

Real-Time Pandemic Planning, Prediction and Response



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References

Papers:

- *CACM 2014*: Computational Epidemiology
- *Nature 2004*: Modelling disease outbreaks in realistic urban social networks.
- *PNAS 2008*: Modeling targeted layered containment of an influenza pandemic in the United States.
- *PNAS 2014*: Opinion: Mathematical models: A key tool for outbreak response.

Tutorials (KDD'14, AAI'16, ICSB'17): <https://covid19.biocomplexity.virginia.edu/publications>

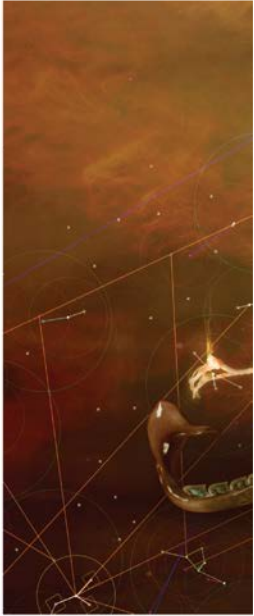
A Tutorial on Generating Synthetic Populations for Social Modeling, [IJCAI 2016](#), & [AAMAS 2016](#).

COVID-19 resource page: <https://covid19.biocomplexity.virginia.edu/>

New NSF Expeditions Project: <https://computational-epidemiology.org>

New NSF Virtual Organization Project: <https://prepare-vo.org>

review articles



DOI:10.1145/2493852.2493871

The challenge of developing and using computer models to understand and control the diffusion of disease through populations.

BY MADHAV MARATHE AND ANIL KUMAR S. VULLIKANTI

Computational Epidemiology

AN EPIDEMIC IS said to arise in a community or region when cases of an illness or other health-related events occur in excess of normal expectancy. Epidemics are considered to have influenced significant historical events, including the plagues in Roman times and Middle Ages, the fall of the Han empire in the 3rd century in China, and the defeat of the Aztecs in the 1500s, due to a smallpox outbreak.⁹ The 1918 flu pandemic in the U.S. was responsible for more deaths than those due to World War I. The last 50 years have seen epidemics caused by HIV/AIDS, SARS, and influenza-like illnesses. Despite significant medical advances, according to the World Health Organization (WHO), infectious diseases account for more than 13 million deaths a year.⁴⁴

Societal interest in controlling outbreaks is probably just as old as the diseases themselves. Interestingly, it appears the Indians and Chinese knew the idea of variolation to control smallpox as early as the 8th century A.D. Epidemiology is a formal branch of science focusing on the study of space-time patterns of illness in a population and the factors that contribute to these patterns. It plays an essential role in public health by

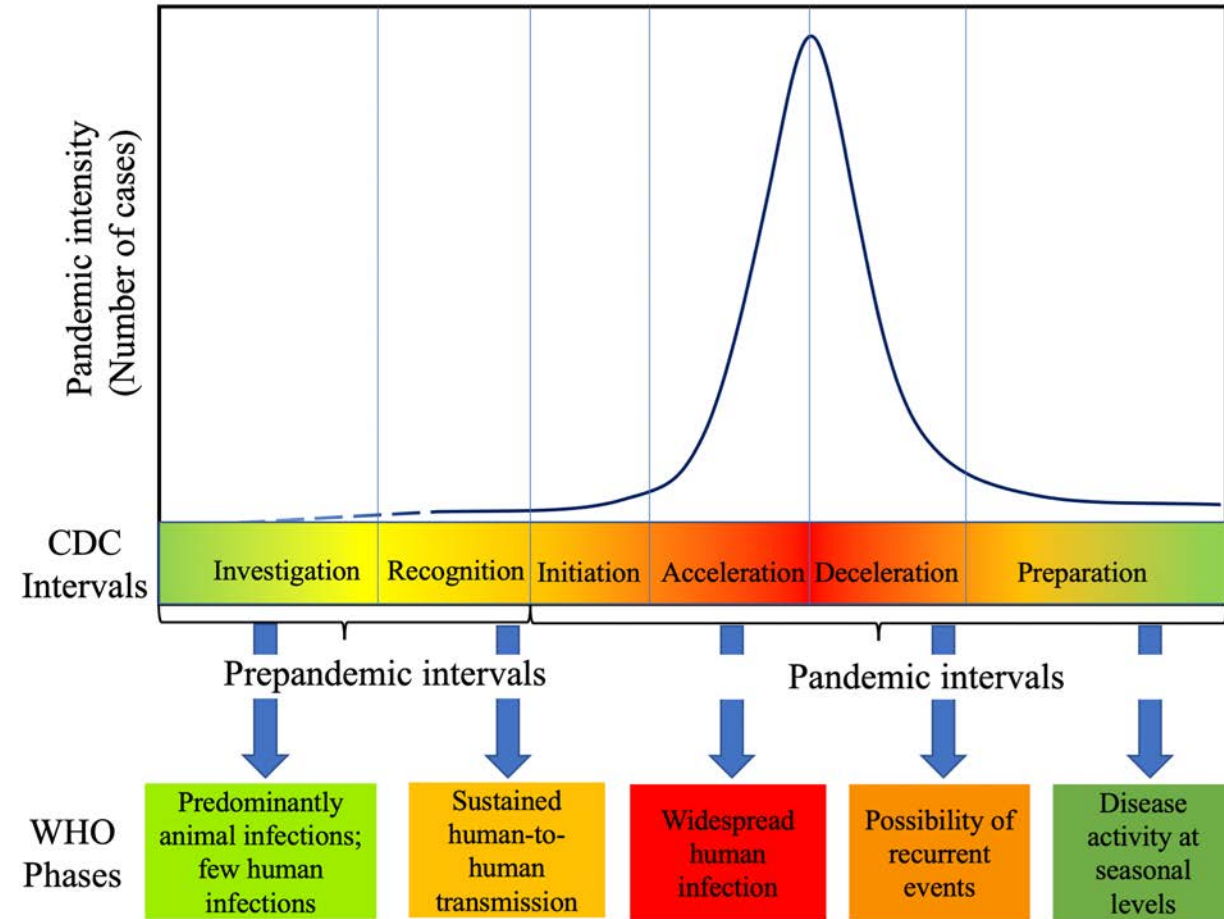
key insights

- Controlling and responding to future pandemics will be challenging due to a number of emerging global trends including increased and denser urbanization, increased local as well as global travel, and a generally older and immuno-compromised population.
- Public health epidemiology is a complex system problem. Epidemics, social-contact networks, individual and collective behavior, and public policies coevolve during a pandemic—a system-level understanding must represent these components and their coevolution.
- Mathematical and computational models of social networks and epidemic spread and methods to analyze them are critical in public health epidemiology.
- Advances in computing, big data, and computational thinking have created entirely new opportunities to support real-time epidemiology.

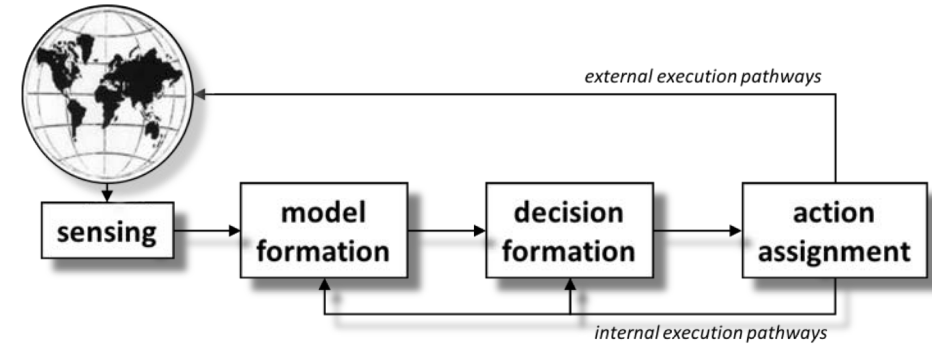
88 COMMUNICATIONS OF THE ACM JULY 2013 | VOL. 56 | NO. 7

Modeling for integrated reasoning about situations & actions

- **VISION:** Real-time Computational Epidemiology
- **GOAL:** Build a flexible suite of models that go beyond prediction and in *real-time*
 - *Synthesize* available data to produce consistent and meaningful representations of the underlying system
 - *Provide* a range of interpretations of incoming measurements
 - *Evaluate* a range of response actions and behaviors
 - *Monitor* effect of intervention responses
 - *Coordinate* understanding among diverse stakeholders
 - *Usable* by analysts and not just computing experts

