



University of Pittsburgh

# The New FRED (the FRamework for Epidemiological Dynamics)

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PHDL Seminar

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FRED

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## Outline

- Overview of Modeling in FRED
- User-defined Conditions and States
- Duration of States
  - distributions
  - absorbing, dormant and transient states
- State Transitions
  - logic-driven
  - probabilistic
  - event-driven
- Examples
- Future tutorial topics
  - web interface
  - plotting and visualization

## What is FRED?

- **FRED** is a **F**ramework for **E**pidemiological **D**ynamics
- **Framework:** FRED is not a model. FRED is a tool for building epidemiological models
- **Epidemiology:** the study and analysis of the patterns, causes, and effects of health and disease conditions in defined populations (Wikipedia)
  - Infectious and noninfectious diseases
  - Health-related conditions such as obesity, drug use, violence, vaccine acceptance, etc
- **Dynamics:** FRED is designed to study how patterns of health conditions in defined populations *vary over time*

*FRED is designed to build agent-based (individual-based) models*

## Agent-based Models

- Include each person in the model, along with social contacts and interactions with the environment
- Include individual responses and behaviors in the model
- Investigate interactions between public health dynamics and spatially distributed resources such as health care facilities

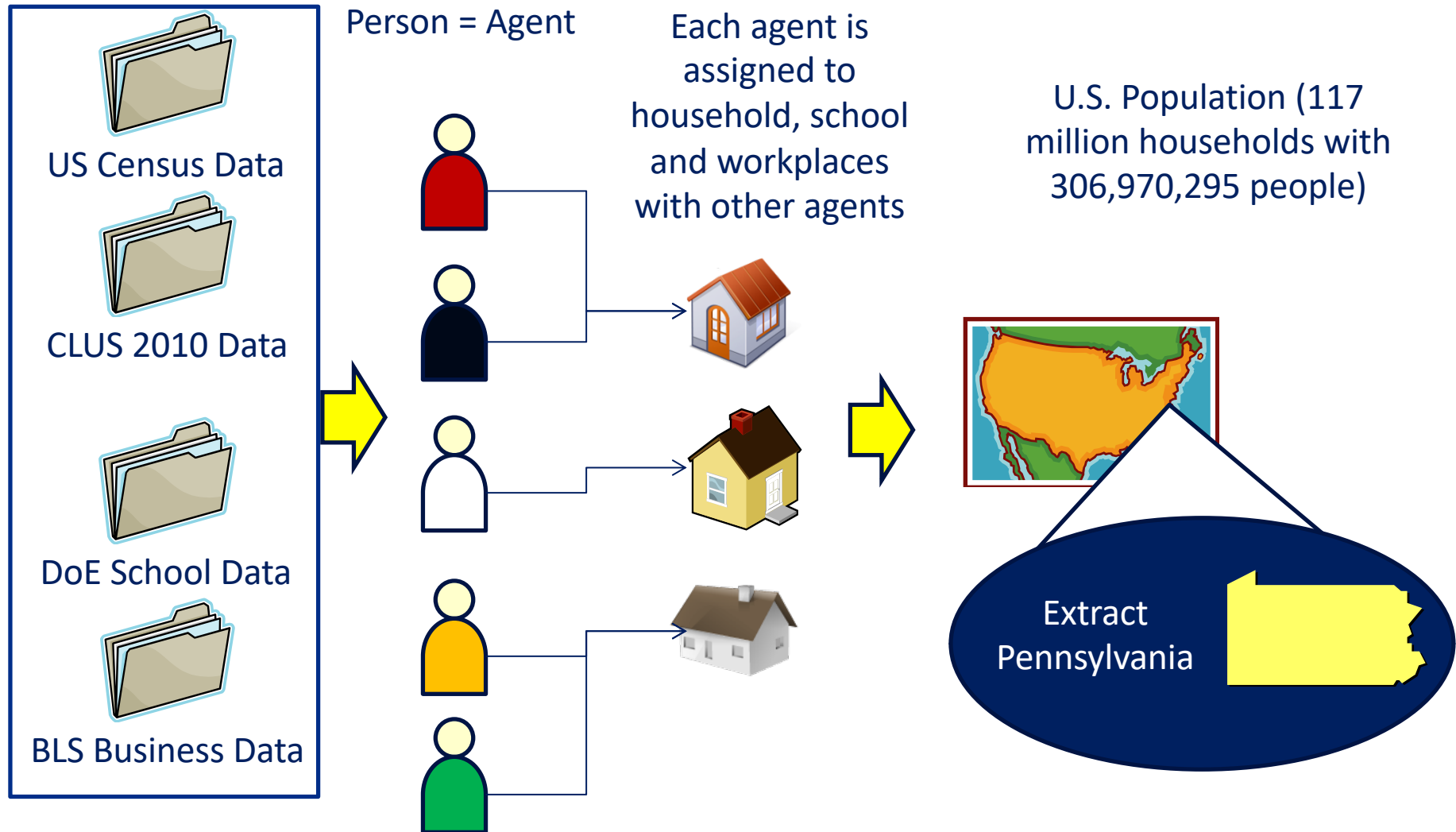
### Purposes

- Study how interactions among *individuals and their environment* can result in patterns of *population behavior*
- Study the *impact of policy and programs* on public health

## Foundational Concepts in FRED

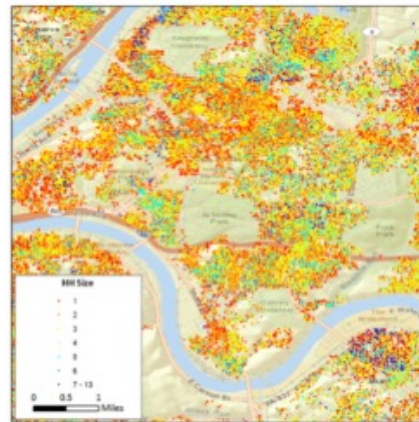
- Space
  - Two-dimensional geography based on actual locations
  - Future: 3-dimensional landscape (e.g. altitude)
- Time
  - Time step = 1 day (agents have multiple serial activities per day)
  - Duration = 1 day to 100 years
- Agent = individual person
- Places (mixing groups for agents)
  - Households, neighborhoods, workplaces, schools, hospitals
  - Future: other place types
- Population
  - Based on census data and other sources
  - Agents are associated with specific places

## Census-matched Synthetic Population

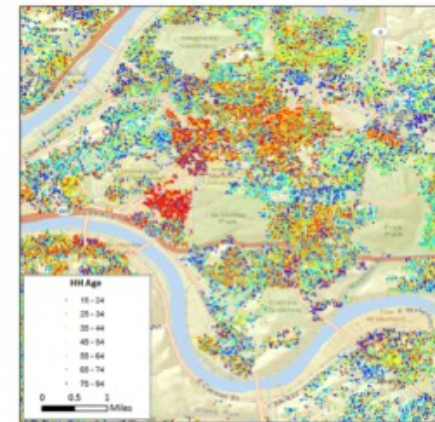


## Matches Actual Demographics

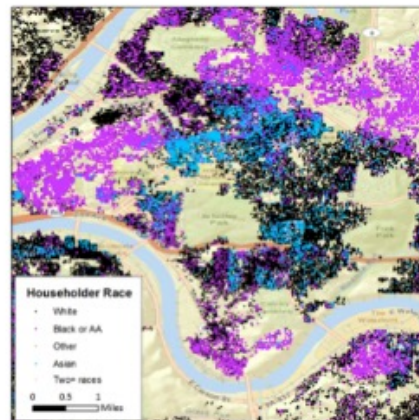
- Iterative proportional fitting assures that synthetic attributes are distributed as real ones are



