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

Estimation of COVID-19 Impact in Virginia

December 9th, 2020

(data current to December 8th) Biocomplexity Institute Technical report: TR 2020-157

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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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Biocomplexity COVID-19 Response Team



Overview

- Goal: Understand impact of COVID-19 mitigations in Virginia
- Approach:
 - Calibrate explanatory mechanistic model to observed cases
 - Project infections 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 growth in Virginia seems to have rebounded following Thanksgiving holiday
- VA mean weekly incidence (38/100K) up (from 28) as national surge also rebounds (to 68/100K from 53/100K).
- Lingering effects from reporting delays and overall testing interruptions continue to impact projections, which are mostly up.
- Recent updates:
 - Ensemble of statistical and Machine Learning models integrated with Adaptive to guide projections
 - Return projection trend window to 2 weeks to minimize holiday effects
 - Planning scenarios remain at Dec 10th and case ascertainment rates updated but are unchanged
- The situation is changing rapidly. Models will be updated regularly.



Situation Assessment



Case Rate (per 100k) by VDH District

Surging Rates continue

- Majority of Districts have increasing rates
- Many districts experiencing highest rates of pandemic •

New River

West Piedmont

date

2020

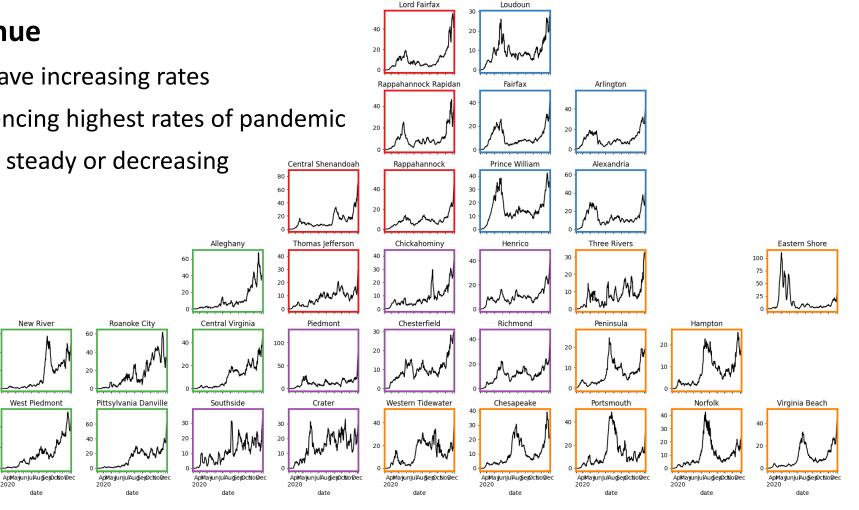
Some Districts remain steady or decreasing

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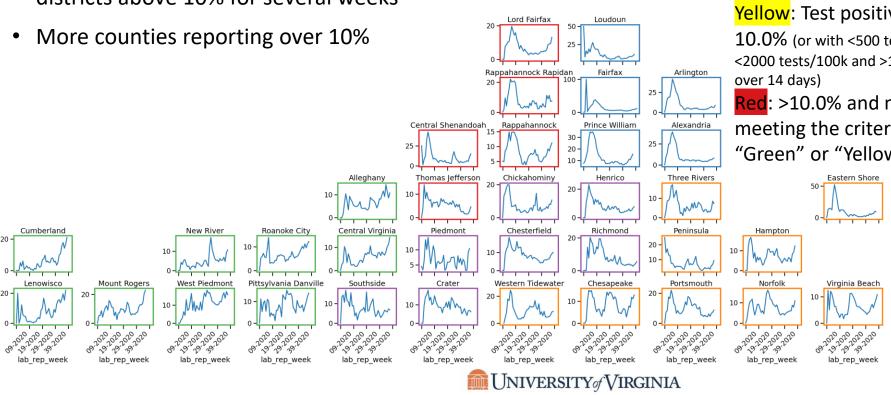
Test Positivity by VDH District

Weekly changes in test positivity by district

 Increasing levels in many districts throughout the commonwealth with many districts above 10% for several weeks

County level test positivity rates for RT-PCR tests.

Green: Test positivity <5.0% (or with <20 tests in past 14 days) Yellow: Test positivity 5.0%-10.0% (or with <500 tests and <2000 tests/100k and >10% positiv over 14 days) **Red**: >10.0% and not meeting the criteria for "Green" or "Yellow"



Oct-28 Nov-04 Nov-11 Nov-18

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https://data.cms.gov/stories/s/q5r5-givu

Yellow

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Count Amherst Count

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Botetourt County Bristol City Buena Vista City Campbell County

Carroll County

Chesapeake City

Clarke County Covington City

Craig County

Culpeper County

Patrick Count

Roanoke City Roanoke Count

Salem Cit

Vise County

Wythe County

Smyth County Stafford County Tazewell Count Washington Count

Rockingham County Russell County

Pittsylvania Count

Prince George Count

Prince William Count Pulaski County

Chesterfield County

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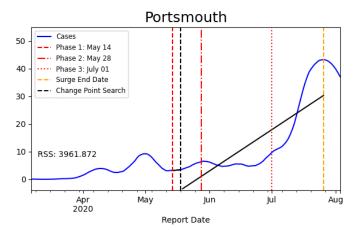
Yellow

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

Hockey stick fit



Trajectory	Description	Weekly Case Rate (per 100K) bounds	# Districts (prev week)
Declining	Sustained decreases following a recent peak	below -0.9	6 (2)
Plateau	Steady level with minimal trend up or down	above -0.9 and below 0.5	1 (3)
Slow Growth	Sustained growth not rapid enough to be considered a Surge	above 0.5 and below 2.5	11 (9)
In Surge	Currently experiencing sustained rapid and significant growth	2.5 or greater	17 (21)



District Trajectories

Status	# Districts (prev week)	Lord Fairfax - Surging Loudoun - Surging 1.5 <= R < 2 *
Declining	6 (2)	$I = \frac{1}{1000} \int_{\frac{1}{1000}} \int_{\frac{1}{1000}} \int_{\frac{1}{1000}} \int_{\frac{1}{1000}} \int_{\frac{1}{1000}} \int_{\frac{1}{1000}} \int_{\frac{1}{10000}} \int_{\frac{1}{100000}} \int_{\frac{1}{1000000000000000000000000000000000$
Plateau	1 (3)	Transition
Slow Growth	11 (9)	Central Shenandoah - Surging Rappahannock - Surging Prince William - Surging Alexandria - Sur
In Surge	17 (21)	
Curve shows smoothe Trajectories of states i Case Rate curve color	in label & chart box	The factor is a second
Curriburiand - Surging	New River - Declining	y - Declining
Lenowisco Slow Growth Countion and the state of the stat		Southside - Plateau Crater - Slow Growth Western Tidewater - Slow Growth Chesapake - Slow Growth Norfolk - Slow Growth • 0 • 0 • 0 • 0 • 0 • 0 • 0 • 0
Report Date Report Date	Heart Date Hear	tour Pepul bit Not Die