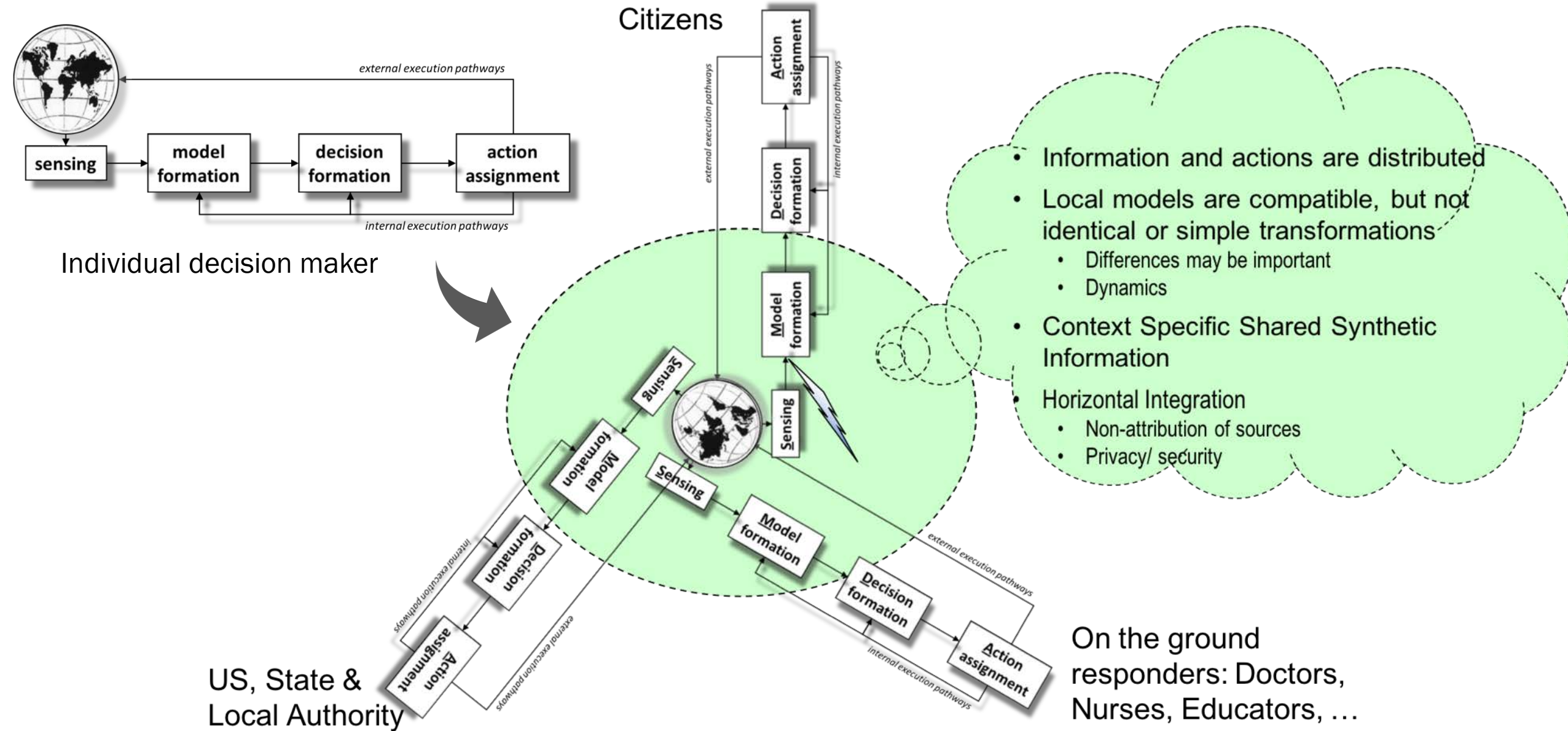


Human-centered model-based distributed decision making

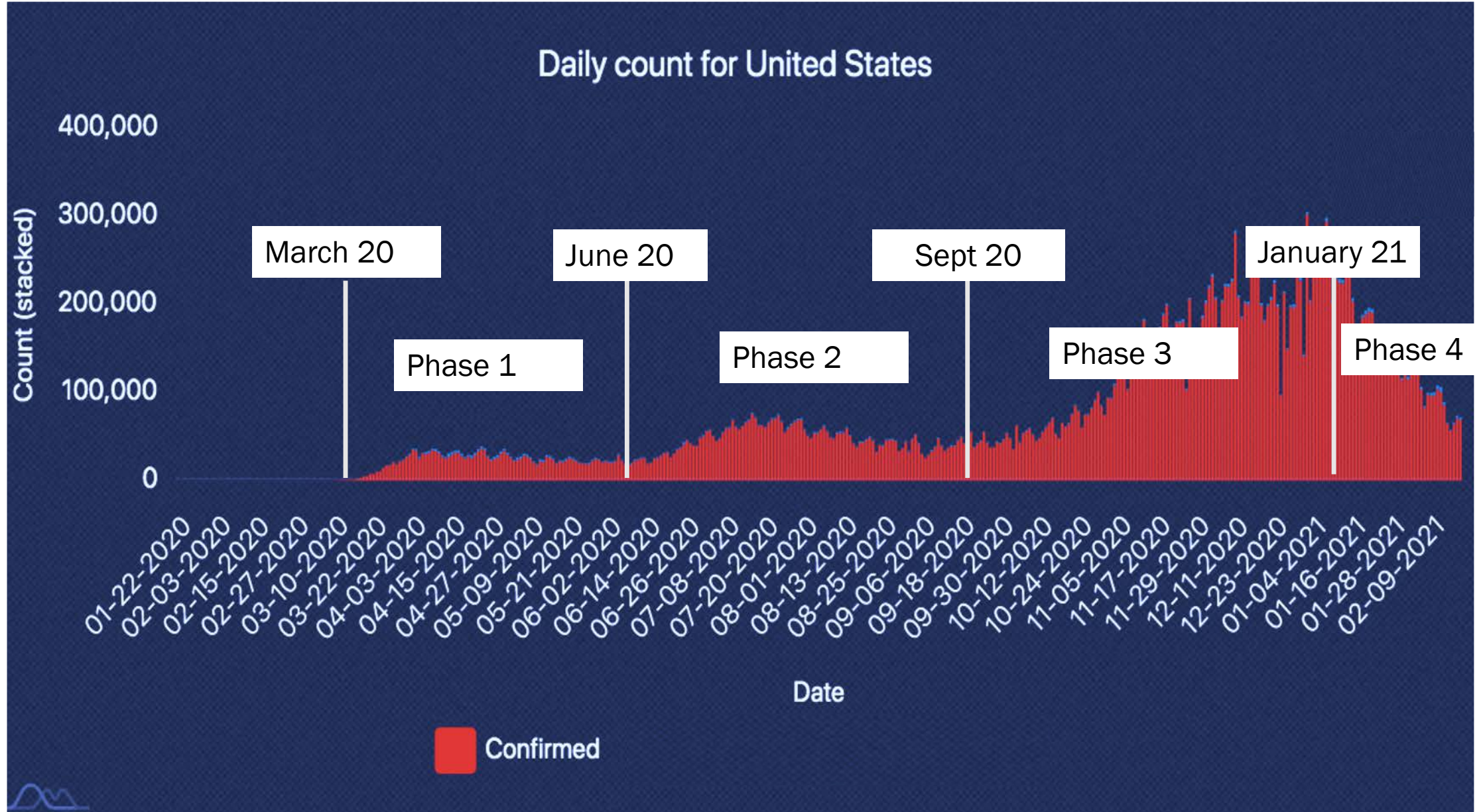


Individual decision maker

Human-centered model-based distributed decision making



The phases of COVID-19 pandemic



Phases and questions

	Phase 1
Questions	<ul style="list-style-type: none">• Predict risk of importation• Infer disease parameters• Evaluate impacts of social distancing
Data Needs	<ul style="list-style-type: none">• Clinical studies on disease outcomes• Global multimodal traffic• Testing and case surveillance
Research Highlights	<ul style="list-style-type: none">• Model projections (<i>Imperial, IHME, Northeastern</i>)• Undocumented infections (<i>Columbia</i>)

Phases and questions

	Phase 1	Phase 2
Questions	<ul style="list-style-type: none">• Predict risk of importation• Infer disease parameters• Evaluate impacts of social distancing	<ul style="list-style-type: none">• Predict risk of resurgence• Infer the role of mobility & mask use• Evaluate the efficacy of contact tracing
Data Needs	<ul style="list-style-type: none">• Clinical studies on disease outcomes• Global multimodal traffic• Testing and case surveillance	<ul style="list-style-type: none">• Cross-scale intervention measures• Local mobility and mixing• Behavior and compliance
Research Highlights	<ul style="list-style-type: none">• Model projections (<i>Imperial, IHME, Northeastern</i>)• Undocumented infections (<i>Columbia</i>)	<ul style="list-style-type: none">• Collaborative ensemble (<i>UMass</i>)• Symptom Surveys (<i>CMU</i>)• Resource allocation (<i>Yale, UT Austin</i>)• Optimal testing (<i>AIM</i>)

Phases and questions

	Phase 1	Phase 2	Phase 3
Questions	<ul style="list-style-type: none"> • Predict risk of importation • Infer disease parameters • Evaluate impacts of social distancing 	<ul style="list-style-type: none"> • Predict risk of resurgence • Infer the role of mobility & mask use • Evaluate the efficacy of contact tracing 	<ul style="list-style-type: none"> • Predict medical resource demand • Infer effect of seasonality • Evaluate K-12, colleges reopening
Data Needs	<ul style="list-style-type: none"> • Clinical studies on disease outcomes • Global multimodal traffic • Testing and case surveillance 	<ul style="list-style-type: none"> • Cross-scale intervention measures • Local mobility and mixing • Behavior and compliance 	<ul style="list-style-type: none"> • Hospital occupancy statistics • Weather and seasonal factors • College reopening plans
Research Highlights	<ul style="list-style-type: none"> • Model projections (<i>Imperial, IHME, Northeastern</i>) • Undocumented infections (<i>Columbia</i>) 	<ul style="list-style-type: none"> • Collaborative ensemble (<i>UMass</i>) • Symptom Surveys (<i>CMU</i>) • Resource allocation (<i>Yale, UT Austin</i>) • Optimal testing (<i>AIM</i>) 	<ul style="list-style-type: none"> • Digital contact tracing study (<i>Oxford</i>) • Susceptibility and climate (<i>Princeton</i>) • Mobility reduction impact (<i>Stanford</i>)

Phases and questions

	Phase 1	Phase 2	Phase 3	Phase 4
Questions	<ul style="list-style-type: none"> • Predict risk of importation • Infer disease parameters • Evaluate impacts of social distancing 	<ul style="list-style-type: none"> • Predict risk of resurgence • Infer the role of mobility & mask use • Evaluate the efficacy of contact tracing 	<ul style="list-style-type: none"> • Predict medical resource demand • Infer effect of seasonality • Evaluate K-12, colleges reopening 	<ul style="list-style-type: none"> • Predict variant dominance • Infer current seroprevalence • Evaluate vaccine rollouts
Data Needs	<ul style="list-style-type: none"> • Clinical studies on disease outcomes • Global multimodal traffic • Testing and case surveillance 	<ul style="list-style-type: none"> • Cross-scale intervention measures • Local mobility and mixing • Behavior and compliance 	<ul style="list-style-type: none"> • Hospital occupancy statistics • Weather and seasonal factors • College reopening plans 	<ul style="list-style-type: none"> • Seroprevalence surveys • Vaccine administration • Genomic sequencing
Research Highlights	<ul style="list-style-type: none"> • Model projections (<i>Imperial, IHME, Northeastern</i>) • Undocumented infections (<i>Columbia</i>) 	<ul style="list-style-type: none"> • Collaborative ensemble (<i>UMass</i>) • Symptom Surveys (<i>CMU</i>) • Resource allocation (<i>Yale, UT Austin</i>) • Optimal testing (<i>AIM</i>) 	<ul style="list-style-type: none"> • Digital contact tracing study (<i>Oxford</i>) • Susceptibility and climate (<i>Princeton</i>) • Mobility reduction impact (<i>Stanford</i>) 	<ul style="list-style-type: none"> • B.1.1.7 US prevalence (<i>Scripps</i>) • Vaccine by serostatus (<i>U. Colorado, Harvard</i>)

