago projected continued decline but cases plateaued
- Projection from 4 weeks ago projected slower decline better capturing recent plateau

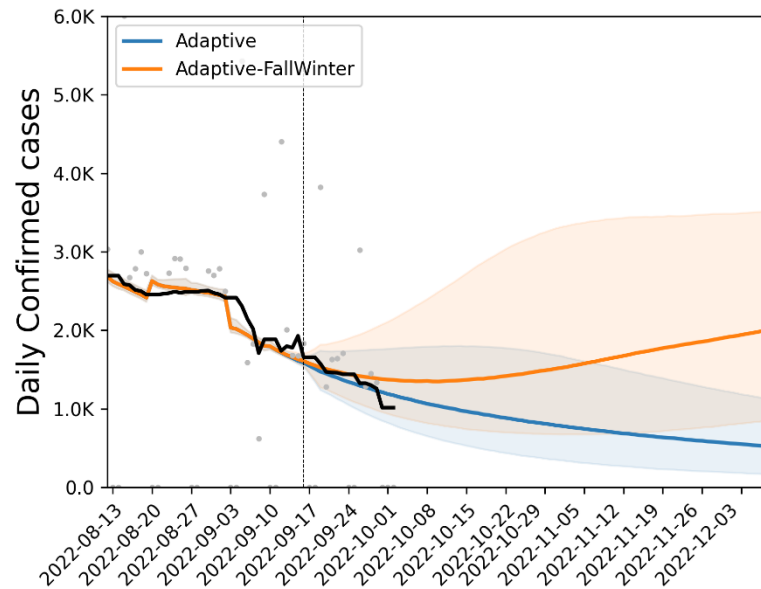
This week's projection

Virginia Daily Confirmed - Comparison 2022-09-30



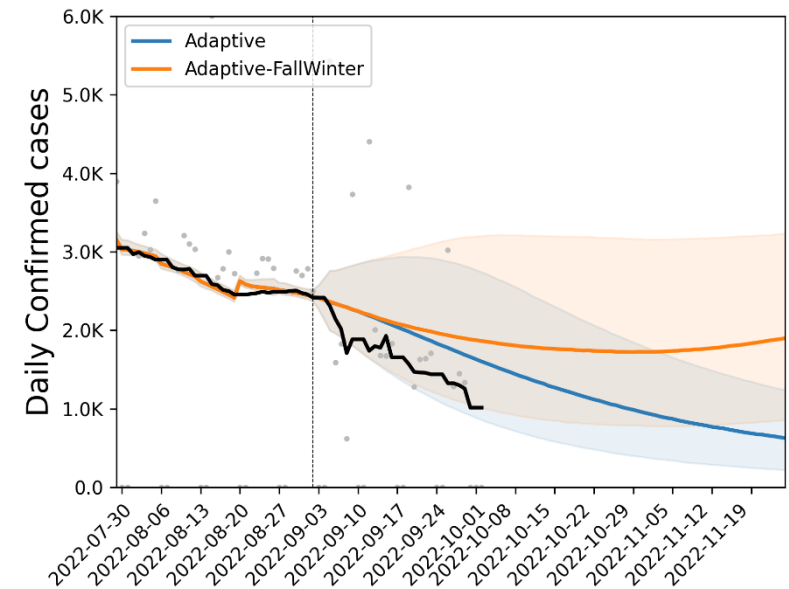
Projection from 2 weeks ago

Virginia Daily Confirmed - Comparison 2022-09-16



Projection from 4 weeks ago

Virginia Daily Confirmed - Comparison 2022-09-02

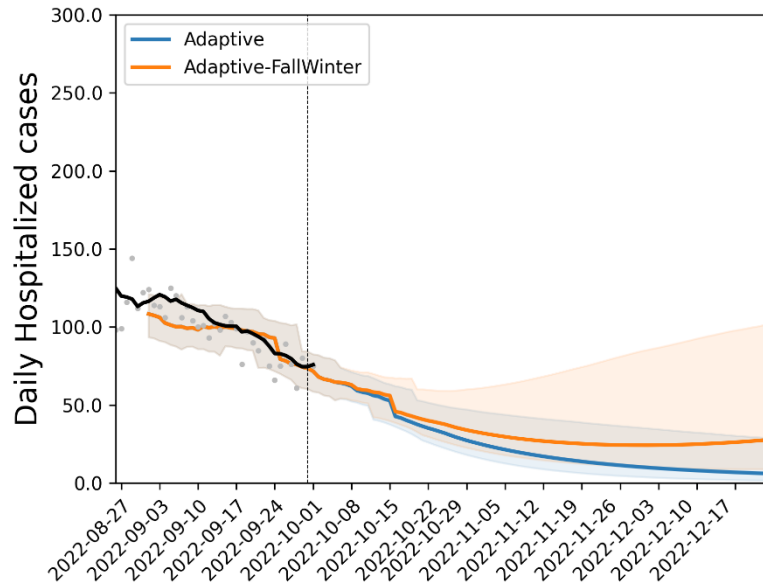


Previous projections comparison - Hospitalizations

- Previous projections have tracked observed hospitalizations well
- Projection from 2 weeks ago projected continued decline
- Projection from late July anticipated a plateau giving way to gentle decline

