Using Mathematical Models for Health Economic Analyses

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Outline

- Introduction to modeling
- Infectious disease modeling
- How models can be used to estimate health outcomes
  - Example: Potential impact of ART for prevention
  - What study data can you use to parameterize models
- When to use which model
- Primer on Decision

An introduction to Mathematical Models

- Framework for understanding and communicating infectious disease
- Explicit assumptions help delineate which parameters are based on evidence
- Quantitative or qualitative results are compared with observed or experimental data
- Validated models can be used to estimate the potential impact of interventions (e.g. ART for prevention)

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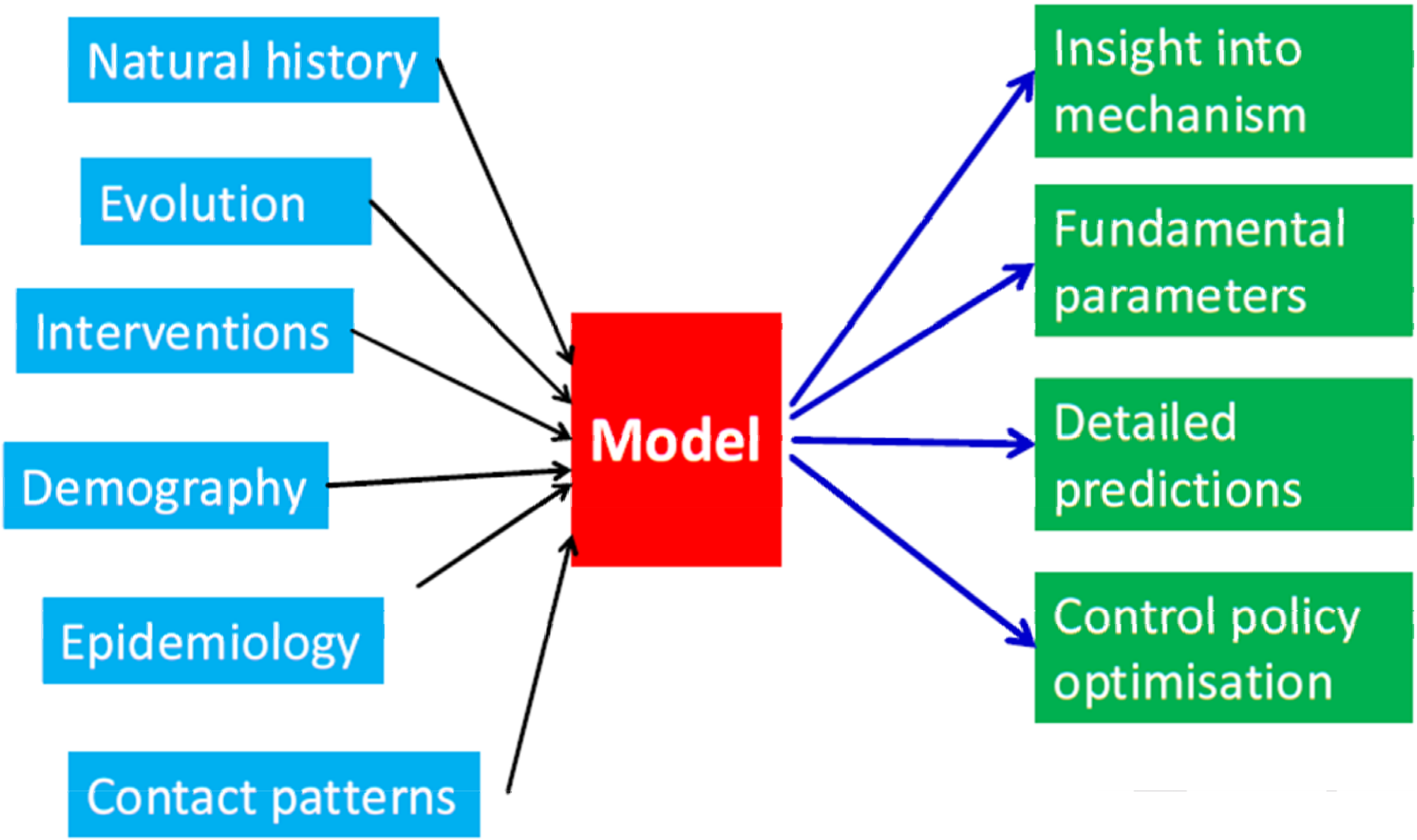
Garnett, G. P. (2002). Sex Transm Infect 78(1): 7-12.

Models in health economic analyses

- Used to structure the economic question and compare all relevant alternatives
- Extrapolate beyond observed data
- Link intermediate and final endpoints
- Generalize results to other settings/patient groups
- Synthesize evidence to simulate comparisons where RCTs don't exist
- Indicate the need for further research

Synthesizing knowledge



Not all models are mathematical!

Why can't we use classical epidemiology to answer questions about infectious disease dynamics?

Why use a (potentially complicated) model?

- Infectious diseases are different because they are caused by **transmission** of a pathogen
- **Prevalence** of infection is the key risk factor for **incidence**.
- Prevalence reflects incidence at earlier time-point—leads to circularity and non-linear dynamics of infectious disease epidemics
- Classic regression of risk factors on disease risk do not incorporate prevalence.
  - ln(disease)= b0 + b1AGE + b2SEX … + b3Income

Population Attributable Fraction (PAR)

- PAR is a useful epidemiologic measure.
- It is the proportion of disease in a population that is attributable to the presence of the risk factor in the population.

$$(Inc_{\text{total pop}} - Inc_{\text{unexposed}})/Inc_{\text{total}} x100$$

E.g. PAR for HIV and male circumcision

- Risk factor: not being circumcised
- Expected incidence among 1,000 susceptibles, of which 500 circumcised:

	Infected	Total population
Circumcised	6	500
Not circumcised	10	500

- Using PAR, (16/1000-6/500)/(16/1000) = 25% of infections in population are due to not being circumcised
- Using a mathematical model, we find 32% of infections are attributable to not being circumcised.
- Why?

