

# Understanding socio-structural drivers of HIV transmission using epidemiology and systems science

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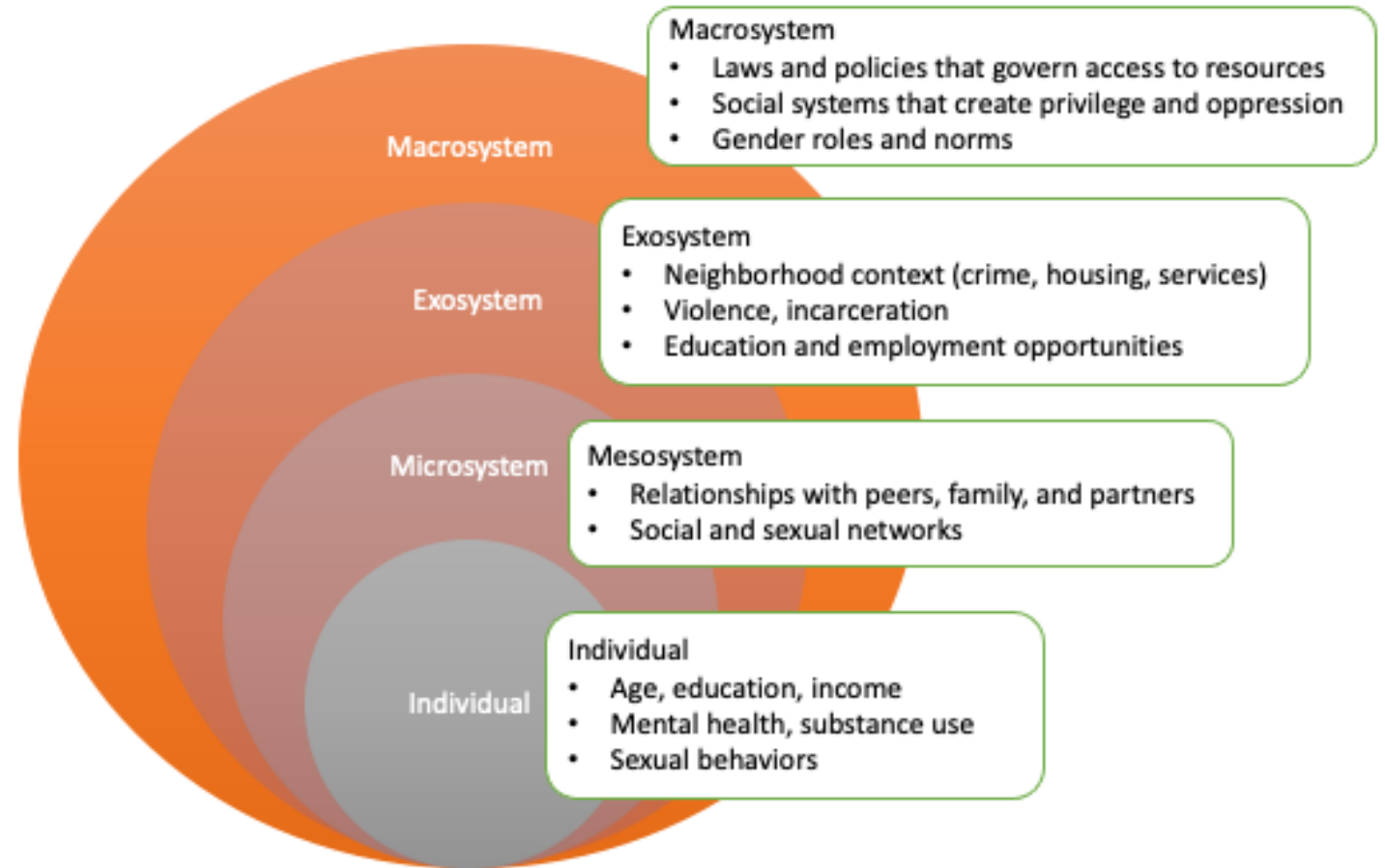


# Outline

- Introduction
- HIV in Chicago
- Brief overview of agent-based modeling (ABM)
- ABM and counterfactual frameworks
- Example: ABM to understand the impact of criminal justice involvement on HIV transmission
- Future research and next steps

# Introduction

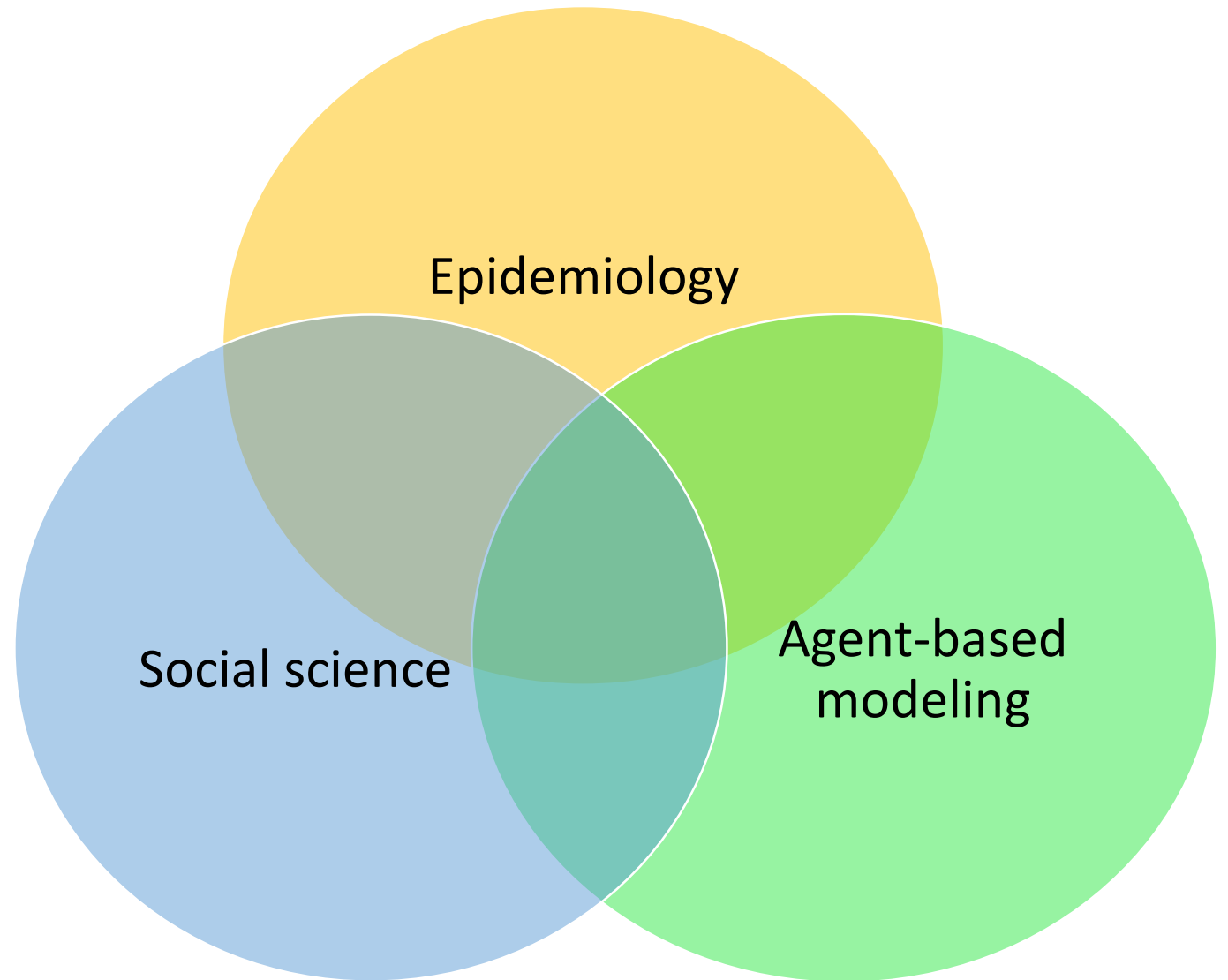
- Socio-structural & contextual influences
  - Dyad-level factors
  - Network influences, environmental context
  - Social determinants of health (SDOH)
- Populations disproportionately affected by HIV/STIs



Adapted from Ecological Systems Theory. Source: Bronfenbrenner, U. (1979). *The Ecology of Human Development: Experiments by Nature and Design*. Cambridge, MA: Harvard University Press.

