

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

# Estimation of COVID-19 Impact in Virginia

October 7<sup>th</sup>, 2020

(data current to October 5<sup>th</sup>)

Biocomplexity Institute Technical report: TR 2020-122



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[biocomplexity.virginia.edu](http://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 infections through December
  - 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:

- **Holding steady with declines outpacing growth.**
- VA weekly incidence (9.8/100K) continues to decline and now well below the national average (16/100K) which has been climbing, fueled by growth in the Plains and Mountain West.
- Projections are mostly downward across commonwealth.
- Recent updates:
  - Adaptive Fitting projection process has been streamlined.
  - Planning Scenarios moved to Nov 1st.
- The situation is changing rapidly. Models will be updated regularly.



# Situation Assessment

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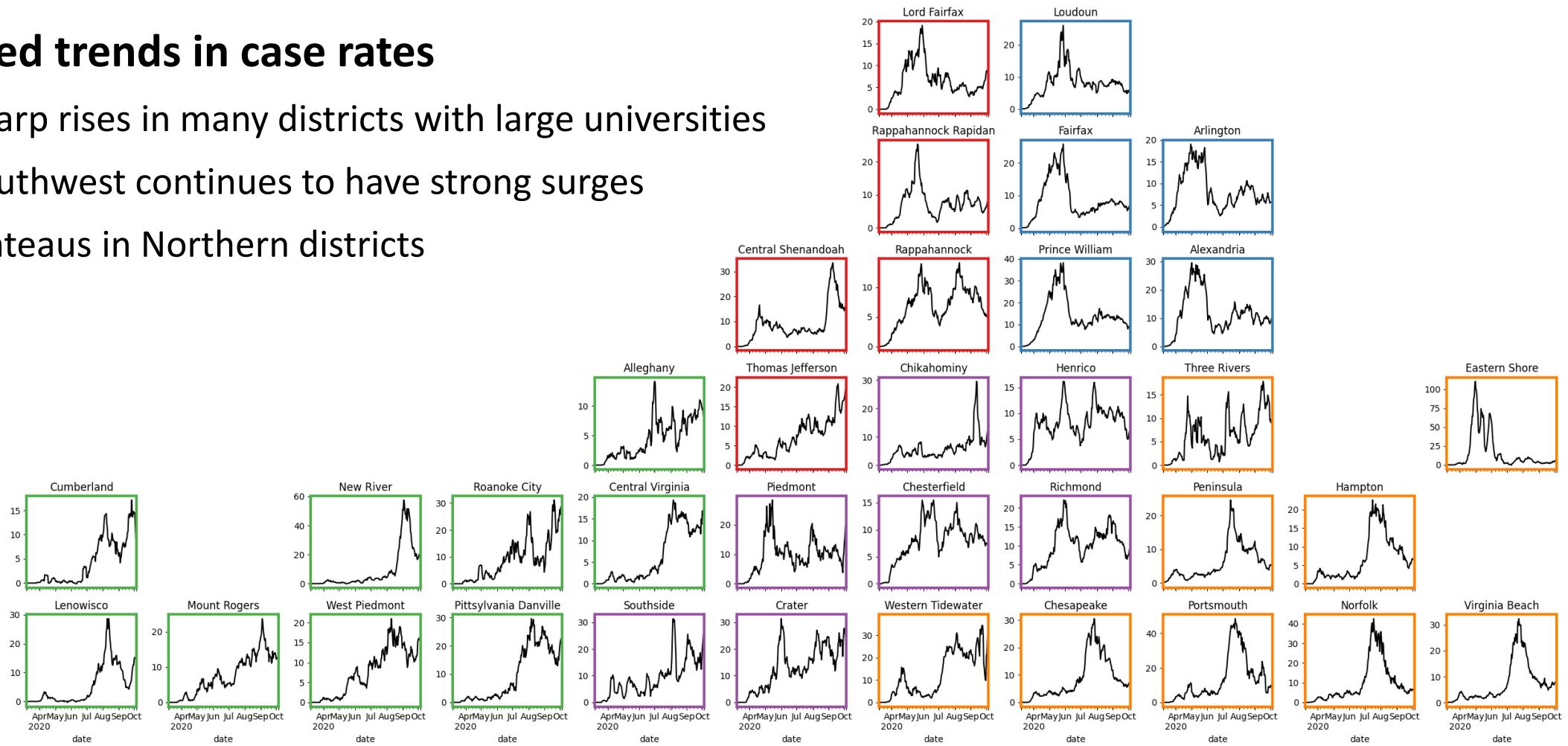
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# Case Rate (per 100k) by VDH District

## Mixed trends in case rates

- Sharp rises in many districts with large universities
- Southwest continues to have strong surges
- Plateaus in Northern districts

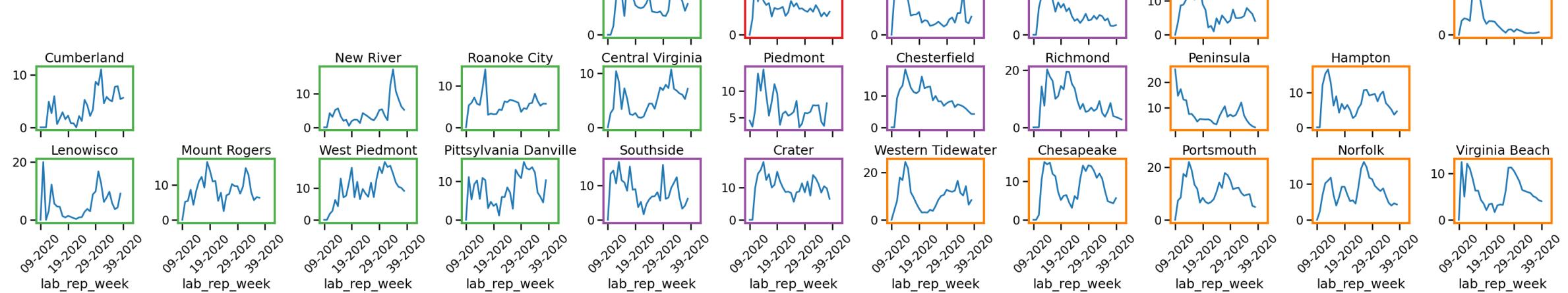


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# Test Positivity by VDH District

## Weekly changes in test positivity by district

- Most districts moving towards lower overall percents
- Areas with most growth also showing high and increasing test positivity, especially in Southwest



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# District Trajectories

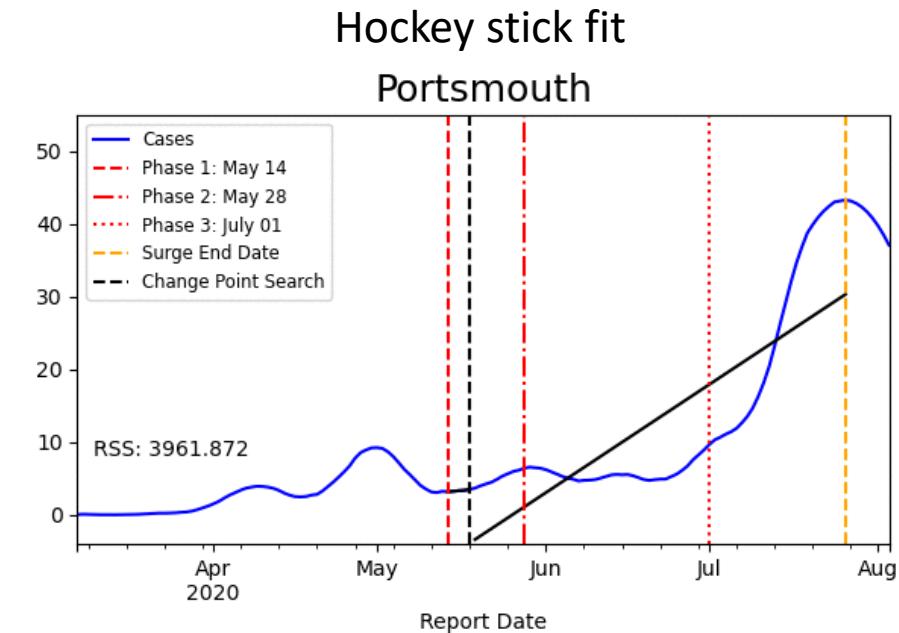
Hockey stick fit used to describe recent growth patterns based on recent trends from last peak or inflection points (based on smoothed case rates per 100k)

**Declining:** Sustained decreases following a recent peak

**Plateau:** Steady level with minimal trend up or down

**Slow Growth:** Sustained growth not rapid enough to be considered a Surge

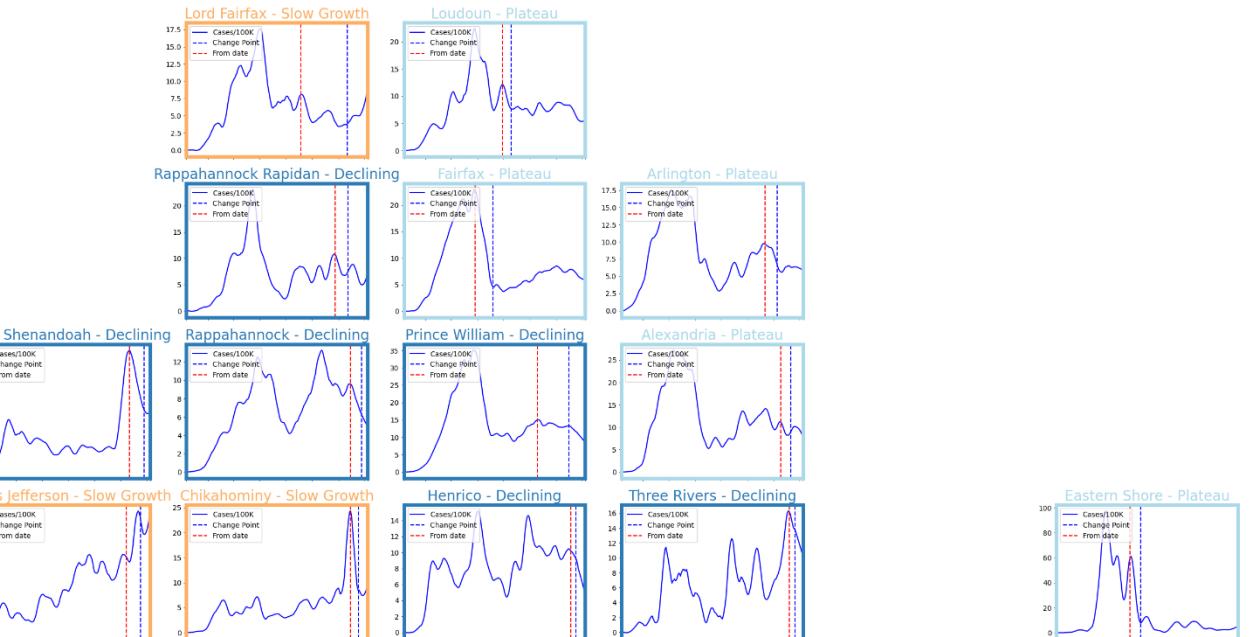
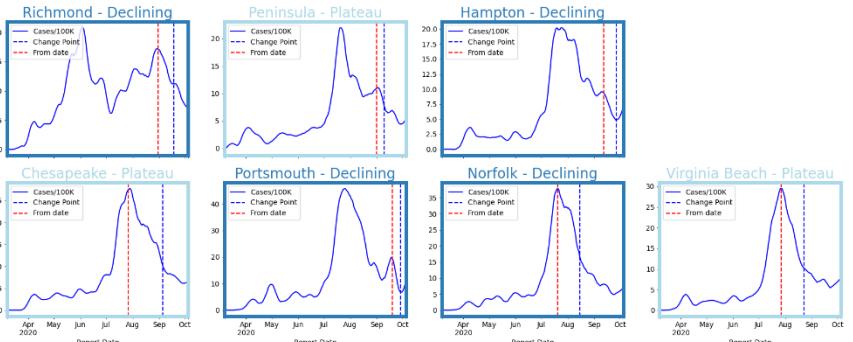
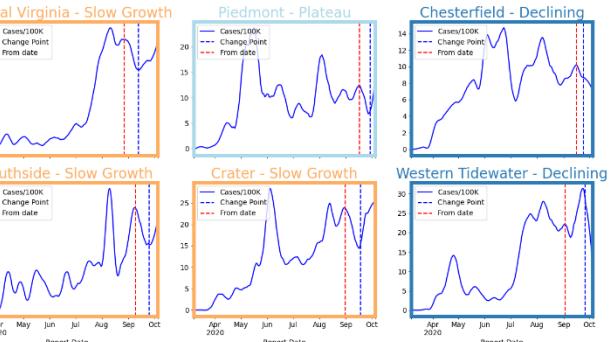
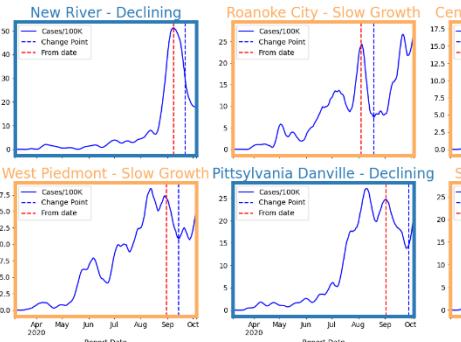
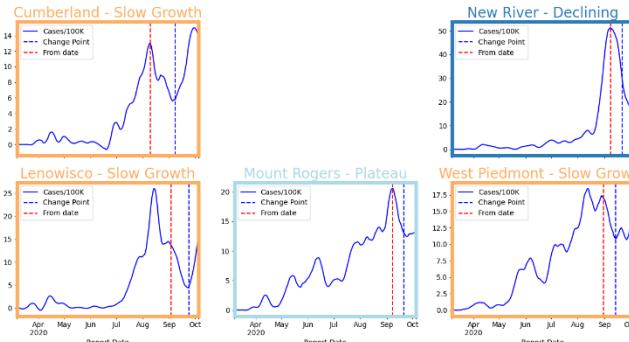
**In Surge:** Currently experiencing sustained rapid growth and exceeds recent inflection points