Estimating Daily Reproductive Number

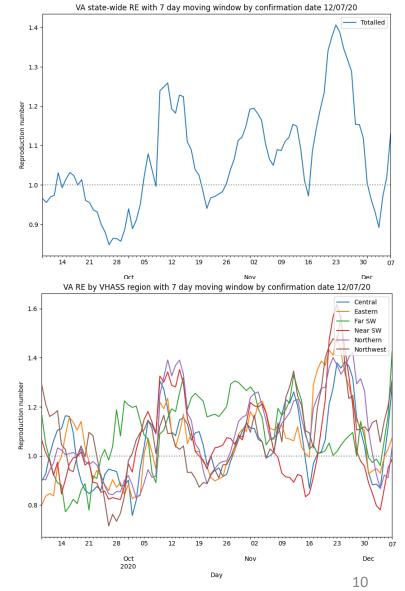
Dec 7th Estimates

Region	Date Confirmed R _e	Date Confirmed Diff Last Week
State-wide	1.130	0.010
Central	1.283	0.242
Eastern	1.073	-0.032
Far SW	1.424	0.349
Near SW	0.987	0.029
Northern	1.034	-0.232
Northwest	1.309	0.190

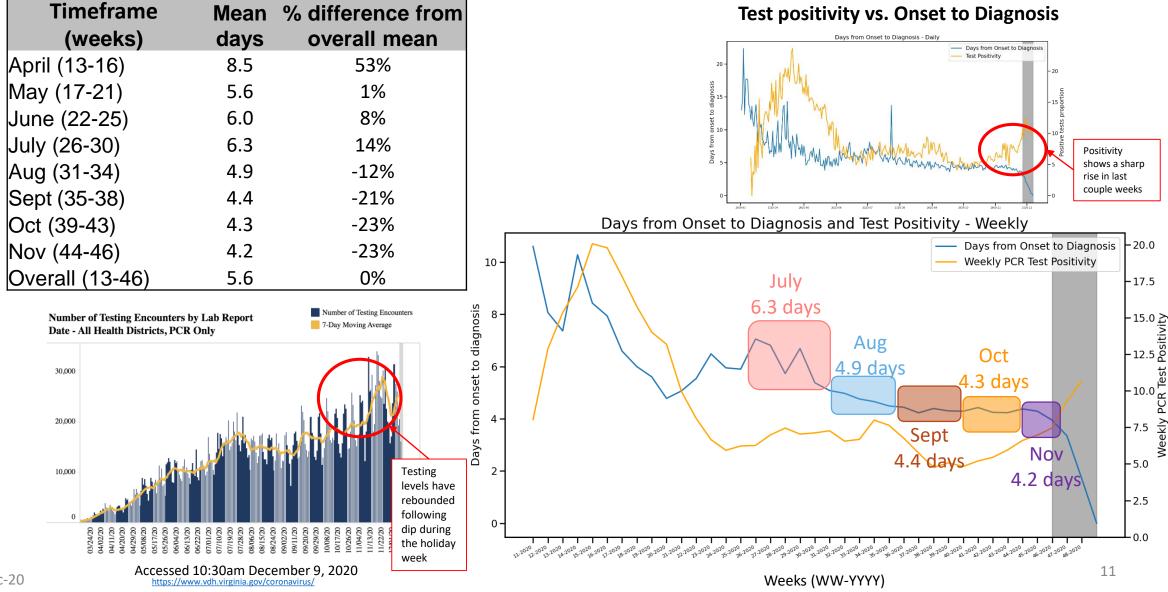
Methodology

- Wallinga-Teunis method (EpiEstim¹) for cases by confirmation date
- Serial interval: 6 days (2 day std dev)
- Using Confirmation date since due to increasingly unstable estimates from onset date due to backfill

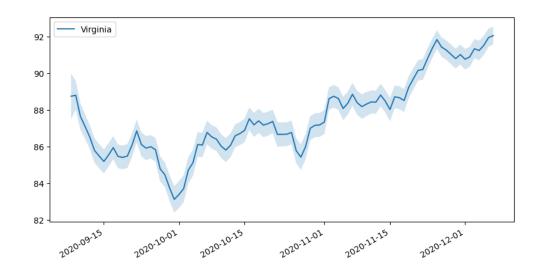
1. Anne Cori, Neil M. Ferguson, Christophe Fraser, Simon Cauchemez. A New Framework and Software to Estimate Time-Varying Reproduction Numbers During Epidemics. American Journal of Epidemiology, Volume 178, Issue 9, 1 November 2013, Pages 1505–1512, <u>https://doi.org/10.1093/aje/kwt133</u>



Changes in Case Detection



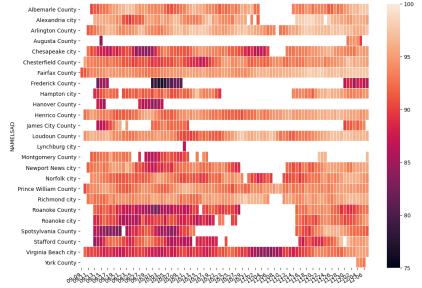
Mask usage in Virginia



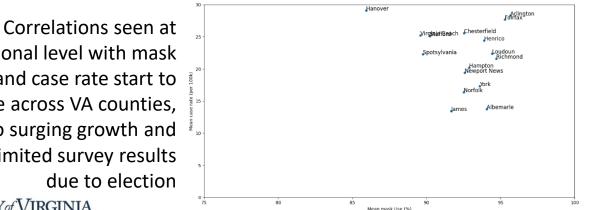
State level mask usage as reported via Facebook surveys over the past month shows ranges from 83% to 89%

- Relatively stable over time •
- Limited variance across the commonwealth •
- ~3000 daily responses from VA ٠

Data Source: https://covidcast.cmu.edu



Some county level fluctuations since beginning of Sept., though data quality may be affected by sample sizes.

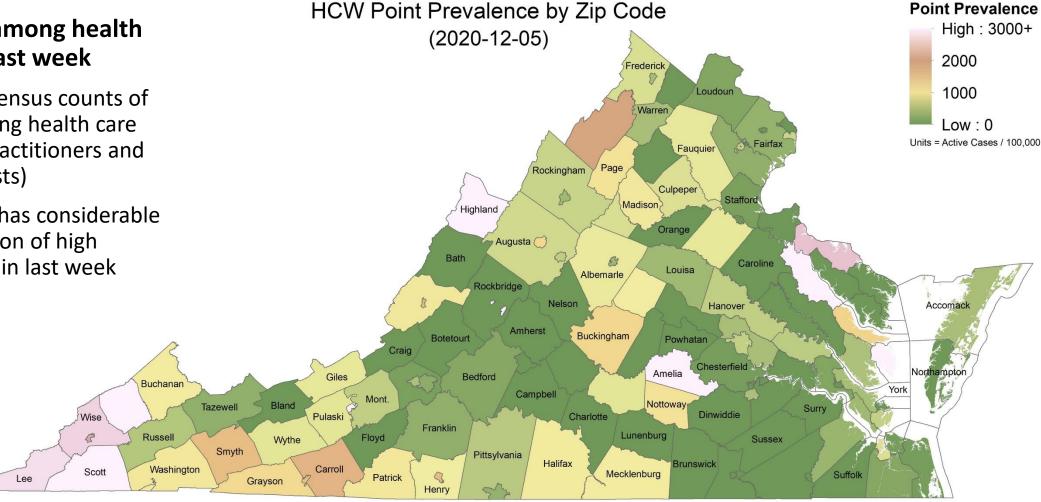


national level with mask use and case rate start to emerge across VA counties, due to surging growth and more limited survey results due to election UNIVERSITY of VIRGINIA

Health Care Worker Prevalence (per 100K)