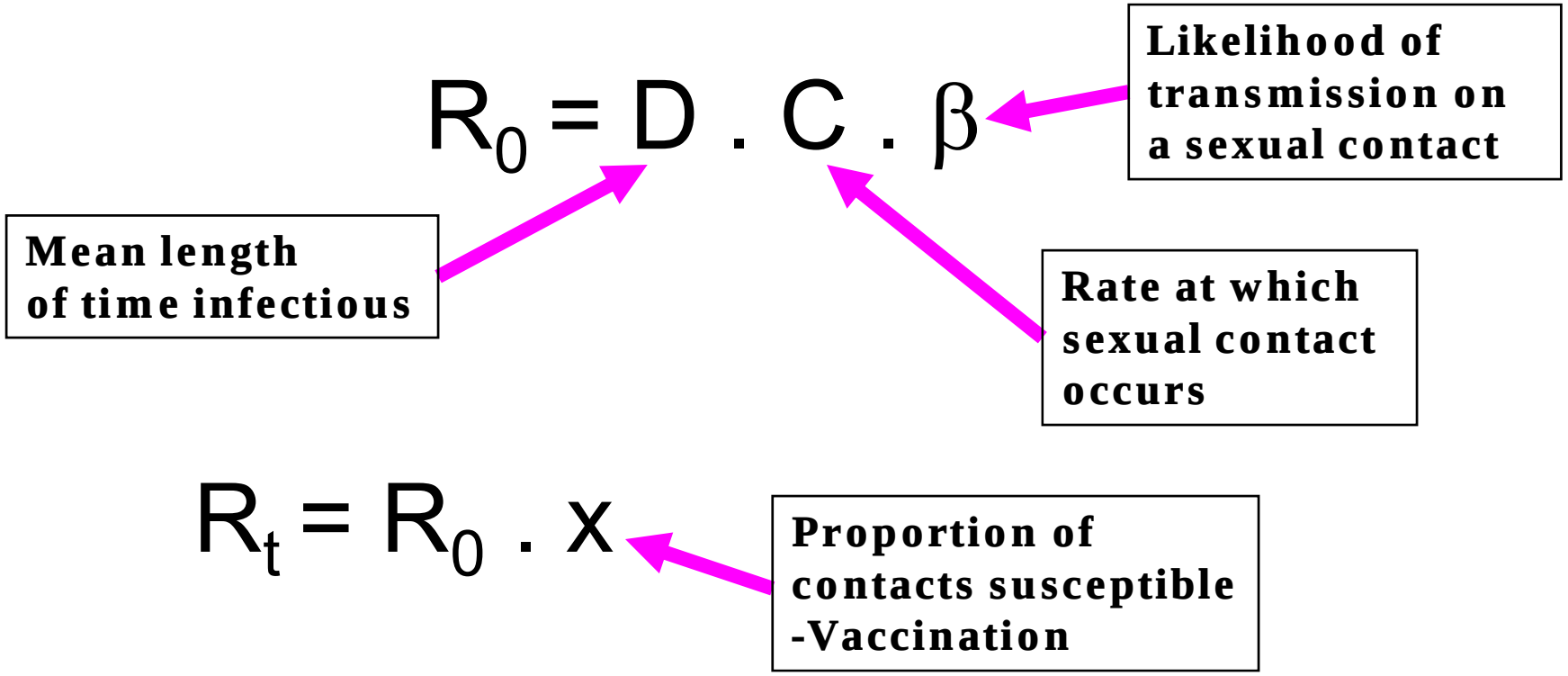
Infectious diseases are non linear

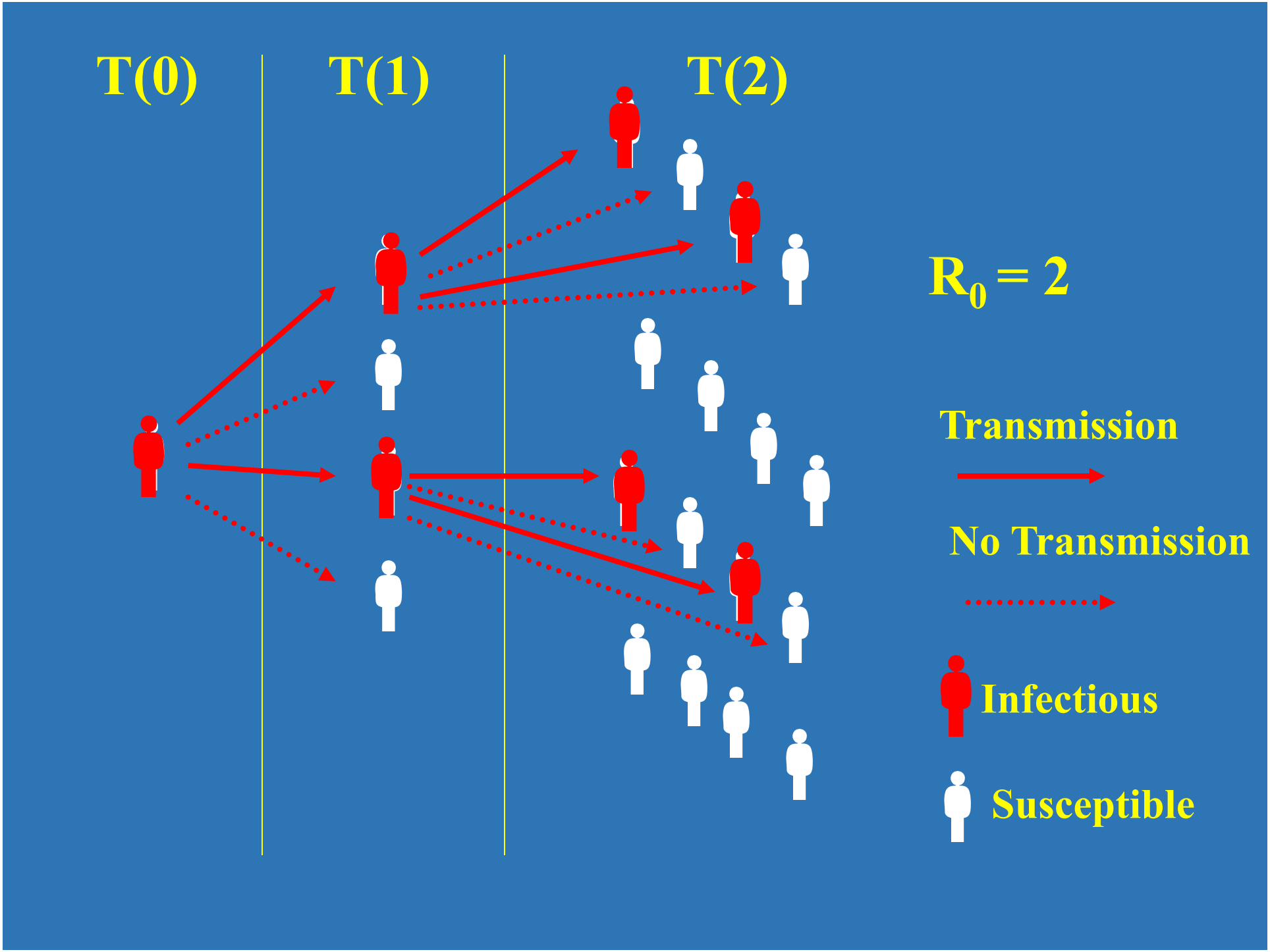
- PAR is a linear calculation
- Infectious diseases depend on non-linear effects
- Herd immunity—circumcision protections other uncircumcised individuals in population
- PAR does not take prevalence into account
- PAR underestimates effects of risk factors on infectious diseases.
- R_0 captures the non-linear dynamics of infectious diseases

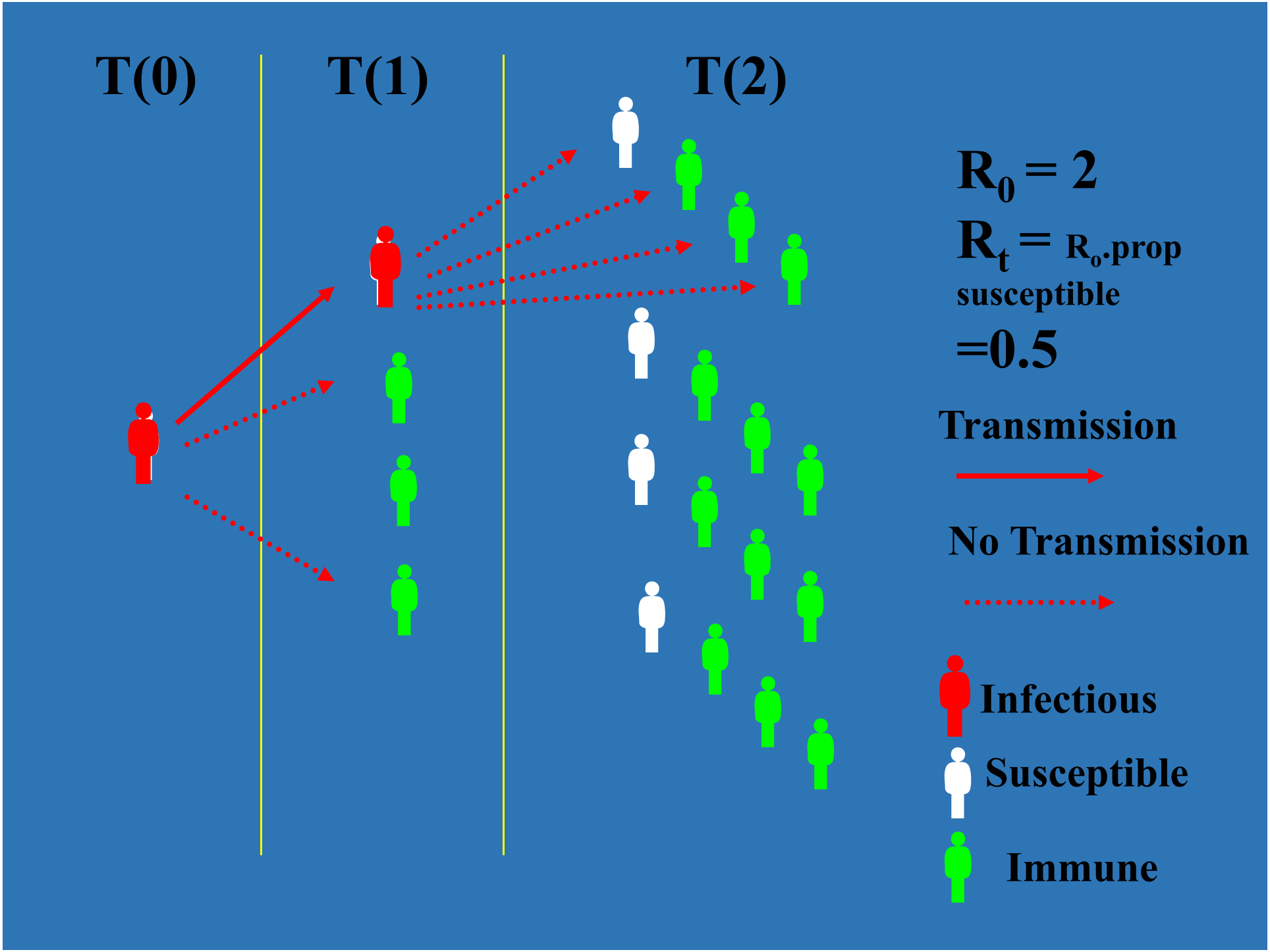
The basic and effective reproductive numbers