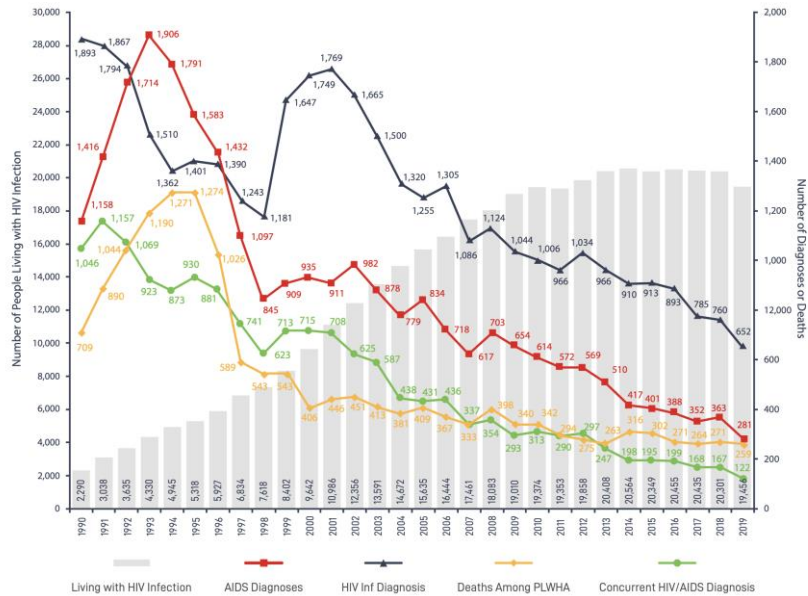
# Research Methods

- Epidemiology
  - Observational research, interventions
  - Causal inference
- Social sciences
  - Conceptual frameworks
  - Social determinants of health
- Systems science & agent-based modeling
  - Useful for studying complex systems
  - Epidemiologic analyses provide input parameters for agent-based models



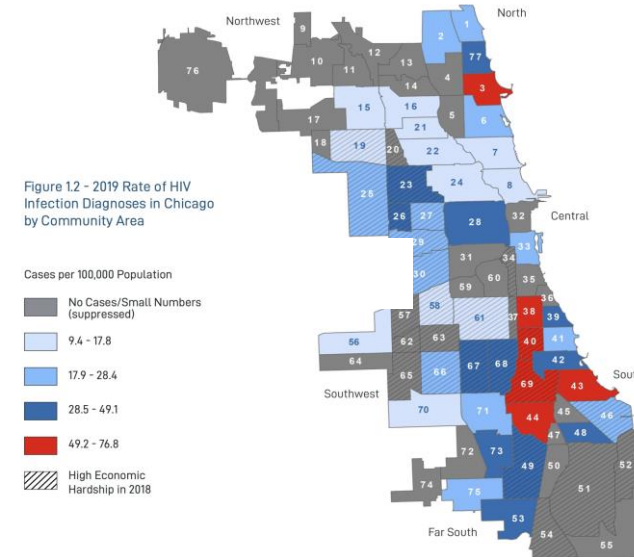
# HIV in Chicago

## Trends in HIV/AIDS Infections, Diagnoses, and Deaths, Chicago, 1990-2019

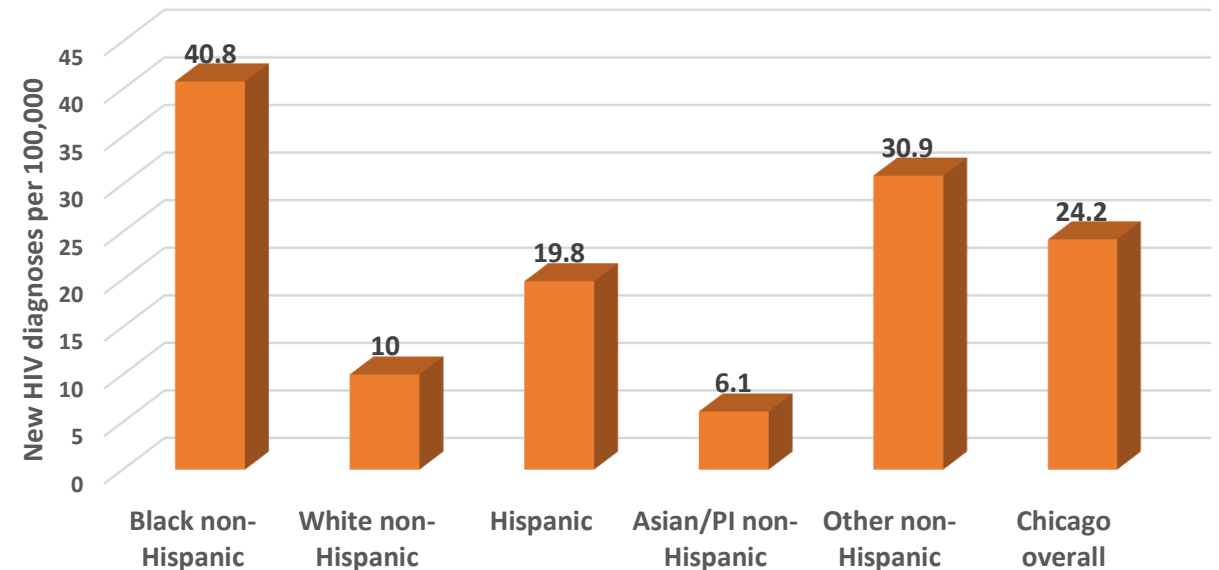


Source: Chicago Department of Public Health. HIV/STI Surveillance Report, 2019. Chicago, IL: City of Chicago; December 2020.