Status	# Districts (last week)
Declining	14 (17)
Plateau	10 (11)
Slow Growth	11 (6)
In Surge	0 (1)

# District Trajectories – Declines outpace Growth

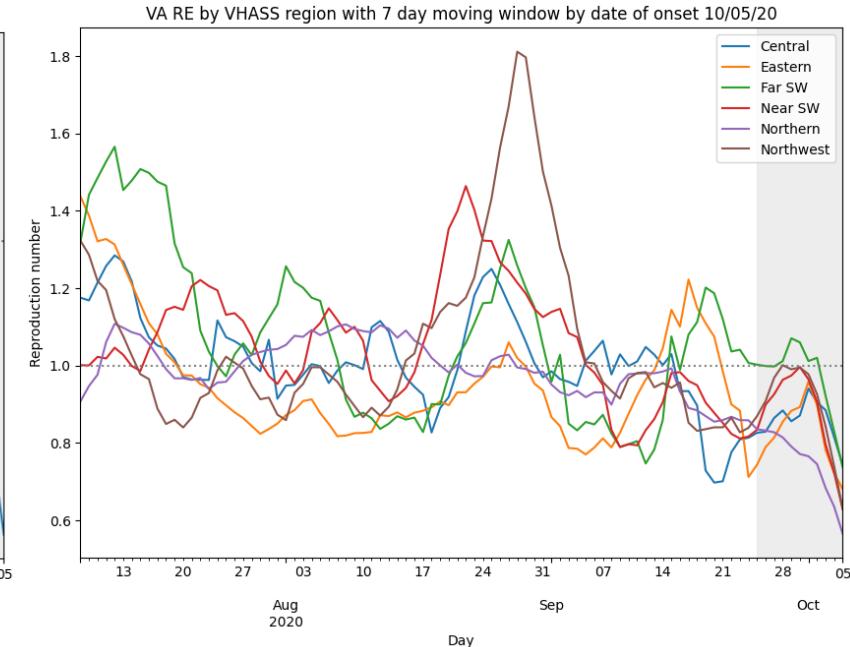
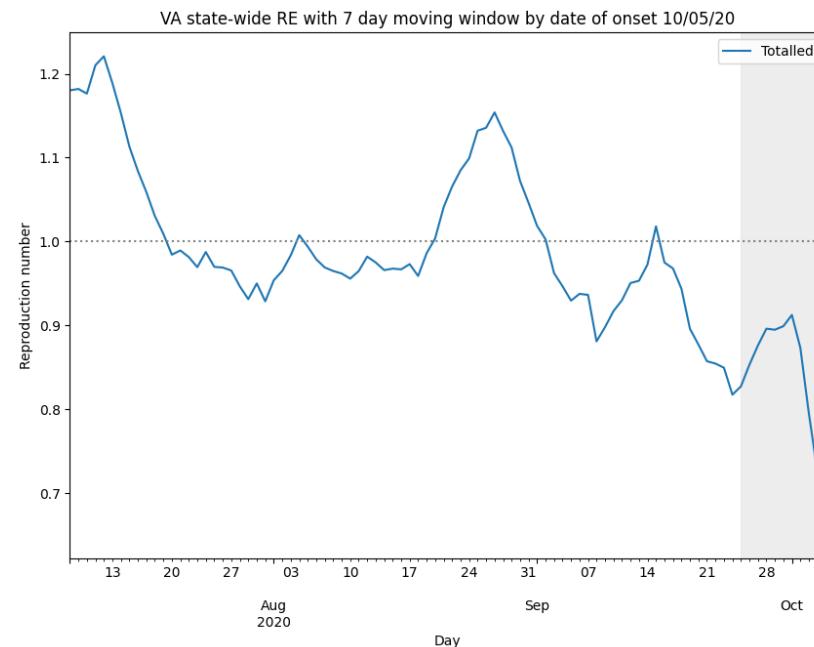
Status	# Districts (last week)
Declining	14 (17)
Plateau	10 (11)
Slow Growth	11 (6)
In Surge	0 (1)



# Estimating Daily Reproductive Number

September 26<sup>th</sup> Estimates

Region	Current $R_e$	Diff Last Week
State-wide	0.853	-0.015
Central	0.829	0.106
Eastern	0.790	-0.298
Far SW	0.999	-0.185
Near SW	0.899	0.041
Northern	0.832	0.008
Northwest	0.909	0.096



## Methodology

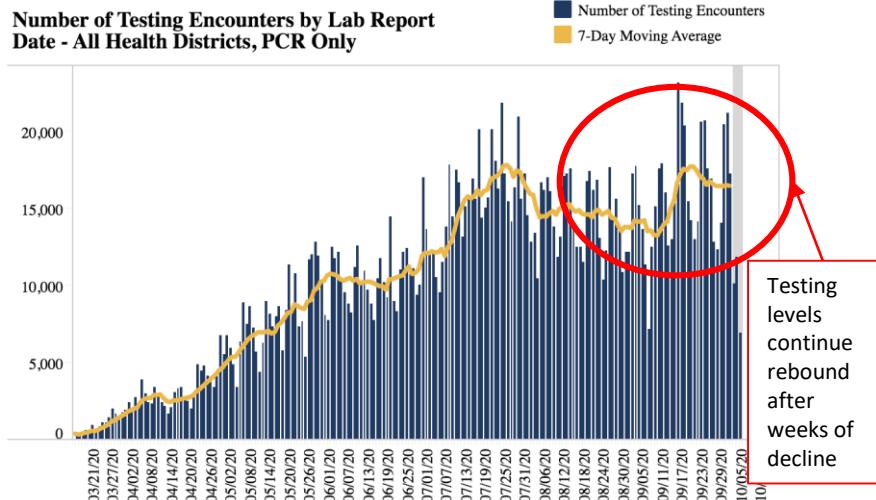
- Wallinga-Teunis method (EpiEstim<sup>1</sup>) for cases by date of onset
- Serial interval: 6 days (2 day std dev)
- Recent estimates may be unstable due to backfill

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>

# Changes in Case Detection

Timeframe (weeks)	Mean days	% difference from overall mean
April (13-16)	8.54	43%
May (17-21)	5.64	-5%
June (22-25)	5.9	-1%
July (26-30)	6.29	5%
Aug (31-34)	4.77	-20%
Sept (35-37)	4.19	-30%
Overall (13-37)	5.96	0%

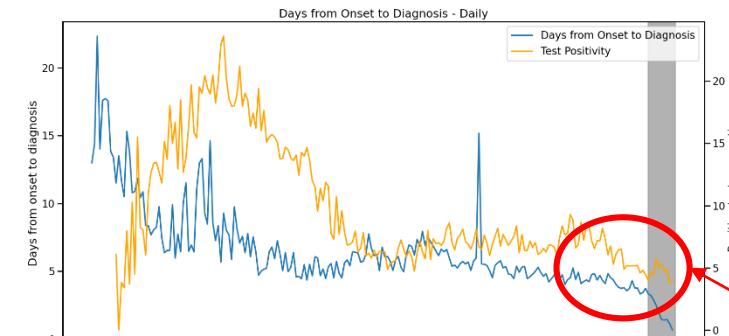
Testing Encounters and test positivity have steadied and increased