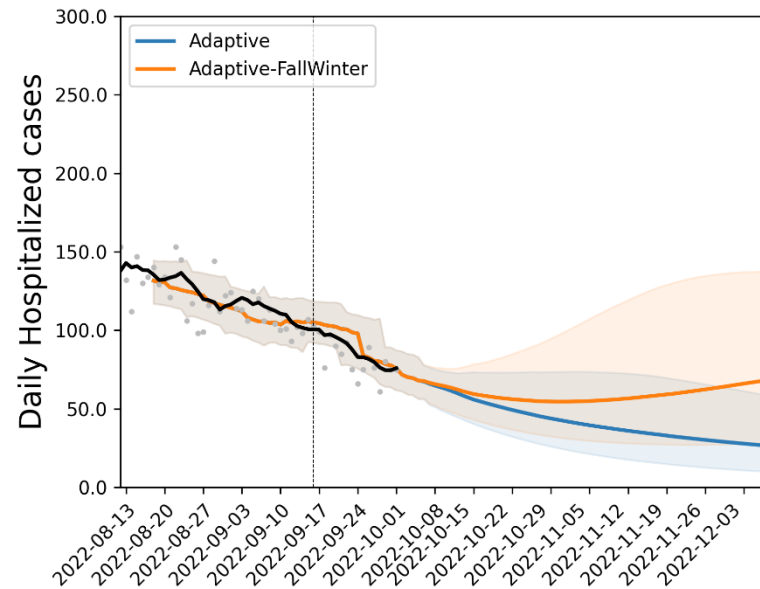
This week's projection

Virginia Daily Hospitalized - Comparison 2022-09-30



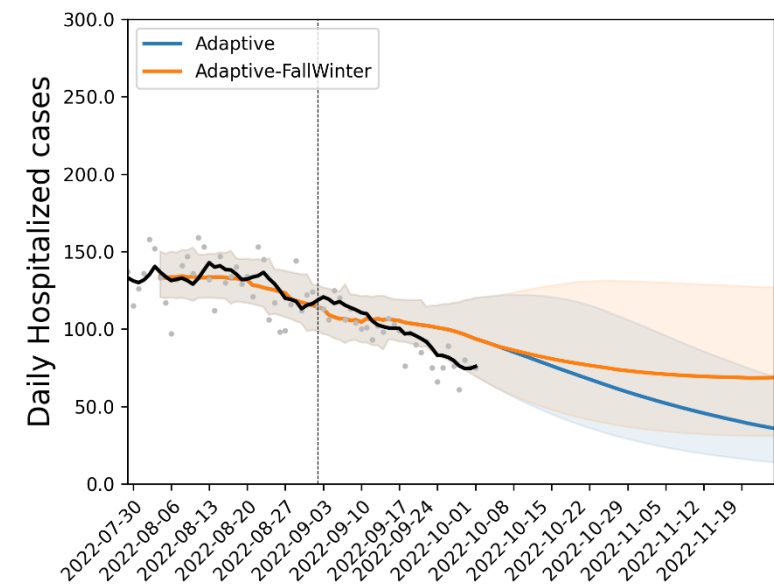
Projection from 2 weeks ago

Virginia Daily Hospitalized - Comparison 2022-09-16



Projection from 4 weeks ago

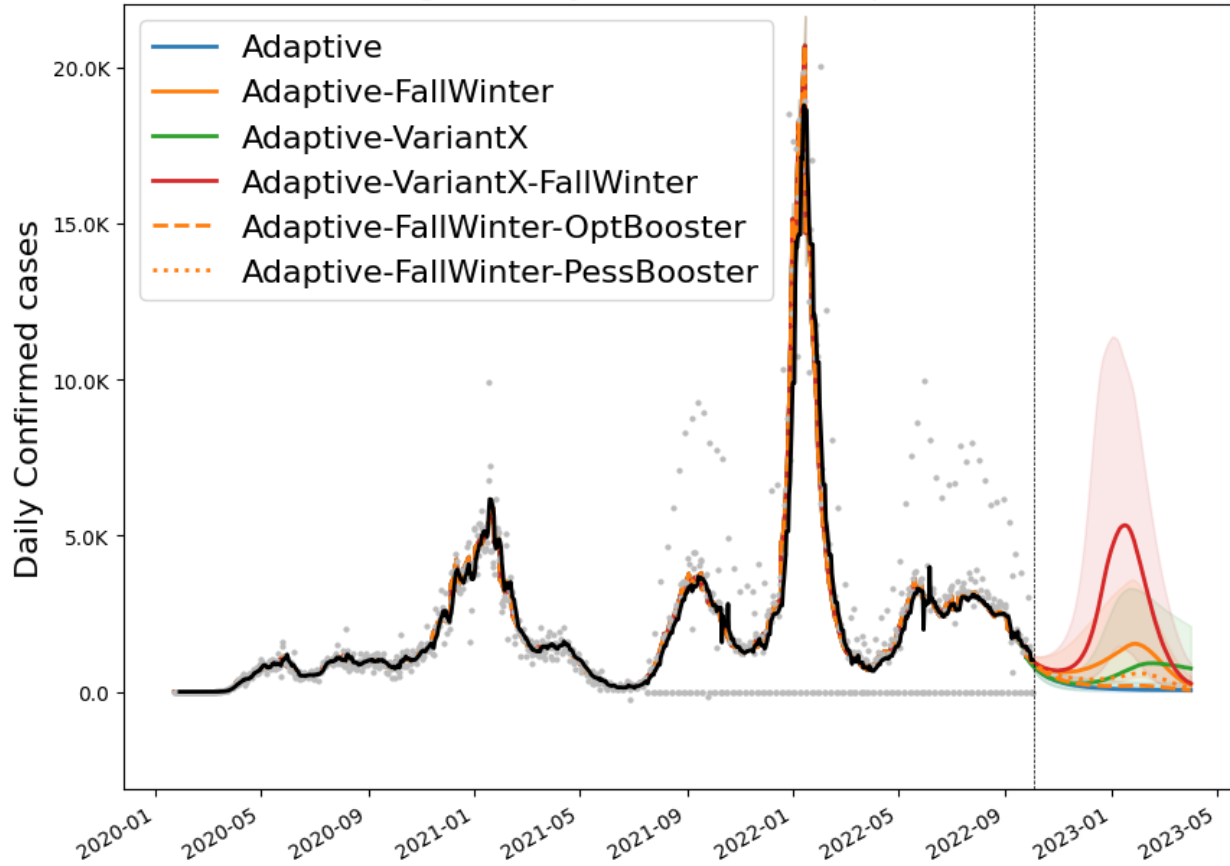
Virginia Daily Hospitalized - Comparison 2022-09-02



Outcome Projections

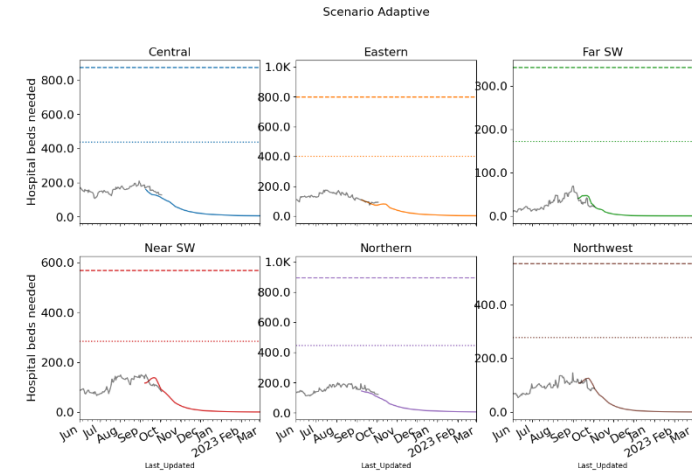
Confirmed cases

Virginia Daily Confirmed - Comparison

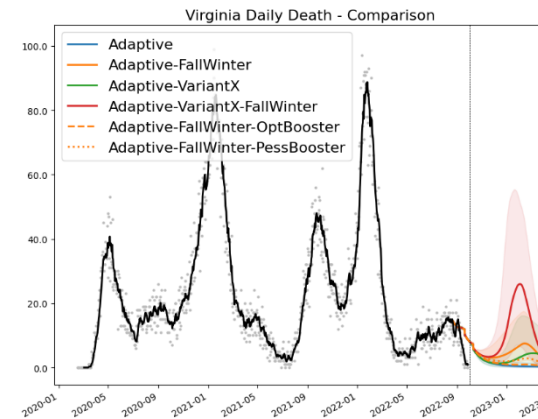


* without surveillance correction VariantBA2 peaked over 10K in July

Estimated Hospital Occupancy

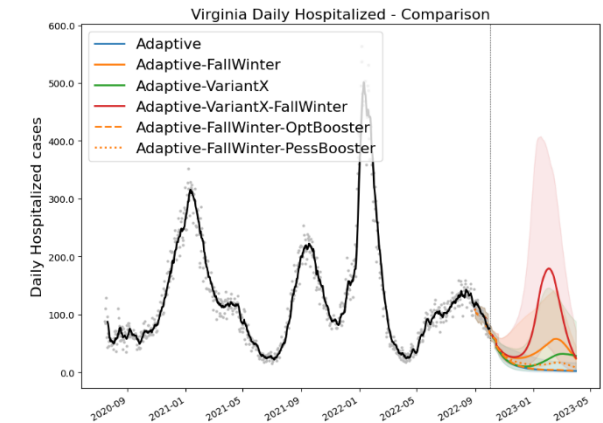


Daily Deaths



Death ground truth from VDH "Event Date" data, most recent dates are not complete