R_0 The Basic Reproductive Number - The number of new infections caused by one infection in an entirely susceptible population

R_t The Effective Reproductive Number - The number of new infections caused by one infection at a given time




T(0)
T(1)
T(2)
$R_0 = 2$
Transmission
No Transmission
Infectious
Susceptible


T(0)
T(1)
T(2)
$R_0 = 2$
$R_t = R_o.prop\ susceptible$
$=0.5$
Transmission
No Transmission
Infectious
Susceptible
Immune

How do we incorporate these important issues into modeling?

The answer depends on:

- the population to be modeled
- data availability
- the scientific question at hand, and
- the modeler's preferences

Types of models

- Static models – Decision Trees
- Dynamic models
  - Force of infection can change over time
  - Includes herd immunity
- Both static and dynamic models can be either deterministic or stochastic (constrained random variables)
- Choice of model depends on scientific question

Model classification

…in a simplified scheme…

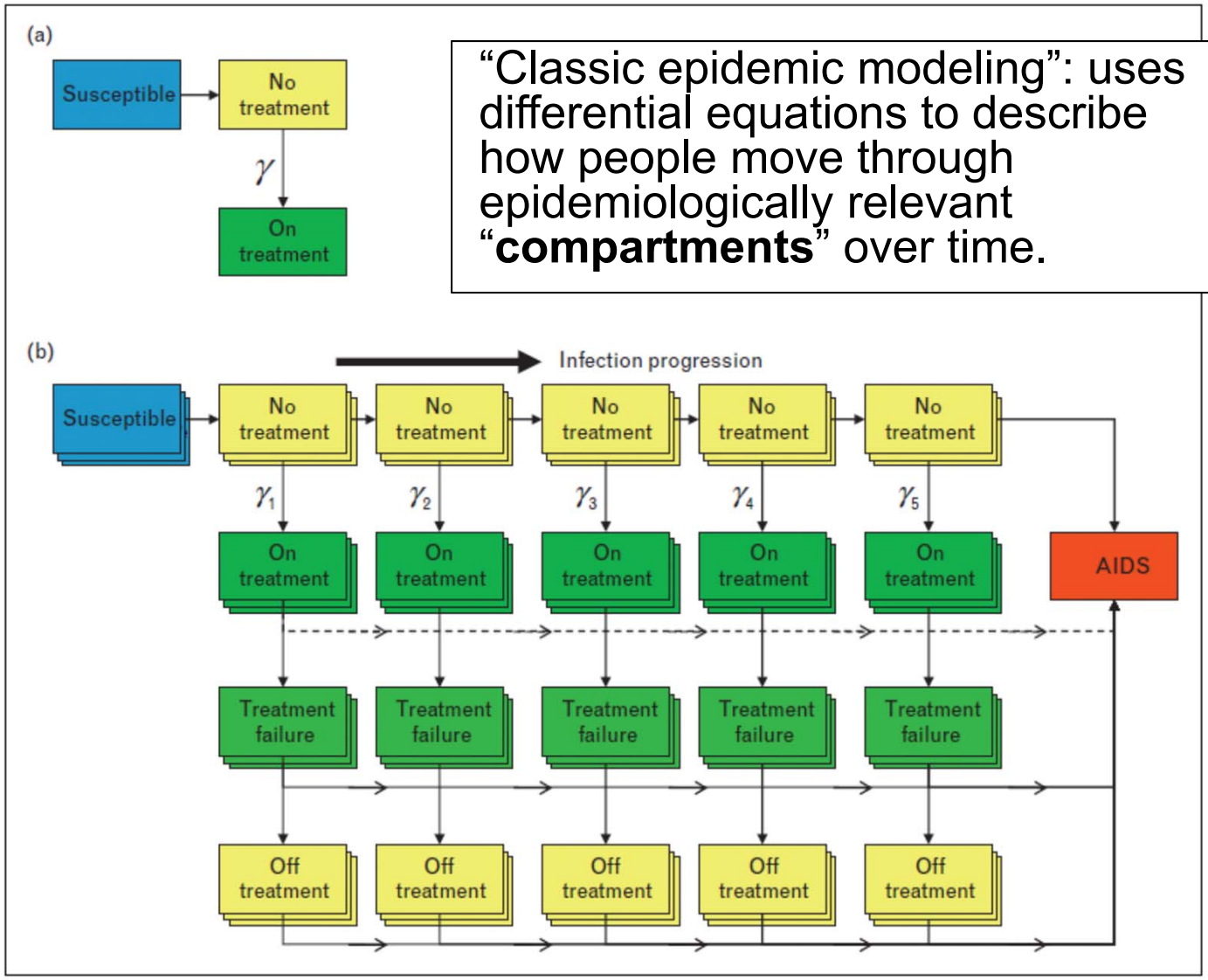
complexity →

Static Deterministic	Dynamic Deterministic
Static Stochastic	Dynamic Stochastic

complexity? ↓

Dynamic, deterministic model

Figure 3 Schematic illustrations of the structure of HIV transmission models incorporating antiretroviral therapy



"Classic epidemic modeling": uses differential equations to describe how people move through epidemiologically relevant "**compartments**" over time.

Baggeley & Fraser. Current Opinion in HIV/AIDS, 2010

How models are like maps

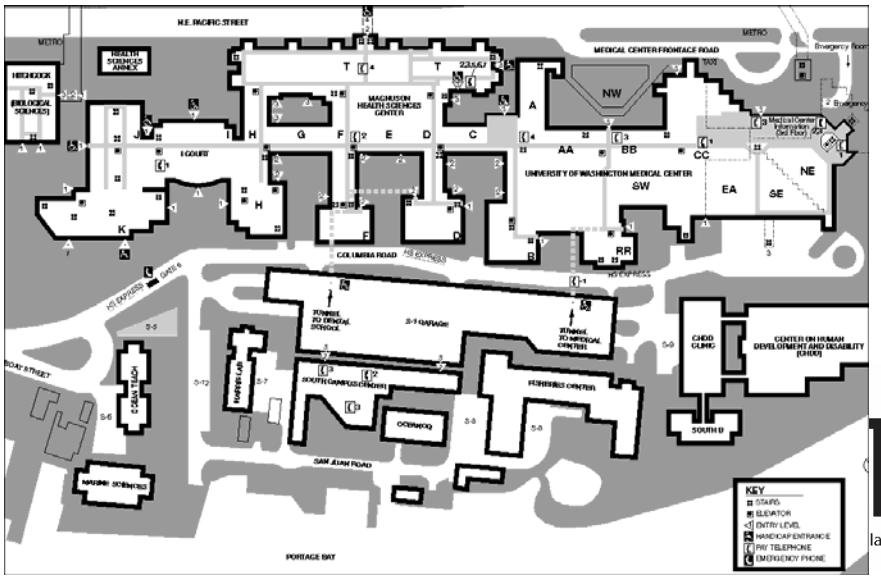
Like maps, models…

… are *abstractions*

… have *scale*

… must trade off *realism* with *generality*

The kind of model you use depends on the question you want to answer.