Challenges for End-to-End Planning, Prediction and Response

From Ecology to Biology to Epidemiology to Sociology

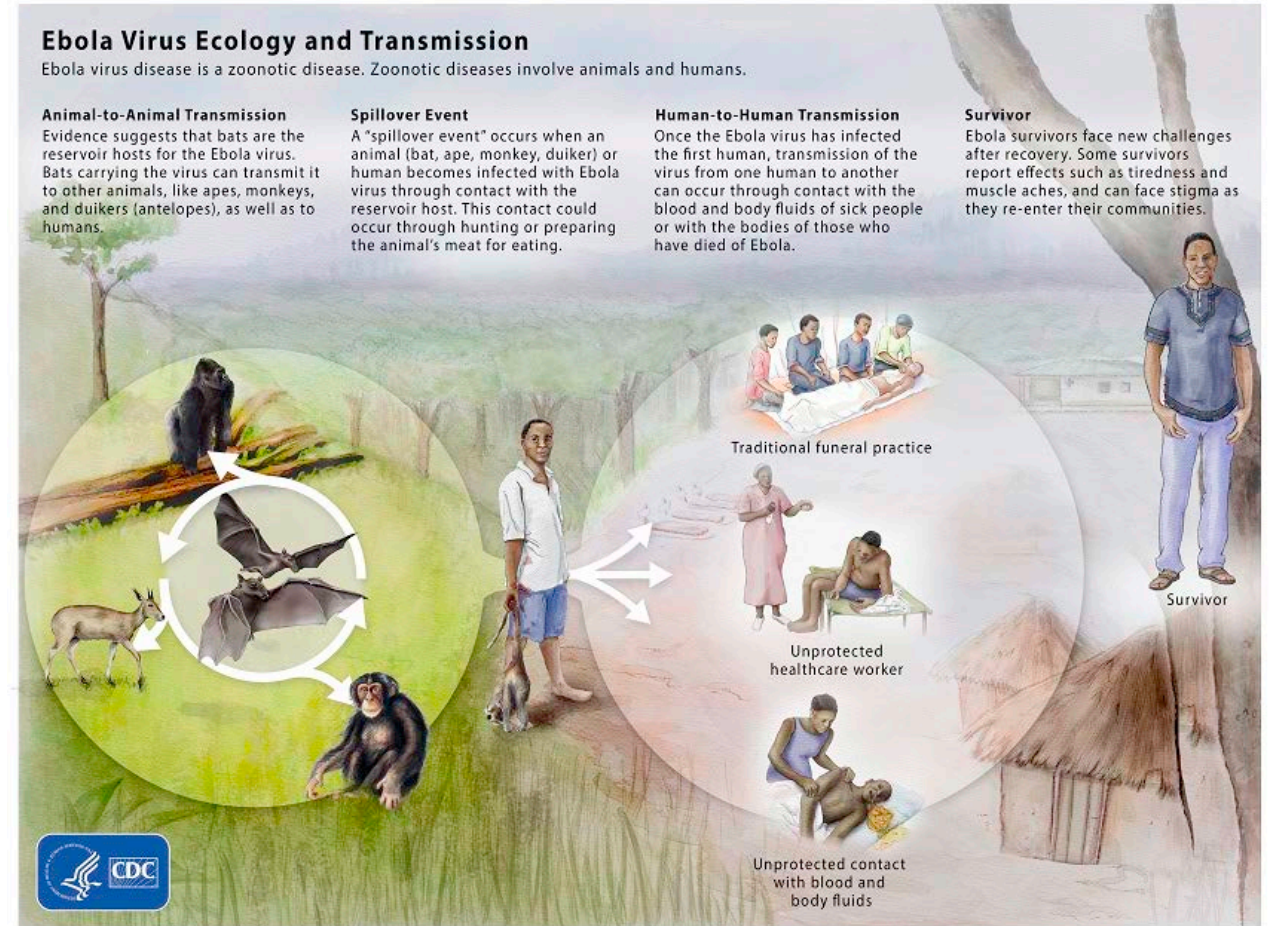
[Zoonosis] Model the evolution of the pathogen in space and time, including its interaction with humans and their immune systems, as well as the effects of interventions.

[Immunological Response] Understand immune response? What is the role of innate and adaptive immunity?

[Vaccine Development and Testing] How effective is the vaccine, how long will the immunity last?

[Viral evolution] How will the virus evolve under selection pressures

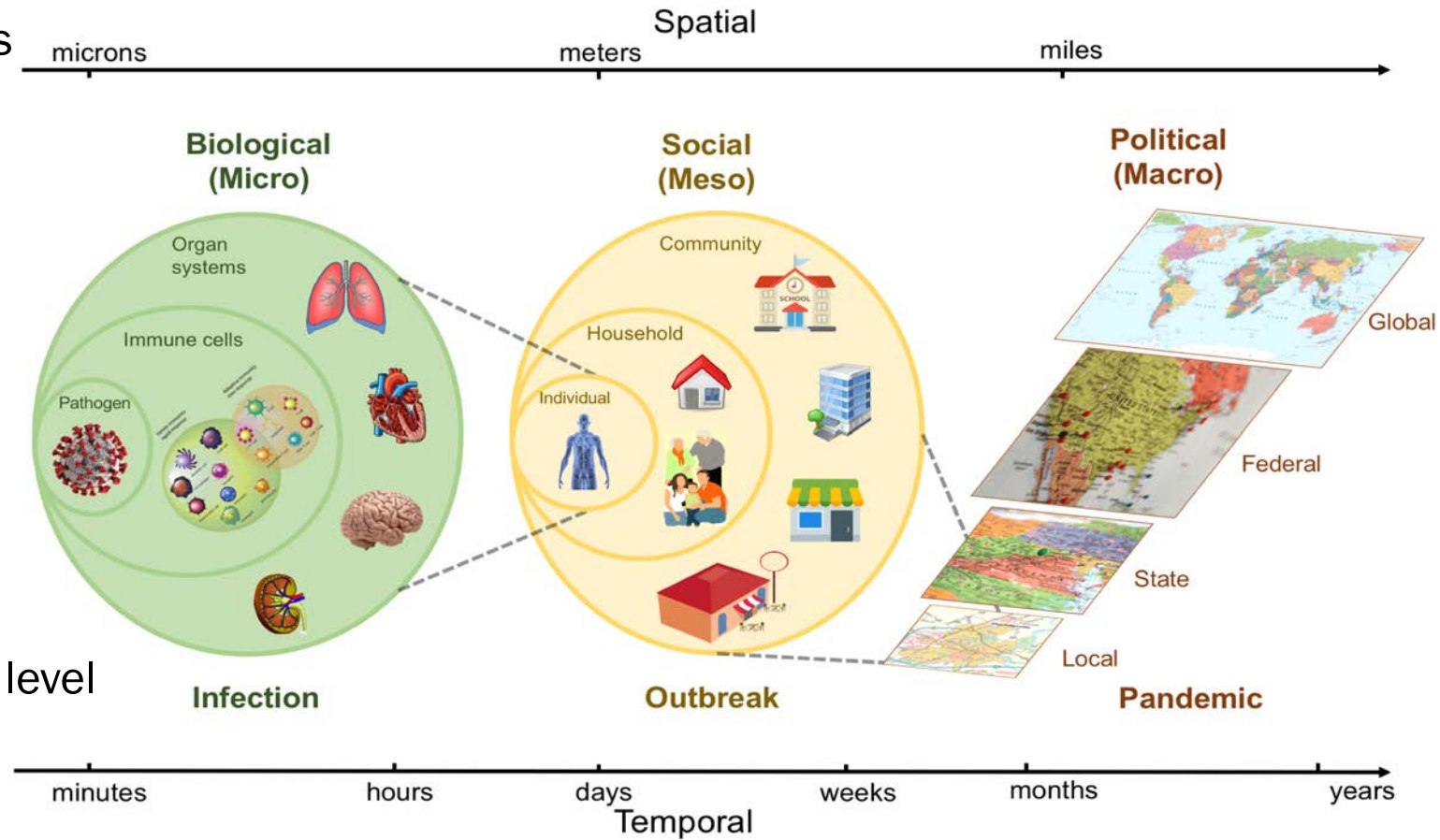
[Socio-economics] How will the pandemic interact with social, political and economic aspects



Need: Link models of viral evolution, human immune system, epidemic spread and socio-economic systems