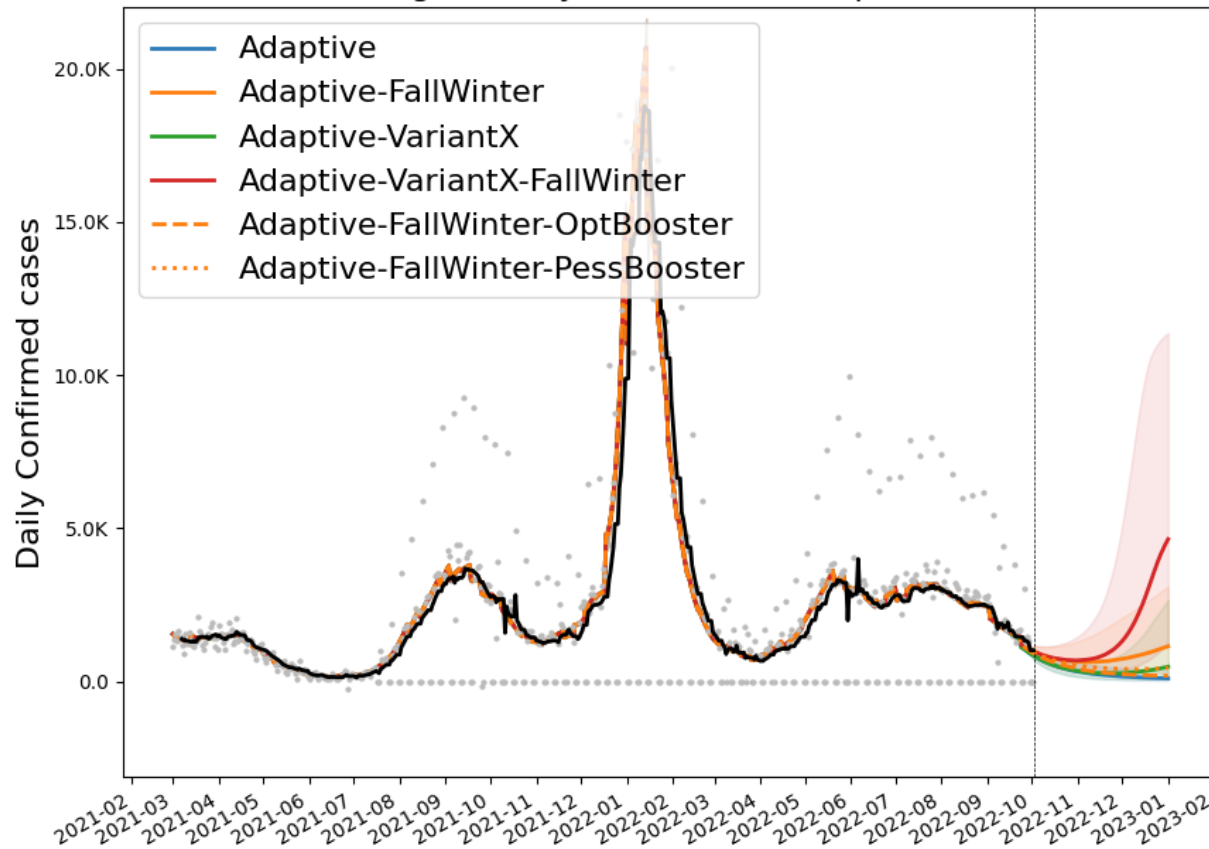
Daily Hospitalized



Outcome Projections – Closer Look

Confirmed cases

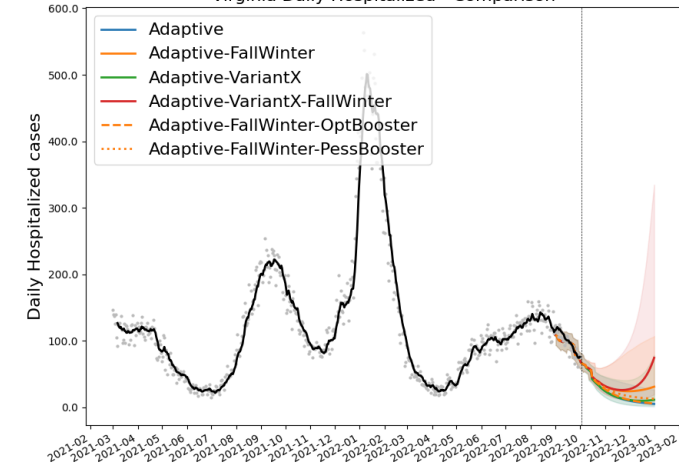
Virginia Daily Confirmed - Comparison



* without surveillance correction VariantBA2 peaked over 10K in July

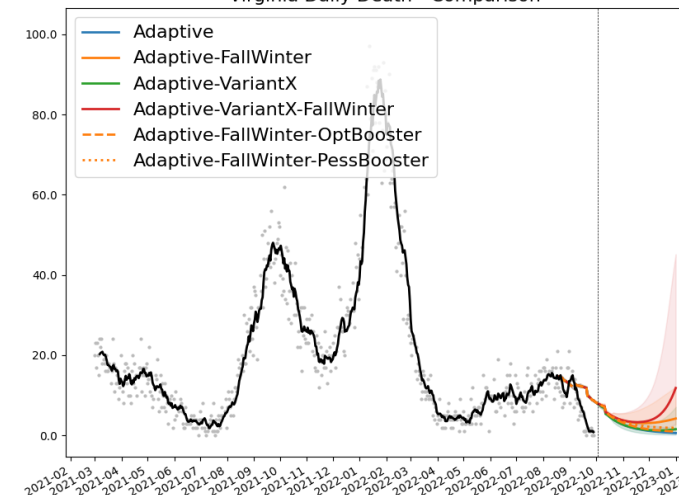
Daily Hospitalized

Virginia Daily Hospitalized - Comparison



Daily Deaths

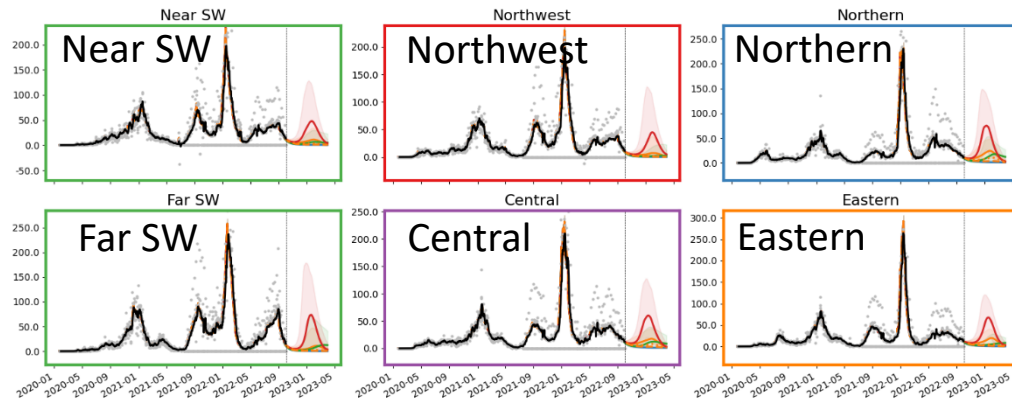
Virginia Daily Death - Comparison



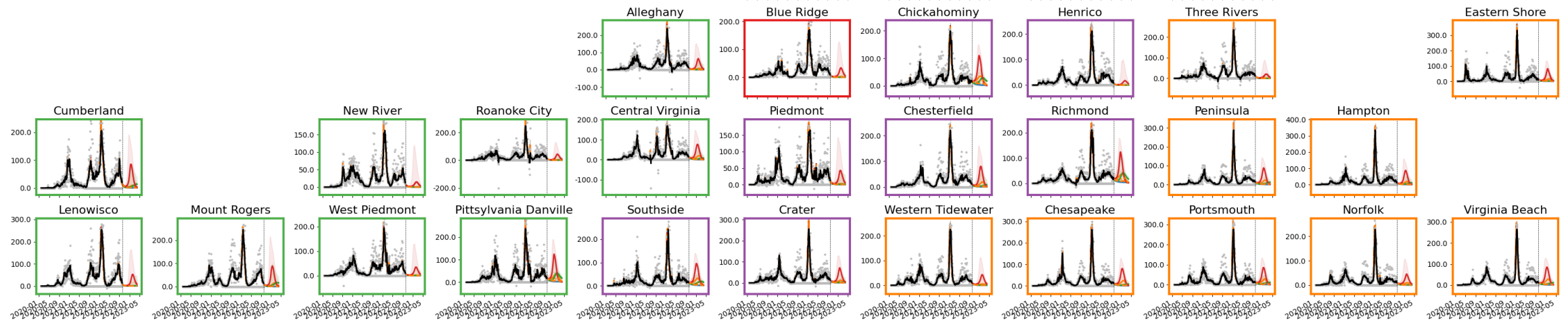
Death ground truth from VDH "Event Date" data, most recent dates are not complete