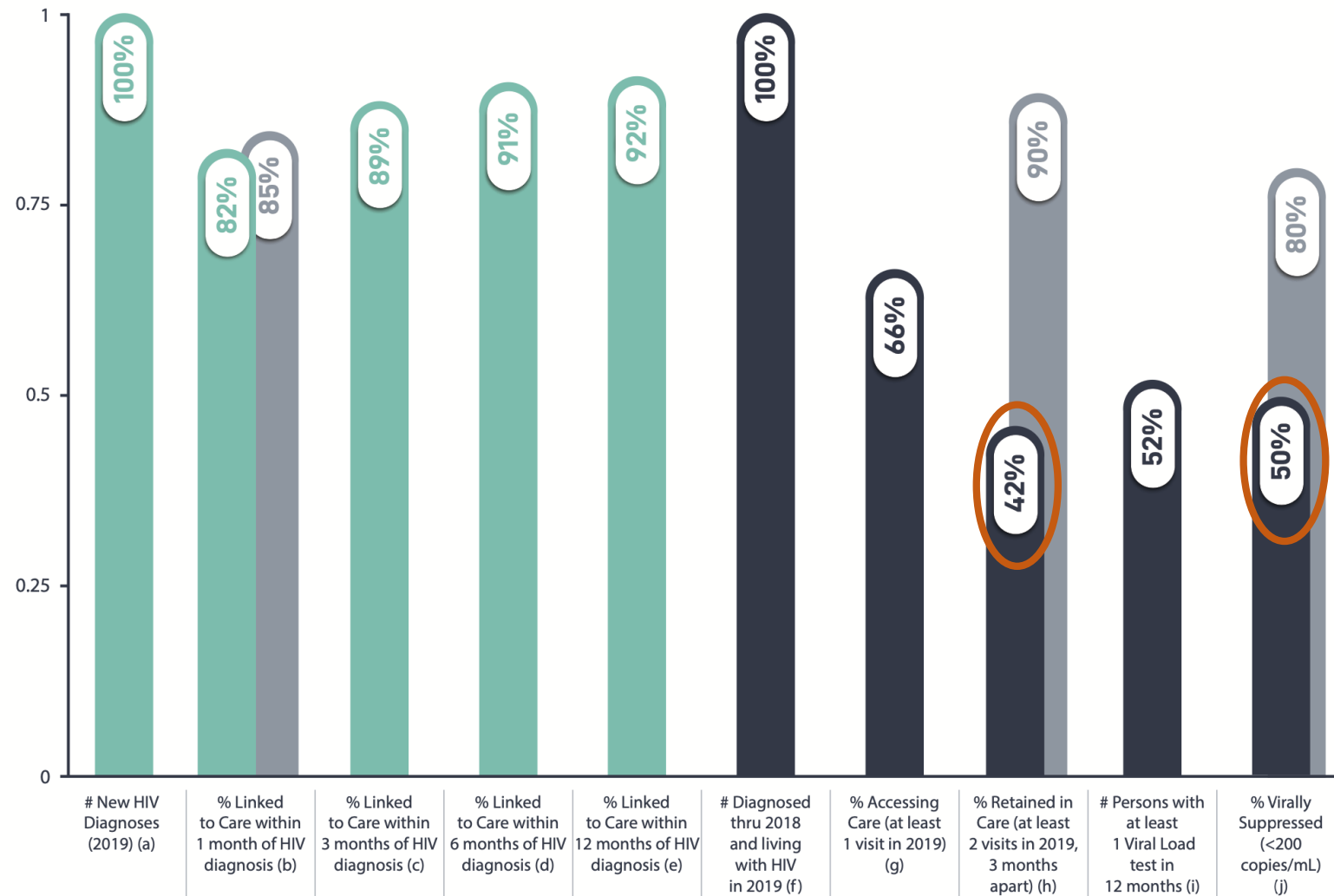
## HIV Infection Rates by Chicago Community Area, 2019



## New HIV diagnoses per 100,000, Chicago, 2019



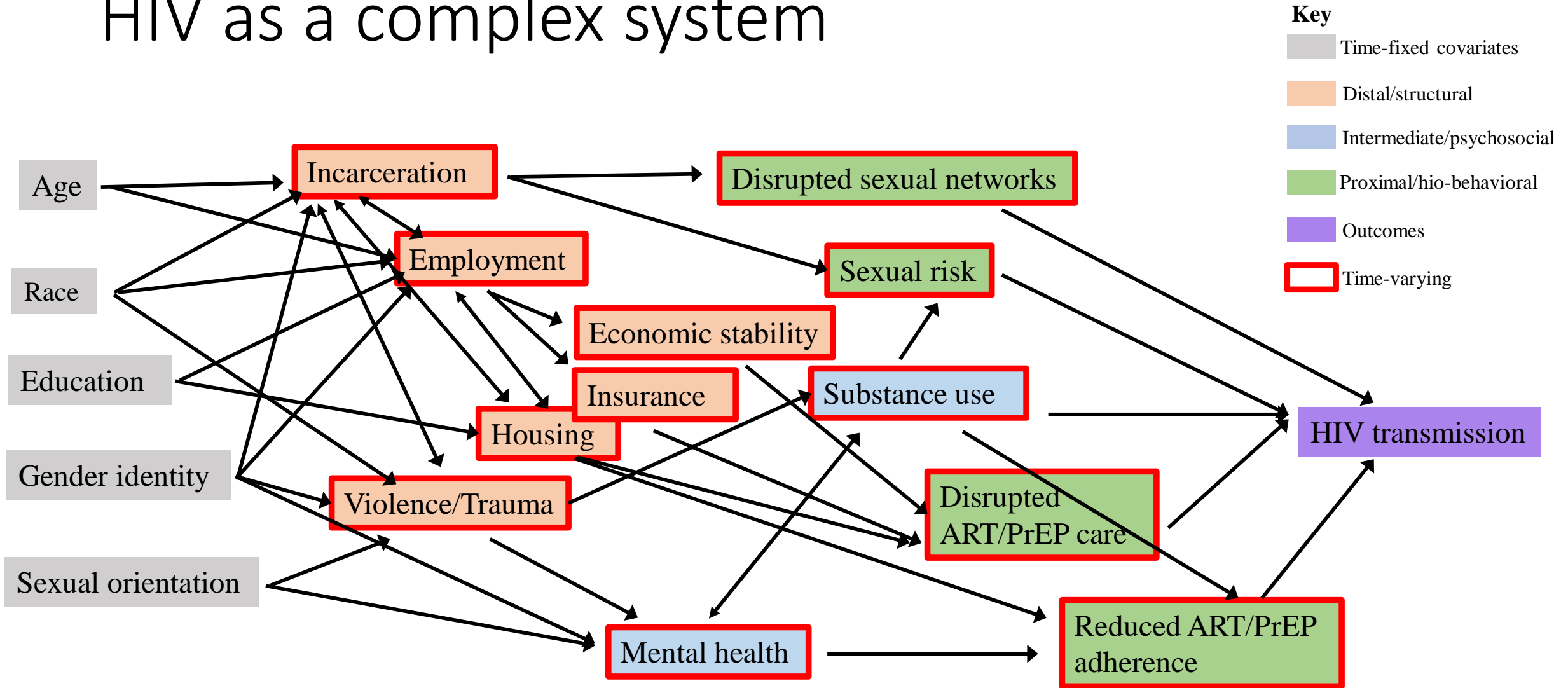
# HIV Continuum of Care Among Persons Aged $\geq 13$ Years, Chicago, 2019



# HIV Elimination goals

- Interventions will need to focus on communities with complex and co-occurring socio-structural barriers to engagement in HIV prevention and care
- Much previous research focused on behavior change at the individual level (sexual risk, substance use), but there is a recognized need to focus on more distal influences on HIV transmission
  - E.g., Housing, employment, incarceration

# HIV as a complex system





# Existing evidence and knowledge gaps

## Evidence

- Associations of SDOH with HIV transmission related factors across multiple observational studies
  - Sexual behaviors
  - Substance use, mental health
  - Engagement in care, viral suppression, PrEP uptake

## Gaps

- Many cross-sectional studies
- Focused on single factors or pathways
- Lacked statistical power or sufficient confounder control; causal associations cannot be inferred
- Effect size magnitude varies widely due to differences in study populations, design, timeframe, confounder control

# Limitations of traditional study designs for understanding complex systems

- Logistical
  - Long duration of follow up required to observe effects
  - Very resource intensive to follow people longitudinally
  - Settings may not be conducive to traditional research designs (e.g., criminal justice settings)
- Ethical
  - Not always feasible or possible to randomize people
  - High participant burden/burnout