Case Rates among health workers in last week

- Based on Census counts of ٠ patient facing health care workers (Practitioners and Technologists)
- Southwest has considerable • concentration of high prevalence in last week



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Age-Specific Attack Rates (per 100K)

Cumulative Age-Specific Attack Rates (per 100k)

- 30-49 year olds bear the highest burden of disease in most districts
 - Major university districts still show a majority of Lord Fairfax Loudour their patients coming from the 0-29 age group 2500 2000 Legend: Age Categories Rappahannock Rapidan Fairfax Arlington 0-29 2500 2000 2500 30-49 Central Shenandoah Rappahannock Prince William Alexandria 50-69 5000 2000 2500 2500 70-79 80 +Thomas Jefferson Chickahominy Henrico Three Rivers Alleghany Eastern Shor 2000 2000 5000 2000 2000 2500 Central Virginia Piedmont Chesterfield Richmond Peninsula Cumberland Roanoke City New River Hampto 5000 5000 5000 2000 2000 2500 2000 2000 2000 Pittsylvania Danville Southside Crater Western Tidewater Chesapeake Portsmouth Norfolk Lenowisco Mount Rogers West Piedmont Virginia Beach 5000 5000 5000 2000 2000 2500 2500 2500 2500 2000 2500 NASAN VERSON NASAN BURGAN 1999 4 Hill Carlor NASAN BUZER NASAN GUEREN NAGAN LIEBER NATA VEREN いまちょうかんだいちょうちょ LESSII UP 18 19 1 19930 UNIVERSITY of VIRGINIA

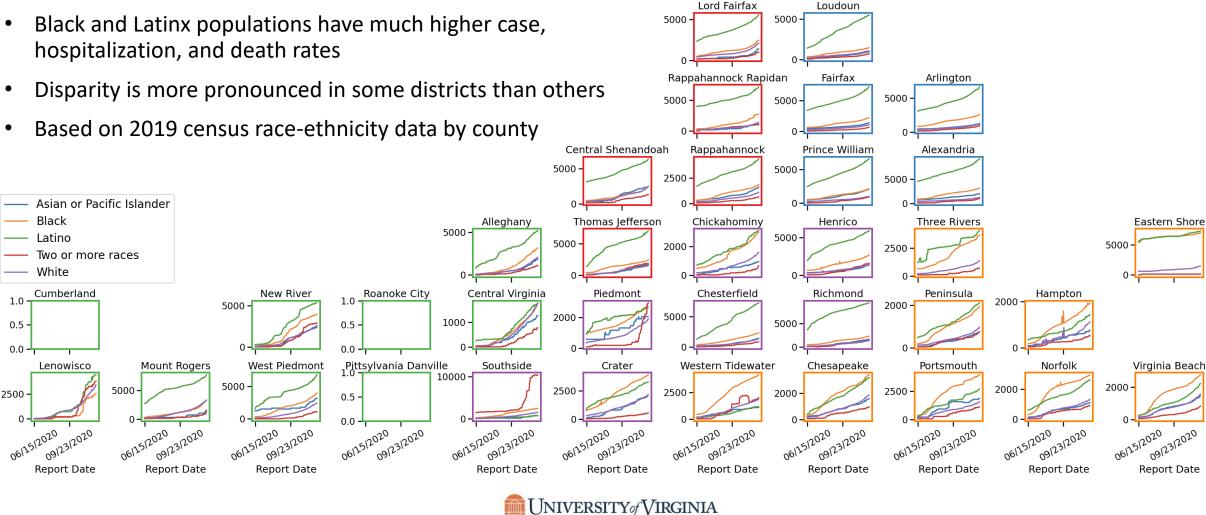
Age-adjusted Cumulative Prevelance Rate Per 100k District Population

•

Race and Ethnicity Attack Rates (per 100K)

Cumulative Race and Ethnicity Attack Rates (per 100k)

- Black and Latinx populations have much higher case, hospitalization, and death rates
- Disparity is more pronounced in some districts than others
- Based on 2019 census race-ethnicity data by county



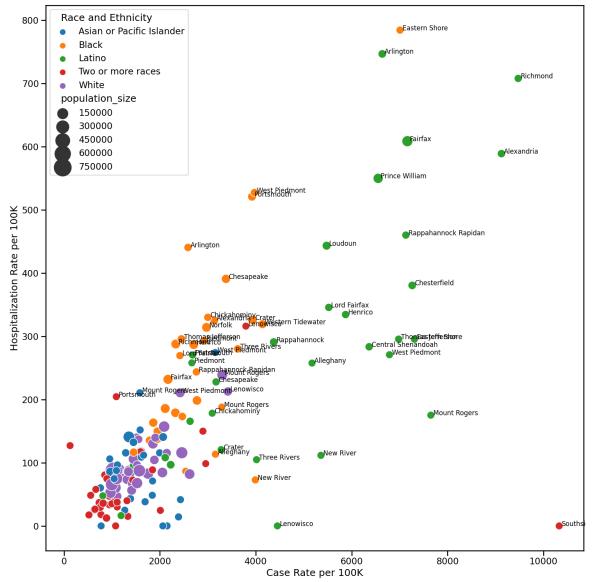
1.0

0.5

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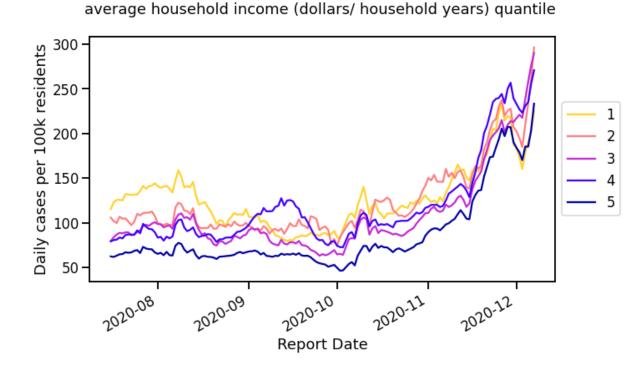
Race and Ethnicity cases per 100K



Rates per 100K of each Racial-Ethnic population by Health District

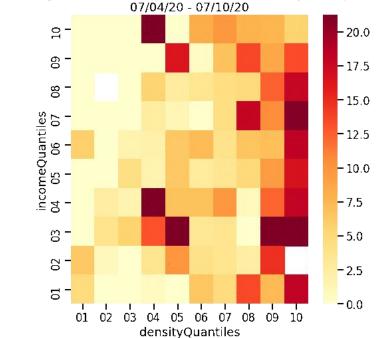
- Each Health District's Racial-Ethnic population is plotted by their Hospitalization and Case Rate
- Points are sized based on their overall population size
- Overlapping labels removed for clarity

Impact across Density and Income



VDH 7-day moving average rate of new COVID-19 cases by zip code

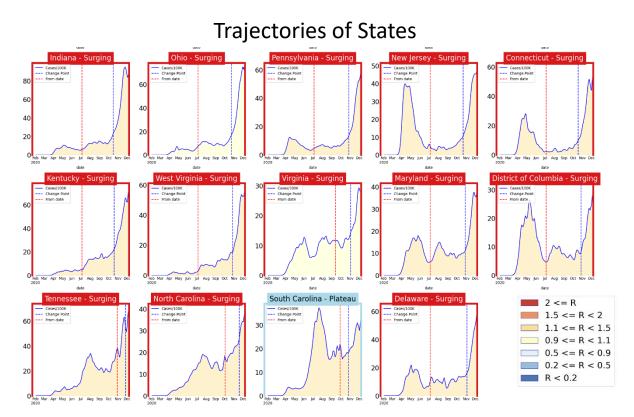
All zip codes show rapid growth and ordering is in flux with the middle quintiles (20th to 80th percentiles) bearing the highest rates VDH mean cases per 100k by zip code population density (person/ sq mile) and average household income (dollars/ household years) quantiles