Detailed Projections: All Scenarios

Projections by Region



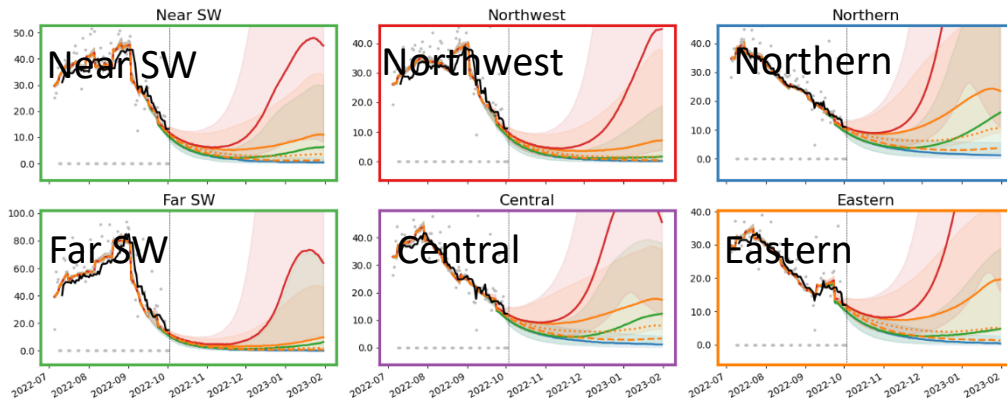
Projections by District



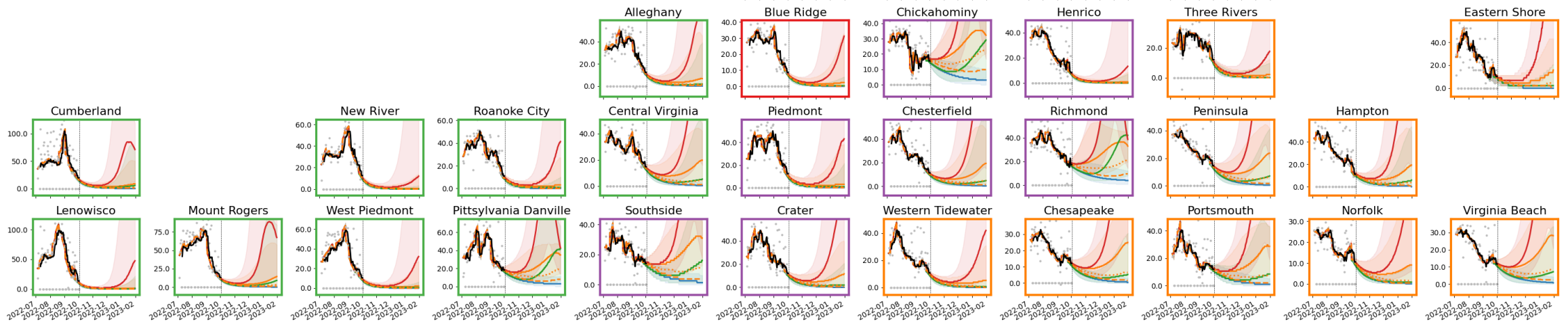
Daily confirmed cases by rate (per 100K) District (grey with 7-day average in black) with simulation colored by scenario

Detailed Projections: All Scenarios - Closer Look

Projections by Region



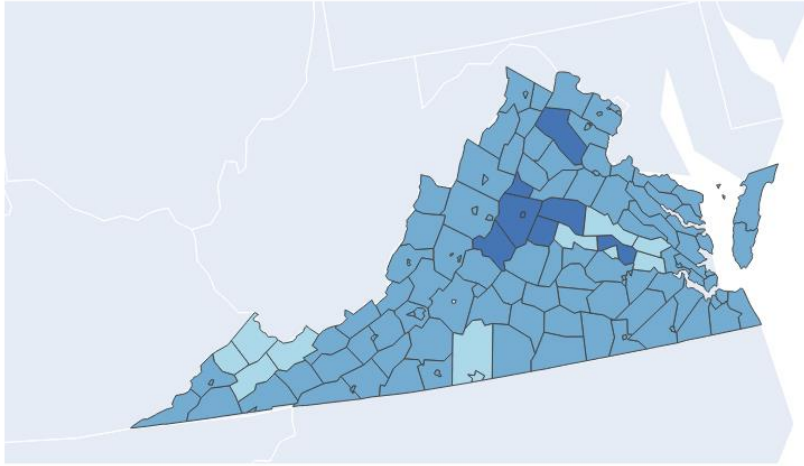
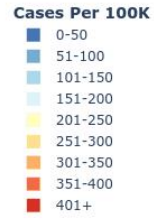
Projections by District



Daily confirmed cases by rate (per 100K) District (grey with 7-day average in black) with simulation colored by scenario

Adaptive

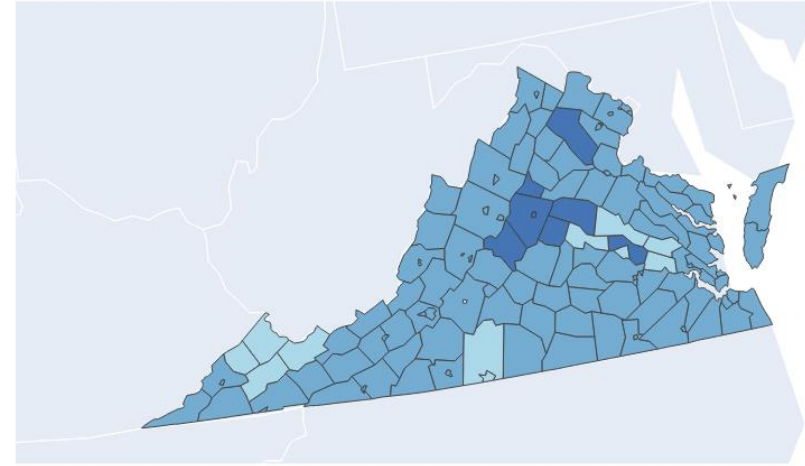
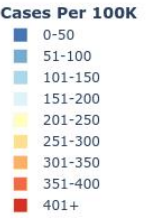
Weekly Projections (Adaptive) 28-Sep-2022



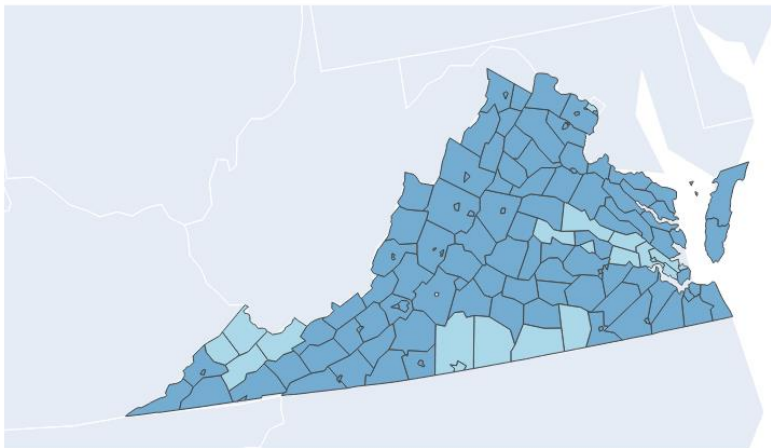
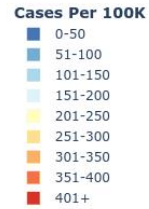
Adaptive