# Statistical challenges

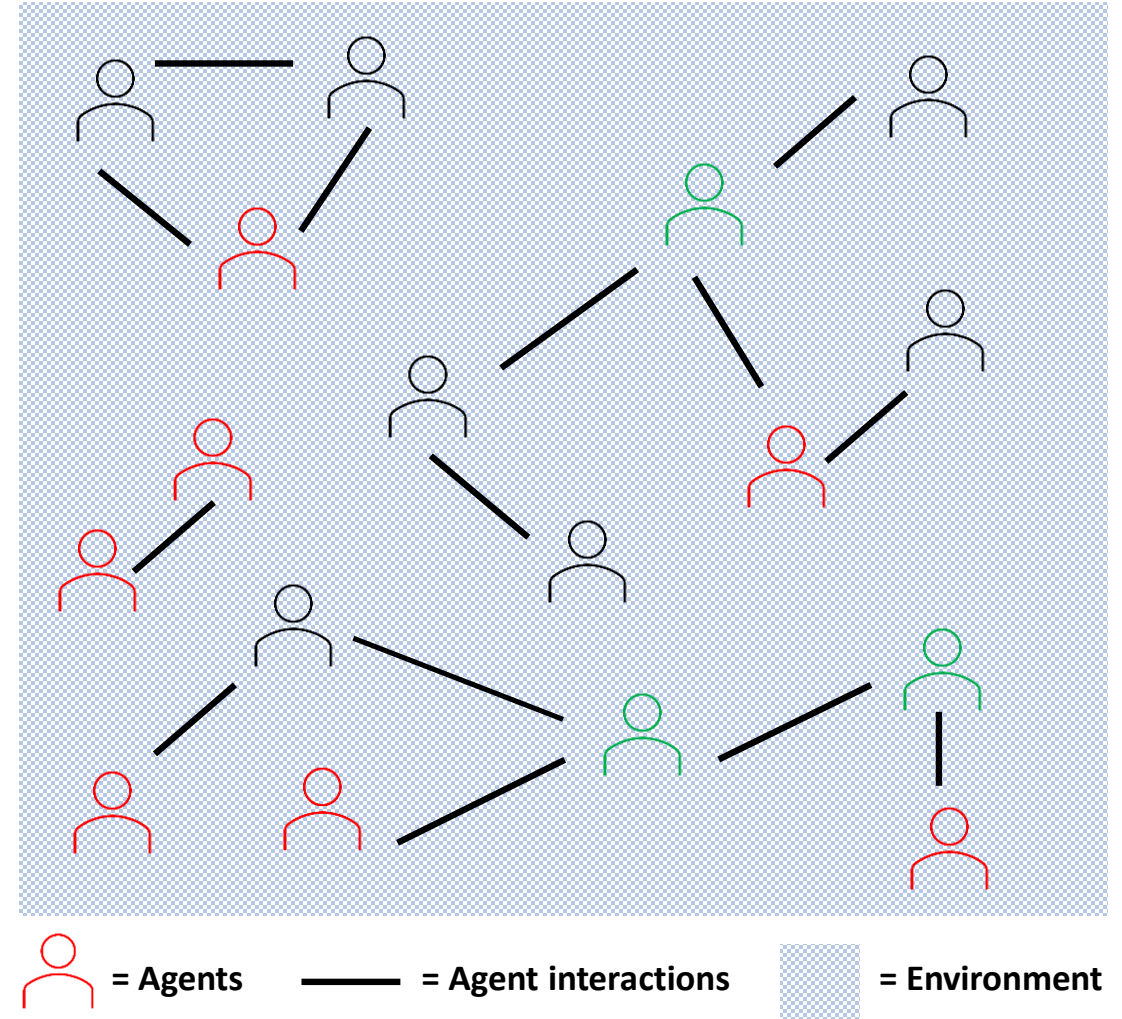
- Standard regression models typically assume relationships between exposures and outcomes are unidirectional, linear, and time-constant and exposures are independent
  - Not well suited to relationships characterized by causal interdependence, non-linearity (e.g., thresholds), feedback loops (magnified or dampening effects), and interference (one person's exposure influences the outcomes of others)
  - Many traditional causal inference frameworks assume these are absent (e.g., unidirectionality, no interference, etc.)
- For rare outcomes (e.g., HIV), very large sample sizes required for sufficient power

# Limitations of randomized controlled trials (RCTs) for evaluating complex interventions

- RCTs – useful for isolating a single intervention effect or component, generally while holding other factors and contexts constant
- Systems science approaches are better suited to studying complex interventions
  - Questions that couldn't be answered with simpler designs or models

# Agent-Based Models (ABMs)

- Computer simulation approach to modeling the dynamics of complex systems
- Models represent social systems composed of agents that interact with and influence each other
- Observe system-level consequences of agent behaviors and interactions
- Effects of interventions can be simulated under various assumptions in a virtual environment



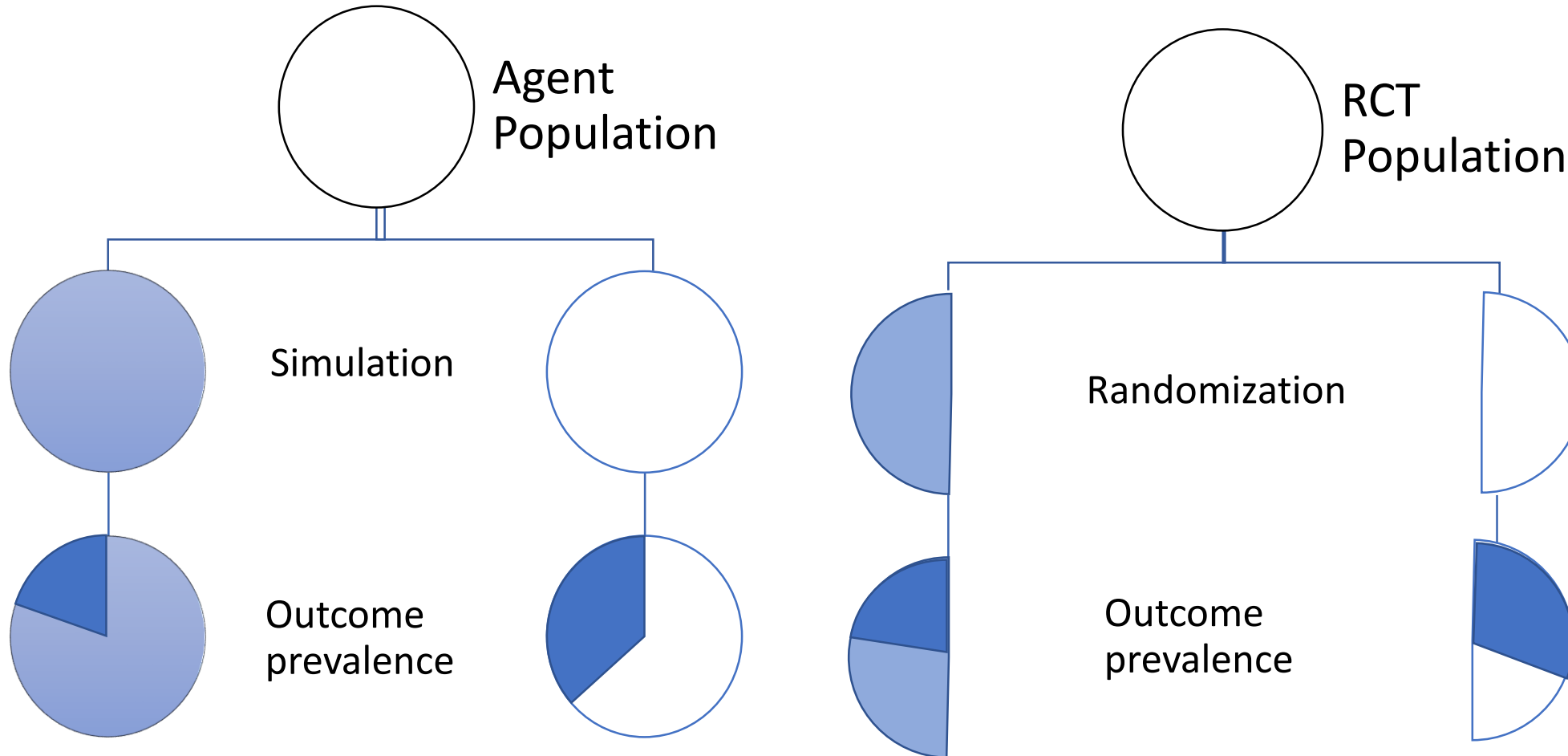
# Agent-based model components

Agents	Agent-Agent interactions	Environments	Agent-environment interactions
<ul style="list-style-type: none"><li>• Attributes (age, sex, race, employment, housing)<ul style="list-style-type: none"><li>• Static or dynamic</li></ul></li><li>• Behaviors<ul style="list-style-type: none"><li>• Based on current information &amp; past history</li></ul></li></ul>	<ul style="list-style-type: none"><li>• Information exchange</li><li>• Disease transmission</li><li>• Contend for resources</li></ul>	<ul style="list-style-type: none"><li>• Social or sexual networks</li><li>• Physical, social, neighborhood environments</li></ul>	<ul style="list-style-type: none"><li>• Take in information on environment</li><li>• Shape environment</li></ul>

# Uses of agent-based modeling in epidemiology

- Understand mechanisms by which exposures (e.g., SDOH) impact population level health outcomes
  - Can show how patterns at the population level arise from exposures that might not be evident in a single study
  - Conduct counterfactual experiments to evaluate hypotheses that may not be possible with standard statistical models
- Evaluate potential interventions
  - Mechanisms by which interventions work
  - How interventions can be most efficiently focused (identify subgroups)
  - Optimal combination/sequence of interventions

# ABM and counterfactual frameworks



Adapted from Marshall BD, Galea S. Formalizing the role of agent-based modeling in causal inference and epidemiology. *Am J Epidemiol.* 2015 Jan 15;181(2):92-9.