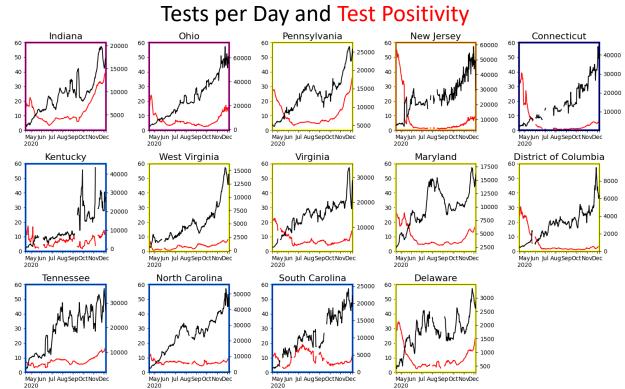
Full evolution of pandemic, shows shifts from denser and wealthier zip codes to poorer and less dense zip codes, followed by a repeat of the pattern. Recently see an uptick across the spectrum of density and income

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Other State Comparisons



- VA and most mid-Atlantic states are in surge (34 total in US)
- Nearly all states maintain highest rates of the pandemic



- VA's test positivity rate continues to rise with many of its neighbors
- Testing volumes have increased for many with recent disruption from the holiday

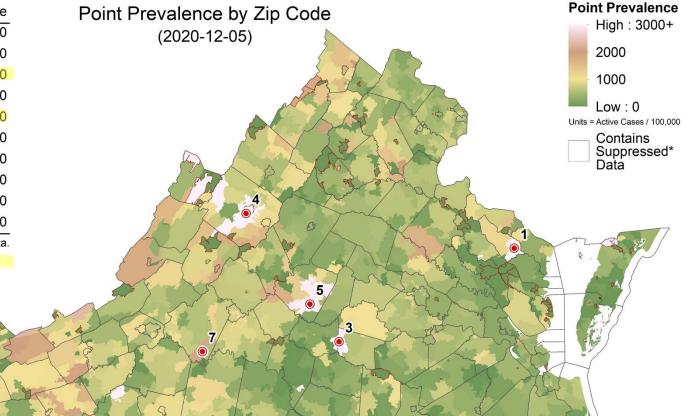


Zip code level weekly Case Rate (per 100K)

Case Rates in the last week by zip code

- Concentrations of very high prevalence in many zip codes
- Several of the top ten zip codes are home to prisons
- Southwest has considerable concentration of high prevalence zips
- Some counts are low and suppressed to protect anonymity, those are shown in white

	Rank	Zip Code	Name	Prevalence	Poi
	1	22460	Farnham	5,220	
	2	24290	Weber City	4,620	
	3	23922	Burkeville *	3,810	
	4	24401	Staunton	3,420	
	5	23936	Dillwyn *	2,950	
	6	24243	Dryden	2,720	
	7	24104	Huddleston	2,090	
	8	24266	Lebanon	2,060	
	9	24228	Clintwood	2,050	
	10	24236	Damascus	2,000	
C	Only inc	ludes zips w	ith pop ≥ 1000 ar	id no supp. data.	
		Denotes zip	codes with state	e prisons.	

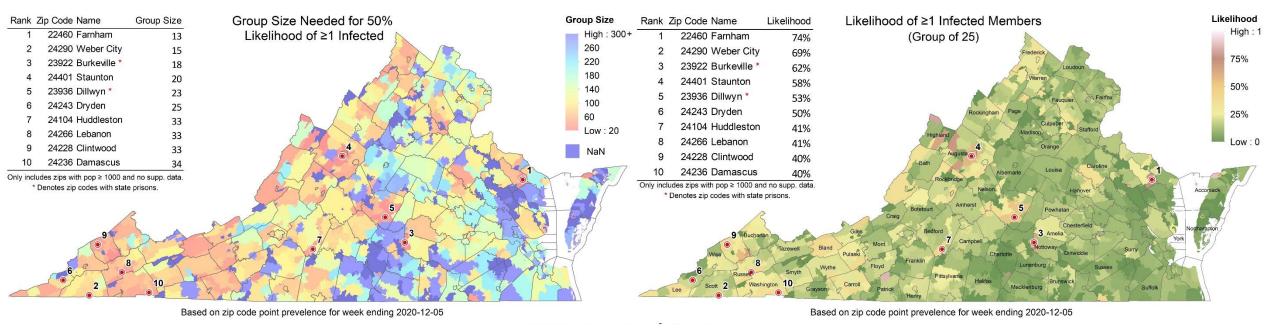




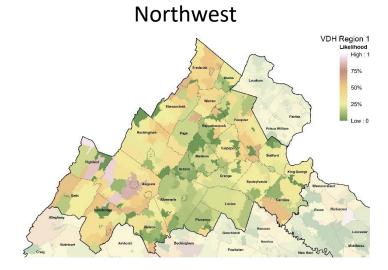
Risk of Exposure by Group Size

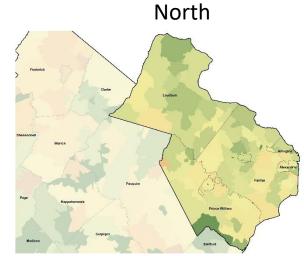
Case Prevalence in the last week by zip code used to calculate risk of encountering someone infected in a gathering of randomly selected people (group size 25)

- Assumes 3 undetected infections per confirmed case (ascertainment rate from recent seroprevalence survey)
- On left, minimum size of a group with a 50% chance an individual is infected by zip code (eg in a group of 20 in Staunton, there is a 50% chance someone will be infected)
- Some zip codes have high likelihood of exposure even in groups of 25