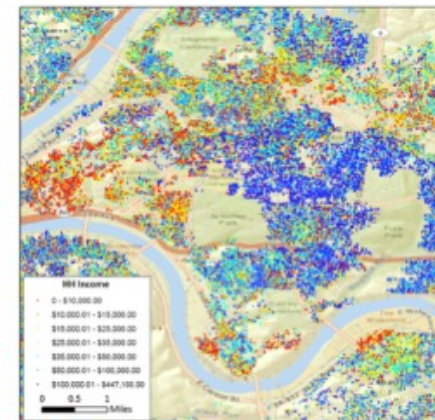
Household Size



Age



Race



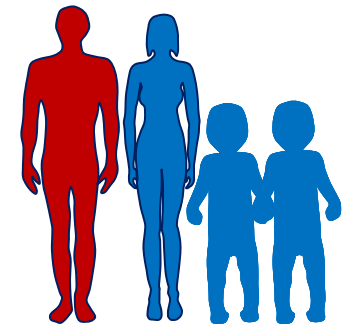
Income



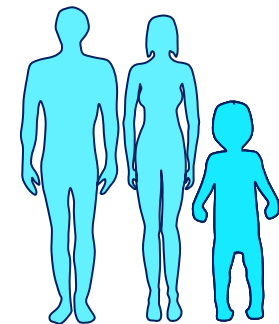
Location and size of  
each school



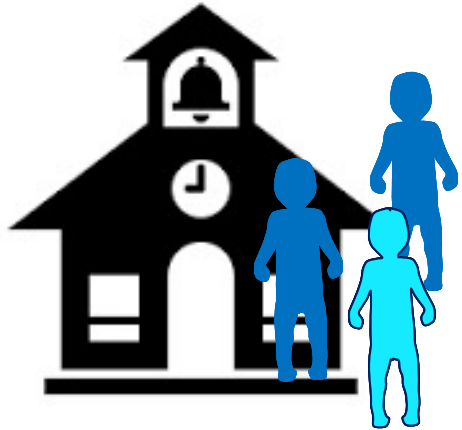
Household size,  
ethnicity, ages, income



Location and size of  
each workplace



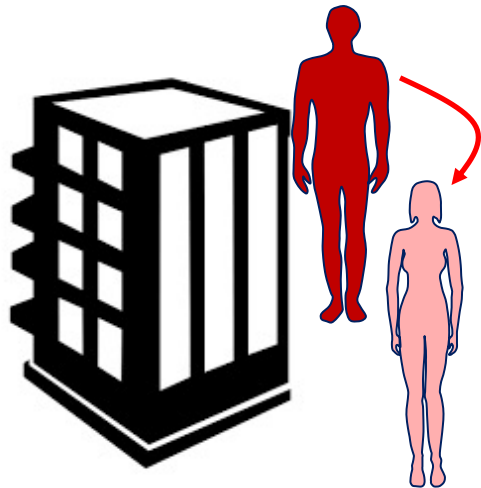




Location and size of  
each school



Household size,  
ethnicity, ages, income



Location and size of  
each workplace

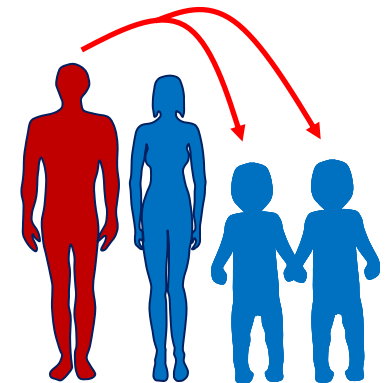




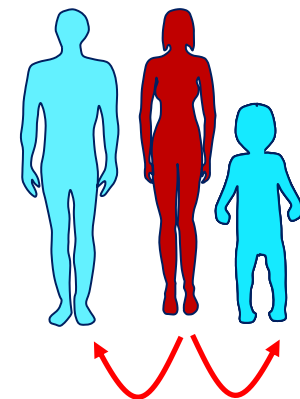
Location and size of each school



Household size, ethnicity, ages, income



Location and size of each workplace



## The New FRED: Easier Population Modeling

- No computer programming required
- You can focus on scientific effort (e.g. data collection, conceptual modeling, experimental design)
- You define (declare) all concepts of interest, rules for agents, environmental resources, initial conditions, and simulation parameters
- FRED processes the declarations, sets up the population, applies initial conditions, simulates the activities and interactions of the agents, and tracks all user-defined conditions within the population
- FRED outputs reports, charts, and visualizations
- FRED provides a simple workflow environment for you and manages all the data produced by the simulation and associated metadata

## How to Model with FRED

1. Decide if FRED is suited to your research problem
2. Conceptual modeling
3. Create rules for individuals
4. Create and run FRED Model
5. Revise model and repeat

## Decide if FRED is suited to the research problem

- FRED is well-suited for
  - Population-level phenomena arising from individual interactions
  - Phenomena with a straightforward "Natural History"
  - Phenomena with strong social determinants
  - Phenomena with strong spatial characteristics
  - Problem dependent on longitudinal demographic projections
- FRED is less well-suited for
  - Detailed physiological modeling
  - Problems associated with detailed interactions among small groups (e.g. resolution of domestic conflicts)
  - Problems depending on high-resolution motion (e.g. traffic models)

## Decide if FRED is suited to the research problem

Do you need this?



FRED

Or this?