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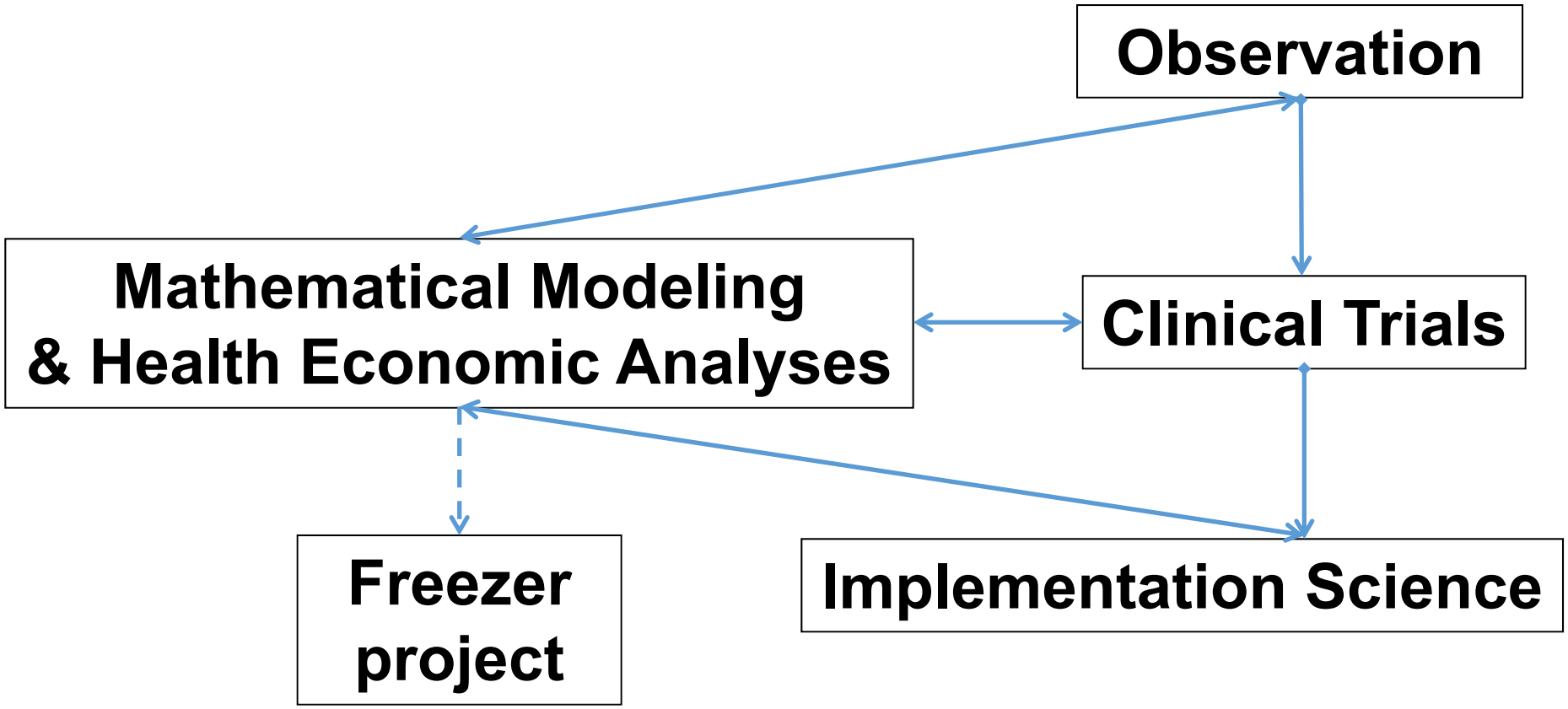
Where do models fit in the path from discovery to implementation?



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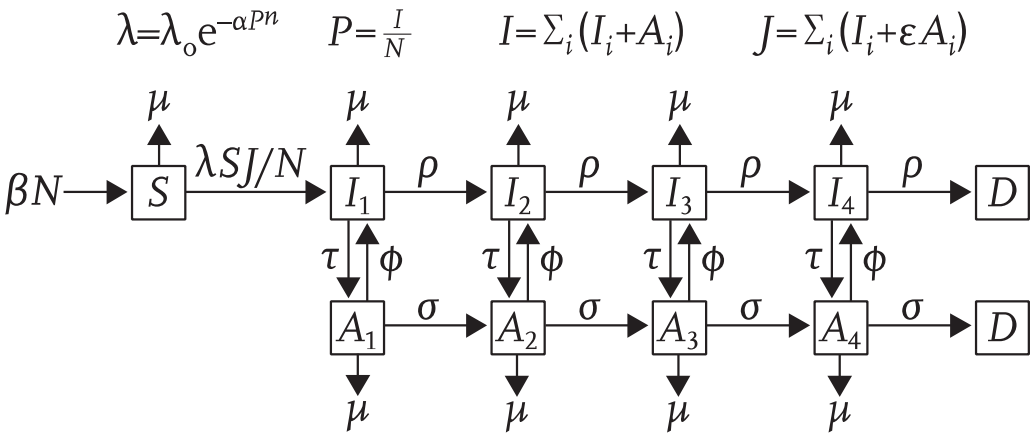
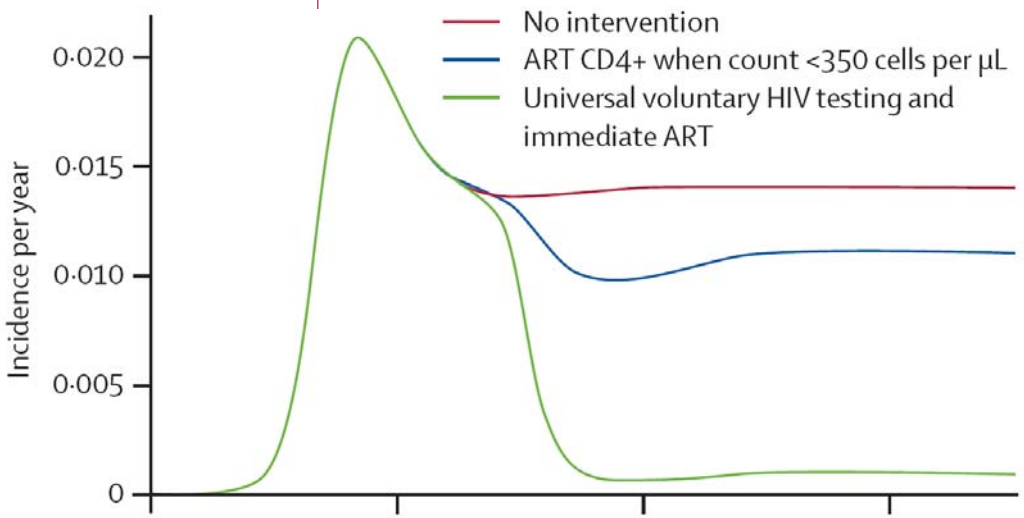
Universal voluntary HIV testing with immediate antiretroviral therapy as a strategy for elimination of HIV transmission: a mathematical model

Reuben M Granich, Charles F Gilks, Christopher Dye, Kevin M De Cock, Brian G Williams



Lancet 2008



Models can estimate potential impact of health programs

Treating our way out of the HIV pandemic: could we, would we, should we? *Geoffrey P Garnett, Rebecca F Baggaley