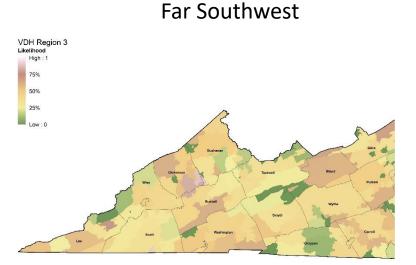


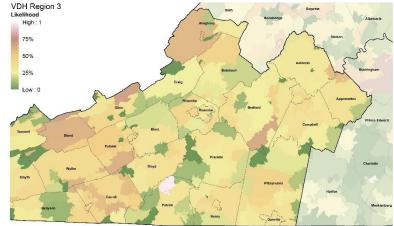
Risk of Exposure in Groups of 50

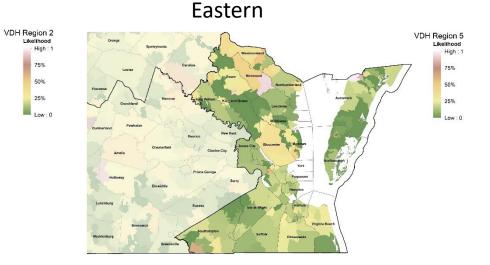




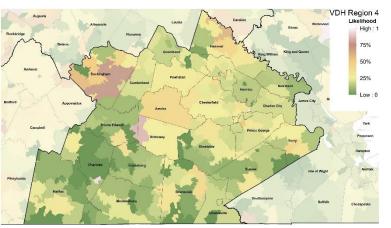
Near Southwest









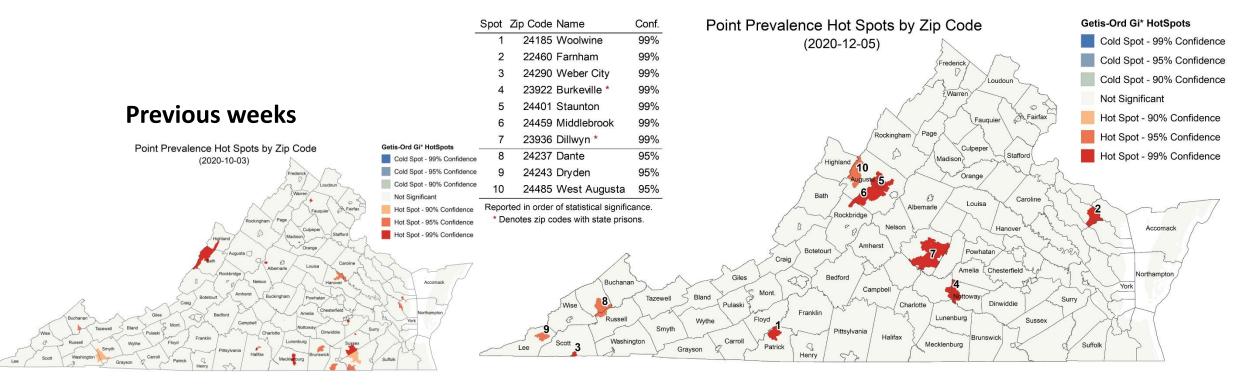




Zip Code Hot Spots

Hotspots across commonwealth

- More spread out but remain concentrated in the Southwest
- Captures some very high prevalence rates in some zips



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Model Update – Adaptive Fitting



Adaptive Fitting Approach

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

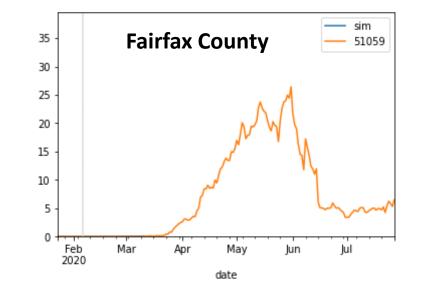
• Allows history to be precisely captured, and used to guide bounds on projections

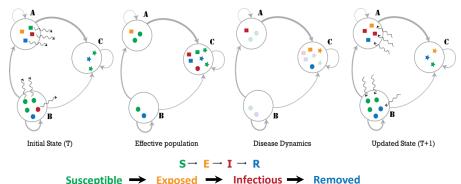
Model: An alternative use of the same meta-population model, PatchSim

- 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



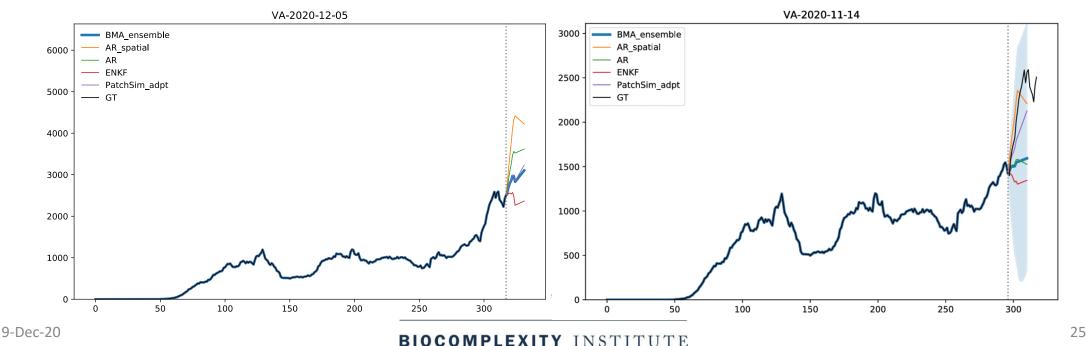




Using Ensemble Model to Guide Projections

An ensemble methodology that combines the Adaptive Fitting and machine learning and statistical models has been developed and refined

- Models: Adaptive Fitting, ARIMA, LSTM, AR, spatially driven AR, Kalman Filters (ENKF)
- This approach facilitates the use of other data streams (weather, mobility, etc.)
- Ensemble provides scaffolding for the Adaptive Fitting's short-term projections



Seroprevalence updates to model design

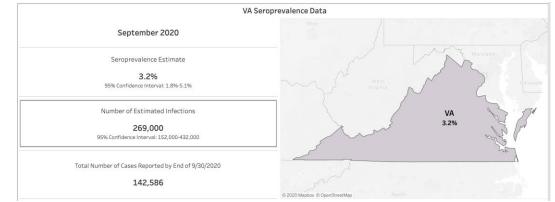
Several seroprevalence studies provide better picture of how many actual infections have occurred

- Virginia Serology Study estimated 2.4% of Virginians estimated infected (as of Aug 15th)
- CDC Nationwide Commercial Laboratory Seroprevalence Survey estimated 3.2% [1.8% – 5.1%] seroprevalence as of Sept 10th-23rd down from 3.9% a month earlier

These findings are equivalent to an ascertainment ratio of ~3x, with bounds of (1x to 7x)

- Thus for 3x there are 3 total infections in the population for every confirmed case
- Uncertainty design has been shifted to these bounds (previously higher ascertainments as was consistent earlier in the pandemic were being used)

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https://covid.cdc.gov/covid-data-tracker/#national-lab

Virginia Coronavirus Serology Project Interim findings by region and statewide - July 22, 2020

			Crude	Weighted prevalence*	
Region	Number of participants	Number antibody positive	prevalence per 100 participants	per 100 population	(95% CI)
Central	400	8	2.0	3.0	(0.5, 5.5)
East	707	9	1.3	1.5	(-0.2, 3.2)
Northern	819	36	4.4	4.2	(2.5, 5.9)
Northwest	756	11	1.5	0.9	(0.2, 1.6)
Southwest	431	3	0.7	1.0	(-0.2, 2.1)
Virginia	3,113	67	2.2	2.4	(1.6, 3.1)

* Weighted prevalence is reweighted by region, age, sex, race, ethnicity, and insurance status to match census population.

https://www.vdh.virginia.gov/content/uploads/sites/8/2020/08/VDH-Serology-Projects-Update-8-13-2020.pdf

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 14 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



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A IS IN OU HANDS Do your pa stop the sp	• irt,	Dashboard Upd Data entered by 5:0		••		
	c	ases, Hospitaliza	tions and Deaths	i		
Total Cases* 267,128		Total Hospitalizations**			Total Deaths	
(New Case	s: 4.398)^	15,	592	4,2	81	
Confirmed [†] 232,940	Probable† 34,188	Confirmed† 15,204	Probable† 388	Confirmed† 3,894	Probable† 387	
Hospitalization status a ew cases represent the /DH adopted the updat e: <u>https://wwwn.cdc.ge</u>	th a positive test (Confirme at time case was investigate number of confirmed and p ed CDC COVID-19 confirm vvnndss/conditions/corona Electronic Disease Surveilla	d by VDH. This underre robable cases reported to ned and probable surveil virus-disease-2019-covic ance System (VEDSS), o	presents the total number o VDH in the past 24 hou lance case definitions on l-19/case-definition/2020 data entered by 5:00 PM to	of hospitalizations in Virgi rs. August 27, 2020. Found /08/05/	nia.	
		Outbi			~	
1	otal Outbreaks* 1,626		Outb	reak Associated (35,999	Cases	
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-		Testing (F	PCR Only)			
Testin	g Encounters PCR O	nly*	Current 7-	Day Positivity Rate P	CR Only**	
3,573,299 10.9%						
	e transcriptase polymerase o ave been received yet. Perce			nplete data.		
		Multisystem I Syndrome				
	Total Cases*			Total Deaths		
	12			0		