VariantX

Weekly Projections (Adaptive-VariantX) 28-Sep-2022

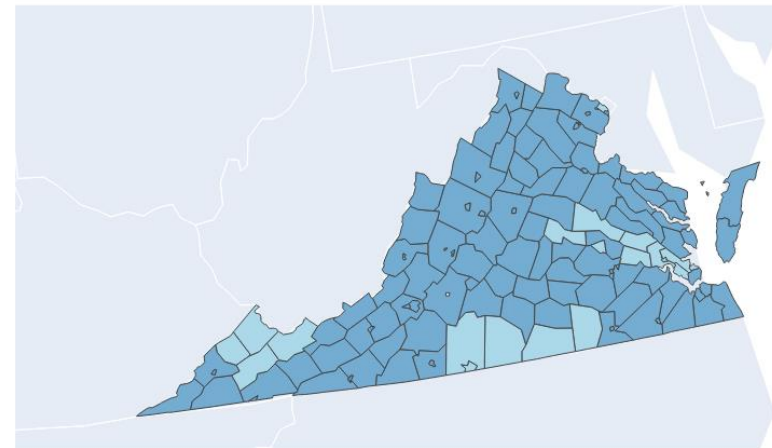
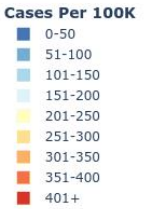


Weekly Projections (Adaptive-FallWinter) 28-Sep-2022



Adaptive-Fall-Winter

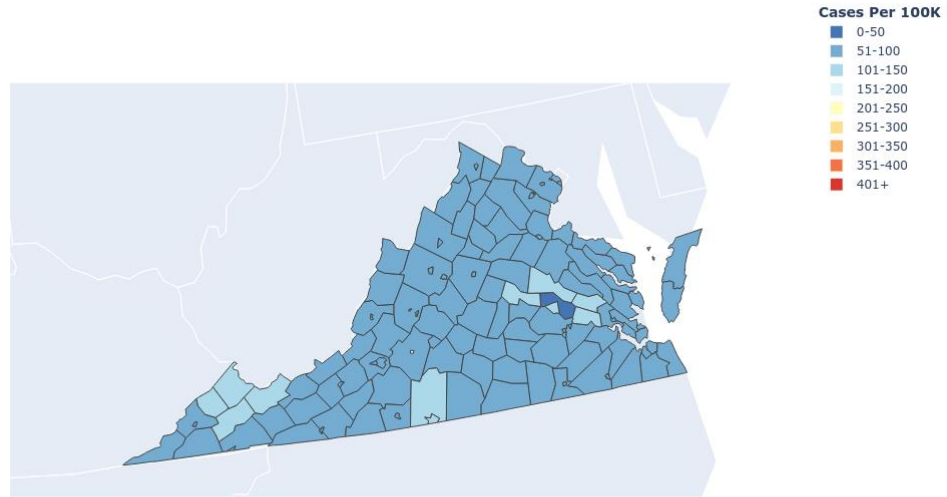
Weekly Projections (Adaptive-VariantX-FallWinter) 28-Sep-2022



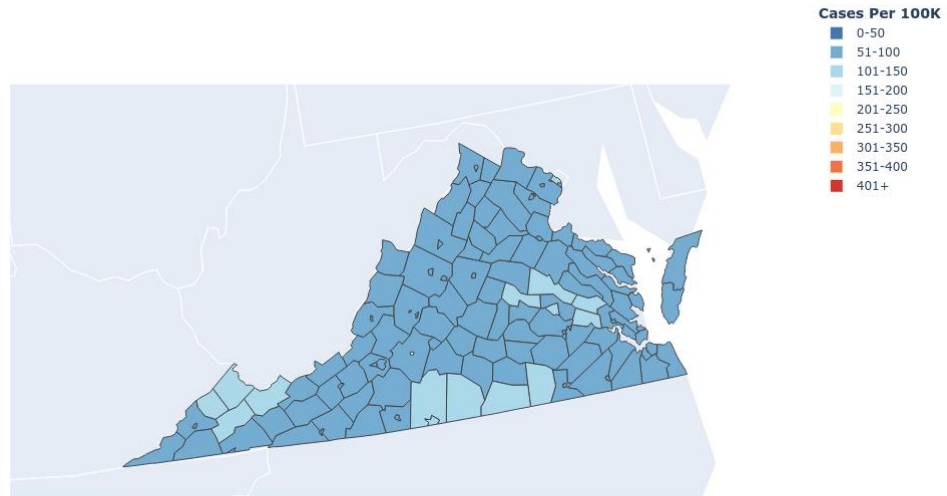
Y
Y

Impact of Optimistic vs. Pessimistic Booster Distribution

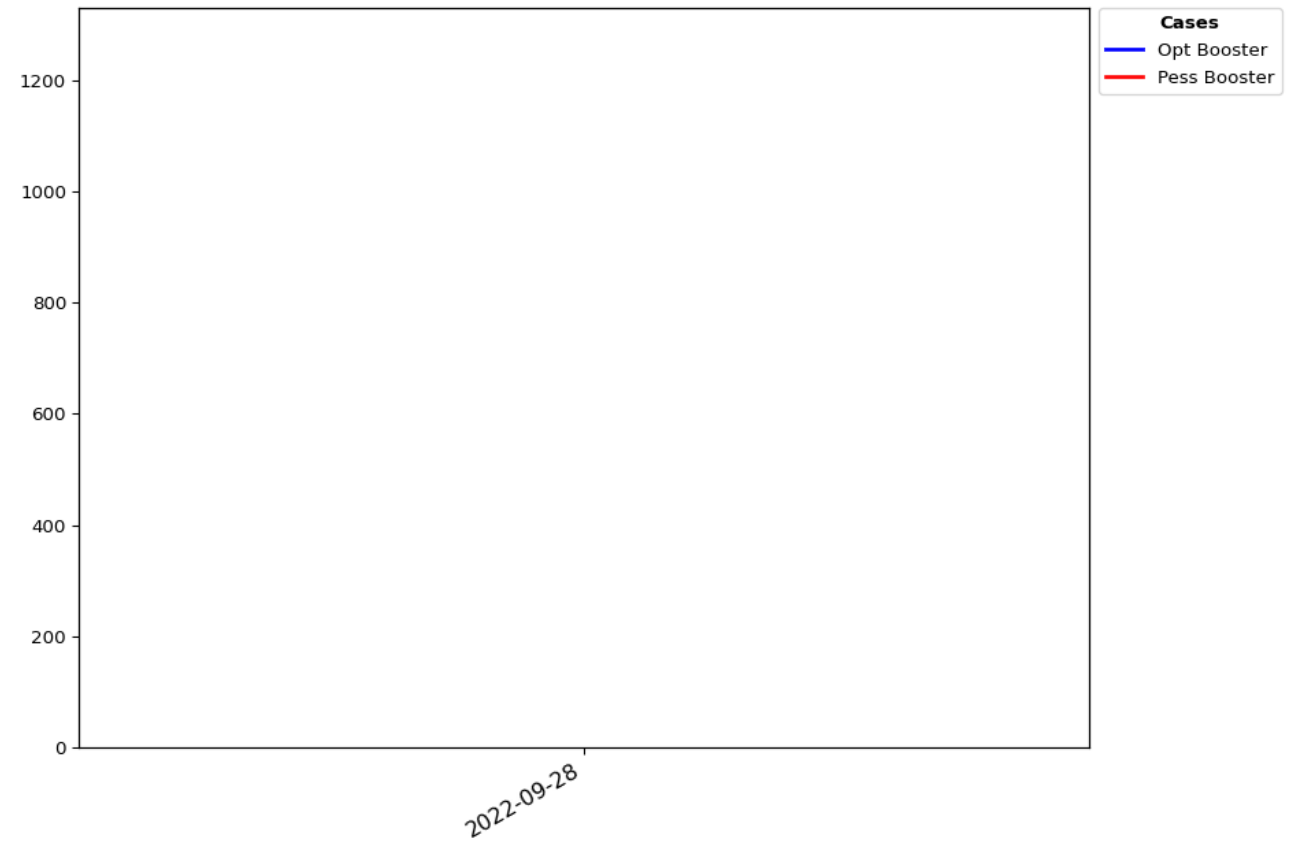
Weekly Projections (Optimistic Booster) 28-Sep-2022