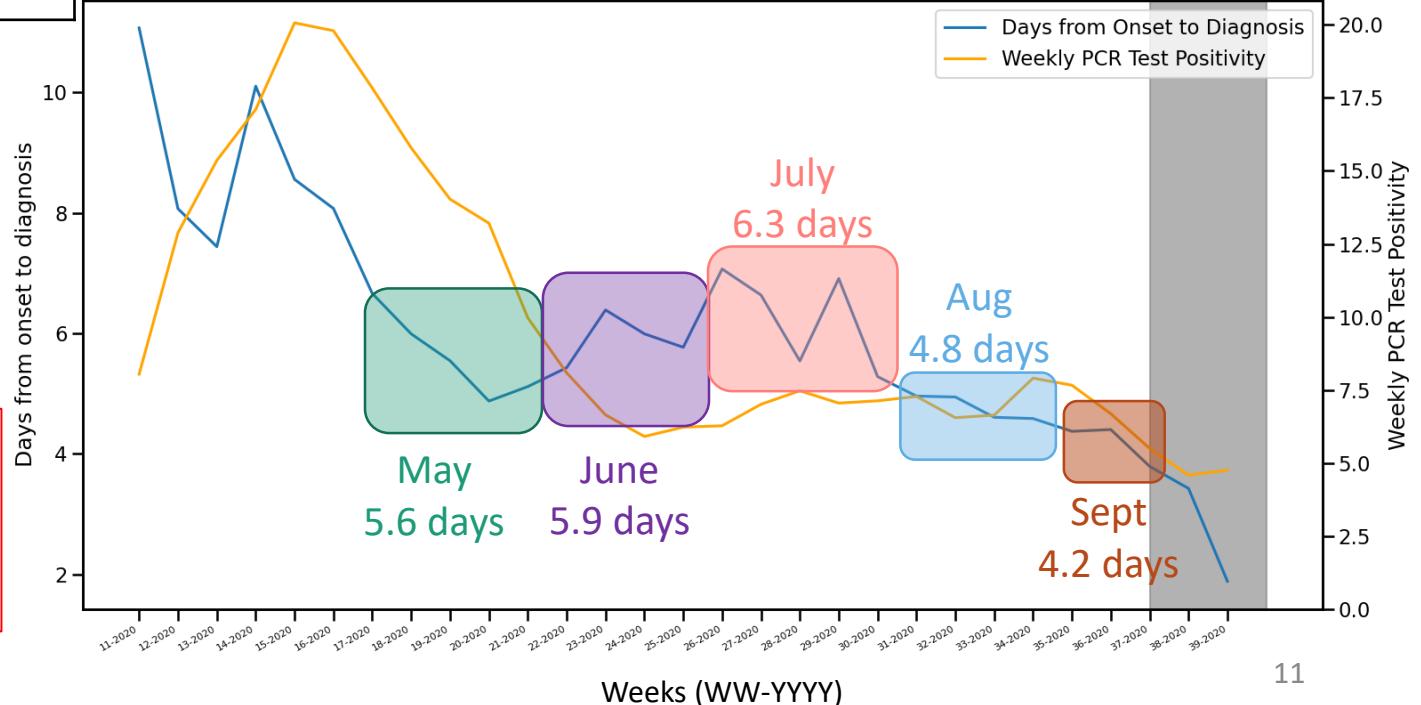
7-Oct-20

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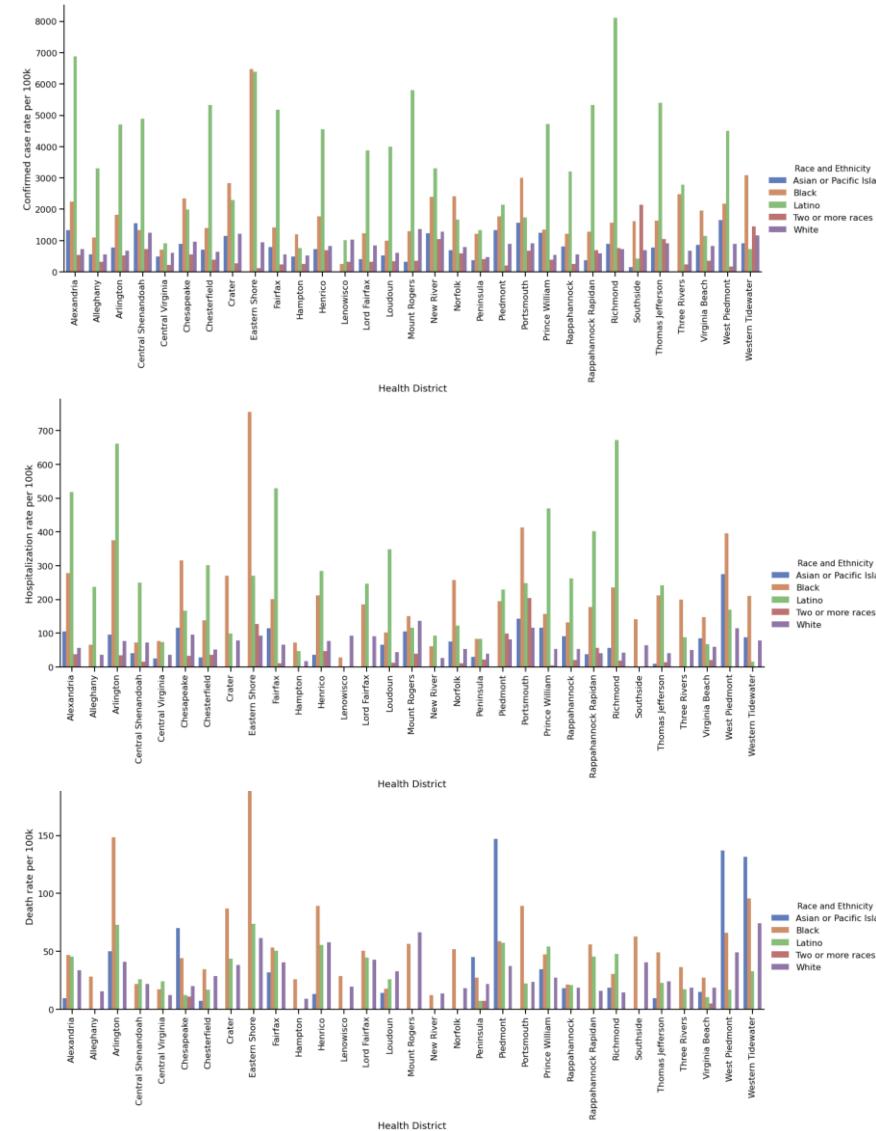
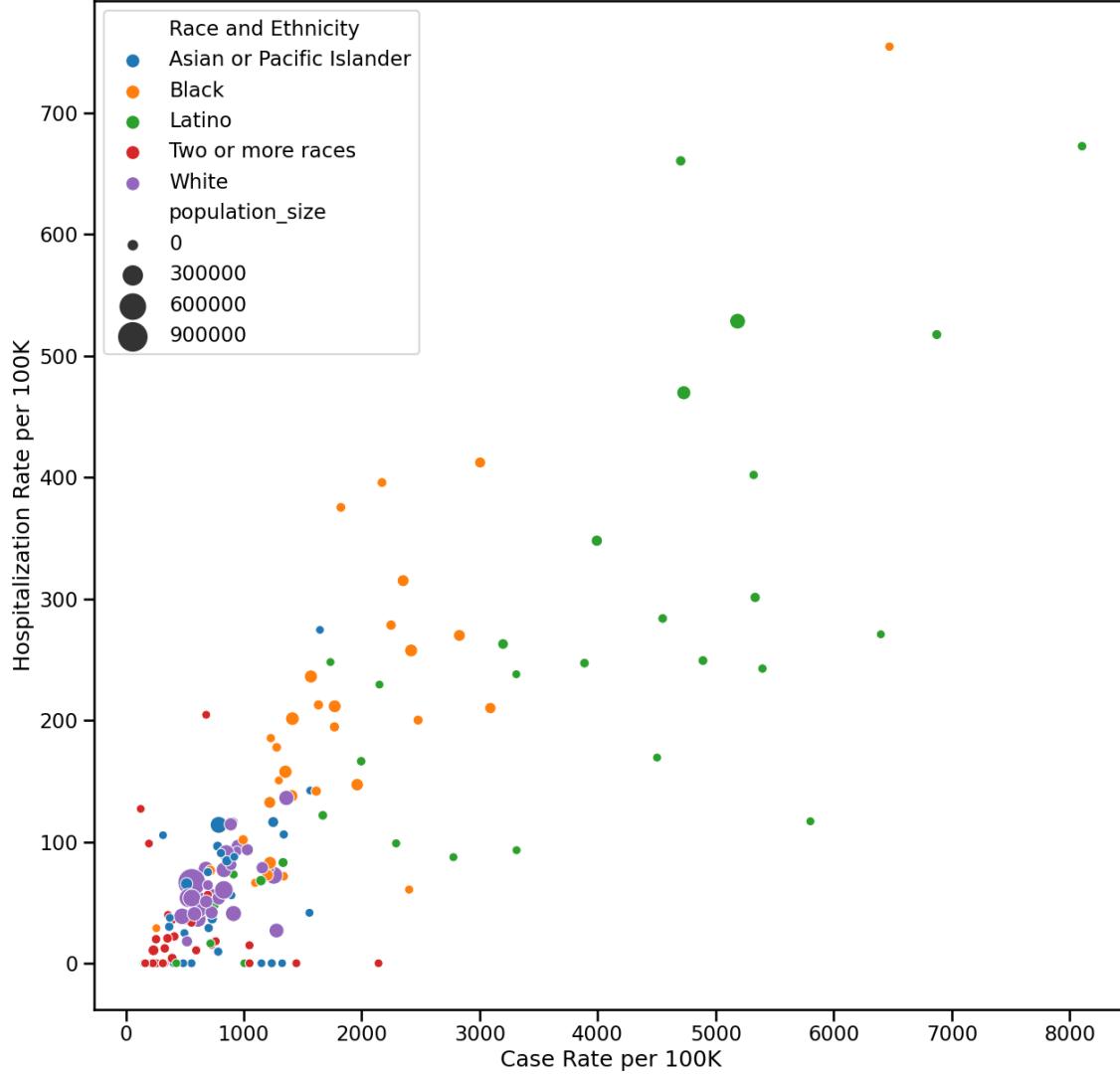
## Test positivity vs. Onset to Diagnosis



## Days from Onset to Diagnosis and Test Positivity - Weekly



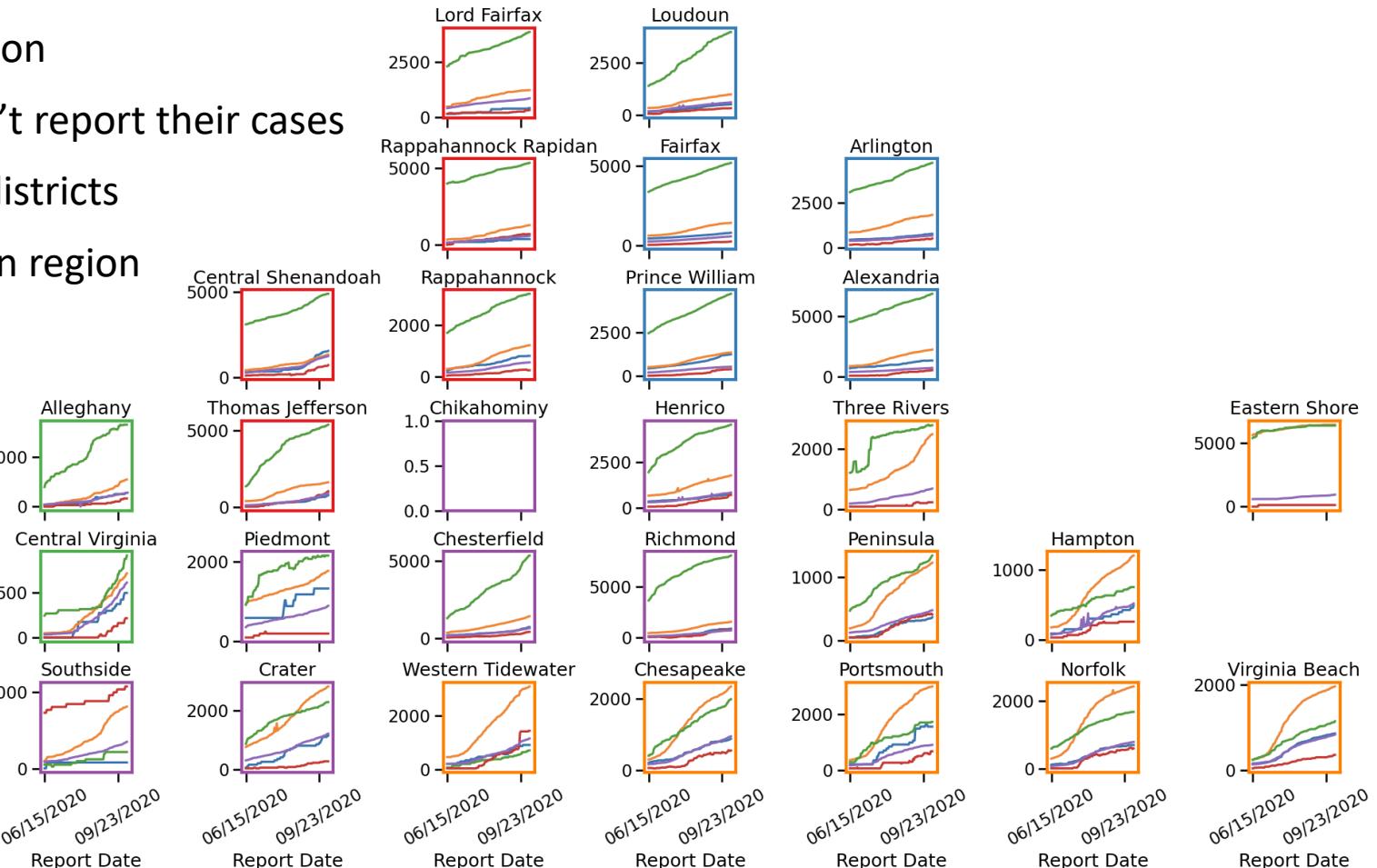
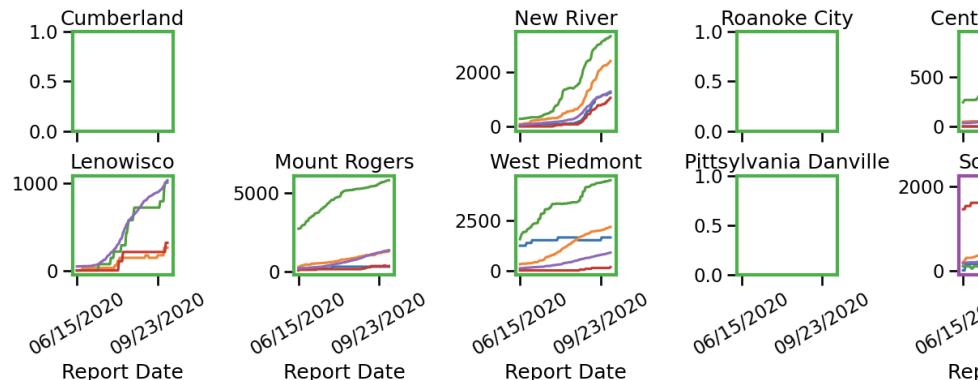
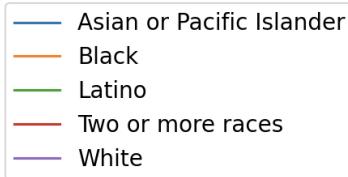
# Race and Ethnicity cases per 100K



# Race and Ethnicity Attack Rates (per 100K)

## Cumulative Race and Ethnicity Attack Rates (per 100k)

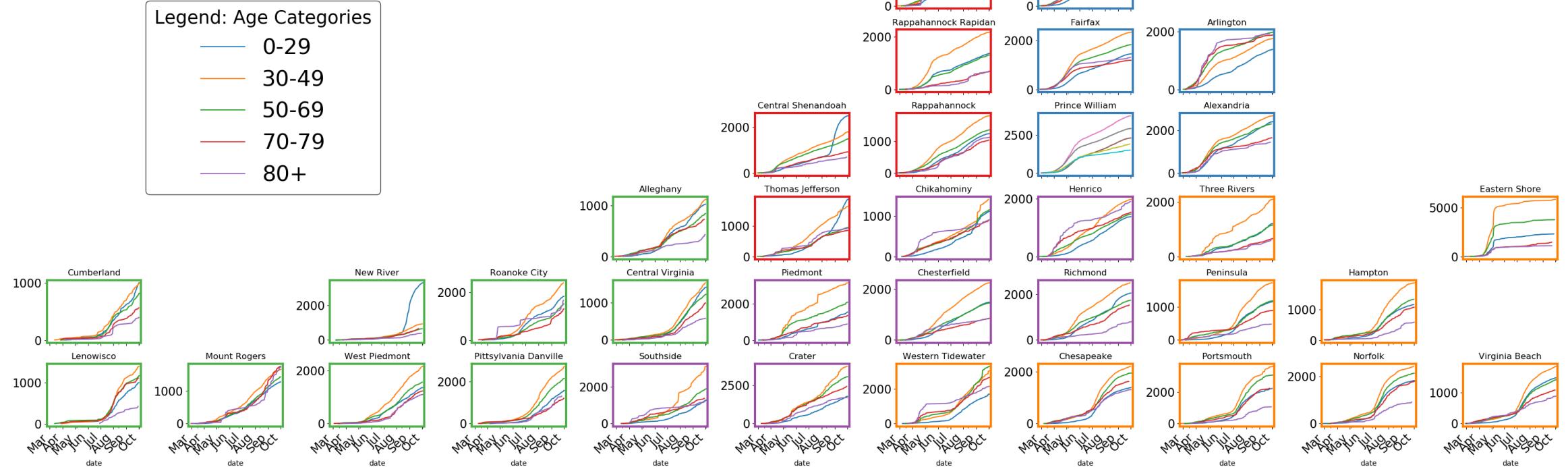
- Case Rates in different groups vary by location
- Some districts have small numbers and don't report their cases
- Latinx population has highest rate in most districts
- Black population has highest rates in Eastern region