# Basic notation: Agents

- At each time step  $t$  ( $t = 1, \dots, T$ ), each agent  $i$  ( $i = 1, \dots, N$ ) has a set of  $m$  ( $m = 1, \dots, M$ ) internal traits that can be described by the matrix  $\mathbf{S}^t$

$$\mathbf{S}^t = \begin{bmatrix} s_{1,1}^t & s_{1,2}^t & \dots & s_{1,M}^t \\ s_{2,1}^t & s_{2,2}^t & \dots & s_{2,M}^t \\ \vdots & \vdots & \vdots & \vdots \\ s_{N,1}^t & s_{N,2}^t & \dots & s_{N,M}^t \end{bmatrix}$$

- Traits:
  - Continuous, nominal, dichotomous
  - Can represent sociodemographics, exposures, behavioral proclivity, etc.

- Analogously, agents can be placed in  $p$  ( $p = 1, \dots, P$ ) environments where  $\mathbf{E}^t$  represents an environmental state matrix

# Agent-agent interactions

- At each time step  $t$  ( $t = 1, \dots, T$ ) each agent  $i$  interacts with a subset of the population  $\{1, \dots, i-1, i+1, \dots, N\}$ 
  - Described by agent-agent interaction matrix  $\mathbf{K}^t$  where each element  $k_{i,j}^t$  indicates whether agent  $i$  interacts with agent  $j$  during timestep  $t$  where  $i$  and  $j = 1, \dots, N$
  - Can be symmetric or asymmetric (information or disease transmission can flow one way or bidirectionally)

# Steps

- Initialize ABM by populating agent trait matrix, environmental state matrix, and interaction matrix with values from pre-defined probability distributions and functions
- Define set of rules **Z** for updating of agent traits, agent-agent interactions, and movement between or interaction with environments
  - Rules **Z** are defined by functions
- At each time step: update the model based on previous values and pre-defined rules **Z**

# Steps

- Monte Carlo simulations to obtain outcomes (e.g., disease incidence, prevalence, mortality) from runs  $r$  ( $r = 1, \dots, R$ ) at time  $T$  for counterfactual scenarios of interest:
  - Scenario A ( $[\mathbf{Z}, \mathbf{S}_A^T, \mathbf{K}_A^T, \mathbf{E}_A^T]$ ) vs. Scenario B ( $[\mathbf{Z}, \mathbf{S}_B^T, \mathbf{K}_B^T, \mathbf{E}_B^T]$ )
- Compute point estimates from scenario A vs. scenario B by averaging across runs for each outcome of interest

$$\hat{\mu}_{o,A}^{T,R} = \frac{\sum_{r=1}^R y_{r,o,A}^T}{R} \text{ vs. } \hat{\mu}_{o,B}^{T,R} = \frac{\sum_{r=1}^R y_{r,o,B}^T}{R}$$

Example: Agent-based modeling to study the impact of criminal justice involvement (CJI) on HIV transmission among young Black sexual and gender minorities (SGM)

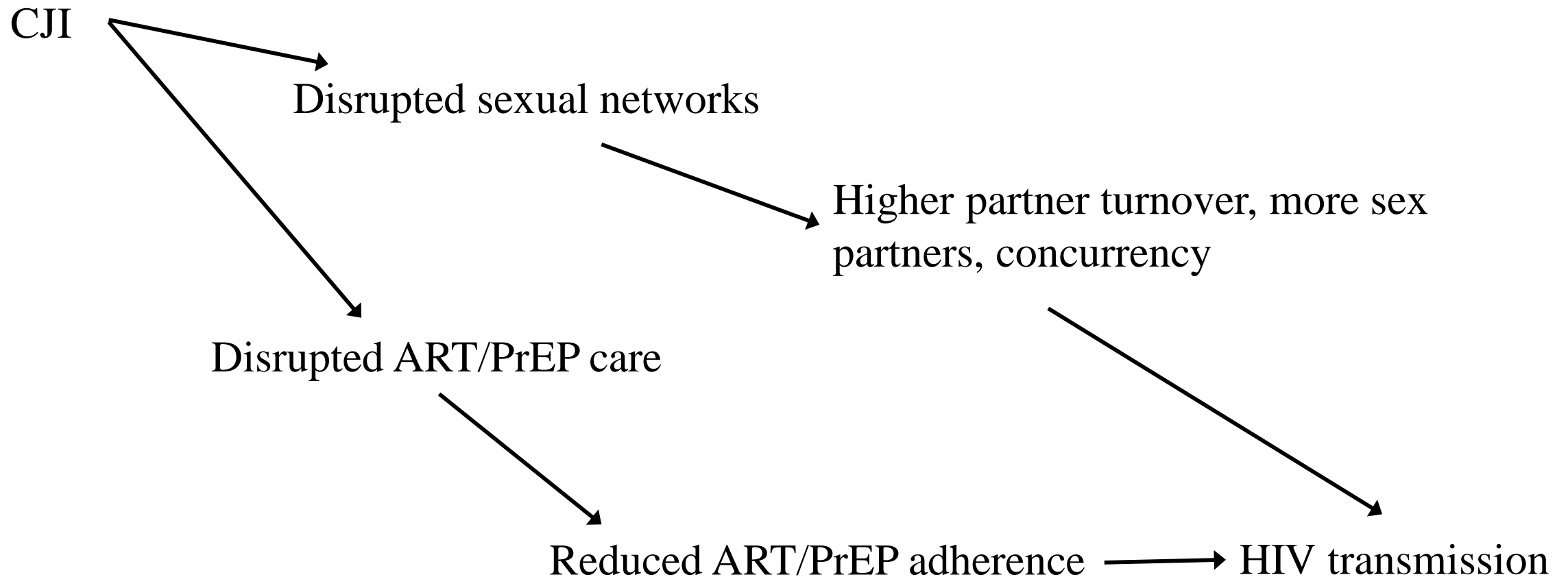
R01DA033934 (Fujimoto, Harawa, & Schneider, PIs)