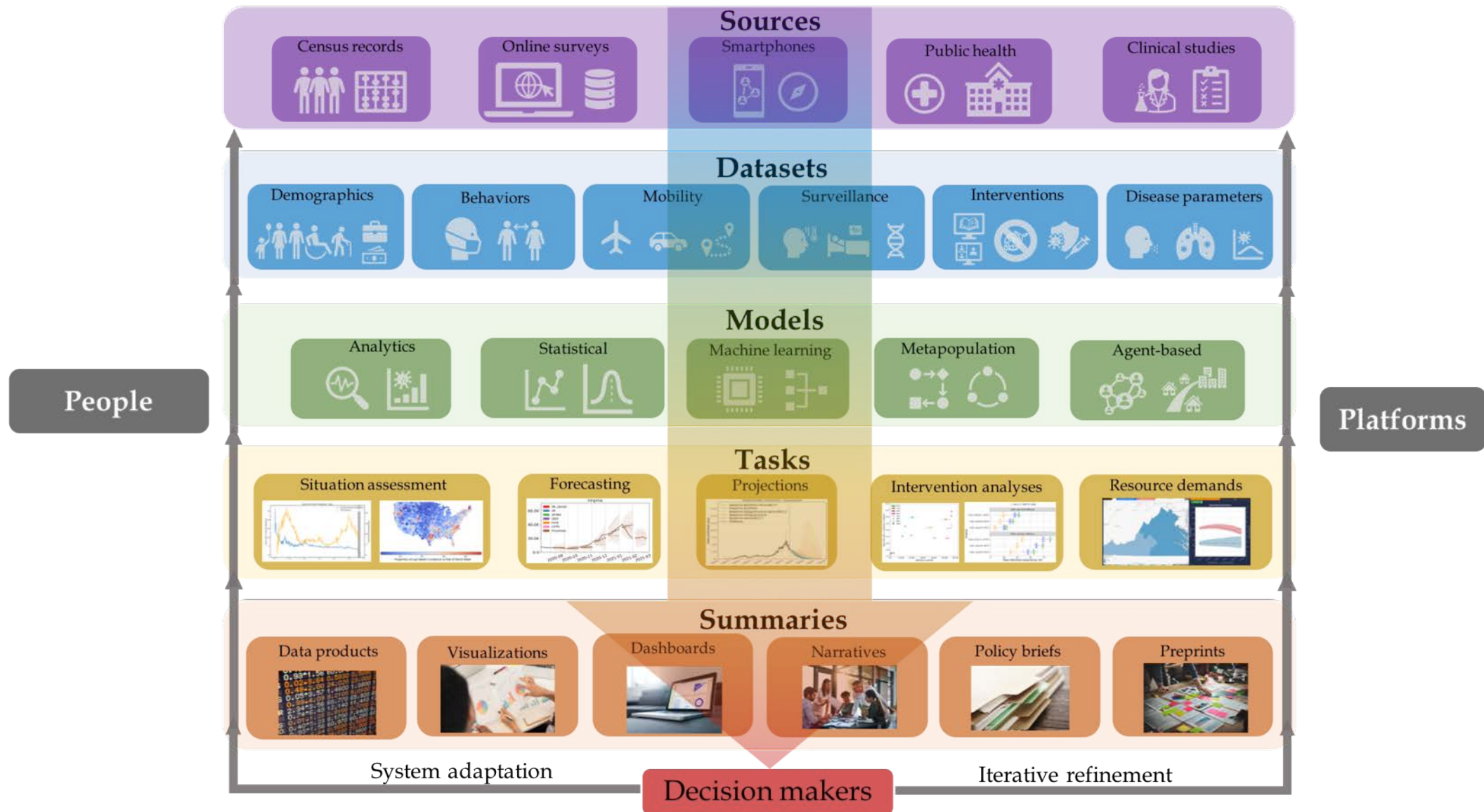
Spatial, temporal and social scales

- Processes unfolding across temporal scales
 - Within host disease progression
 - Between host transmission
 - Individual behavioral change
 - Community outbreak response
 - Public health control measures
 - Global response coordination
 - Seasonality and waning immunity
- Spatial and Social Interaction Scale
 - House (building), neighborhood (block group), city; state, country and region
 - Household, neighborhood and country level contact network
- Individual and collective behavior
 - Organizations (months), Community (days) and individual (days)



Need: Multi-scale, multi-theory, multi-level network representations and simulations

Data to decisions and communication



NEED: Context-specific, decentralized information integration and decision making across socio-political scales

Validation and uncertainty quantification (trust and adequacy)

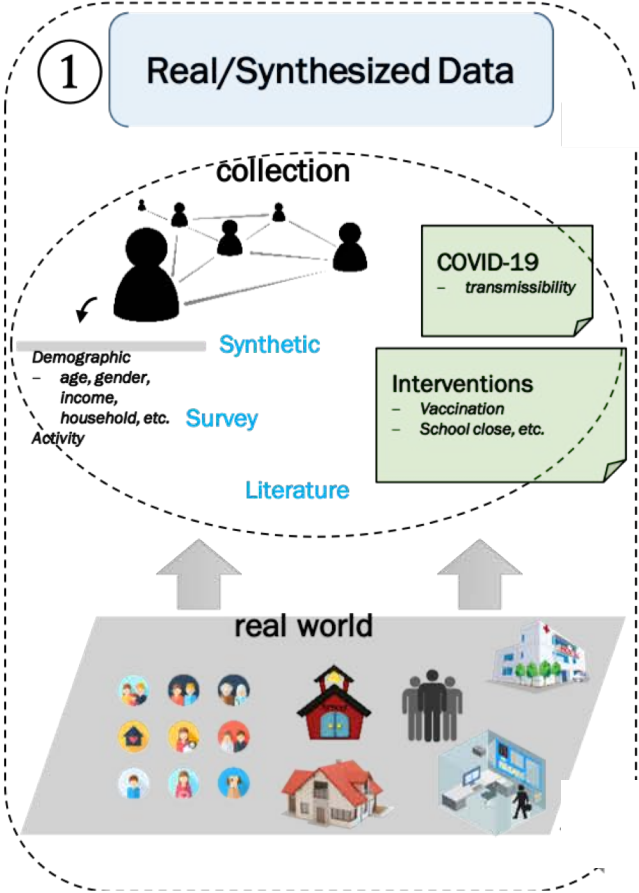
- Retrospective and predictive validity are not as useful in crisis situations when data is limited.
- External Validation
 - validate past predictions
 - update future projections
- Internal Validation
 - ensure structurally correct
- How do we gather and incorporate relevant data in real-time to:
 - actively learn when modeling assumptions cause models to fail to capture real-world dynamics?



Need: New approaches for V&V, UQ and model adaptation for co-evolving networks

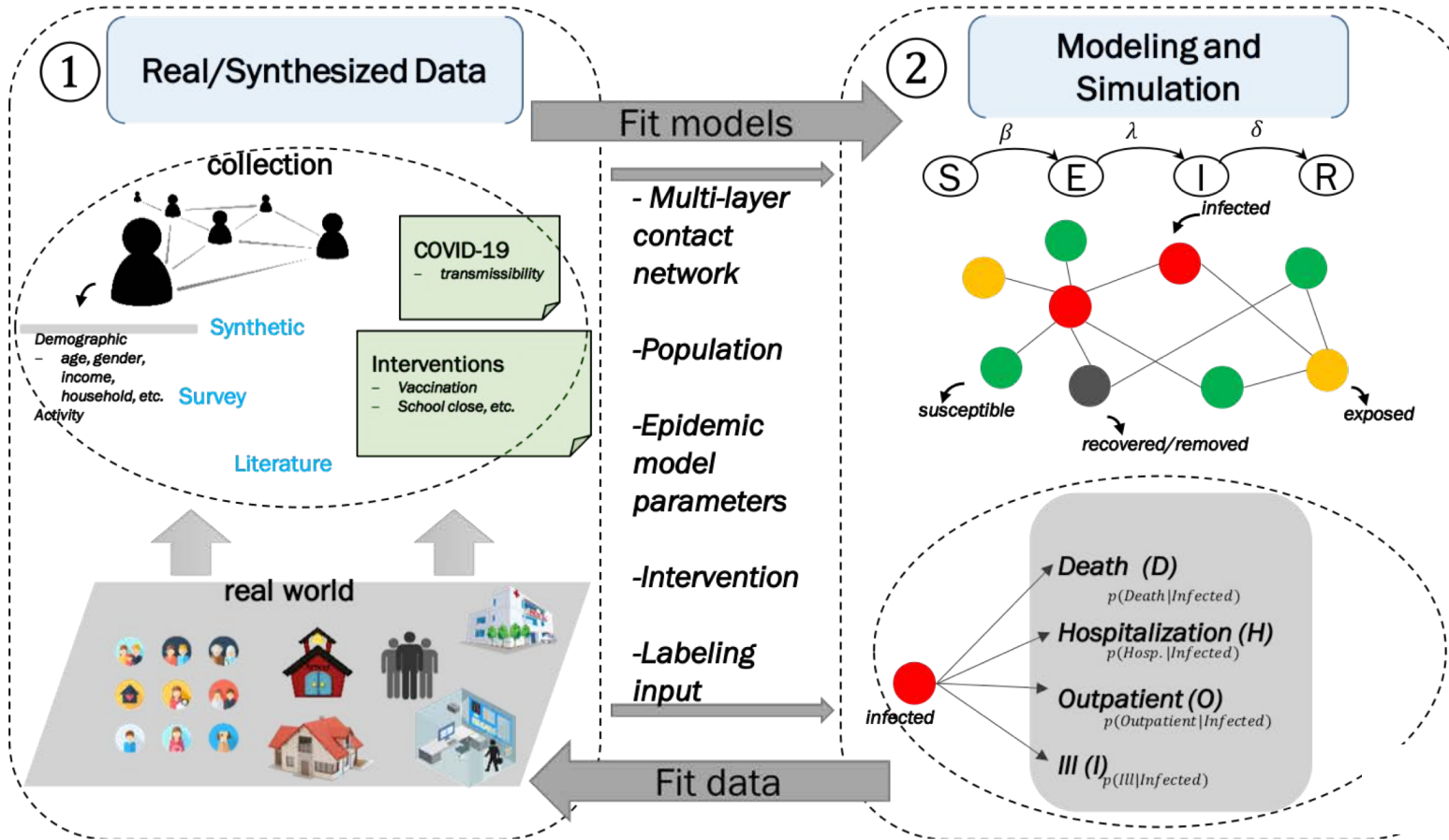
AI and High-Performance
Computing-Enabled Prediction and
Decision Informatics for Real-time
Epidemic Science