# Age-Specific Attack Rates (per 100K)

## Cumulative Age-specific Attack Rates (per 100k)

- Younger age groups outpace older in many districts
- Some districts with previous surge in young cases now show a spillover from 0-29 to 30-49 (eg. Alleghany)

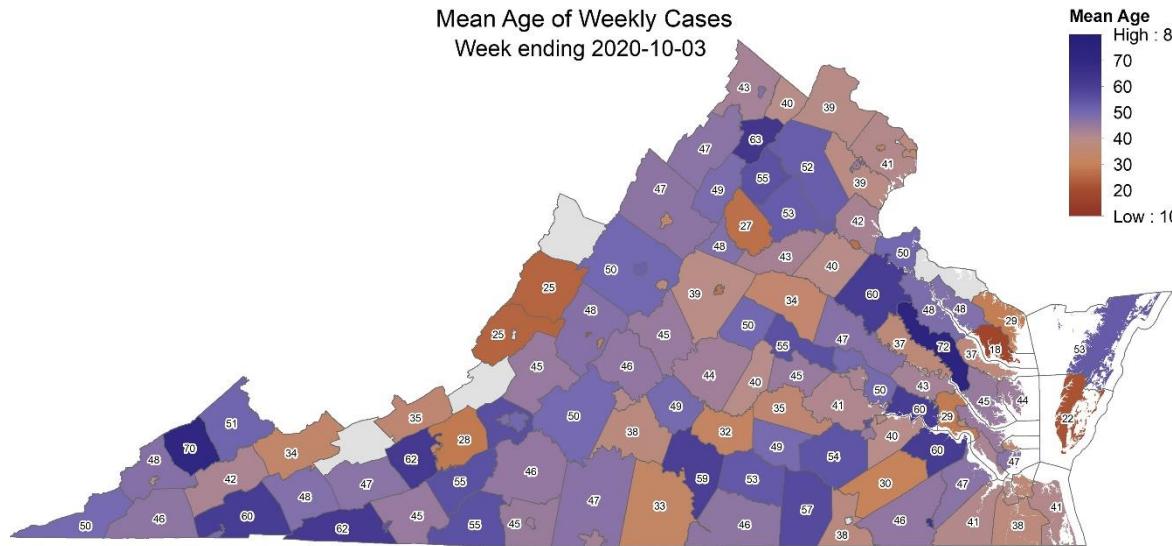


# Age-Specific Case Prevalence

## How different is this from the Population?

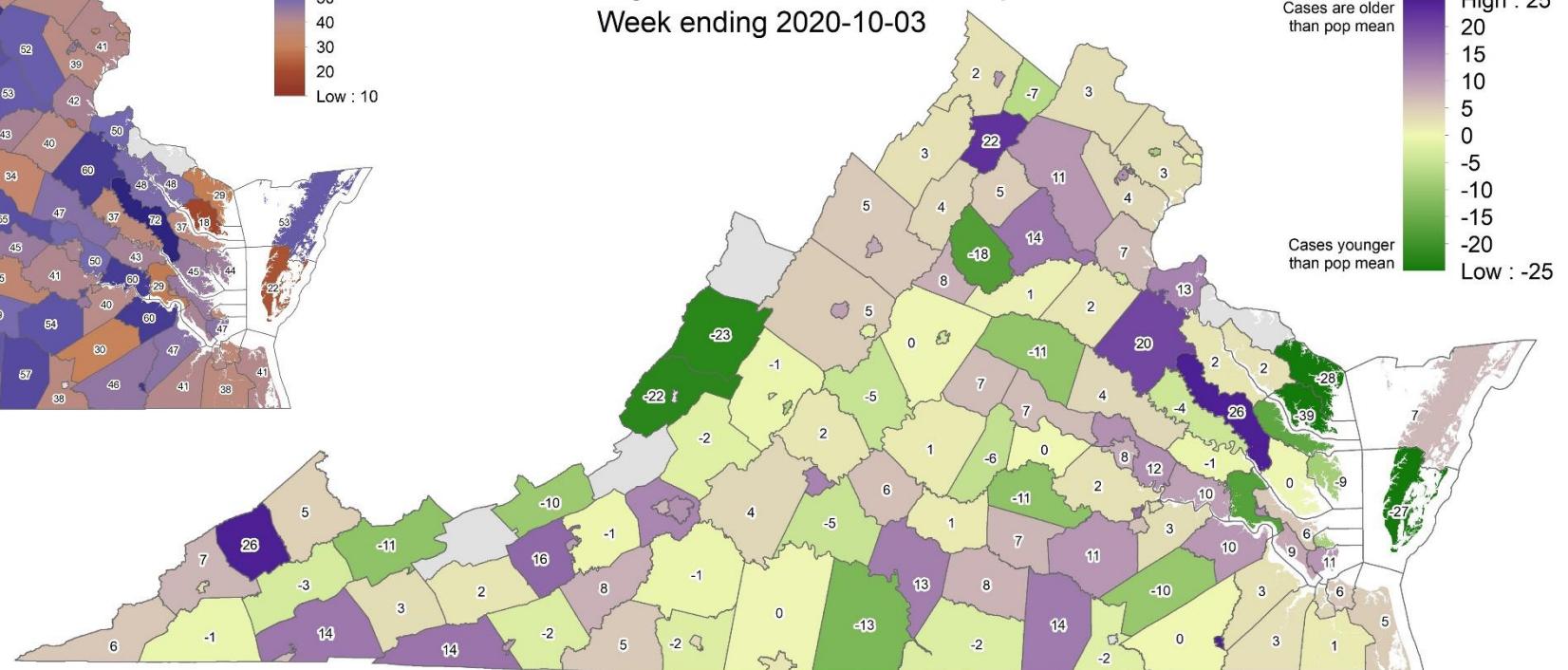
Difference in mean age of cases vs. population as a whole

Purple = Cases are older than pop; Green = Cases are younger than pop



## Difference in Age between Cases and Population

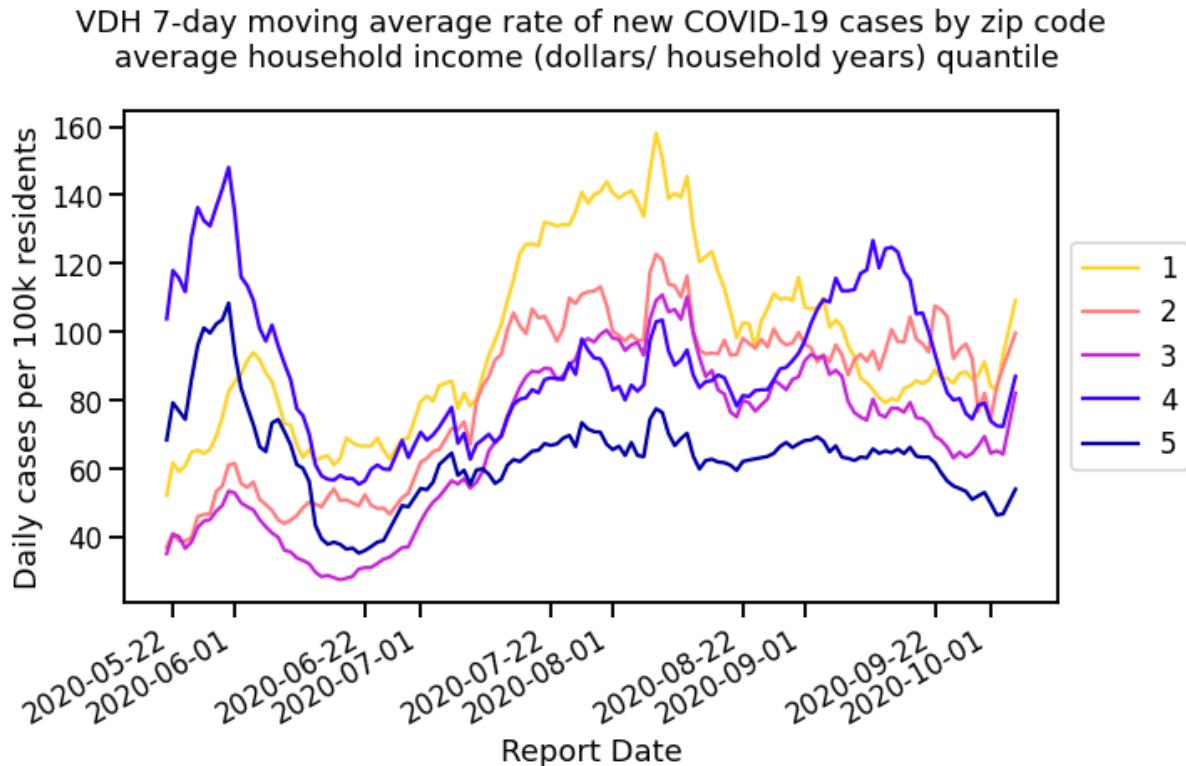
Week ending 2020-10-03



## What is the average age of the cases by county?

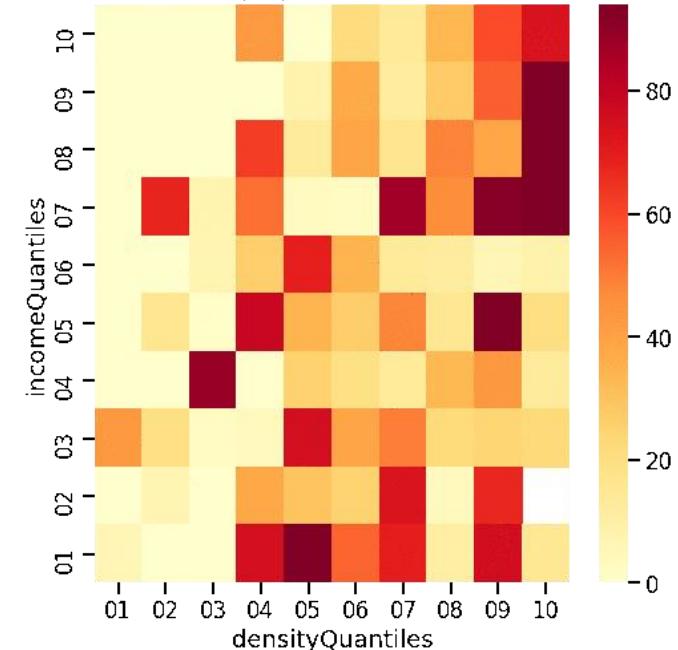
Younger cases in Northern VA, Far SW, Tidewater, and around universities

# Impact across Density and Income



Shift back to higher income zip codes partially driven by surges in areas surrounding universities