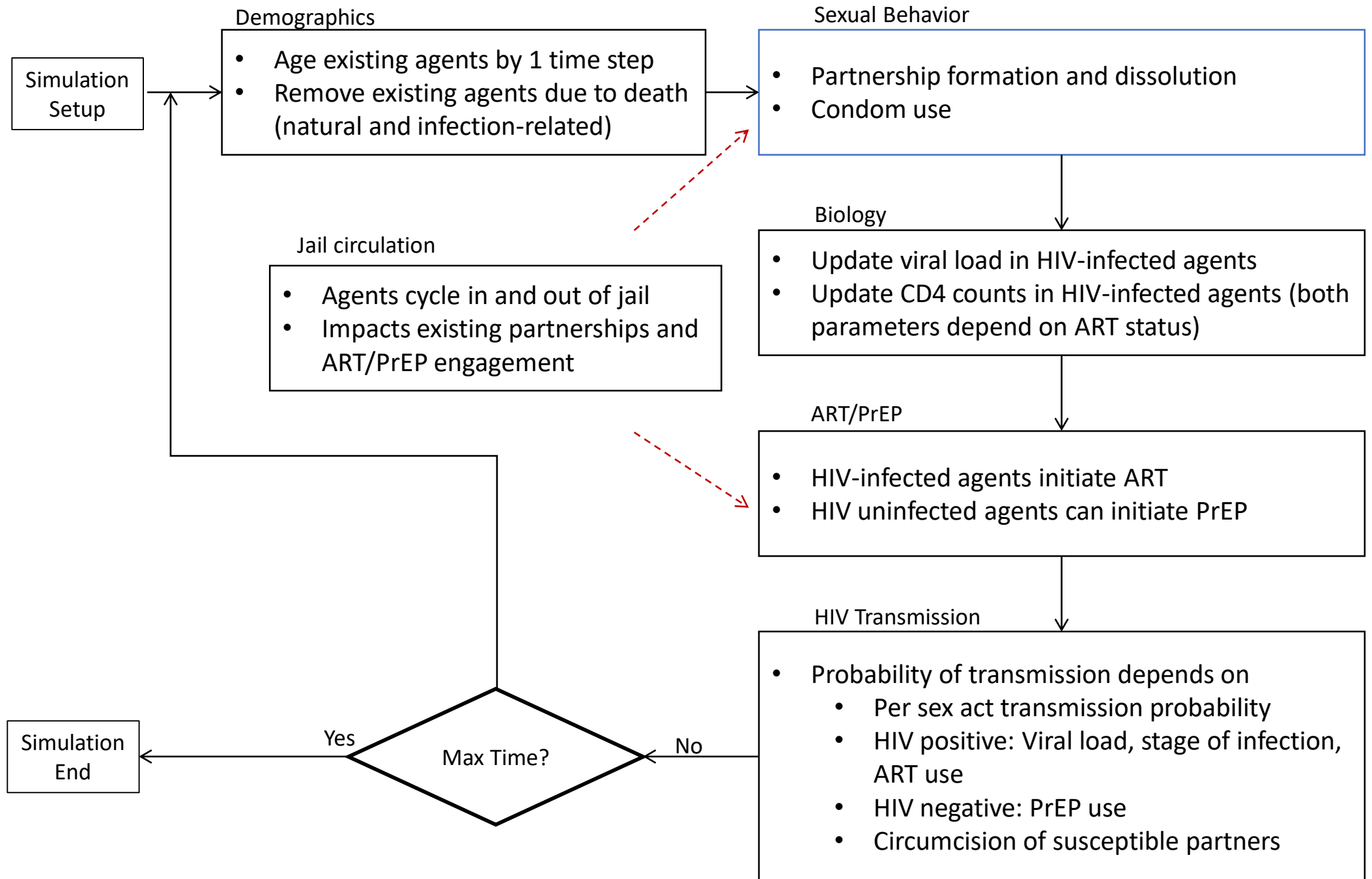
R21MH128116 (Hotton, PI)

# Intersection of HIV and criminal justice involvement

- Black SGM are disproportionately impacted by HIV and criminal justice involvement
  - Frequent cycling between communities and criminal justice settings
- CJI can impact:
  - Employment and housing opportunities
  - Access to medical care
  - Social and sexual network stability
- Agent-based models can be used to:
  - Provide insights to understand how CJI impacts HIV transmission
  - Evaluate interventions for criminal justice involved individuals and their networks

# Hypothesized mechanisms by which justice involvement impacts HIV transmission







# Model population and data sources

- Model population: 10,000 agents representing Black SGM ages 18-34 in the city of Chicago
- Data sources: Local cohort studies, clinical data, and public health surveillance
- Outcomes: HIV incidence and prevalence (average annualized estimates computed over 10 years)
- Calibration: local HIV incidence and prevalence estimates (CDPH surveillance and local studies), incarceration incidence and prevalence
- Model components: Developed with Repast HPC ABM toolkit using C++
- Network formation and dissolution dynamics modeled with exponential random graph models using the *statnet* suite of packages in R

# Experiments

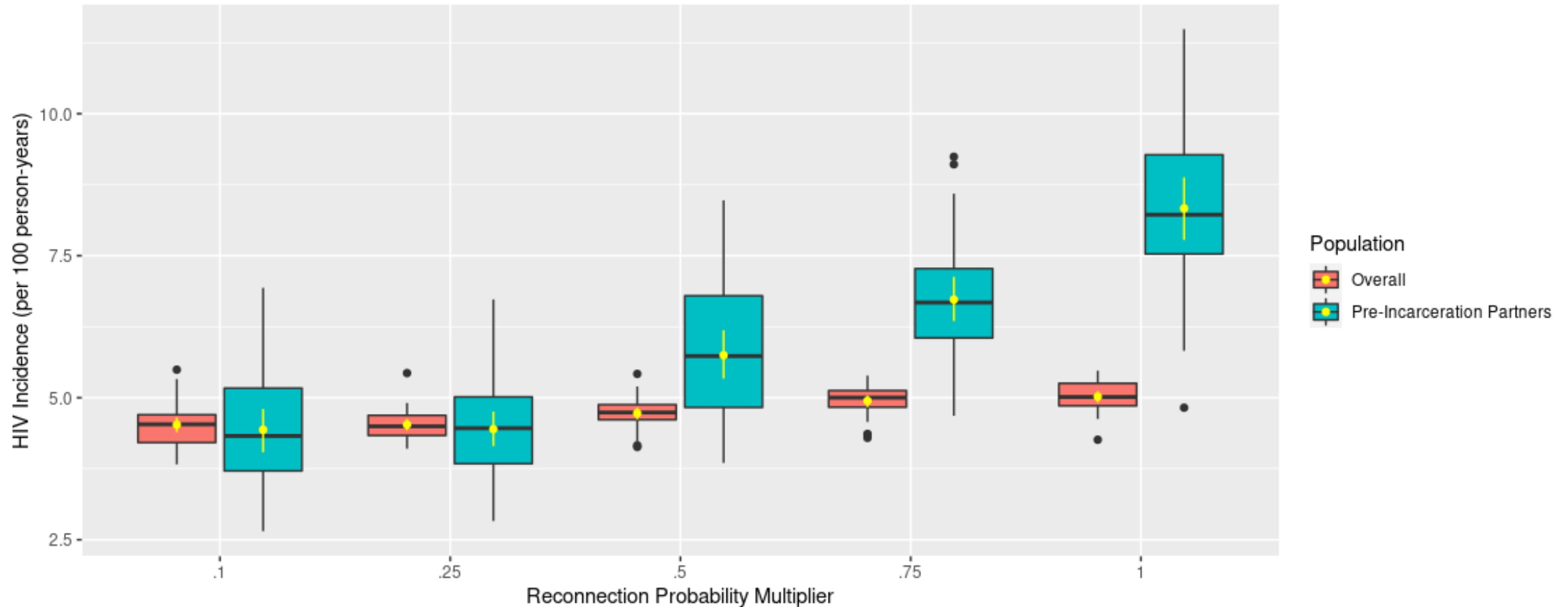
- Quantify the impact of criminal justice involvement:
  - Population level HIV incidence
  - HIV incidence among justice involved individuals and their networks
- Evaluate interventions to reduce the impact of justice involvement on HIV transmission in different sub-populations
  - Examples:
    - Reduce post-release disruption in HIV/PrEP care (e.g., interventions to facilitate care engagement by reducing insurance, housing, or employment barriers)
    - Focused or enhanced PrEP and ART interventions for justice involved individuals and their networks
- Each scenario was repeated across 30 runs to incorporate stochasticity and outcomes were averaged across runs