How do we do it: Data driven-networked epidemiology



Step 1: Build a digital twin of a city/country

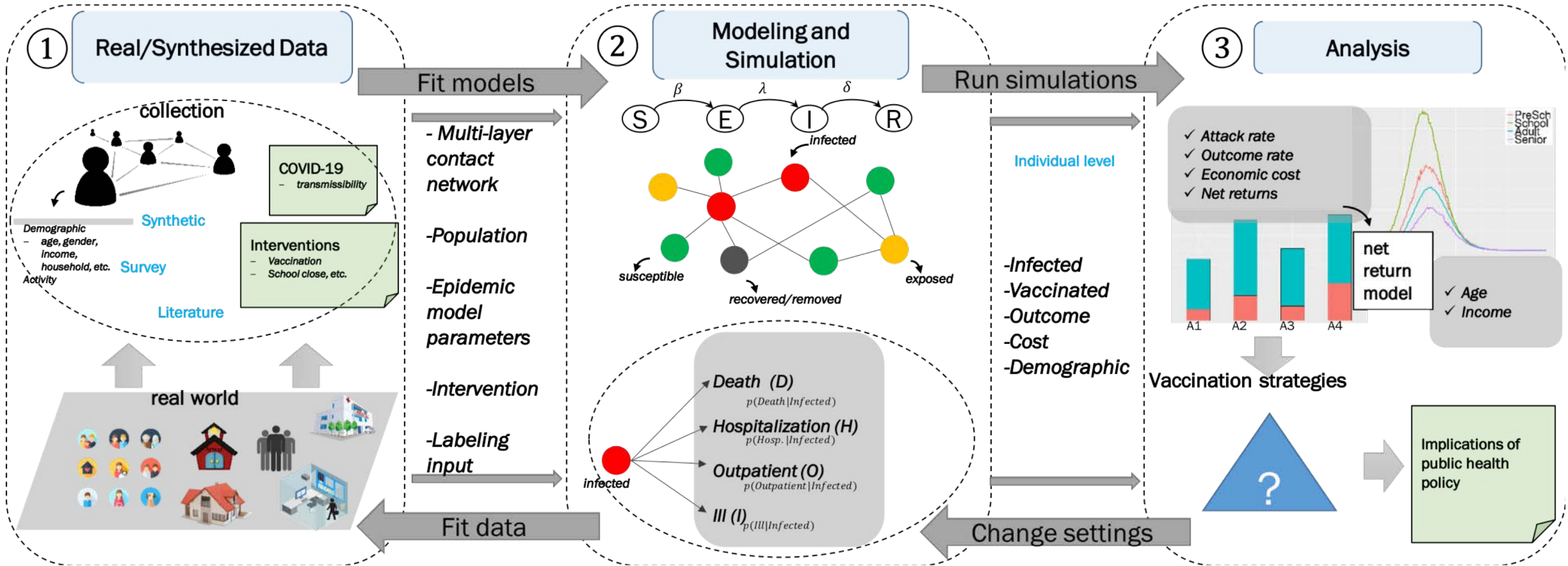
How do we do it: Data driven-networked epidemiology



Step 1: Build a digital twin of a city/country

Step 2: Build agent-based simulations of disease propagation

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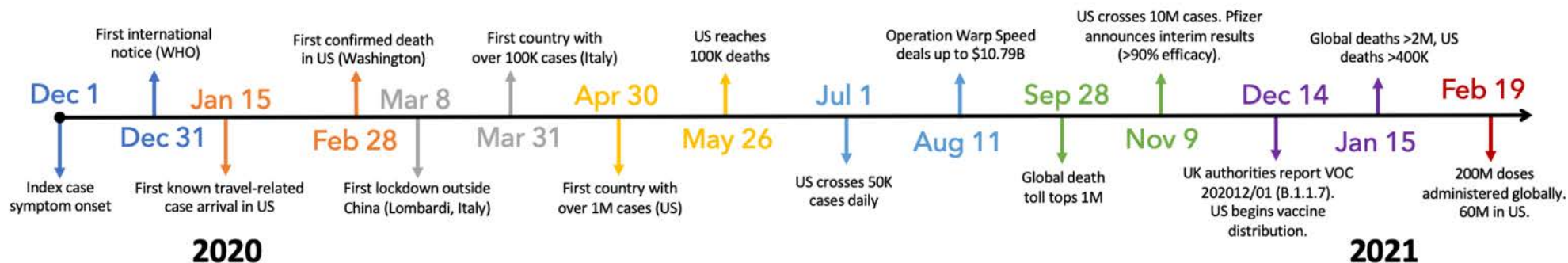
Step 3: Epidemiological workflows using simulation and ML

Who do we support

80+ scientists, staff, and students at UVA and collaborating institutions



50+ weeks of modeling briefs to public health agencies

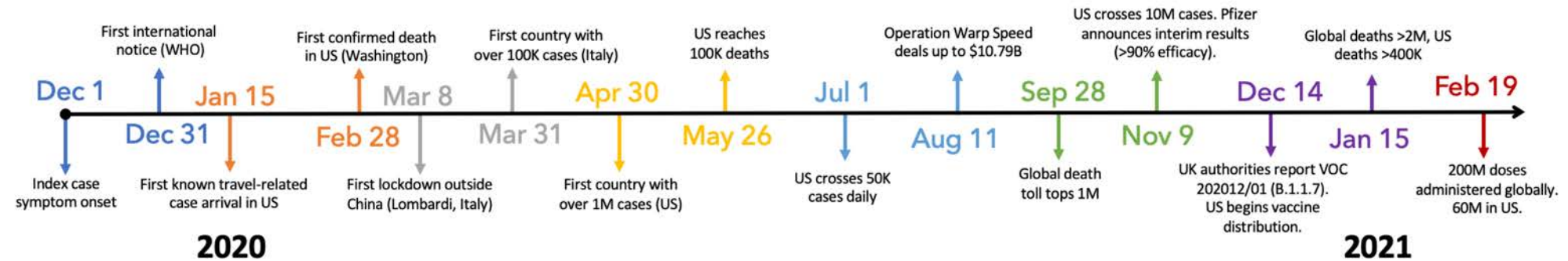


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Federal agencies



Commonwealth of Virginia

- Primary modeling for planning and response efforts



Local agencies

- Virginia Hospital and Healthcare Association (comprised of 27-member health systems and 110 community and specialty hospitals)



Weekly updates to state and federal agencies since February 2020

Network Systems
Science & Advanced
Computing
Biocomplexity Institute
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University of Virginia

Estimation of COVID-19 Impact in Virginia

February 17th, 2021

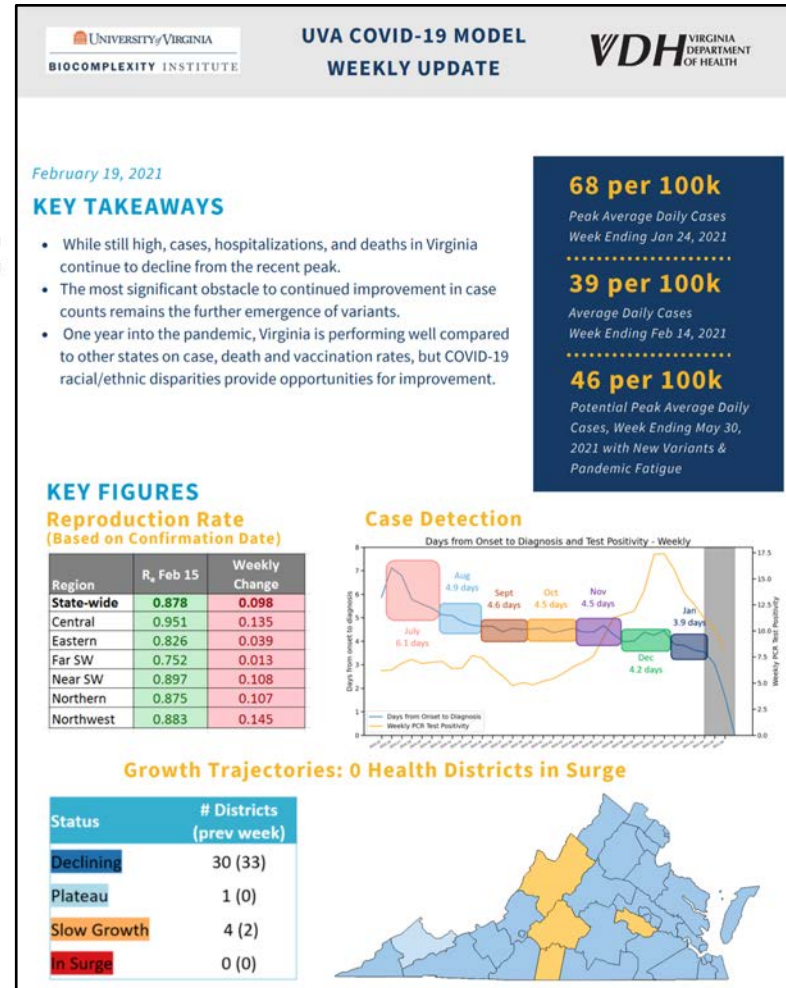
(data current to February 15th – 16th)