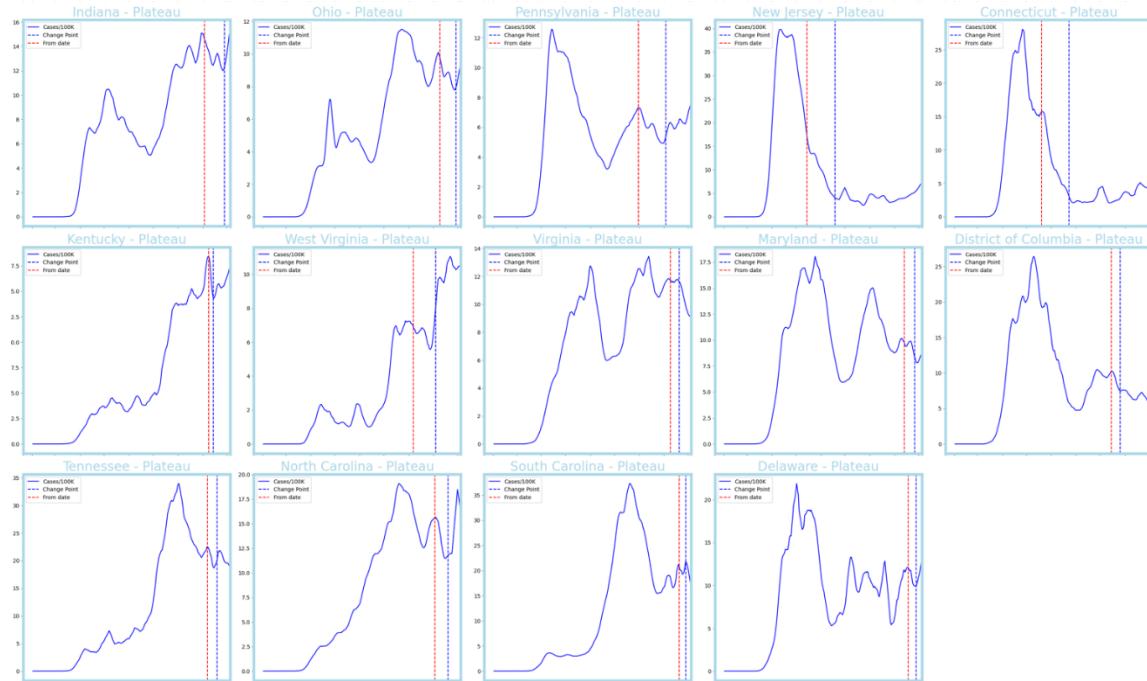
VDH mean cases per 100k by zip code population density (person/ sq mile)  
and average household income (dollars/ household years) quantiles  
05/15/20 - 05/21/20



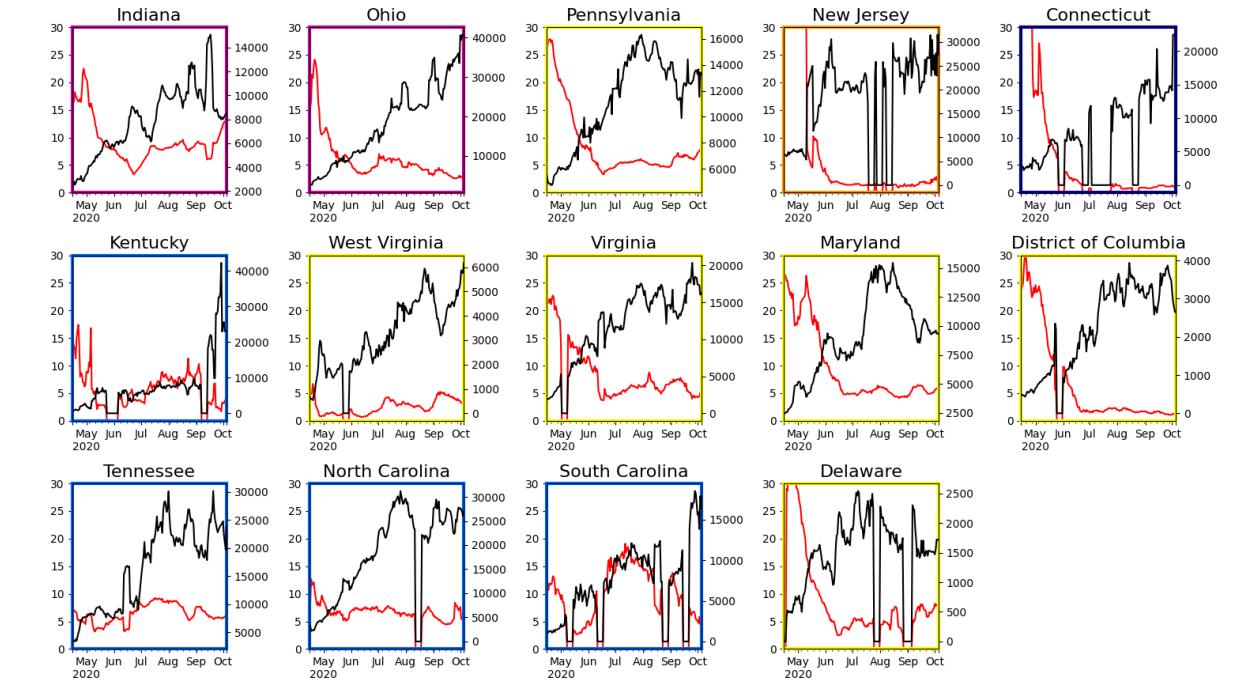
Can see the evolution from denser and wealthier zip codes to poorer and less dense zip codes, then recently back to denser wealthier zip codes

# Other State Comparisons

## Trajectories of States



## Tests per Day and Test Positivity



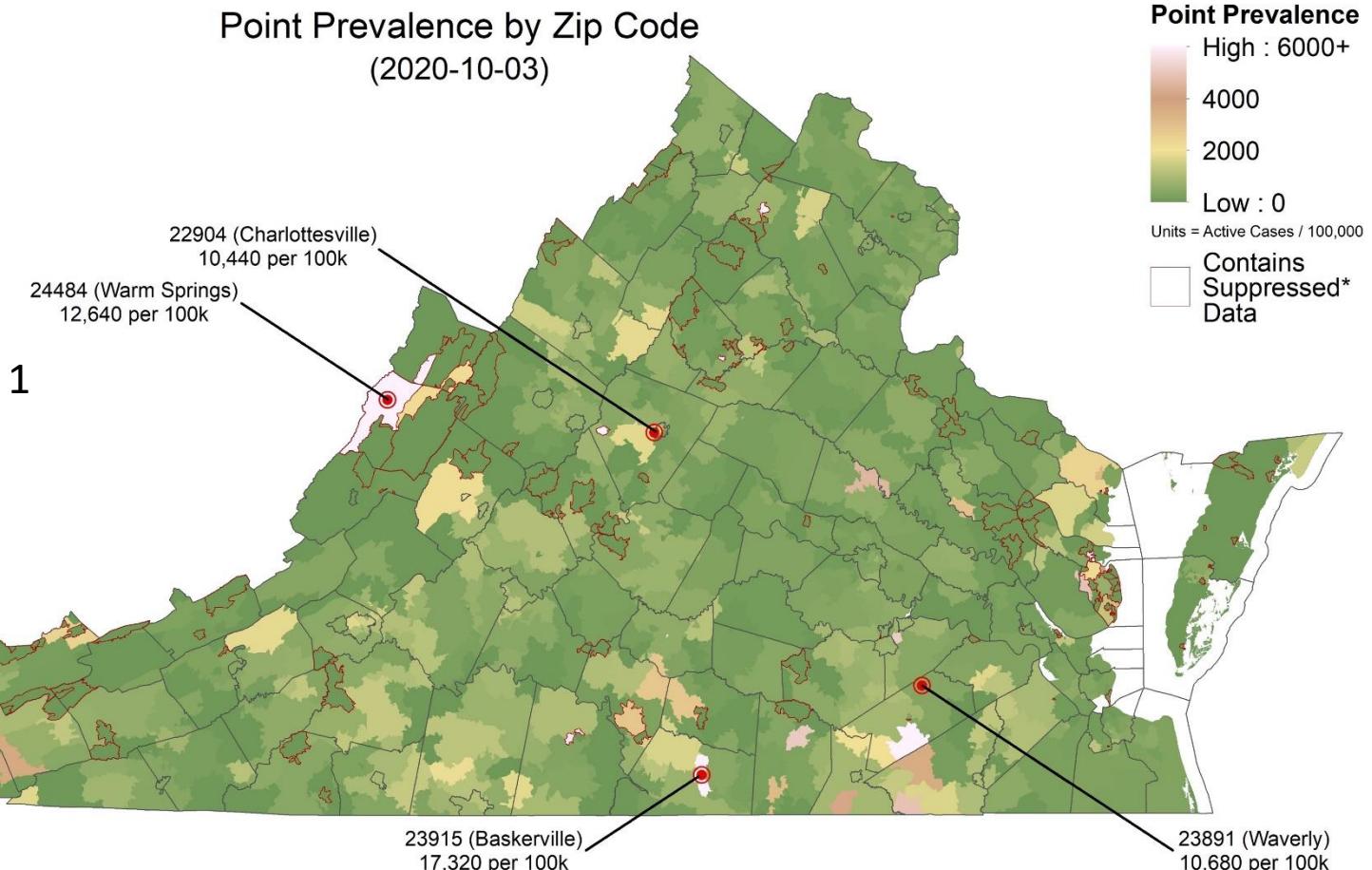
- VA and all neighbors plateauing
- Slight mixed trends but within the bounds of steady
- Case rates remain over 10/100K in TN, NC, SC, DE, and WV

- Test positivity mixed, VA's declining rate has slowed.
- Testing volumes steady in most states

# Zip code level weekly Case Rate (per 100K)

## Case Rates in the last week by zip code

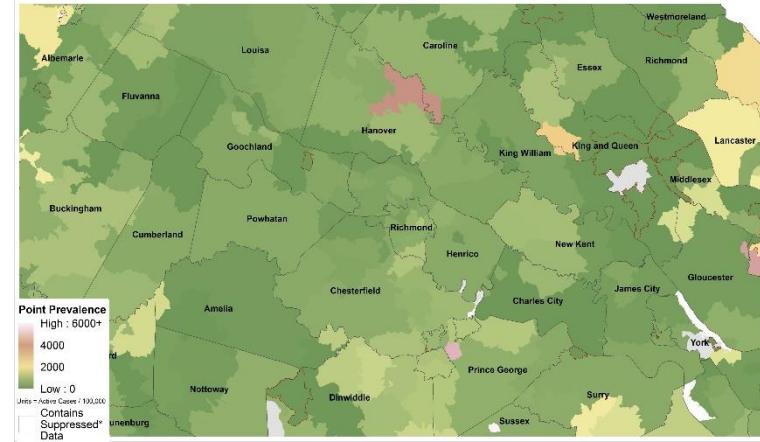
- Concentrations of very high prevalence in some zip codes, some fueled by low population size
- High prevalence zips well distributed across the commonwealth
- Many counts are low and suppressed to protect anonymity, those are assumed to be 1 case (per zip per day) and shown in white



# Zip code level weekly Case Rate (per 100K)

## Richmond

Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



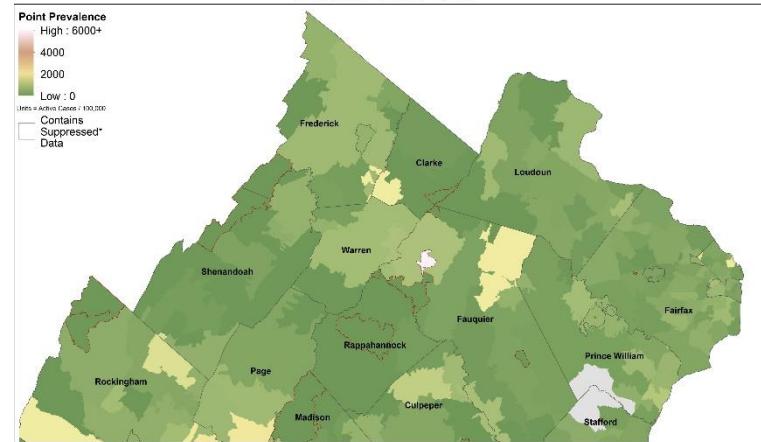
## Albemarle

Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



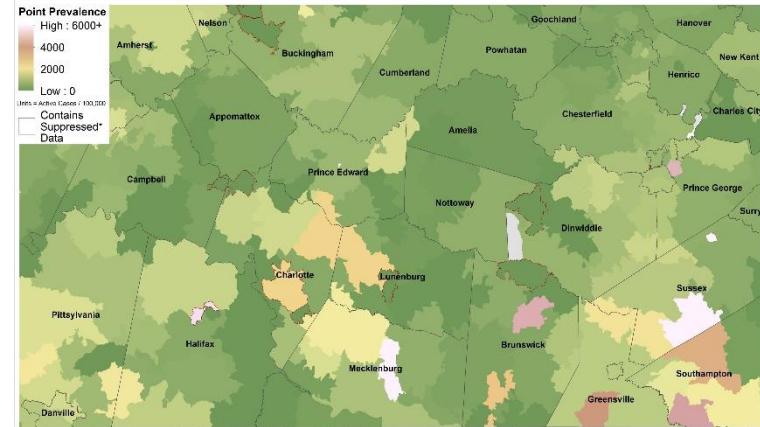
## Northern Virginia

Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



## Southside

Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



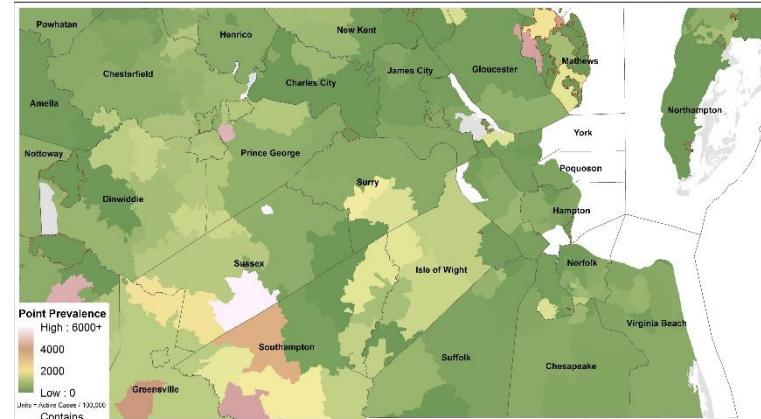
## Three Rivers

Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



## Tidewater

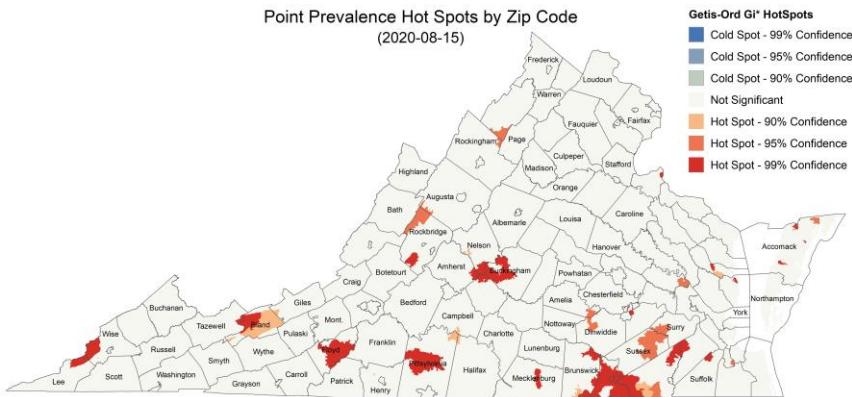
Point Prevalence by Zip Code  
2020-09-27 to 2020-10-03