# Annual HIV incidence by subpopulation

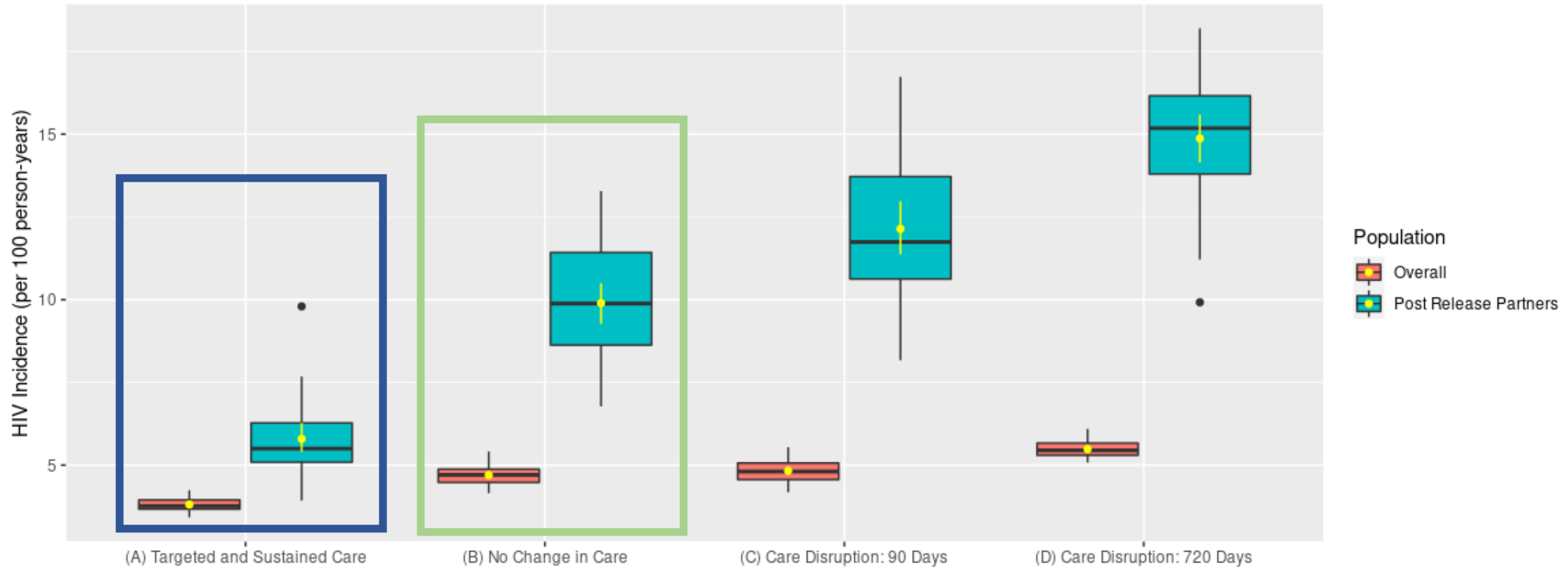
Population	HIV incidence per 100 person-years	95% CI*
<i>Individuals</i>		
Ever incarcerated	5.72	5.52 – 5.92
Never incarcerated	4.73	4.65 – 4.83
<i>Partners of individuals with CJI</i>		
Pre-Incarceration Partners	6.83	6.31 – 7.40
Post-Release Partners	12.14	11.4 – 13.0

\*Confidence intervals obtained via bootstrapping

# HIV incidence by probability of partner reconnection after release from jail



# HIV incidence under various levels of post-release care disruption and intervention



# Average HIV incidence under different care disruption scenarios

	<b>Post-Release Partners</b>	<b>Overall</b>
	<b>Incidence per 100 person- years (95% CI)</b>	<b>Incidence per 100 person- years (95% CI)</b>
Intervention: Targeted and sustained care	5.80 (5.40, 6.28)	3.81 (3.74, 3.89)
Standard: No change in care	9.90 (9.27, 10.50)	4.70 (4.59, 4.82)
Incidence rate ratio	0.59 (0.53, 0.65)	0.86 (0.84-0.89)
Incidence rate difference	-4.10 (-4.85, -3.35)	-0.89 (-1.02, -0.76)

# Summary

- Identified a subgroup who could benefit from targeted PrEP interventions (partners of those recently released from jail) which may have been hard to identify using observational research designs
  - Can give ideas about where to target limited public health resources
- Suggests need for interventions to increase ART and viral suppression among HIV-positive individuals with CJ *and* increase PrEP/ART use in their networks
- Next steps
  - Evaluate impact of interventions to distribute PrEP to networks of released individuals
  - Explicitly model interventions to reduce post-release disruption in care by reducing insurance, housing, or employment barriers and recidivism
  - Combinations of interventions applied simultaneously or sequentially

# Limitations & open questions

- Estimates used as input parameters for agent-based models are often uncertain or potentially biased – need for sensitivity analysis
  - Model results may be dependent on parameter inputs for which the true magnitude of effect is often unknown
  - Transportability – estimates of effect from one population may not generalize to another
- Can provide a range of effect estimates as priors and use computational techniques to refine estimates – ongoing work in this area
  - Large-scale sensitivity analyses and model exploration with high-performance computing
  - Identify variables that have the most impact on model (system) behavior



# Extensions and ongoing work

- Ongoing work: extend existing model to incorporate additional social determinants of health (housing, employment) and evaluate their impact
  - Increase the granularity of the synthetic population in order to represent HIV transmission with sufficient realism to examine more nuanced research questions
- Develop formal methods for evaluating assumptions needed for valid inference with agent-based modeling
  - Counterfactual frameworks and high-dimensional sensitivity analyses to assess the impact of varying causal or mechanistic assumptions
  - Quantify the impact of incomplete or imprecise empirical data
- G-computation
  - Methods to estimate the causal effect of a time-varying exposure in the presence of time-varying confounders affected by the exposure; also applies to settings with feedback loops
  - Extensions can address interference (auto g-computation)
  - May complement agent-based modeling to triangulate information if adequate longitudinal data are available
  - Can provide causal effects estimates as starting parameters for agent-based models to be refined using computational approaches within the ABM

# Collaborators and funding

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# Technical resources and documentation

## **Repast for High performance computing:**

Homepage: [https://repast.github.io/repast\\_hpc.html](https://repast.github.io/repast_hpc.html)

Repast documentation, including Repast4Py (Python implementation) and lots of tutorials: <https://repast.github.io/docs.html>

Collier N, North M. Parallel agent-based simulation with Repast for High Performance Computing. *SIMULATION*. 2013;89(10):1215-1235.

## **Estreme-scale Model Exploration with Swift:** <https://emews.github.io/>

Ozik J, Collier NT, Wozniak JM, Macal CM, An G. Extreme-Scale Dynamic Exploration of a Distributed Agent-Based Model With the EMEWS Framework. *IEEE Transactions on Computational Social Systems*. 2018;5(3):884-895.