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

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Accessed 10:30am December 9, 2020 https://www.vdh.virginia.gov/coronavirus/

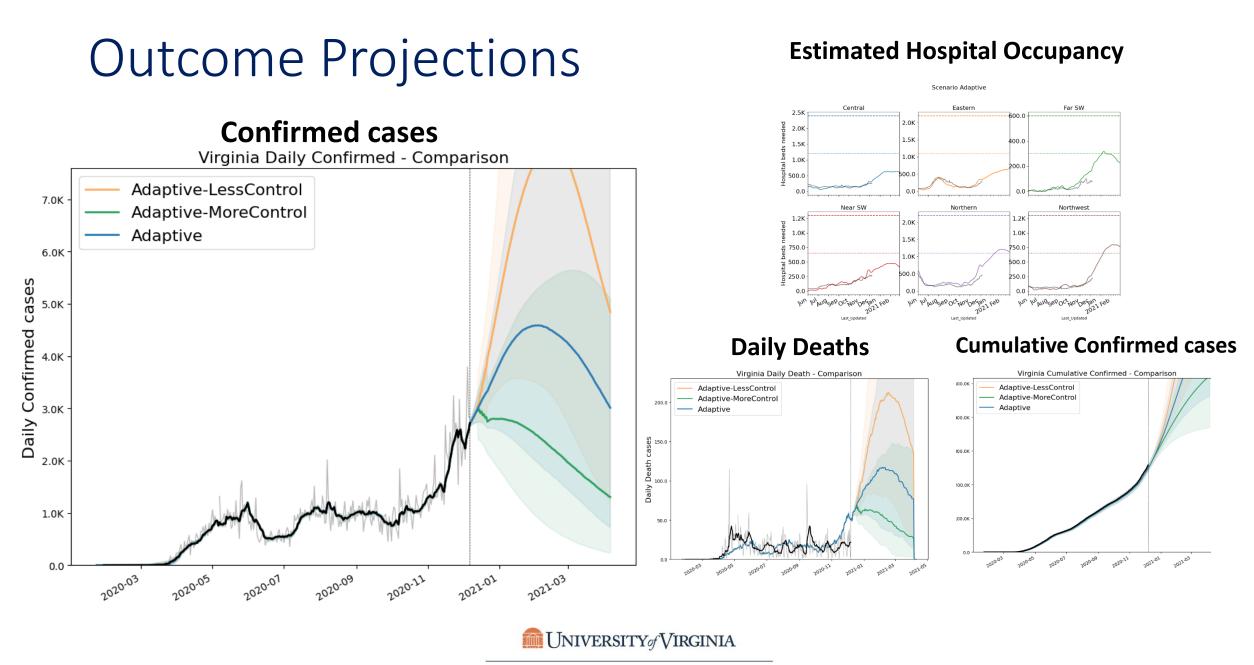
Scenarios – Seasonal Effects

- Societal changes in the past month have led to an increase in transmission rates, these could continue to drive transmission
 - Seasonal impact of weather patterns
 - More interactions at places of learning
 - Travel related to holidays and traditional large family gatherings
 - Fatigue with infection control practices
- Population's behaviors determine the level of control of transmission we can achieve
- Three scenarios capture possible trajectories starting Dec 10th, 2020
 - Adaptive: No change from base projection
 - Adaptive-MoreControl: 15% decrease in transmission starting Dec 10th, 2020
 - Adaptive-LessControl: 15% increase in transmission starting Dec 10th, 2020