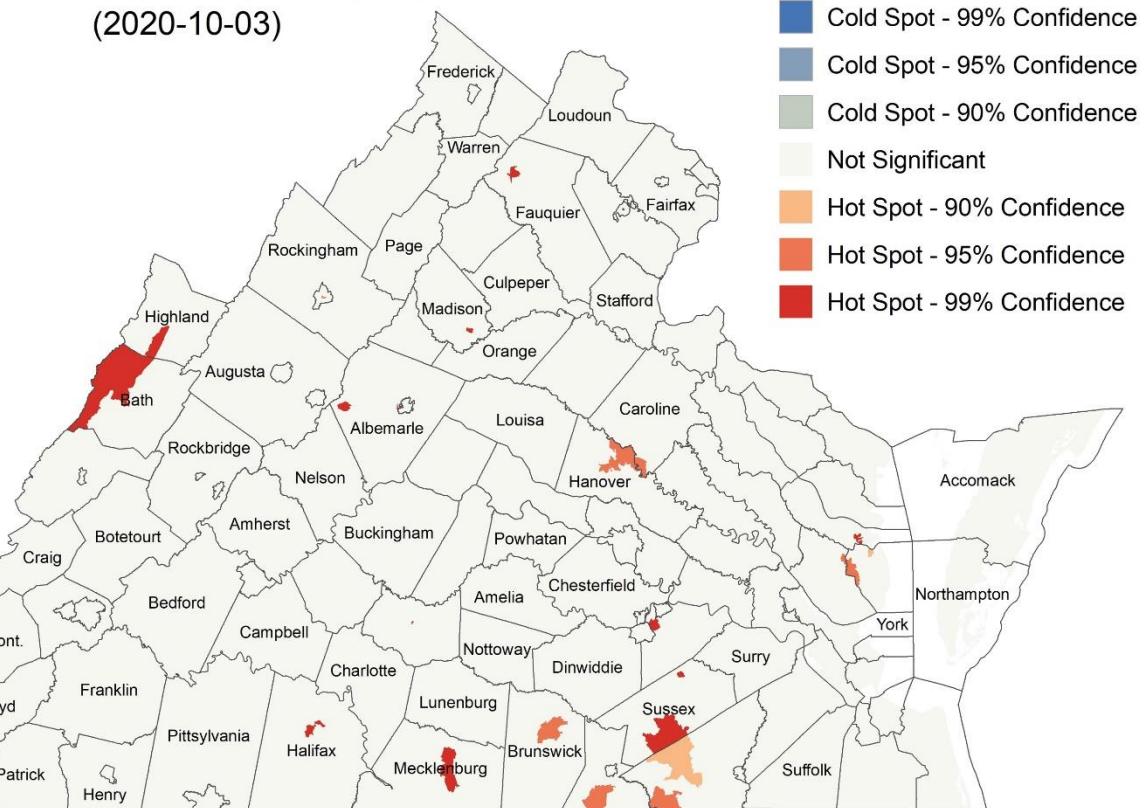
7-Oct-20

# Zip Code Hot Spots

## Previous weeks



Point Prevalence Hot Spots by Zip Code  
(2020-10-03)



## Hotspots across commonwealth

- General trend towards fewer hotspots over the last month
- Most university hotspots now less significant



# Model Update – Adaptive Fitting

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# Adaptive Fitting Approach

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

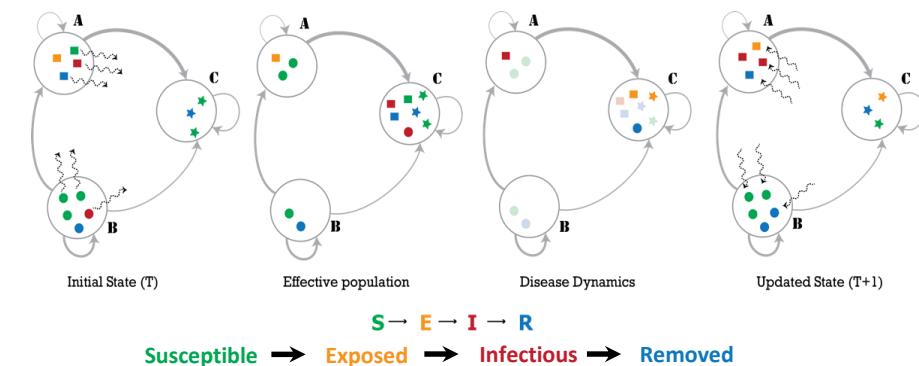
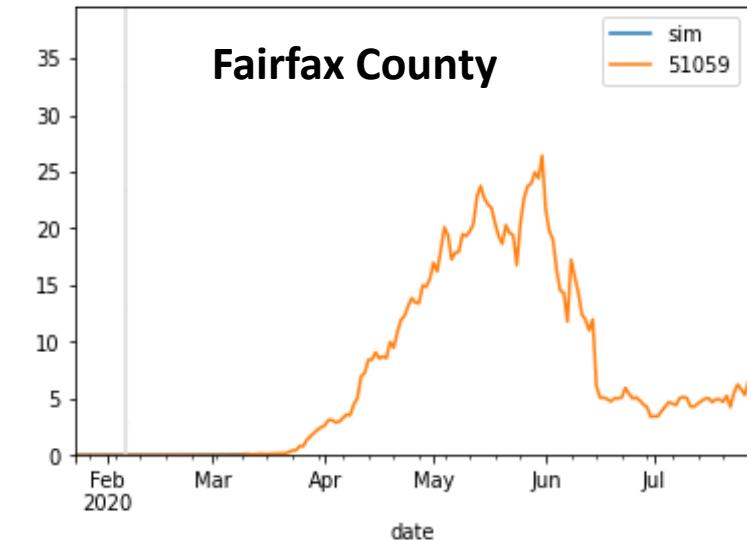
- Allows history to be precisely captured, and used to guide bounds on projections

**Model:** An alternative use of the same meta-population model, PatchSim

- Allows for future “what-if” Scenarios to be layered on top of calibrated model
- Eliminates connectivity between patches, to allow calibration to capture the increasingly unsynchronized epidemic

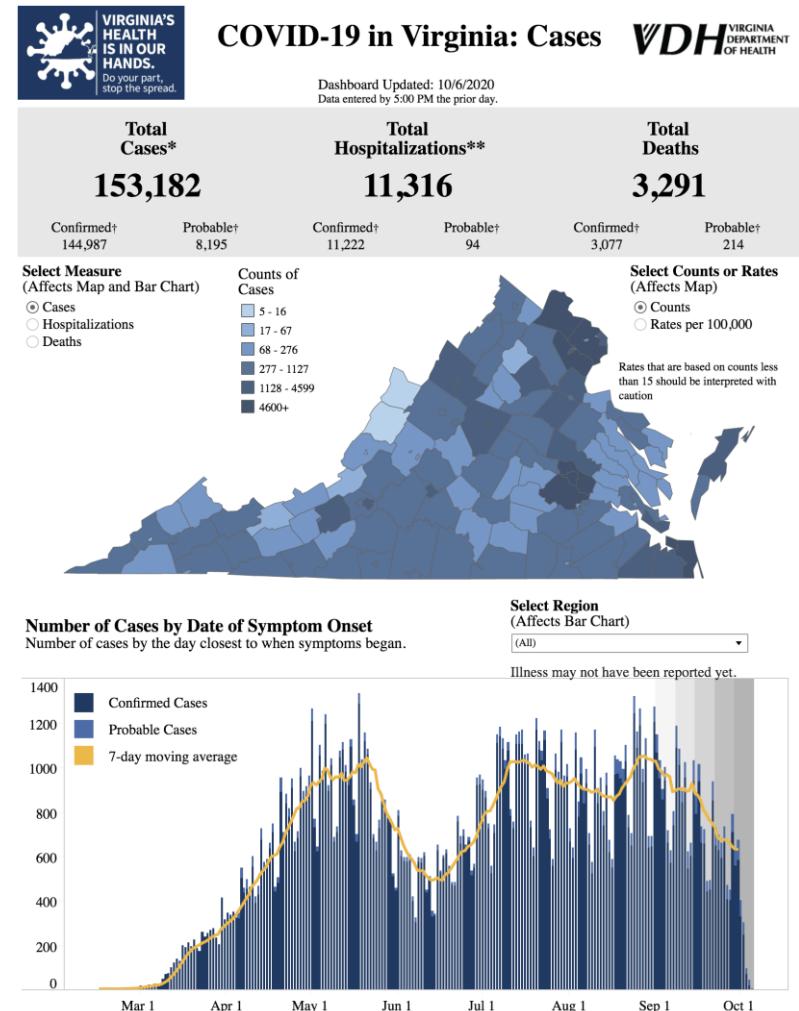
**External Seeding:** Steady low-level importation

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



# Calibration Approach

- **Data:**
  - County level case counts by date of onset (from VDH)
  - Confirmed cases for model fitting
- **Calibration:** fit model to observed data
  - Tune transmissibility across ranges of:
    - Duration of incubation (5-9 days), infectiousness (3-7 days)
    - Undocumented case rate (2x to 15x)
    - 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 using the most recent parameters with constraints learned from the history of the fit parameters
  - Mean trend from last 7 days used, adjusted by variances in the previous 3 weeks
  - 1 week interpolation to smooth transitions in rapidly changing trajectories
  - Particles with high error or variance filtered out



Accessed 8:30am October 7, 2020  
<https://www.vdh.virginia.gov/coronavirus/>

# Scenarios – Seasonal Effects

- Societal changes in the coming weeks may lead to an increase in transmission rates
  - Start of in-person school
  - Changes to workplace attendance
  - Seasonal impact of weather patterns
- Three scenarios provided to capture possible trajectories related to these changes starting Nov 1<sup>st</sup>, 2020
  - Adaptive: No change from base projection
  - Adaptive-Low: 10% increase in transmission starting Nov 1<sup>st</sup>, 2020
  - Adaptive-High: 20% increase in transmission starting Nov 1<sup>st</sup>, 2020