- "HIV prevention is easy in theory – the practice is hard."
- Need intensive HIV testing and robust linkages to care, even among people who feel well
- Strategies need to be effective and cost-effective

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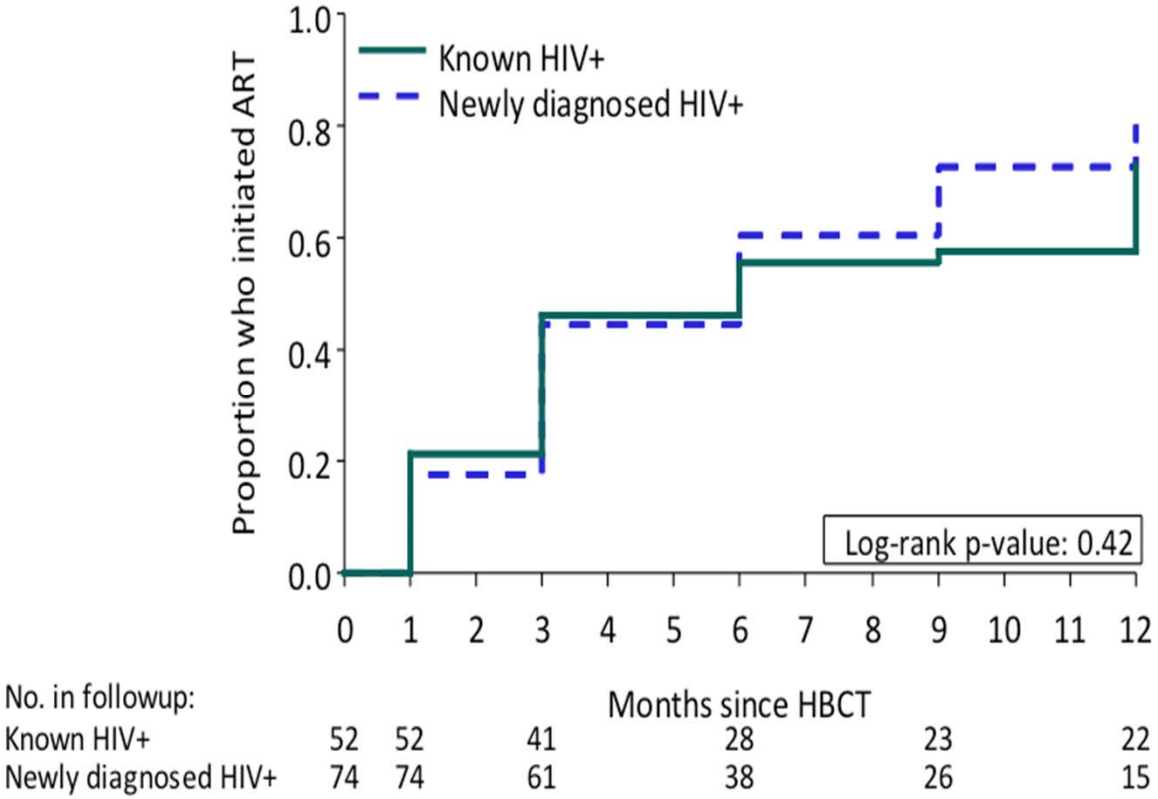
Initiation of antiretroviral therapy and viral suppression after home HIV testing and counselling in KwaZulu-Natal, South Africa, and Mbarara district, Uganda: a prospective, observational intervention study

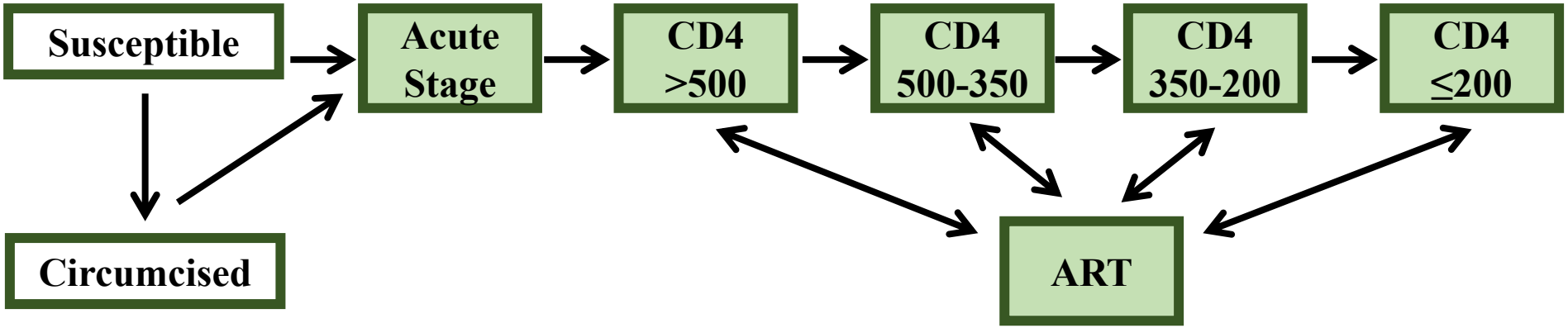
Ruanne V Barnabas, Heidi van Rooyen, Elioda Tumwesigye, Pamela M Murnane, Jared M Baeten, Hilton Humphries, Bosco Turyamureeba, Philip Joseph, Meighan Krows, James P Hughes, Connie Celum

74% ART uptake among eligible participants
(CD4 count ≤350 cells/µL and not on ART at enrollment)



ART Model: Structure

- Mathematical model to evaluate ART scale-up



- Realistic assumptions regarding testing and ART coverage

Using study data for models

- Demographics
- Mixing patterns
- Natural history
- Transmission probability
- Factors that change susceptibility
- Factors that change infectiousness
- Effectiveness of interventions
- Engagement in health care

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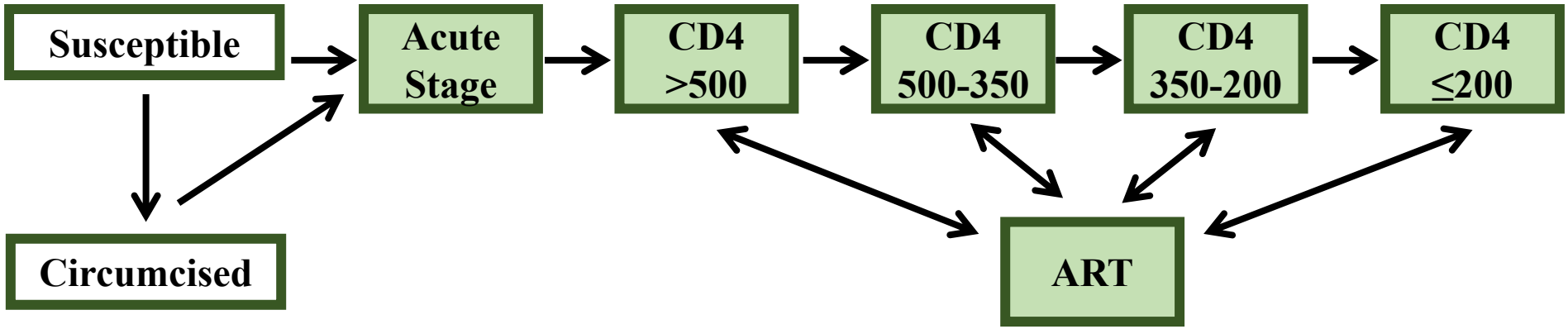
What study data can you use to parameterize models?