Biocomplexity Institute Technical report: TR 2021-020



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Situation assessment

Literature surveys

<https://www.vdh.virginia.gov/coronavirus/covid-19-data-insights/>

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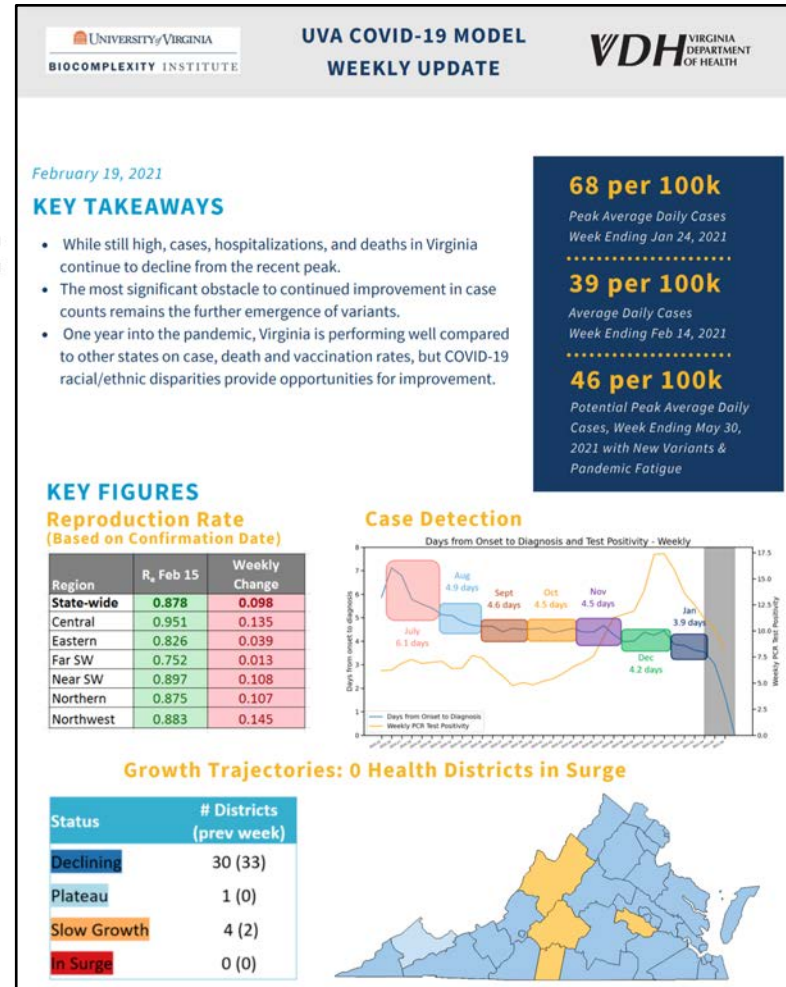
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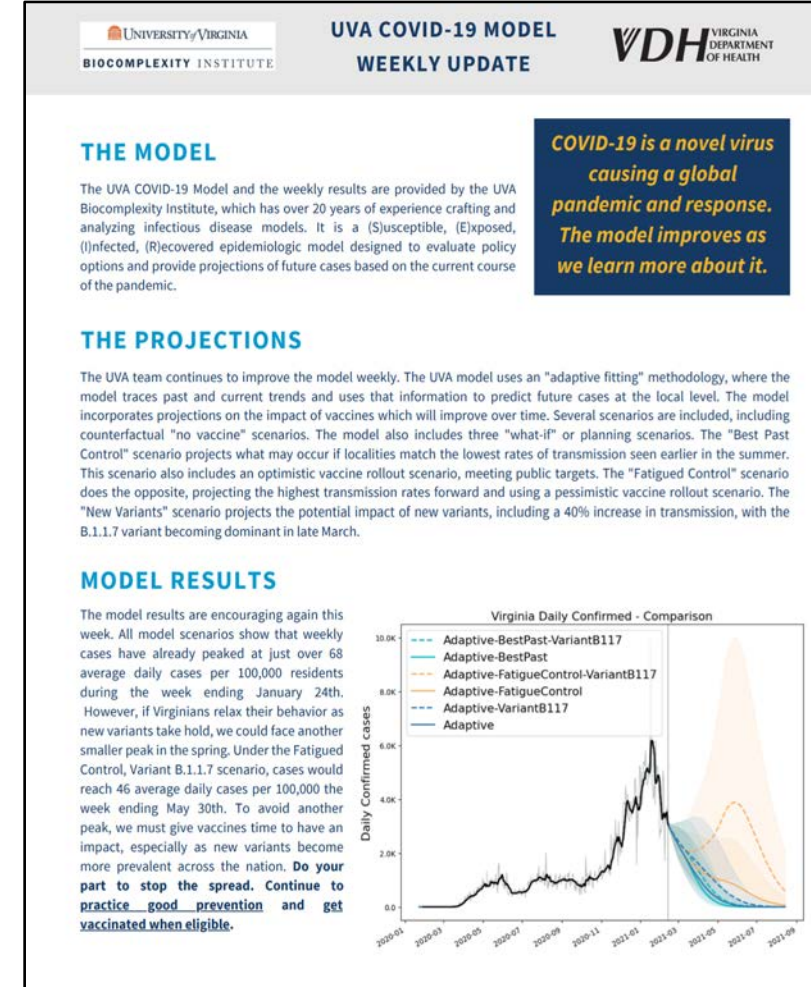
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Situation assessment

Literature surveys



Model projections

Narrative summaries

<https://www.vdh.virginia.gov/coronavirus/covid-19-data-insights/>

Building dashboards

Integrated Solutions:

Accessible and helpful to everyone across the globe, reliable, rich visualizations with easy-to-use interface

COVID-19 Surveillance Dashboard:

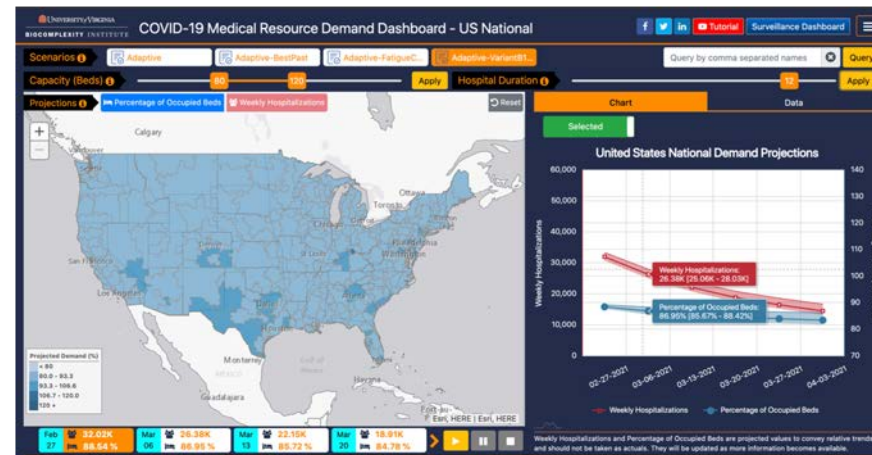
1.2 million users worldwide
3.7 million views since Feb 3, 2020

Spatial: 210 Countries, 450 States, 3,200 Counties (USA)

Temporal: Updated multiple times a day since Jan 22, 2020;

Medical Resource Demand Dashboard

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



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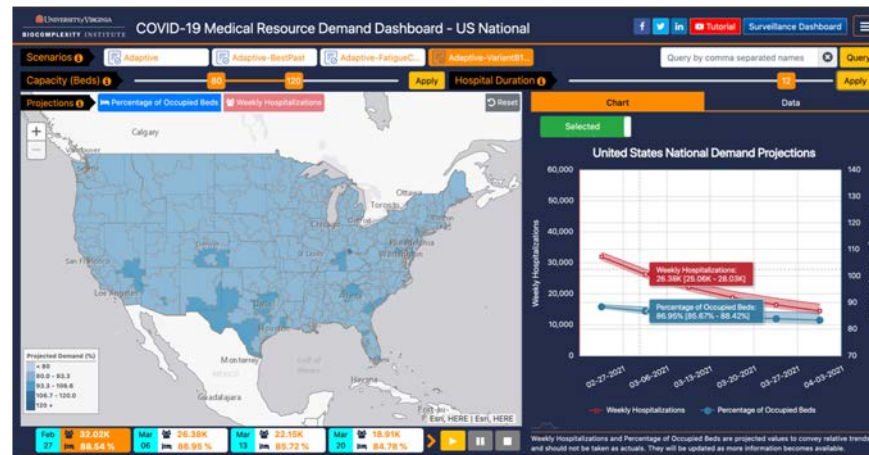
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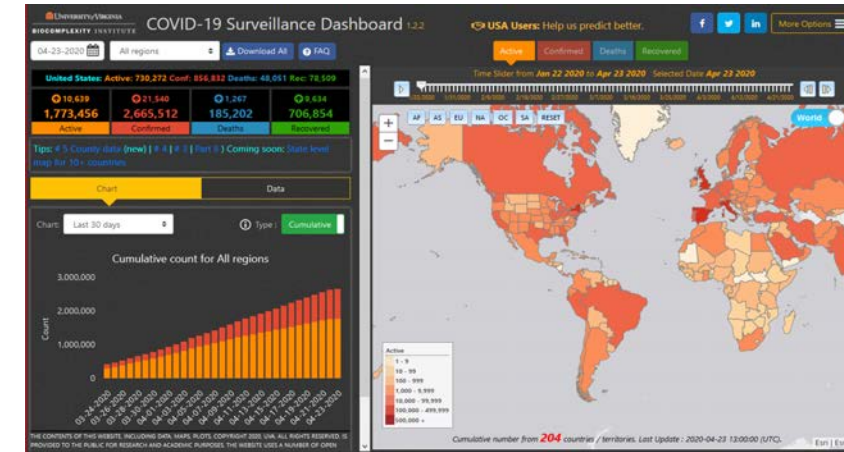
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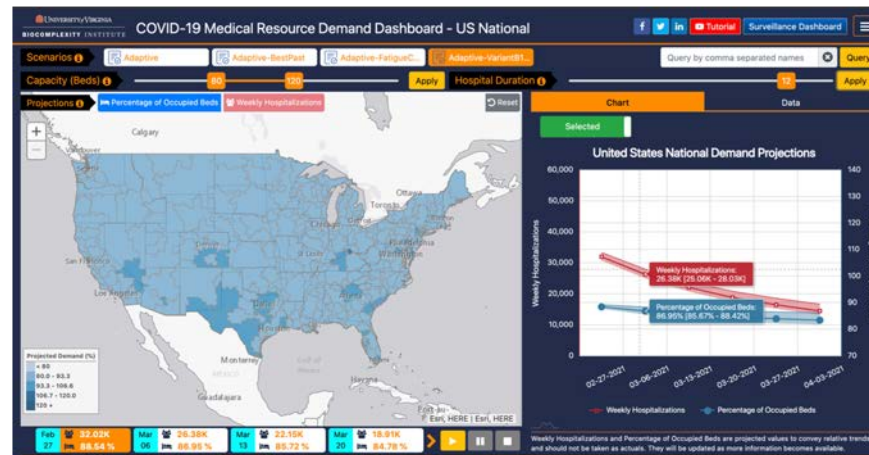
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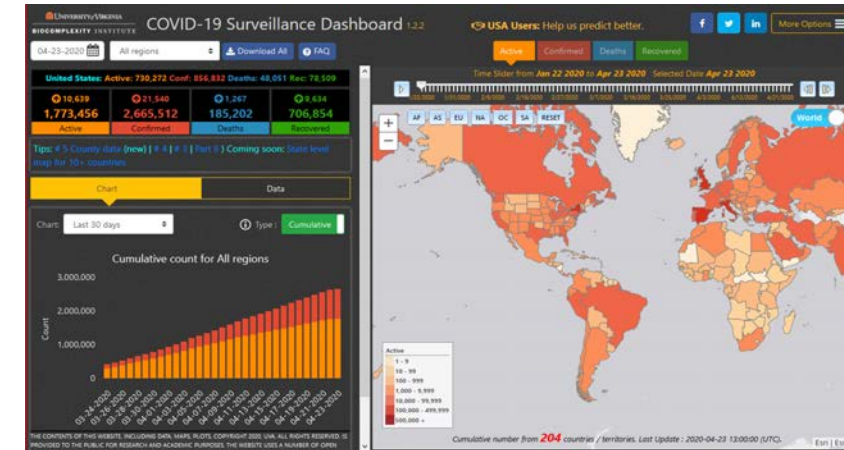
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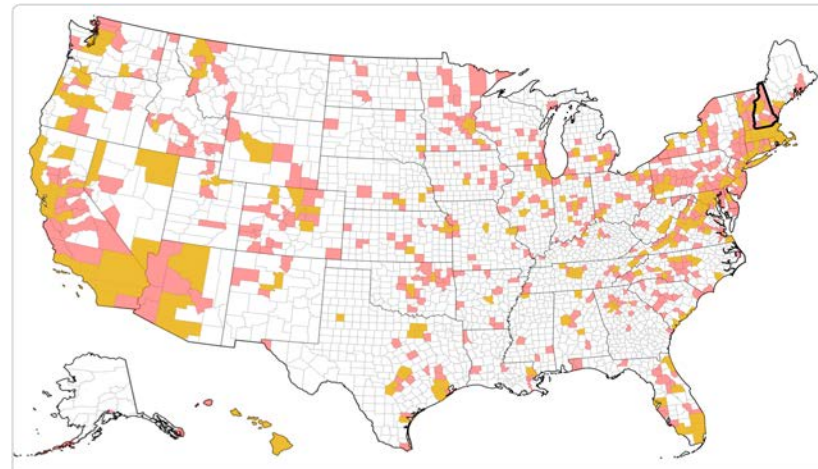
COVID-19 Surveillance Dashboard