# Model Results

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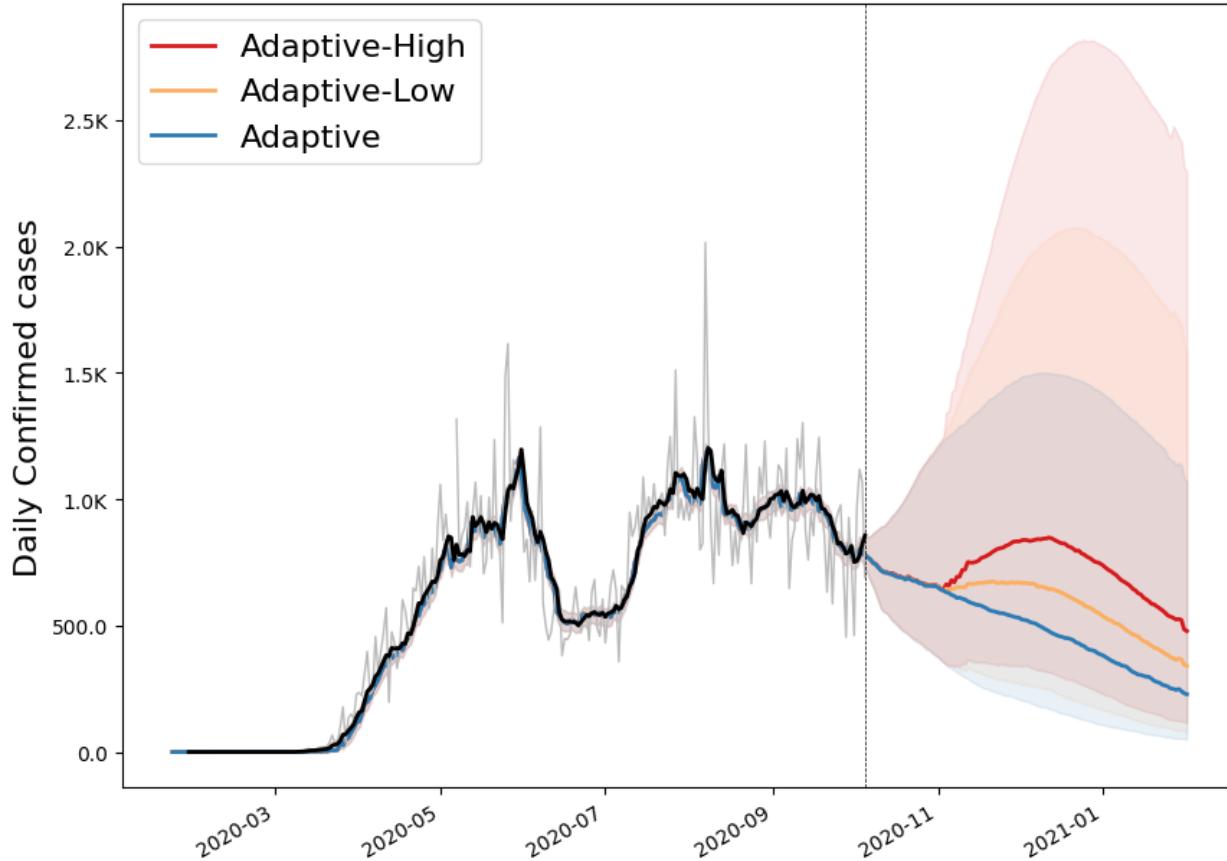


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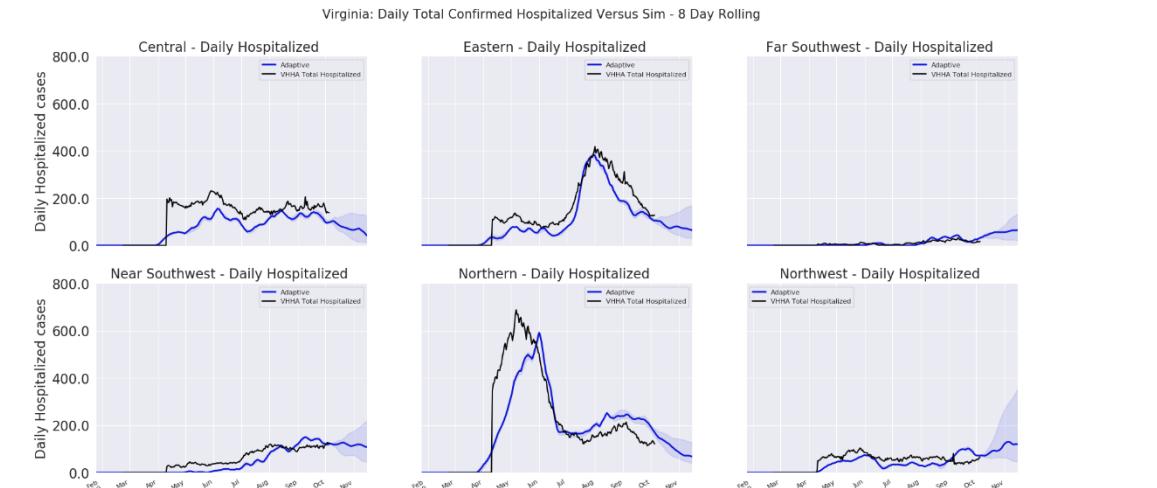
# Outcome Projections

## Confirmed cases

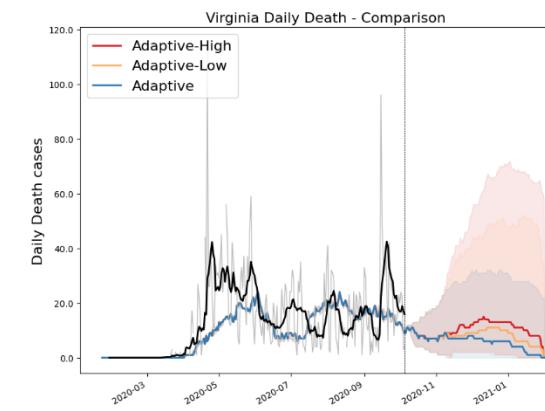
Virginia Daily Confirmed - Comparison



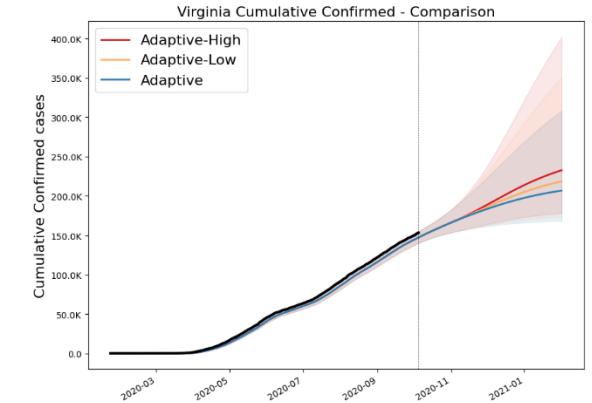
## Estimated Hospital Occupancy



## Daily Deaths



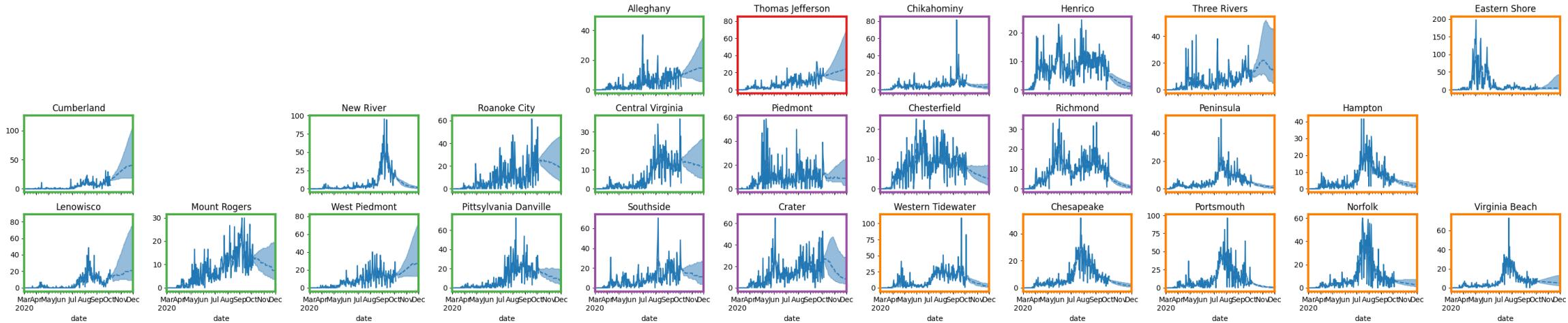
## Cumulative Confirmed cases



# District Level Projections: Adaptive

## Adaptive projections by District

- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (blue solid) with simulation colored by scenario

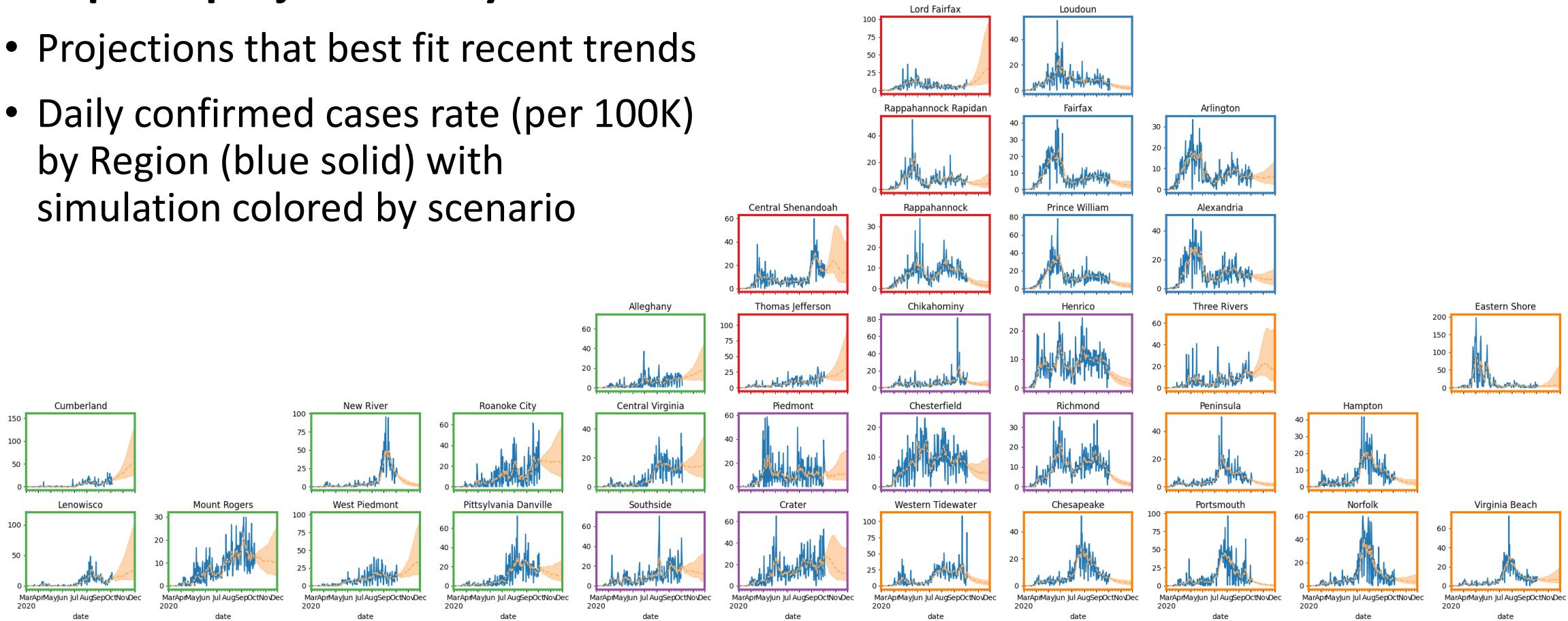


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# District Level Projections: Adaptive-Low

## Adaptive projections by District

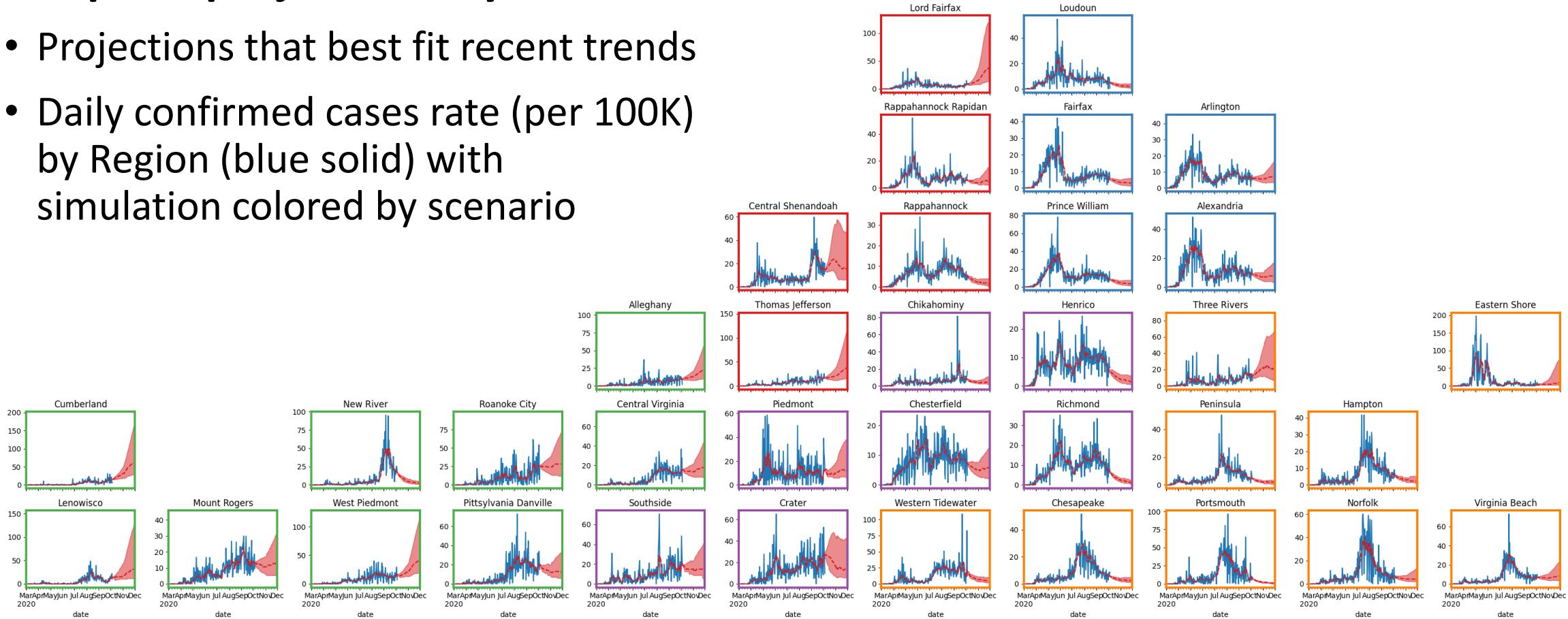
- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (blue solid) with simulation colored by scenario