Weekly Projections (Pessimistic Booster) 28-Sep-2022



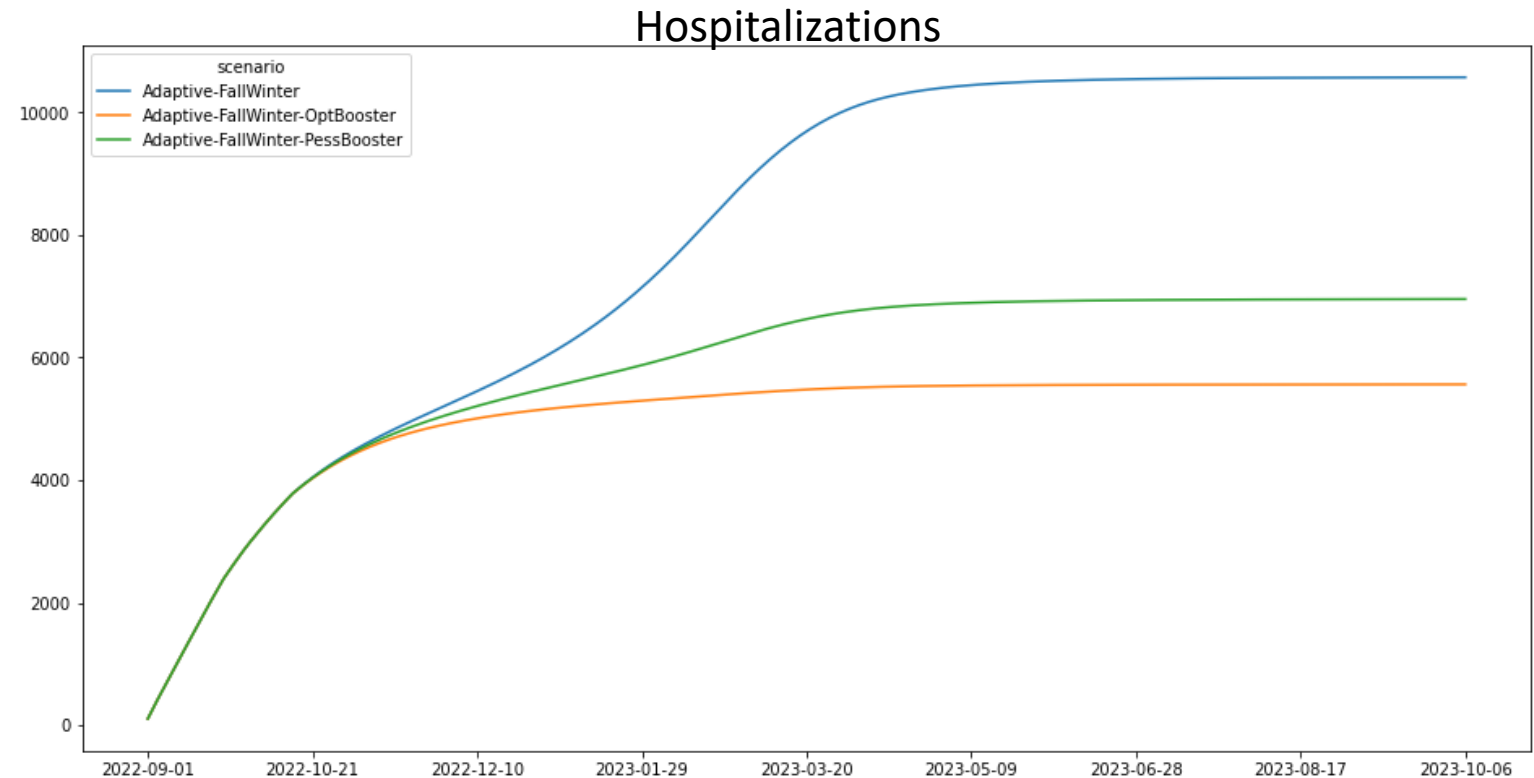
Cases for Optimistic vs. Pessimistic Boosters 28-Sep-2022



Booster Campaign Coverage has impact on future hospitalizations

Booster Campaign can significantly limit future hospitalizations and severe outcomes

- Reduction in future hospitalizations (3600 to 5K) through Spring 2023

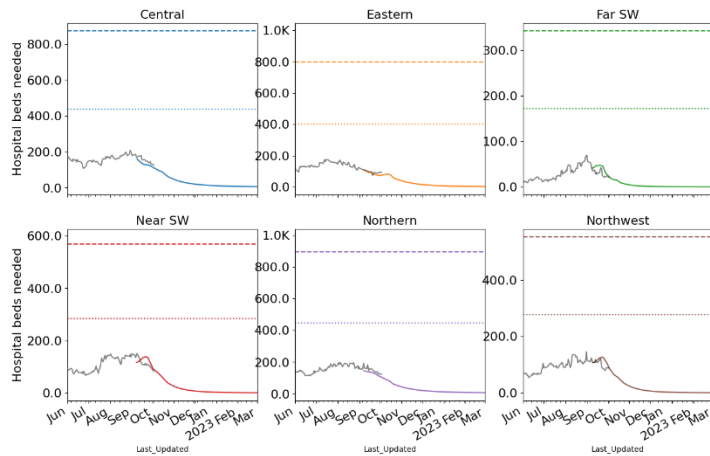


Hospital Demand and Bed Capacity by Region

Capacities by Region

COVID-19 capacity ranges from 80% (dots) to 120% (dash) of total beds

Adaptive



Length of Stay Estimates

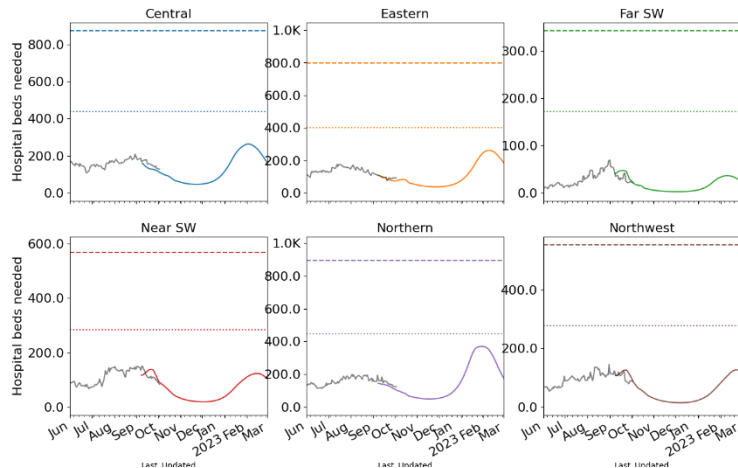
| | |
|--------------|---|
| Central | 8 |
| Eastern | 6 |
| Far SW | 4 |
| Near SW | 9 |
| Northern | 5 |
| Northwestern | 9 |

Length of Stay more variable with Omicron, occupancy projections may vary as a result, ad-hoc estimation performed per region

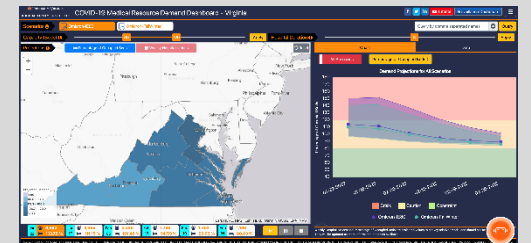
Estimated LOS shortened slightly to better fit observed data