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**EpiModel** (R package for mathematical models of infectious disease): <https://www.epimodel.org/>

## **Tutorials on agent-based modeling:**

C M Macal (2016) Everything you need to know about agent-based modelling and simulation, *Journal of Simulation*, 10:2, 144-156, DOI: [10.1057/jos.2016.7](https://doi.org/10.1057/jos.2016.7)

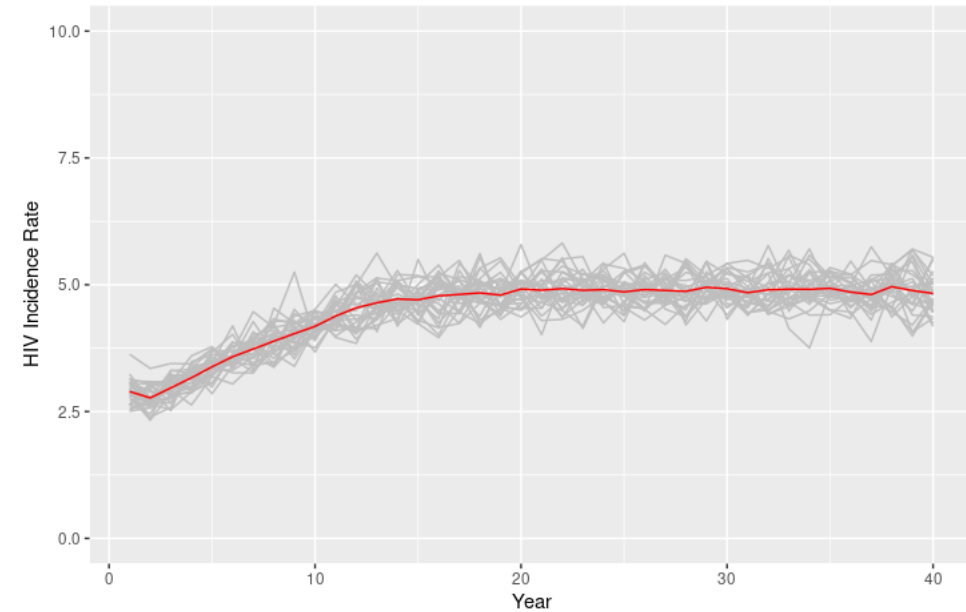
Macal & North: Introductory tutorial on agent-based modeling and simulation:  
<http://simulation.su/uploads/files/default/2014-macal-north.pdf>

Extra slides

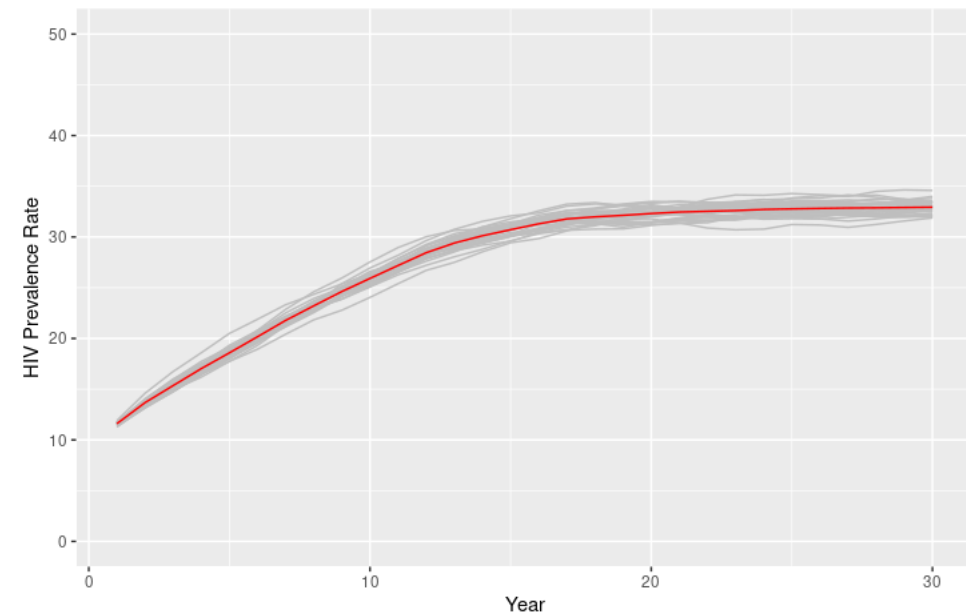
# Model calibration

- Initial set of 270 calibration runs
- Calibration targets: annual HIV incidence (5-7%), HIV & incarceration prevalence
  - Examined differences by age and prior incarceration
- Tested a range of scenarios using empirical estimates from local data as inputs
  - Probability & duration of incarceration
  - PrEP & ART care continuum disruption
  - Network tie retention probabilities
- Refined estimates after initial examination of model output
- Selected the set of parameters that produced outputs consistent with empirical calibration targets for further experimentation

HIV incidence



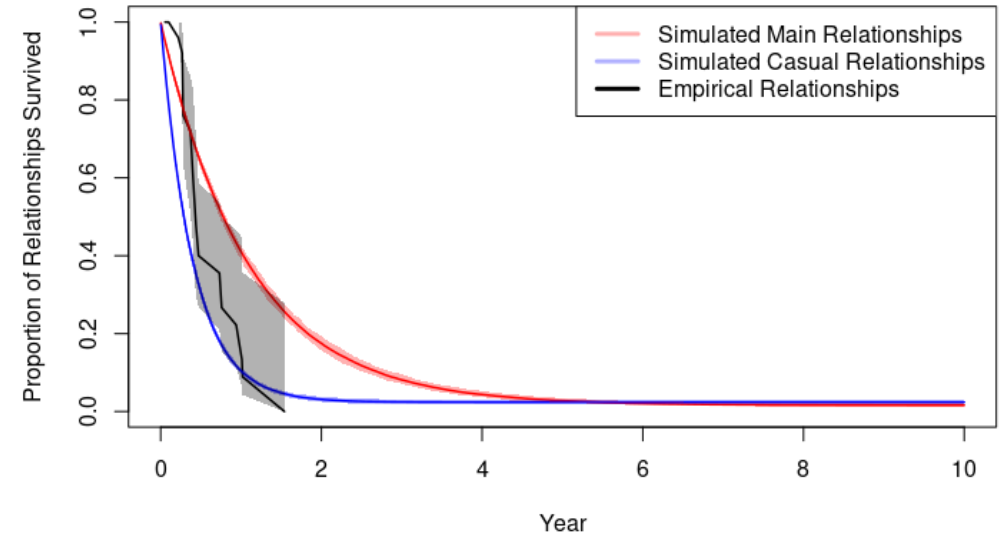
HIV prevalence





# Network tie retention

- When agents go to jail existing network ties are broken with a probability of reconnecting after release
- Determine the status quo survival rate of relationship ties over time using the existing ABM with no incarceration processes implemented
- Apply multiplier to represent the impact of incarceration on probability of reconnecting ties after release
  - Results in a shift in the status quo distribution
    - Multiplier = 1: no impact on tie retention
    - Multiplier < 1: probabilities of retained ties less than the status quo



# Local data sources

Source	Year	Description	Parameter categories
<b>UConnect (R01DA033875)</b>	2013-2015	Cohort study of Black MSM & transwomen ages 16-29, RDS recruitment (n=618)	Sociodemographics, networks, substance use, risk/prevention behavior, HIV/STI prevalence Chicago
<b>National HIV Behavioral Surveillance (NHBS)</b>	June-December 2017 (MSM cycle V)	Time location venue sampling of White, Black, Hispanic/Latino MSM (transwomen not eligible) of all ages (n ~ 500)	Sociodemographics, substance use, risk/prevention behavior, HIV prevalence, PrEP & ART use, partner by partner characteristics/sex behaviors for up to 3 partners
<b>CDPH HIV surveillance</b>	Ongoing	HIV surveillance records	HIV incidence and prevalence, retention in care, viral suppression
<b>US Census Bureau</b>	Ongoing	Demographics of Chicago population	Age-specific mortality rates, population growth rates, population size overall and by subgroup

# Agent-based models for evaluating adaptive interventions

- Adaptive interventions modify intervention or intervention components based on participants' initial response
- ABM can provide insights about the potential impact of adaptive interventions
  - Observe predicted patient trajectories to inform and adjust dose or other intervention components
  - Adjustments to service-level factors, such as provider training
- Can model dynamic, time-varying processes, multicomponent interventions

# Agent-based models for implementation science research

- What-if scenarios can be used to evaluate questions at a conceptual level at the beginning of the implementation process even if empirical data are limited
  - Rerun models under different implementation strategies or policy scenarios
  - Can identify barriers and strengths early in the implementation process
- Incorporate behavioral rules at the individual level and organizational-level interactions
- Individual interventions; combinations of interventions applied simultaneously or sequentially; multi-level interventions; cost-benefit analysis