# District Level Projections: Adaptive-High

## Adaptive projections by District

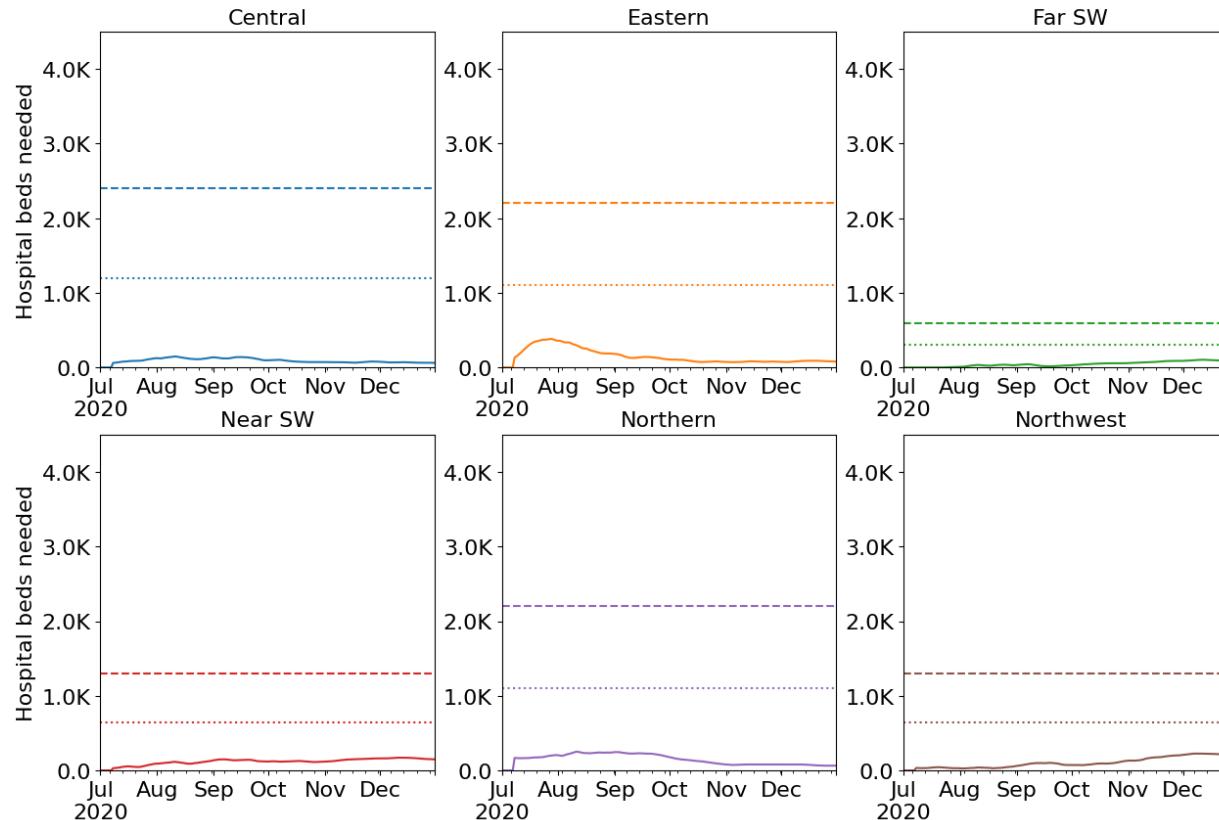
- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (blue solid) with simulation colored by scenario



# Hospital Demand and Capacity by Region

## Capacities by Region – Adaptive-High

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



\* Assumes average length of stay of 8 days

Week Ending	Adaptive	Adaptive-High
10/4/20	5,506	5,506
10/11/20	5,246	5,250
10/18/20	4,906	4,933
10/25/20	4,760	4,790
11/1/20	4,604	4,634
11/8/20	4,390	4,651
11/15/20	4,177	5,136
11/22/20	3,975	5,440
11/29/20	3,810	5,709
12/06/20	3,654	5,852
12/13/20	3,430	5,904
12/20/20	3,188	5,774

Based on Adaptive-High scenario

- No regions forecast to exceed capacity

# Key Takeaways

Projecting future cases precisely is impossible and unnecessary.

Even without perfect projections, we can confidently draw conclusions:

- **Holding steady with declines outpacing growth.**
- VA weekly incidence (9.8/100K) continues to decline and now well below the national average (16/100K) which has been climbing, fueled by growth in the Plains and Mountain West.
- Projections are mostly downward across commonwealth.
- Recent updates:
  - Adaptive Fitting projection process has been streamlined.
  - Planning Scenarios moved to Nov 1st.
- The situation is changing rapidly. Models will 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.
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- Biocomplexity Institute. COVID-19 Surveillance Dashboard. <https://nssac.bii.virginia.edu/covid-19/dashboard/>
- Google. COVID-19 community mobility reports. <https://www.google.com/covid19/mobility/>
- Biocomplexity page for data and other resources related to COVID-19: <https://covid19.biocomplexity.virginia.edu/>

# Questions?

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# Supplemental Slides



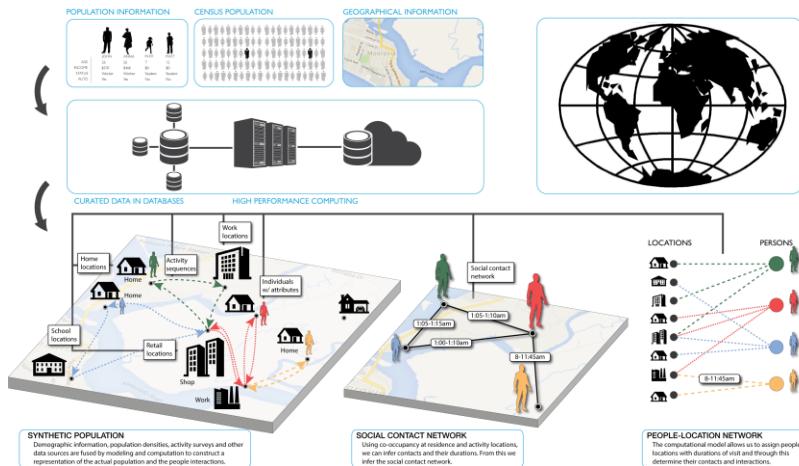
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# Agent-based Model (ABM)

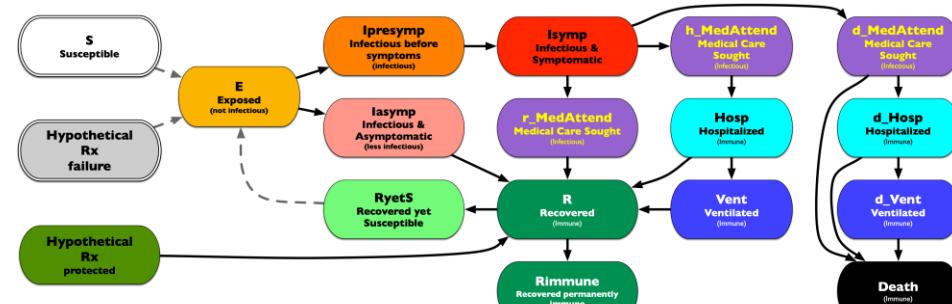
## EpiHiper: Distributed network-based stochastic disease transmission simulations

- Assess the impact on transmission under different conditions
- Assess the impacts of contact tracing



### Synthetic Population

- Census derived age and household structure
- Time-Use survey driven activities at appropriate locations



### Detailed Disease Course of COVID-19

- Literature based probabilities of outcomes with appropriate delays
- Varying levels of infectiousness
- Hypothetical treatments for future developments

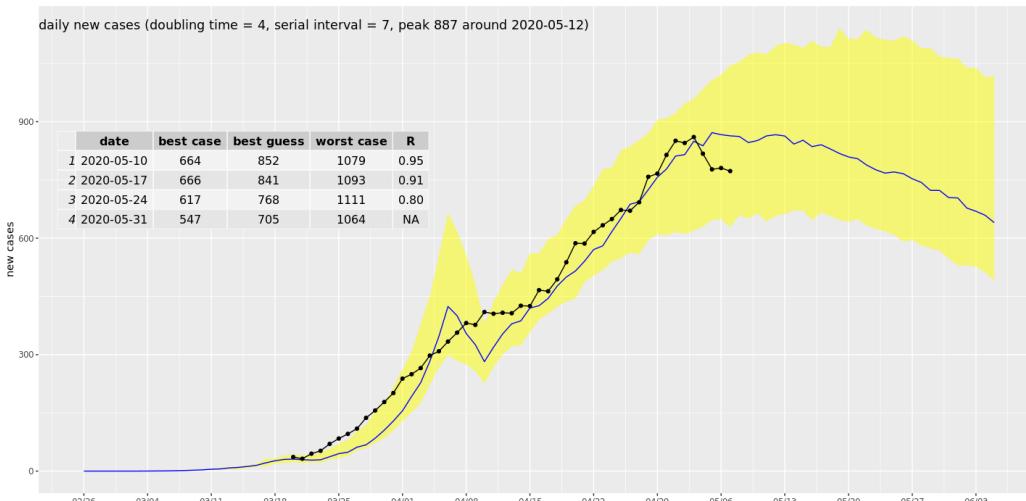


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# ABM Social Distancing Rebound Study Design

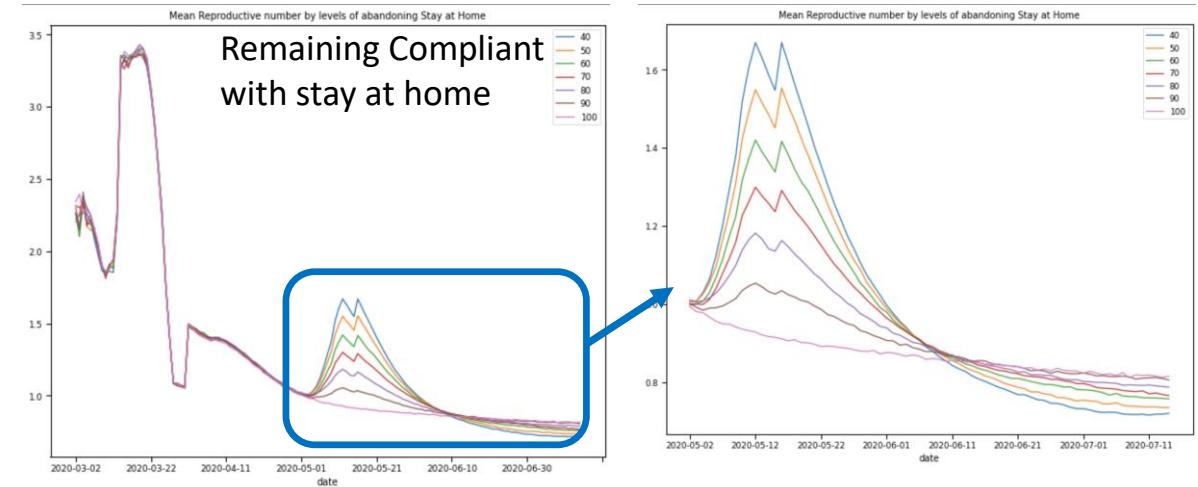
## Study of "Stay Home" policy adherence

- Calibration to current state in epidemic
- Implement “release” of different proportions of people from “staying at home”



### Calibration to Current State

- Adjust transmission and adherence to current policies to current observations
- For Virginia, with same seeding approach as PatchSim



### Impacts on Reproductive number with release

- After release, spike in transmission driven by additional interactions at work, retail, and other
- At 25% release (70-80% remain compliant)
- Translates to 15% increase in transmission, which represents a 1/6<sup>th</sup> return to pre-pandemic levels

# Medical Resource Demand Dashboard

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