- Country specific demographics
- Distribution of CD4 count and viral load
- Intervention (including treatment) coverage and efficacy
- Factors that impact on HIV transmission: viral load, gender, circumcision status, co-infection status

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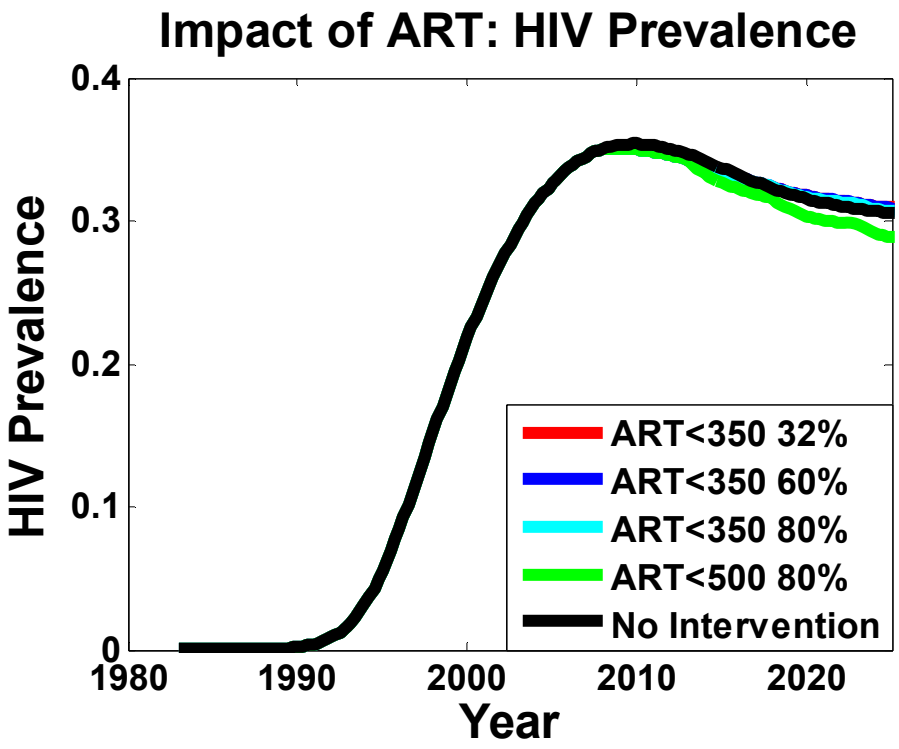
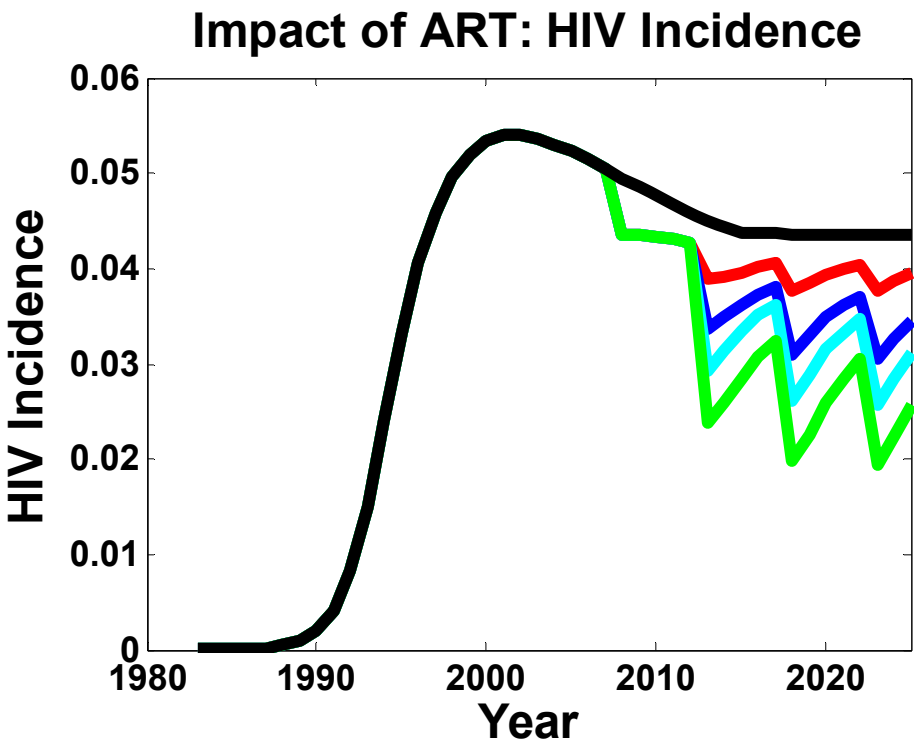
ART Model: Results





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Roger Ying, ISSTDR 2013

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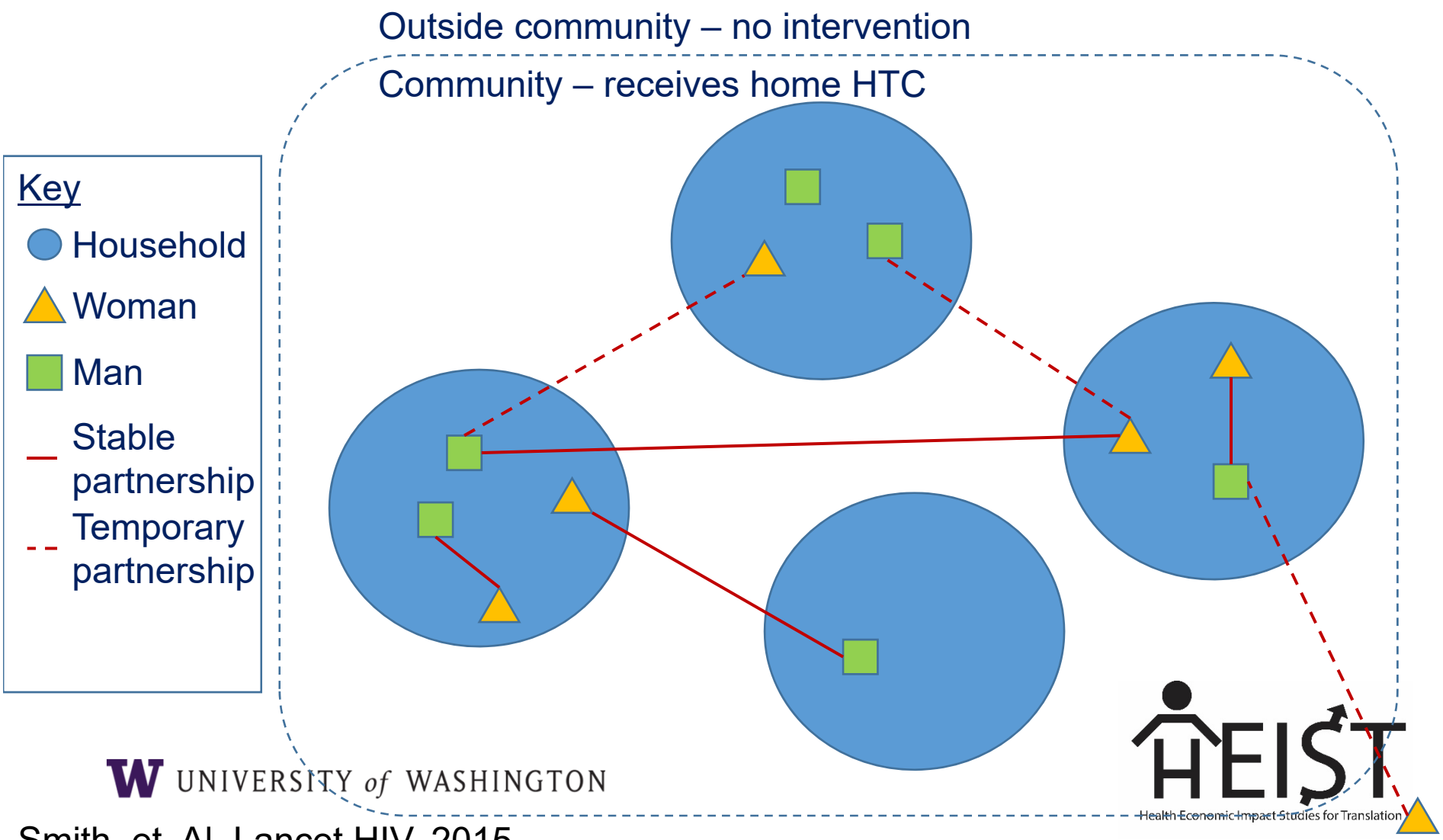
Cost-effectiveness

	Change in HIV incidence*	Change in HIV prevalence*	ICER per infection averted	ICER per death averted	ICER per QALY gained
Home HTC: ART for 48% of all HIV-positive people	−33·8%	−4·7%	Dominated†	$3290	$860
Home HTC plus CD4: additional ART for CD4 count 350–500 cells per μL	−40·6%	−6·7%	Dominated†	$4070	$900
Home HTC plus high viral load: additional ART for viral load >10 000 copies per mL	−51·6%	−12·1%	$2960	$5020	$1710

Results are shown for a 10 year time period of 2015–25 with a 6% annual dropout from ART care. Costs and effectiveness are discounted by 3% annually. ICER=incremental cost-effectiveness ratio. QALY=quality-adjusted life-year. HTC=home HIV testing and counselling. ART=antiretroviral therapy. *Relative to a "no ART" counterfactual. †A dominated strategy is more costly and less effective or more costly and less cost-effective than a combination of other interventions.