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



Social distancing survey

<https://socialdistancing.stanford.edu/>



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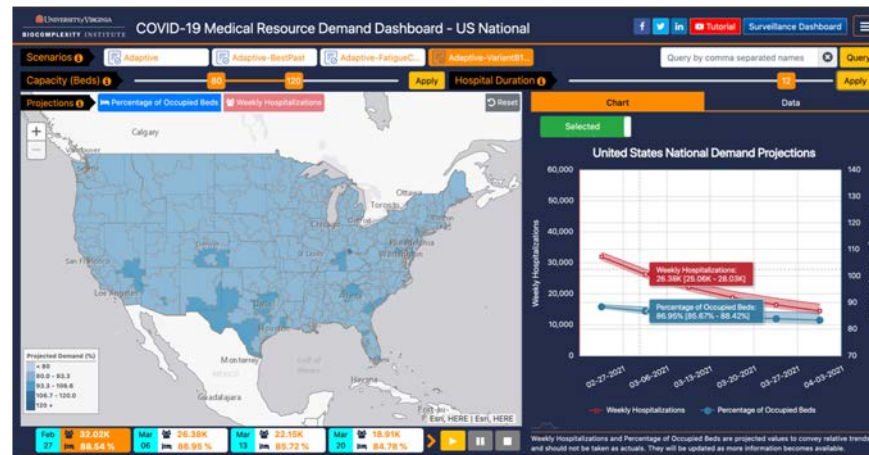
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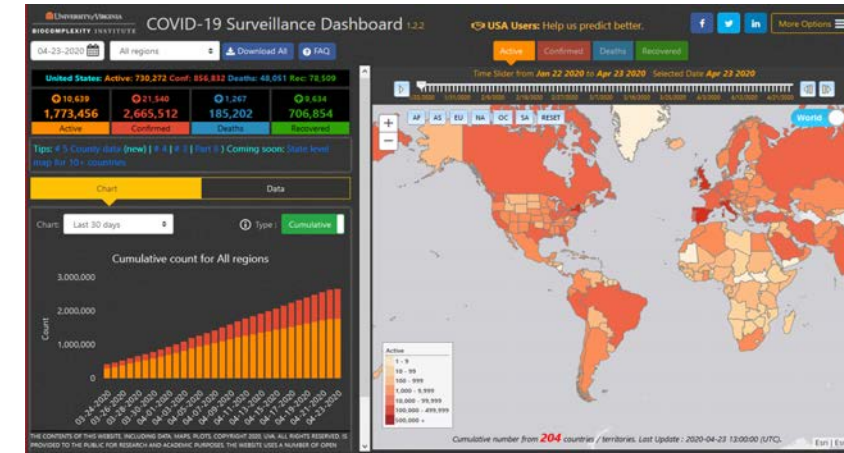
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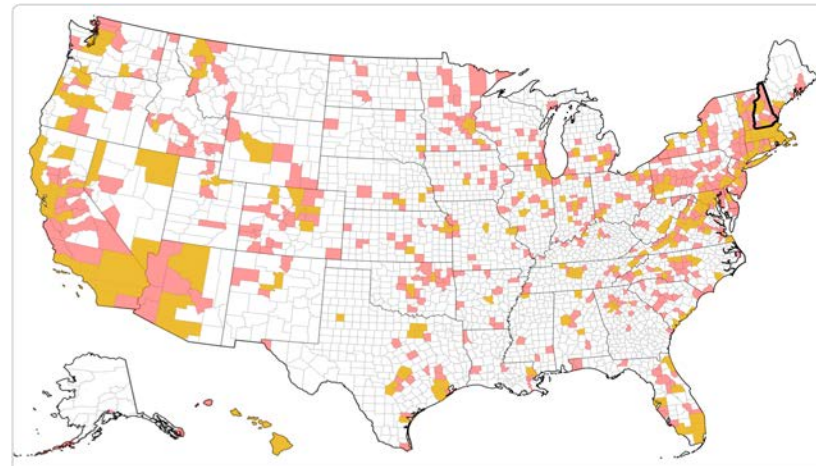
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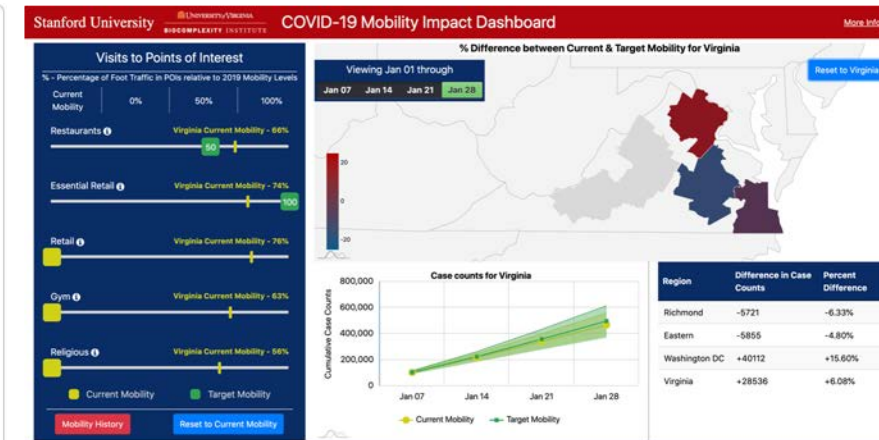