NetLogo

## Conditions in FRED

- *Conditions* are patterns of interest defined over a population

### Examples

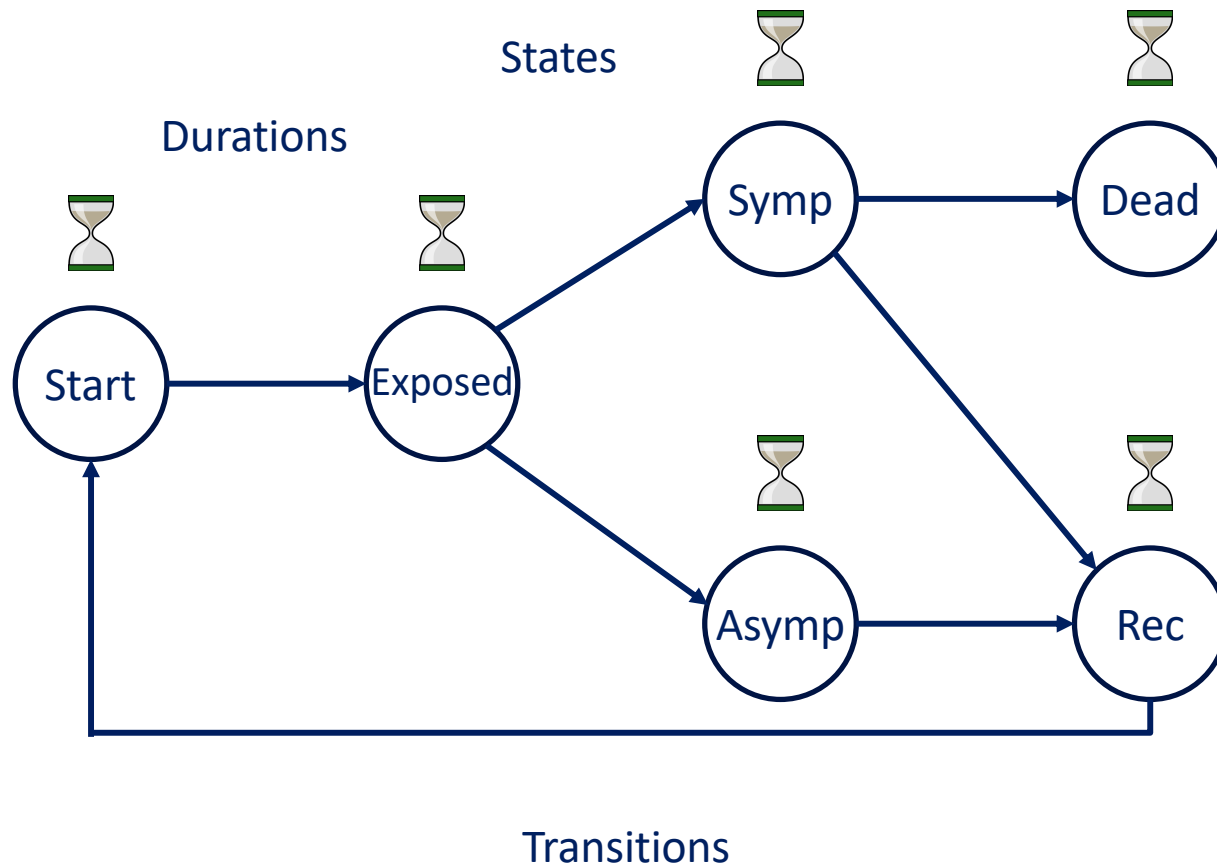
- Demographic conditions
  - maternity, mortality
- Health-related behaviors
  - vaccine acceptance and uptake, condom use, personal hygiene, exercise, violence, compliance, care-giving
- Social conditions
  - income inequality, access to insurance, gang activity, community activism, gun ownership
- Infectious diseases
  - flu, measles, dengue, chikungunya, HIV, HepC
- Noninfectious diseases
  - CVD, asthma, alzheimers, drug use disorder, obesity

## Conditions and States

- You can define any number of conditions
- Each condition has a user-defined set of *states*
  - Conditions and states can have whatever names you choose
- Each agent is in exactly one state associated with each condition
  - The first named state is the initial state for each agent
- You define the duration of each state
  - Selected from statistical distribution or changed by side effect
- You provide rules for changing states
  - Logistic regression for probability of changing to any other state
- State can have health side effects
  - Transmissibility, susceptibility, symptoms, mortality, maternity
- States can have social contact side effects
  - Absent from school or work, isolation, hospitalization, etc.
- States can have side effects on other conditions



## Condition = Influenza



## Duration of States

- For every condition, each agent has a *current state*
- For every condition, each agent has at most one *scheduled state transition*
- When a state transition time arrives, the *transition matrix* is used to select the agent's *next state*
- You define the state durations and the state transitions

## Duration of States

- How long before an agent changes state?