Table 3: **Effectiveness and cost-effectiveness of ART uptake from home HTC with varying ART initiation guidelines**

Model: community structure & partnerships



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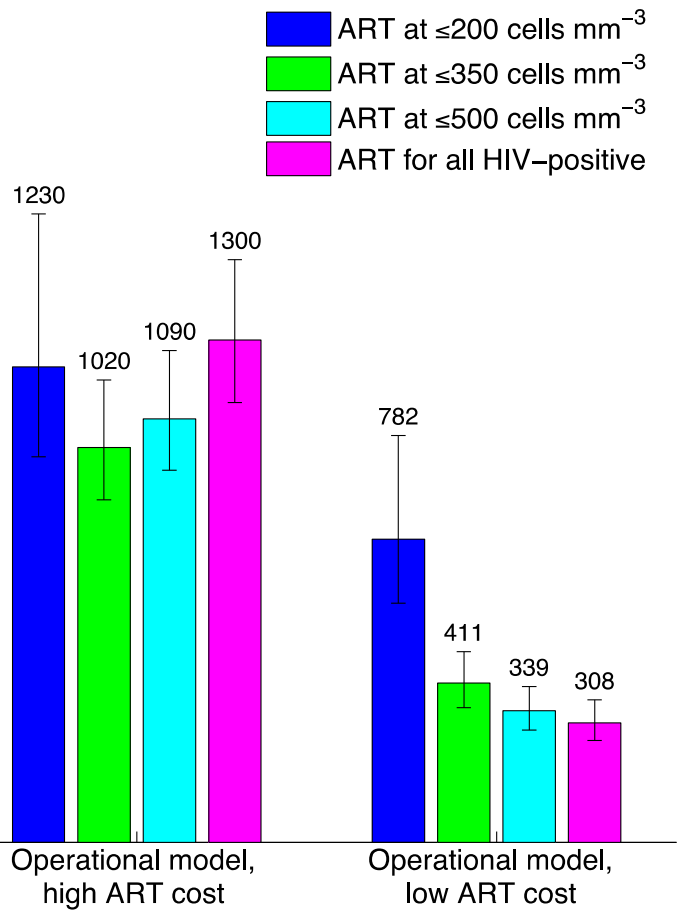
Smith, et. Al, Lancet HIV, 2015

Incremental cost per DALY averted

- **All ICERs per DALY averted are <20% of South African GDP per capita (2012), which by WHO standards are very cost-effective**
- **Reducing ART cost to CHAI target reduces ICER per DALY averted by 36-76%**



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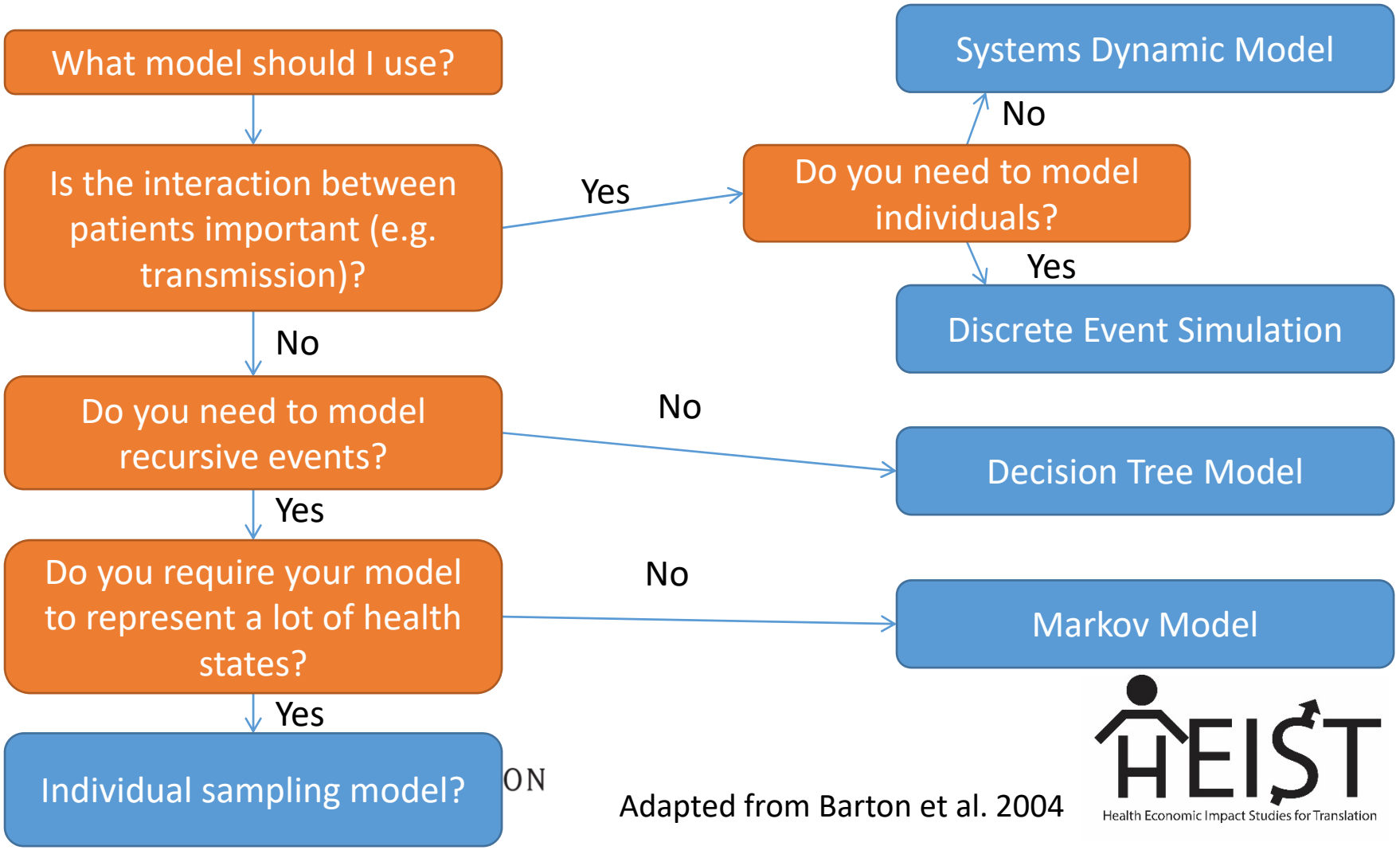
How to choose the appropriate model for health outcomes



Adapted from Barton et al. 2004

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Summary

- Infectious disease modeling is a useful tool
- Modeling can be used to estimate health outcomes
- Study data can be used to parameterize models

Primer on Decision Trees

James G. Kahn, MD, MPH

Professor of Health Policy, Epidemiology, & Global Health

University of California, San Francisco

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Overview

- What is a decision tree?
- When should I use a decision tree?
- How to construct a decision tree
- How to analyze a decision tree
- Software options

What is a decision tree?