Model Results

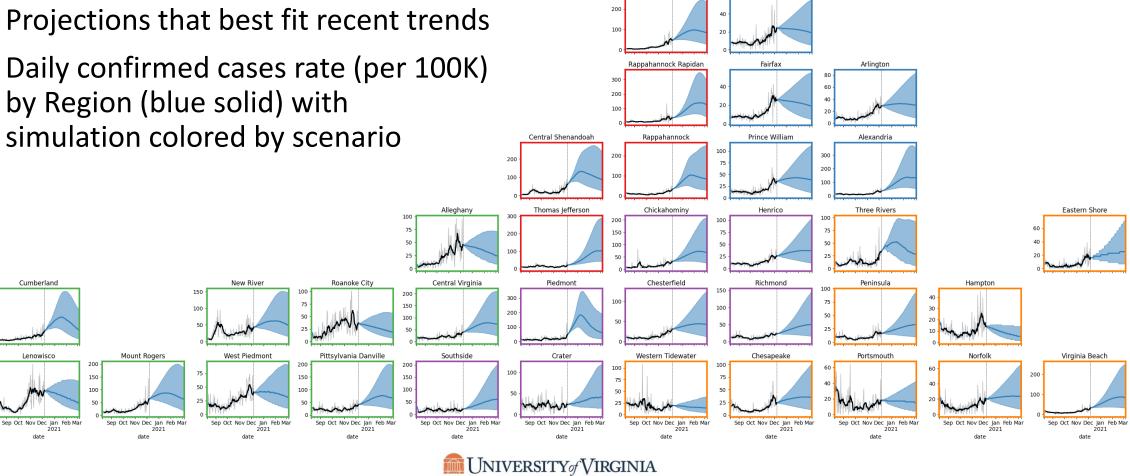




District Level Projections: Adaptive

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



Lord Fairfax

Loudoun

300

200

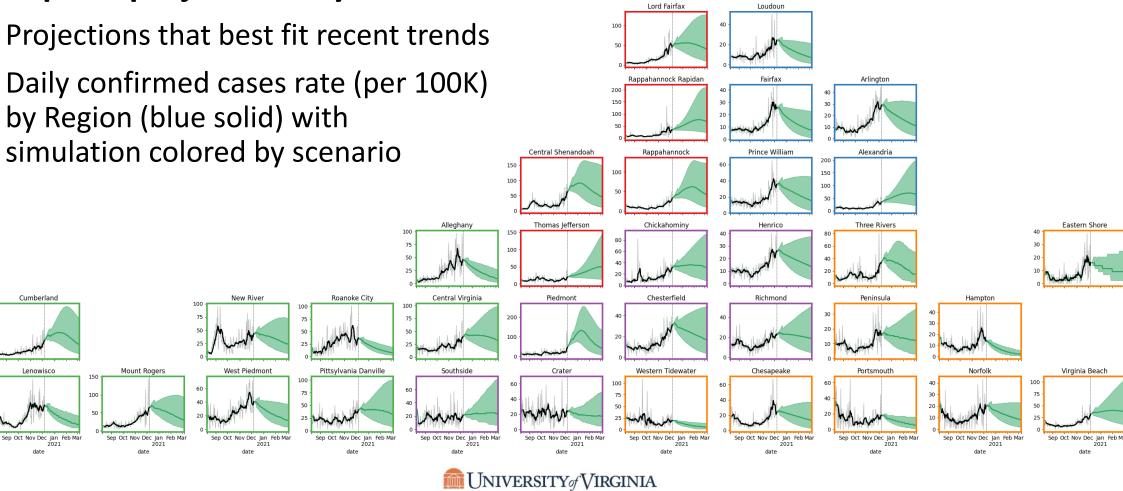
100

100

District Level Projections: Adaptive-MoreControl

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

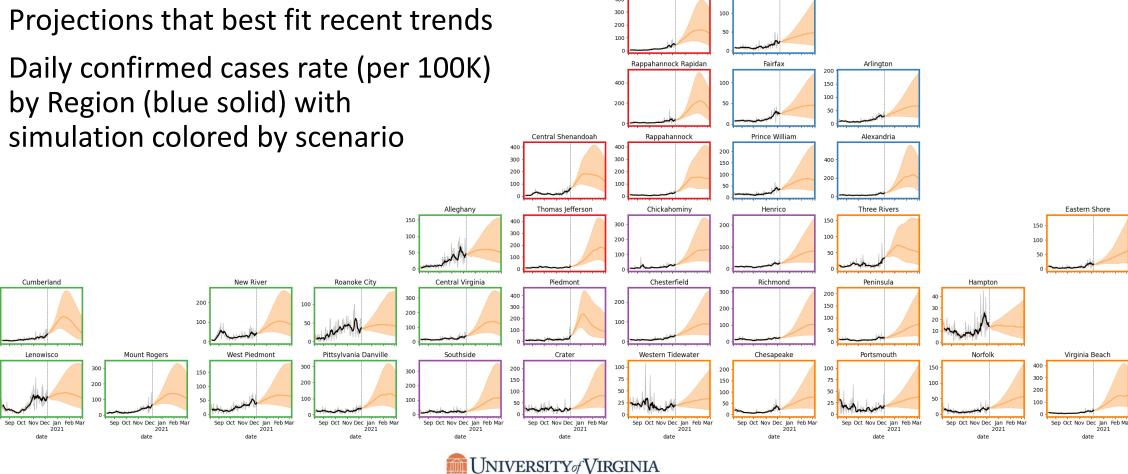


150 100

District Level Projections: Adaptive-LessControl

Adaptive projections by District

- Projections that best fit recent trends
- by Region (blue solid) with simulation colored by scenario



Lord Fairfax

Loudoun

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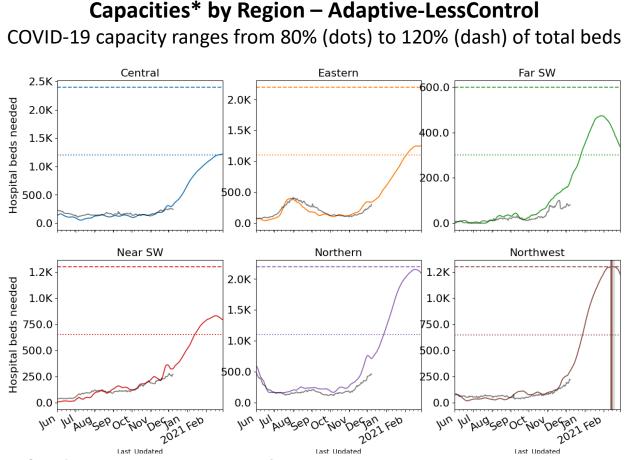
200

100

150

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Hospital Demand and Bed Capacity by Region



Week Ending	Adaptive	Adaptive- LessControl	
11/29/20	17,405	17,405	
12/06/20	16,823	16,823	
12/13/20	19,811	19,826	
12/20/20	21,987	23,587	
12/27/20	24,055	29,112	
1/3/20	26,459	34,782	
1/10/20	28,636	40,422	
1/17/20	30,276	45,443	
1/24/20	31,391	49,562	
1/31/20	31,941	52,748	
2/7/20	32,081	54,887	
2/14/20	31,753	56,209	

If Adaptive-LessControl scenario persists:

- All regions approach initial bed capacity this winter, starting mid-Dec through mid-Feb
- Over bed capacity possible in Northwest in mid February.
- * Assumes average length of stay of 8 days 9-Dec-20

Key Takeaways

Projecting future cases precisely is impossible and unnecessary. Even without perfect projections, we can confidently draw conclusions:

- Case growth in Virginia seems to have rebounded following Thanksgiving holiday
- VA mean weekly incidence (38/100K) up (from 28) as national surge also rebounds (to 68/100K from 53/100K).
- Lingering effects from reporting delays and overall testing interruptions continue to impact projections, which are mostly up.
- Recent updates:
 - Ensemble of statistical and Machine Learning models integrated with Adaptive to guide projections
 - Return projection trend window to 2 weeks to minimize holiday effects
 - Planning scenarios remain at Dec 10th and case ascertainment rates updated but are unchanged
- The situation is changing rapidly. Models will be updated regularly.



References

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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. <u>https://github.com/NSSAC/PatchSim</u> (Accessed on 04/10/2020).

Virginia Department of Health. COVID-19 in Virginia. <u>http://www.vdh.virginia.gov/coronavirus/</u> (Accessed on 04/10/2020)

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

Google. COVID-19 community mobility reports. <u>https://www.google.com/covid19/mobility/</u>

Biocomplexity page for data and other resources related to COVID-19: <u>https://covid19.biocomplexity.virginia.edu/</u>



Questions?

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Supplemental Slides



Estimating Daily Reproductive Number

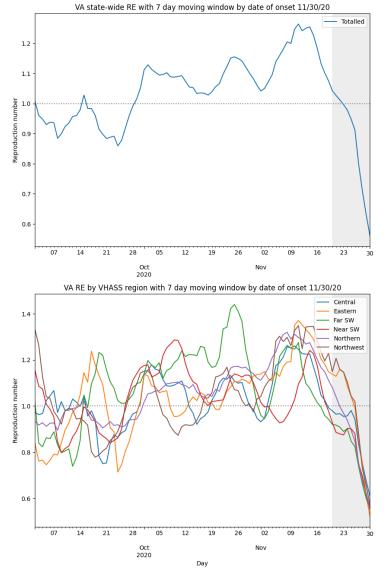
Nov 30th Estimates

Region	Date of Onset R _e	Date Onset Diff Last Week	
State-wide	1.030	-0.222	
Central	0.963	-0.264	
Eastern	1.121	-0.188	
Far SW	0.908	-0.166	
Near SW	0.922	-0.311	
Northern	1.069	-0.207	
Northwest	1.135	-0.205	

Methodology

- Wallinga-Teunis method (EpiEstim¹) 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, <u>https://doi.org/10.1093/aje/kwt133</u>

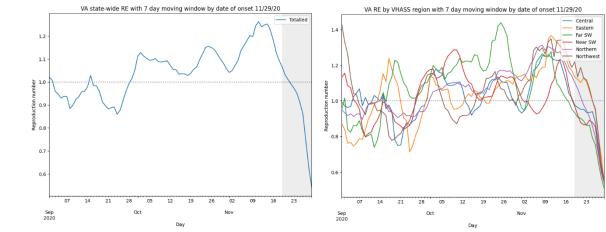


Estimating Daily Reproductive Number

Region	Date of Onset R _e	Confirmed Date R _e	Range of R_e	Onset Diff Last Week	Confirmed Diff Last Week	Range of Diff from Last Week
State-wide	1.030	1.234	[1.03-1.234]	-0.222	0.143	[-0.22 to 0.14]
Central	0.963	1.064	[0.963-1.064]	-0.264	-0.035	[-0.26 to -0.03]
Eastern	1.121	1.369	[1.121-1.369]	-0.188	0.332	[-0.19 to 0.33]
Far SW	0.908	1.022	[0.908-1.022]	-0.166	-0.204	[-0.17 to -0.2]
Near SW	0.922	1.225	[0.922-1.225]	-0.311	0.307	[-0.31 to 0.31]
Northern	1.069	1.280	[1.069-1.28]	-0.207	0.125	[-0.21 to 0.12]
Northwest	1.135	1.399	[1.135-1.399]	-0.205	0.192	[-0.21 to 0.19]