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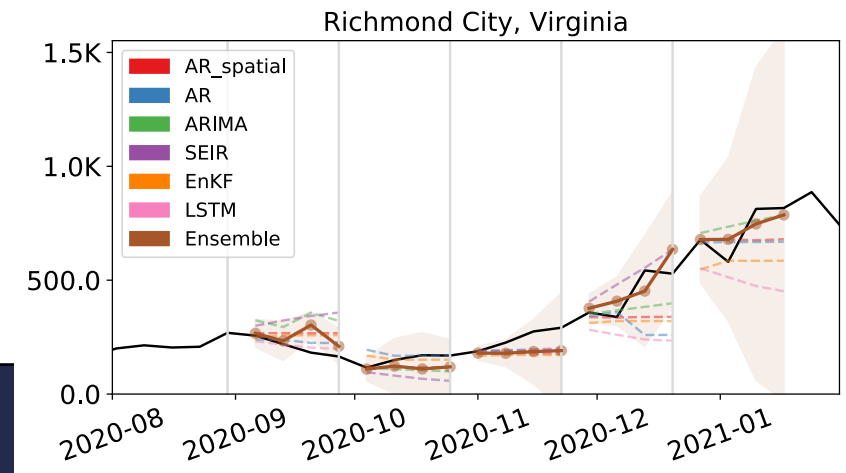
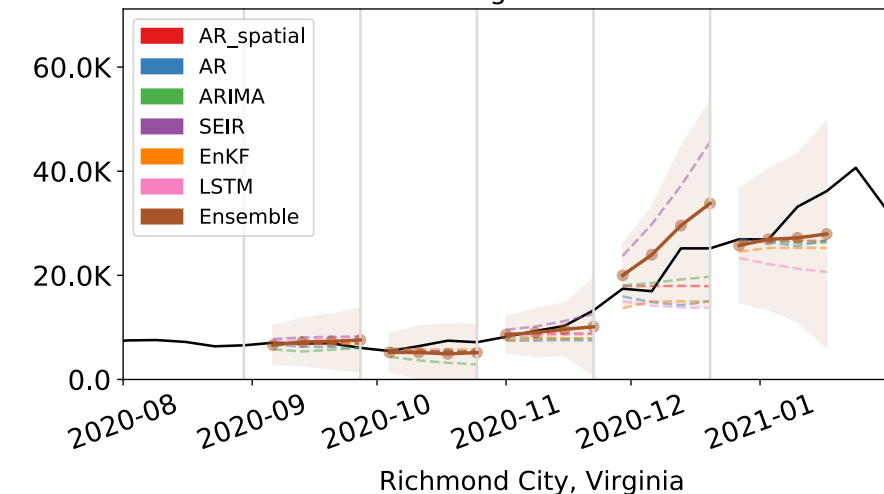
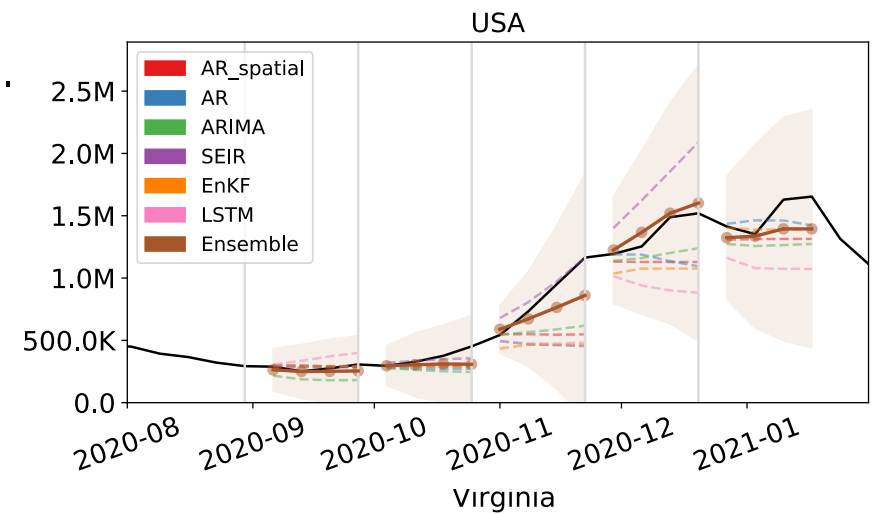


COVID-19 Mobility Impact Dashboard



Multi-scale, multi-method weekly COVID-19 forecasting

- Incorporate multiple classes of models:
 - Statistical methods (AR, Kalman filters), deep learning models (LSTM), and Mechanistic meta-population models (SEIR model)
- Ensemble:
 - Combine forecasts from multiple methods to produce probabilistic forecasts at county level (performance-based ensemble)
 - Bayesian model averaging (instead of model selection) to avoid overconfident inferences & include individual model uncertainty
 - Ensemble forecasts usually perform better than individual forecast
- Key observation: All models are useful.



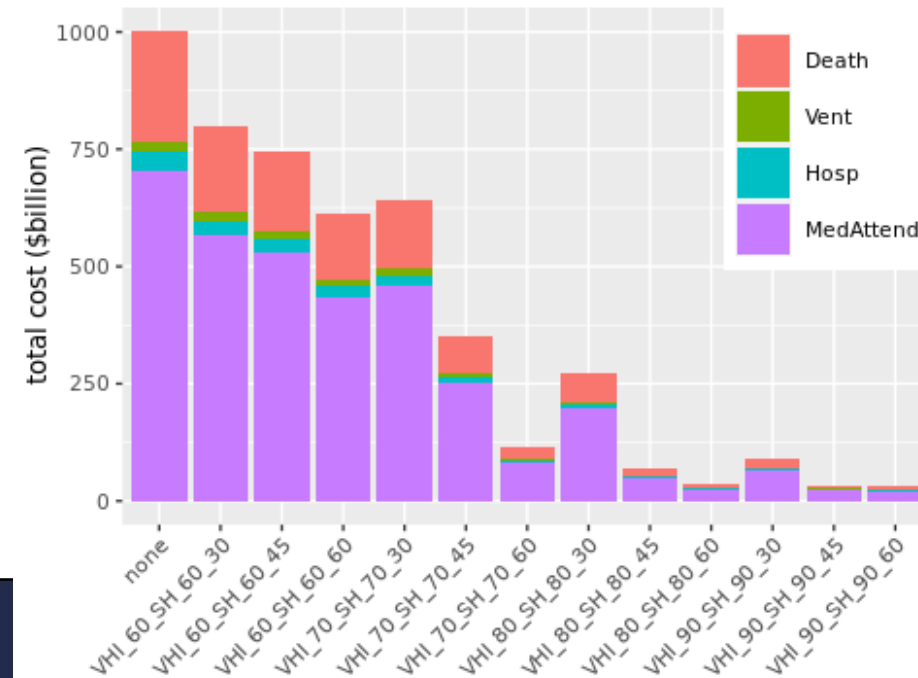
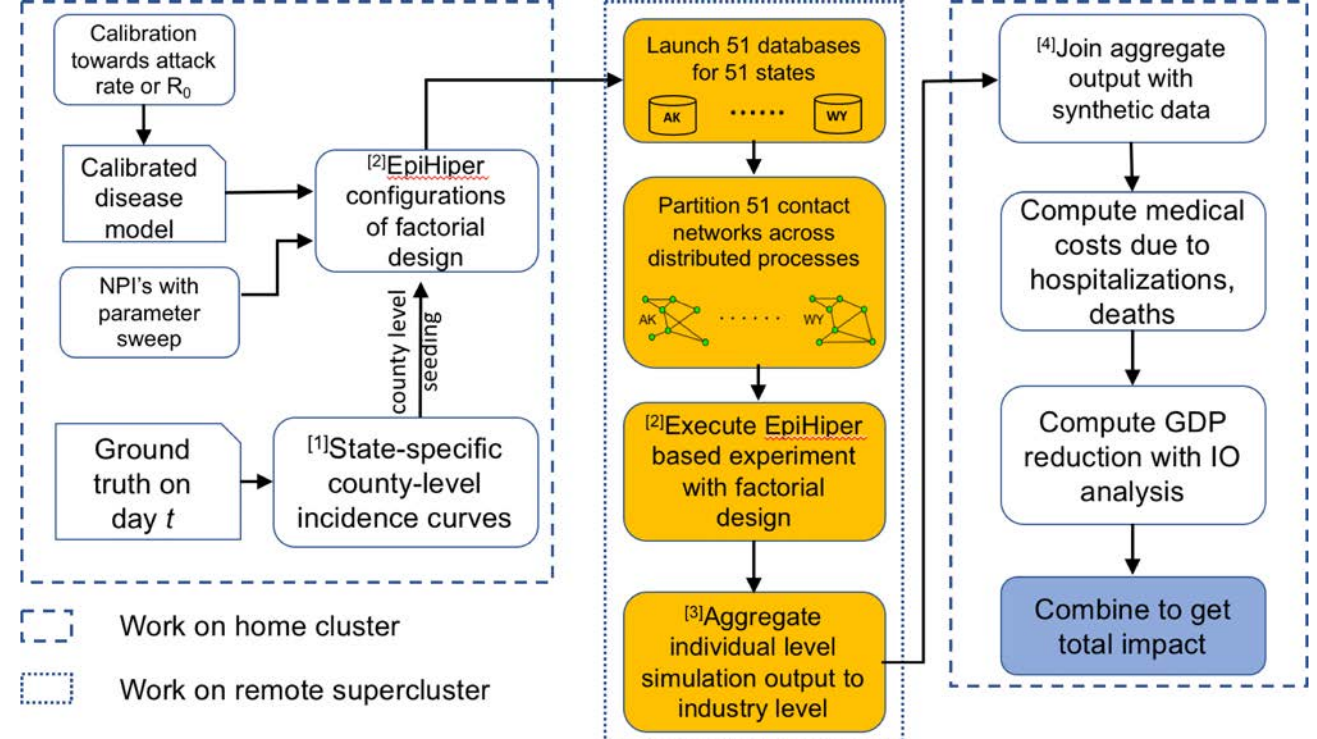
HPC-Grid workflow to compute US medical costs

[1] Daily Incidence data: ~ 3100 counties \times 200 days.

[2] A typical design: 2 VHI compliances \times 3 lockdown durations \times 2 lockdown compliances \times 51 states \times 15 replicates = 9180 simulation instances. Network with 300 million nodes and 7.9 billion edges partitioned across all 50 states

[3] Size of *individual* level output data: 12 cells \times 51 states \times 15 replicates multi-million state transitions = approx. multi-billion entries (3TB).

[4] Size of *aggregate* output data: 12 cells \times 51 states \times 15 replicates \times 365 days \times 90 health states \times 3 counts = ~ 1 billion entries (2.5GB).



Lessons learned

- **Work closely with stakeholders**
 - Build models that are explainable, transparent
- **Be agile and flexible**
 - Each situation is new and comes with unique challenges: requires constant model adaptation
- **Unusual effectiveness of transdisciplinary team science**
 - Working in teams is critical – skills, perspective and collaboration matters
- **Social, political and economic considerations are increasingly important**
- **Communicating scientific results in such situation needs to be thoughtful and deliberate**

Concluding remarks and key takeaways

An effective strategy to reduce the global burden of epidemics must:

- **Detect** timing and location of occurrence.
- **Anticipate** public reaction to an outbreak.
- **Develop** actionable interventions that enable targeted and effective responses.

Needed advances

- **Real-time collection** and updating of data, models in rapidly changing environments.
- **Incorporate social and behavioral** components in the models
- **Models** that are scientifically effective, explainable & operational.

New looming challenges

- Climate Change
- Anti-microbial resistance
- Synthetic pathogens
- Infodemics and role of social media
- Urbanization & increased global transactions
- Expectation of timely information

Some Background Information