- A branching structure that leads from a **choice** (among competing courses of action) through a probability net of possible **consequences** (temporary and final) …
- … in which each path of consequences has an associated **probability** and set of **outcomes** of interest (e.g., cost and health status) such that …
- … each course of action can be assigned **expected values** for the outcomes (as the weighted mean of relevant paths) that can be compared and used to **guide decisions** among the actions.

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Decision tree structure to evaluate interventions

Population	Decision node / Course of Action / Options	Chance nodes / Consequences	Path Probability	DALYs		Costs	
				For terminal nodes	Intermediate product	For terminal nodes	Intermediate product
Young men (smokers)	No intervention	cancer 0.2	die young 0.8	0.16	10	1.6	$20,000	$3,200
			die old 0.2	0.04	1	0.04	$20,000	$800
		no cancer 0.8	die young 0.2	0.16	8	1.28	$5,000	$800
			die old 0.8	0.64	0	0	$5,000	$3,200
				1.00		**2.92**		**$8,000**

Expected value for this option.

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			die old 0.8	0.64	0	0	$5,000	$3,200
				1.00		**2.92**		**$8,000**
	Smoking cessation program	cancer 0.15	die young 0.8	0.12	10	1.2	$21,000	$2,520
			die old 0.2	0.03	1	0.03	$21,000	$630
		no cancer 0.85	die young 0.2	0.17	8	1.36	$6,000	$1,020
			die old 0.8	0.68	0	0	$6,000	$4,080
				1.00		**2.59**		**$6,250**
					Difference	**-0.33**	**Difference**	**$250**
						ICER	**$758**	

Expected value for this option.

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When should I use a decision tree?

- **Conceptualizing: Almost always.** Extremely useful to develop and portray the structure of a cost-effectiveness analysis … clarify thinking, tighten logic, avoid omissions of possible paths. Can be used in conjunction with other visual portrayals of model dynamics.
- **Operationalizing: Often.** Assures that conceptual approach is reflected in implementation. Often used in conjunction with other calculation tools. Balance of tree & other calculation structures is personal preference.
- **Presenting: Sometimes.** Some analyses done with trees are presented with trees, some not.

How to construct a decision tree

- **Population & context**
- **Decision node (square)** – the question under study, 2 or more action options – all plausible (judgment call). Later decisions brought to front.
- **Chance nodes (circles)** – in each node probabilities sum to 100%. Mutually exclusive & exhaustive. Dichotomous easiest to manipulate. Markov can be incorporated.
- **Terminal node utilities = outcomes** – health, costs (direct, time)
- **Expected values** for health and costs, for each action option as weighted mean of paths.
- **Iterative revision** – unlike RCTs, the approach can (and nearly always does) change with early results and better understanding. The trick is knowing when to stop refining, and balancing completeness with transparency.

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How to analyze a decision tree

- **Comparisons across options** – compare expected values for costs and health outcomes … ordered (least to most expensive) & step-wise incremental … then incremental cost-effectiveness ratios (ICERs)

	Cost	Δ Cost	DALYs	Δ DALYs (averted)	ICER	
Option A	$1,000		10			
Option C	$1,500	$500	8	2.0	$250	
Option B	$1,700	$200	8.5	-0.5	Dominated	
Option D	$2,500	$800	7.5	1.0	$2,000	[vs C]

- **Sensitivity analyses** – 1-way, 2-way, scenarios, thresholds, multivariate (eg Monte Carl0).

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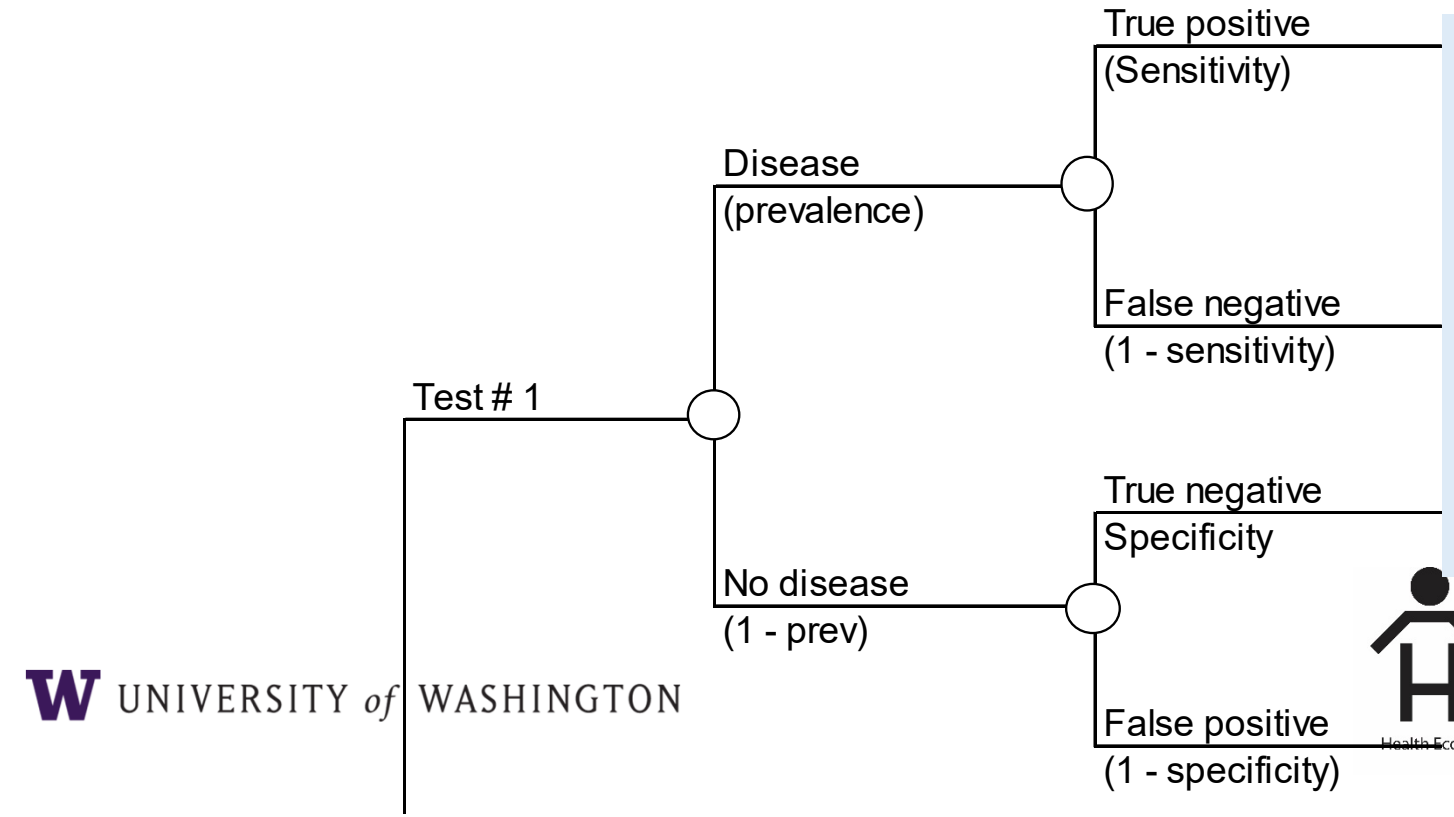
Software

- **Excel** – familiar, generic, flexible (eg incorporate epidemic and cost models), has sensitivity analysis add-ons (Crystal Ball, @Risk). My favorite. Consider starting with template.
- **TreeAge** – new, specialized, efficient for set CEA tasks, less flexible, quirky manual and implementation.
- **@Risk** – newer, specialized, efficient for set CEA tasks, powerful, complex, narrow market.

Extra credit: testing analysis
prevalence before test performance

Tree structure to evaluate diagnostic tests

Population	Decision: Test or not	True disease prevalence (risk of disease)	Test performance*



Why?

Forces equal disease prevalence in all arms, o/w source of error

Avoid ppv, npv, etc which are more complex (prevalence-dependent)

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- Thank you
- Email: rbarnaba@uw.edu

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