Projections show continued declines and with expanded capacities and adjusted length of stay, no capacities exceeded

Adaptive – VariantX & Fall Winter



Interactive Dashboard with regional projections



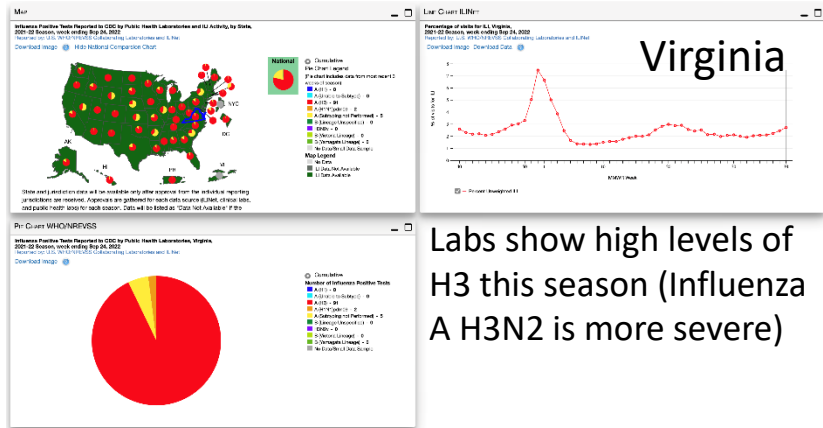
<https://nssac.bii.virginia.edu/covid-19/vmrddash/>

Current Influenza Hospitalization Forecast

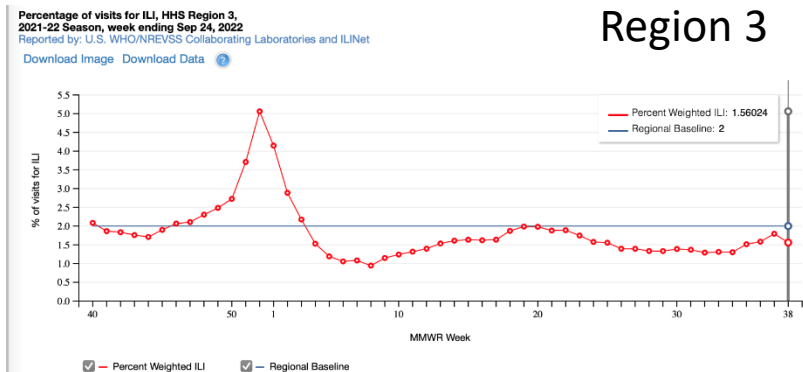
Statistical models for submitting to CDC FluSight forecasting challenge

- Similar to COVID-19 case forecasts, uses a variety of statistical and ML approaches to forecast weekly hospital admissions for the next 4 weeks for all states in the US

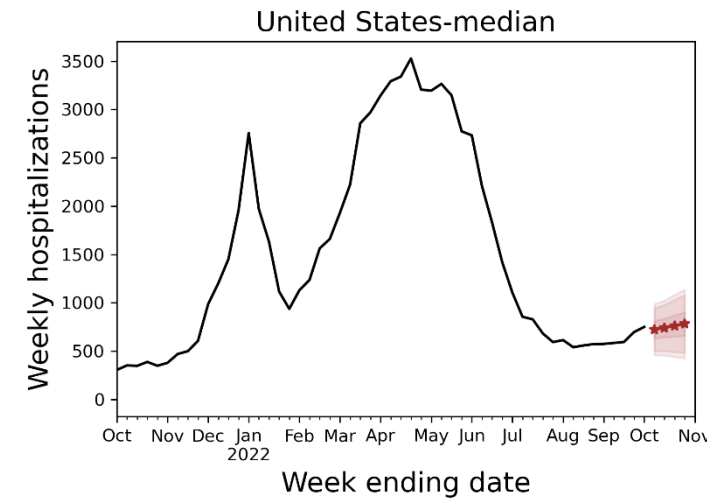
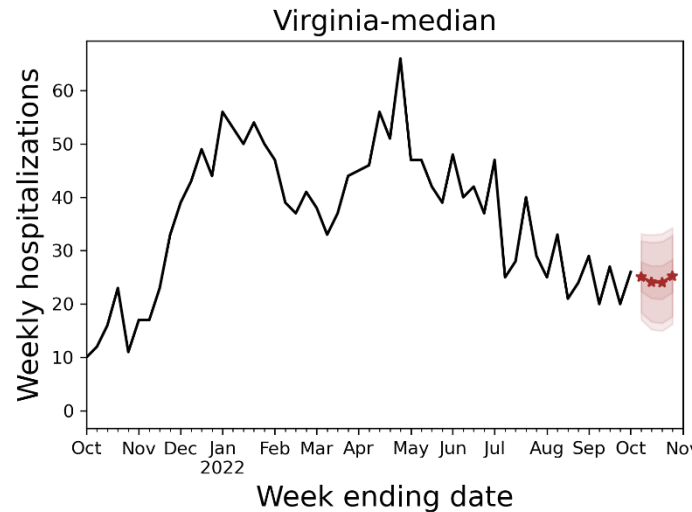
Influenza-like Illness Activity



Labs show high levels of H3 this season (Influenza A H3N2 is more severe)



Hospital Admissions for Influenza and Forecast for next 4 weeks (UVA ensemble)



Initial forecasts have wide uncertainty due to noisiness in data due to low numbers of hospitalizations

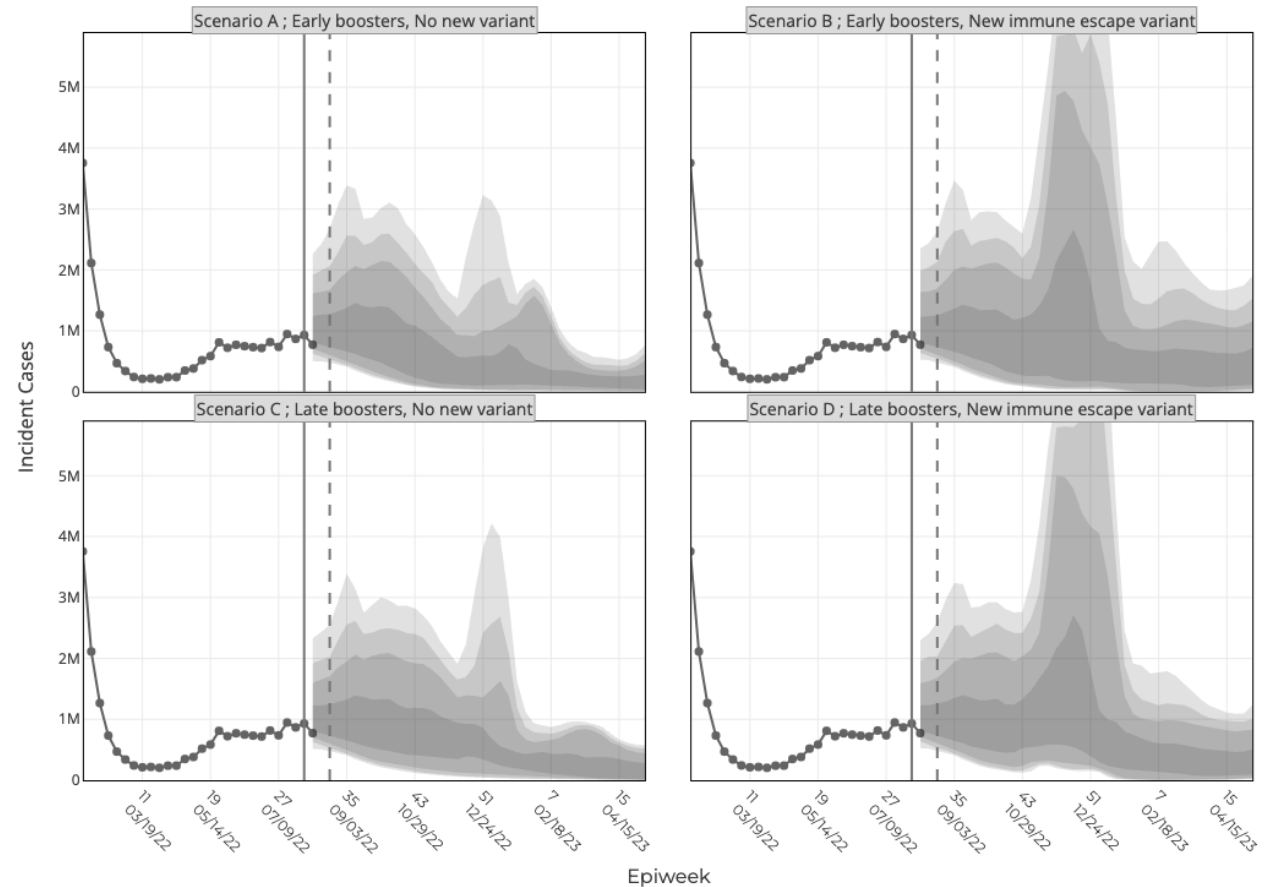
Scenario Modeling Hub – COVID-19 (Round 15)