Nov 20th Estimates

9-Dec-20

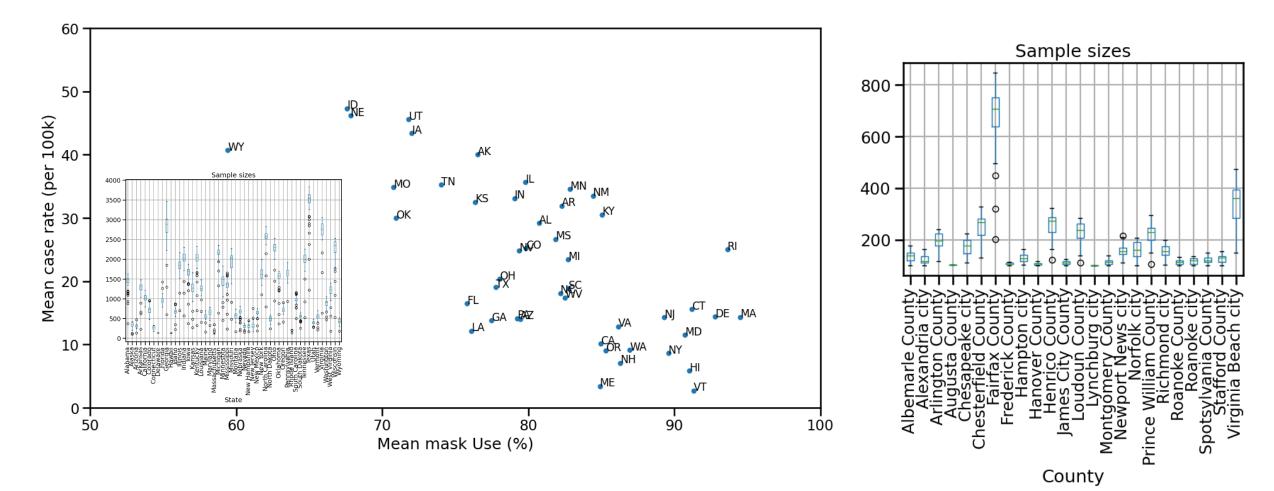


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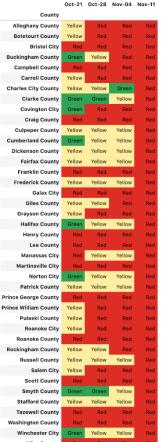
Mask usage sample sizes



Test positivity across VA counties

- CMS weekly summary (used for guiding nursing homes testing protocol)
- Data: COVID-19 Electronic Lab Reporting (CELR); HHS Unified Testing Dataset;
- County level testing counts and test positivity rates for RT-PCR tests.
 - Green: Test positivity <5.0% or with <20 tests in past 14 days
 - Yellow: Test positivity 5.0%-10.0% or with <500 tests and <2000 tests/100k and >10% positivity over 14 days
 - Red: >10.0% and not meeting the criteria for "Green" or "Yellow"

https://data.cms.gov/stories/s/q5r5-gjyu





Red on Oct 21 (4-week back)

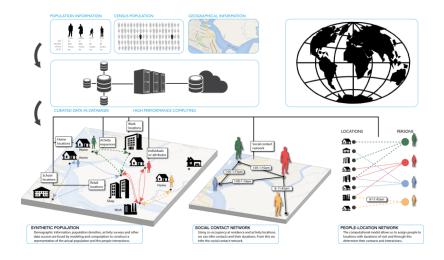
Red on Nov 11 (latest)

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Agent-based Model (ABM)

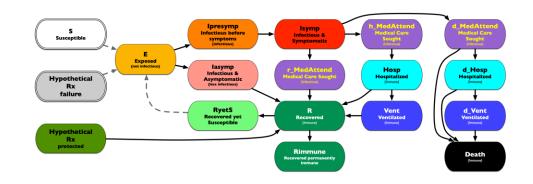
EpiHiper: Distributed network-based stochastic disease transmission simulations

- 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

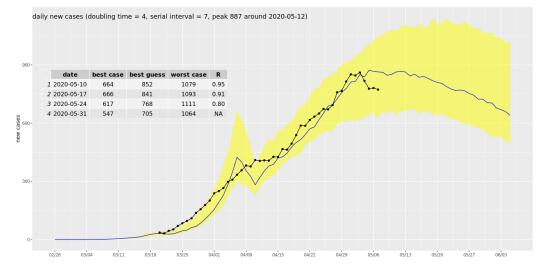
- Literature based probabilities of outcomes with appropriate delays
- Varying levels of infectiousness
- Hypothetical treatments for future developments

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ABM Social Distancing Rebound Study Design

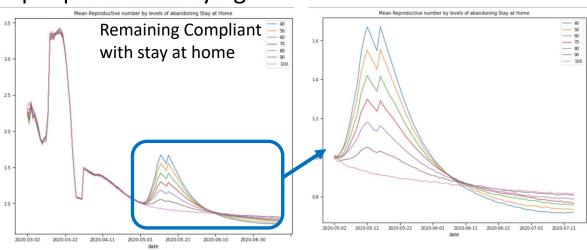
Study of "Stay Home" policy adherence

- 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



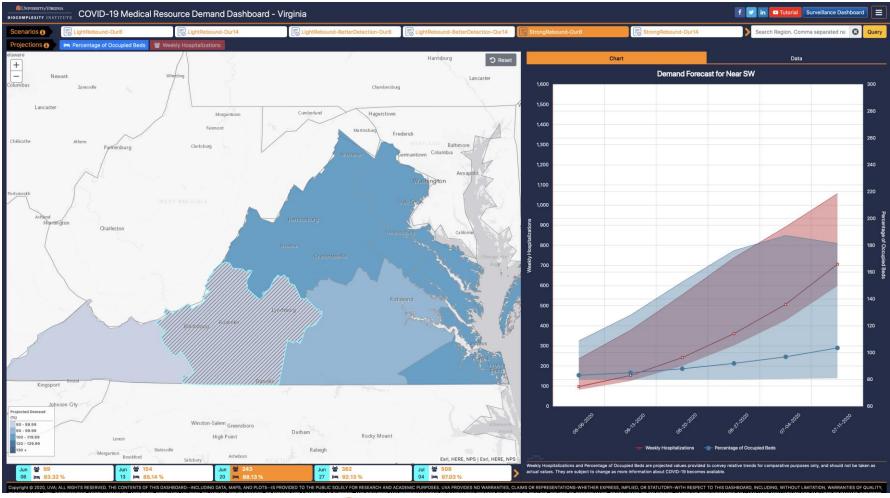
Impacts on Reproductive number with release

- 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 1/6th return to pre-pandemic levels

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Medical Resource Demand Dashboard

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



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