Collaboration of multiple academic teams to provide national and state-by-state level projections for 4 aligned scenarios

- Round 15 results published
 - Scenarios: Test benefits of reformulated fall boosters w/ and w/out a new variant
 - Timing of reformulated boosters is one of the axes

<https://covid19scenariomodelinghub.org/viz.html>

Projected Incident Cases by Epidemiological Week and by Scenario for Round 15 - US
(- Projection Epiweek; -- Current Week)



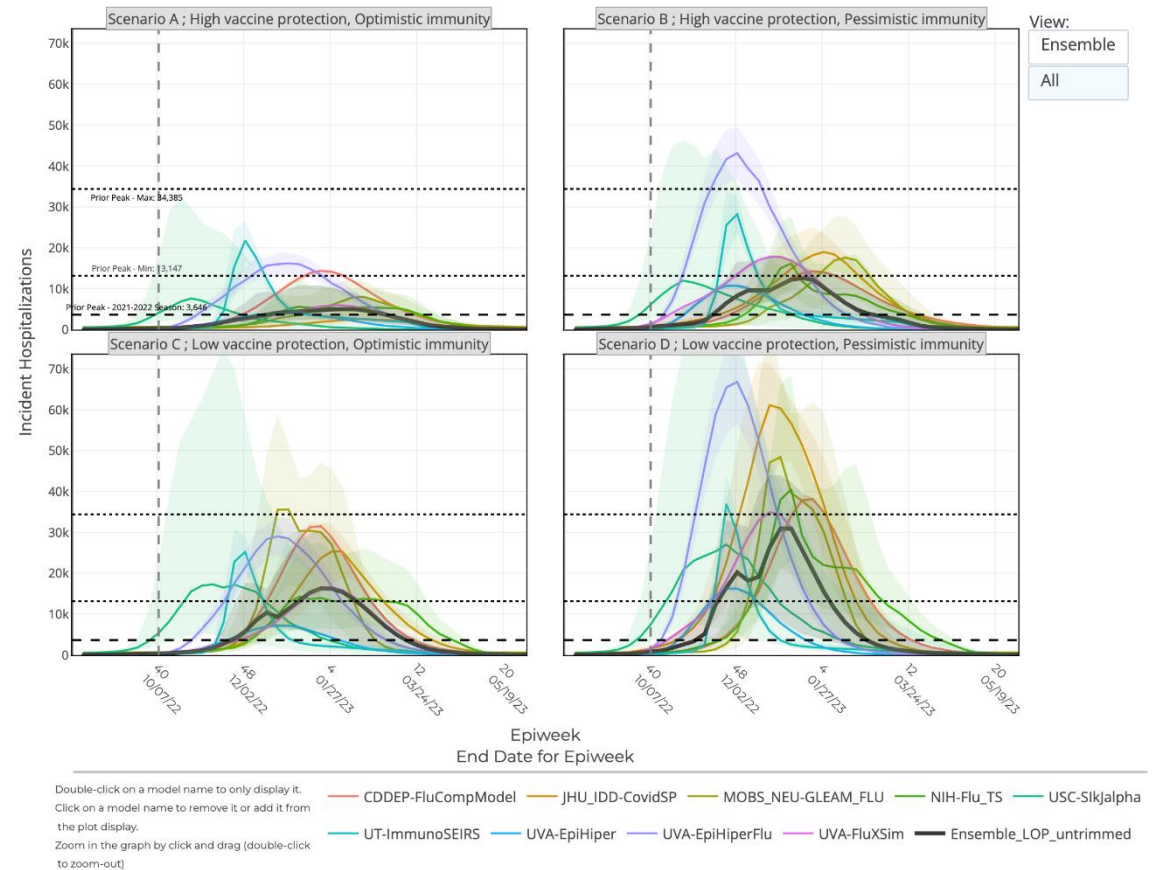
Scenario Modeling Hub – Influenza (Round 1)

Collaboration of multiple academic teams to provide national and state-by-state level projections for 4 aligned scenarios

- Round 1 results recently published
 - Impact of missed flu seasons on pre-season immunity
 - Testing different seasonal vaccine coverage and efficacy
 - Projected from Aug 14th 2022
- High degree of uncertainty as previous 2 seasons have been irregular and there is still limited data for this season available
- Demonstrates importance of good vaccine coverage especially if previous immunity is weak

<https://fluscenariomodelinghub.org/viz.html>

Projected Incident Hospitalizations by Epidemiological Week and by Scenario for Round 1 - US
(- Projection Epiweek; -- Current Week)



Key Takeaways

Projecting future cases precisely is impossible and unnecessary.

Even without perfect projections, we can confidently draw conclusions:

- **Case rates continue their decline, hospitalizations continue steady decline**
- VA weekly case rate continues decline to 99/100K from 109/100K
 - US weekly case rate is down considerably as well at 89/100K from 105/100K
 - VA hospital occupancy (rolling 7 day mean of 545 down from 599 a week ago) has continued to decline
- Projections anticipate continued declines in cases as well as hospitalizations
- Potential for rebounds due to seasonal forces and/or novel sub-variants in the Fall remains
- Model updates:
 - Maintained Booster Scenarios by slowing down the rate of vaccination adopting the rate of 3rd dose rollout
 - Current monitoring still not finding a definite candidate for Variant X, though BQ.1.1 or BA.2.75.2 remain likely, 50% prevalence pushed back to Nov 15th

The situation continues to change. Models continue to be updated regularly.

References

Venkatramanan, S., et al. "Optimizing spatial allocation of seasonal influenza vaccine under temporal constraints." *PLoS Computational Biology* 15.9 (2019): e1007111.

Arindam Fadikar, Dave Higdon, Jiangzhuo Chen, Bryan Lewis, Srinivasan Venkatramanan, and Madhav Marathe. Calibrating a stochastic, agent-based model using quantile-based emulation. *SIAM/ASA Journal on Uncertainty Quantification*, 6(4):1685–1706, 2018.

Adiga, Aniruddha, Srinivasan Venkatramanan, Akhil Peddireddy, et al. "Evaluating the impact of international airline suspensions on COVID-19 direct importation risk." *medRxiv* (2020)

NSSAC. PatchSim: Code for simulating the metapopulation SEIR model. <https://github.com/NSSAC/PatchSim>

Virginia Department of Health. COVID-19 in Virginia. <http://www.vdh.virginia.gov/coronavirus/>

Biocomplexity Institute. COVID-19 Surveillance Dashboard. <https://nssac.bii.virginia.edu/covid-19/dashboard/>

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Questions?

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