Distribution	Parameters	Use
normal	mean, std	produced by many small effects acting additively and independently e.g., human gestation period
lognormal	median, dispersion	produced by multiplication many effects, each of which is positive e.g., duration of illness, length of marriage before divorce
geometric	mean (mean = $1.0/p$ )	time before success of Bernoulli trials with prob $p$ . e.g., time before winning a lottery
uniform	min, max	e.g., value of a dice toss

- Transient states
  - States with duration 0
- Absorbing states have infinite duration
  - examples: Death, Race, Male/Female (in models without Transgenders)
- Dormant states
  - States that FRED doesn't need to track (for efficiency)

## Example: Pandemic Influenza Natural History

The states are: Susceptible, Exposed, Infectious, and Recovered

Rules:

1. People start out susceptible and remain susceptible until they are infected (that is, exposed).
2. People who are exposed become infectious after about 1-3 days.
3. People who are infectious recover after about 3-8 days.
4. Infectious people have moderate symptoms and stay home about 50% of the time.
5. Once recovered, people remain immune.

```
INFLUENZA.states = Susceptible Exposed Infectious Recovered
```

```
# TRANSMISSION
```

```
INFLUENZA.transmission_mode = respiratory
```

```
INFLUENZA.transmissibility = 1.0
```

```
# RULE 1
```

```
INFLUENZA.Susceptible.susceptibility = 1.0
```

```
INFLUENZA.exposed_state = Exposed
```

```
# RULE 2 (from Lessler, 2009)
```

```
INFLUENZA.Exposed.duration_distribution = lognormal
```

```
INFLUENZA.Exposed.duration_median = 1.9
```

```
INFLUENZA.Exposed.duration_dispersion = 1.51
```

```
INFLUENZA.transition[Exposed][Infectious] = 1
```

```
# RULE 3 (Citation?)
```

```
INFLUENZA.Infectious.transmissibility = 1.0
```

```
INFLUENZA.Infectious.duration_distribution = lognormal
```

```
INFLUENZA.Infectious.duration_median = 5.0
```

```
INFLUENZA.Infectious.duration_dispersion = 1.5
```

```
INFLUENZA.transition[Infectious][Recovered] = 1
```

```
# RULE 4
```

```
INFLUENZA.Infectious.symptoms = moderate
```

```
INFLUENZA.Infectious.probability_of_household_confinement = 0.5
```

```
# RULE 5
INFLUENZA.Recovered.is_dormant = 1

#####

# SIMULATION SETUP

# DURATION
days = 100

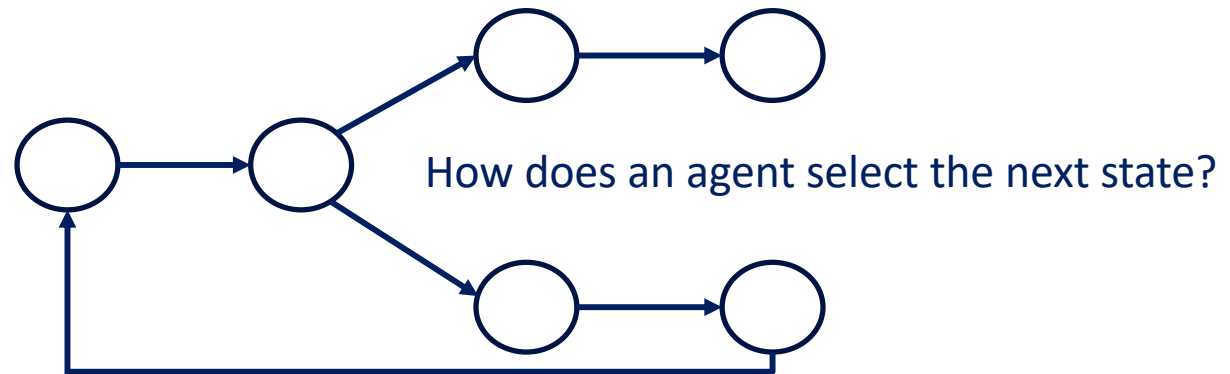
# POPULATION
locations_file = locations.txt

# CONDITIONS
conditions = INFLUENZA

# IMPORT CASES
INFLUENZA.import_file = import_influenza.txt

# VISUALIZATION
enable_visualization_layer = 1
INFLUENZA.Exposed.visualize = 1
INFLUENZA.Infectious.visualize = 1
INFLUENZA.Recovered.visualize = 1
```

## Transition Rules



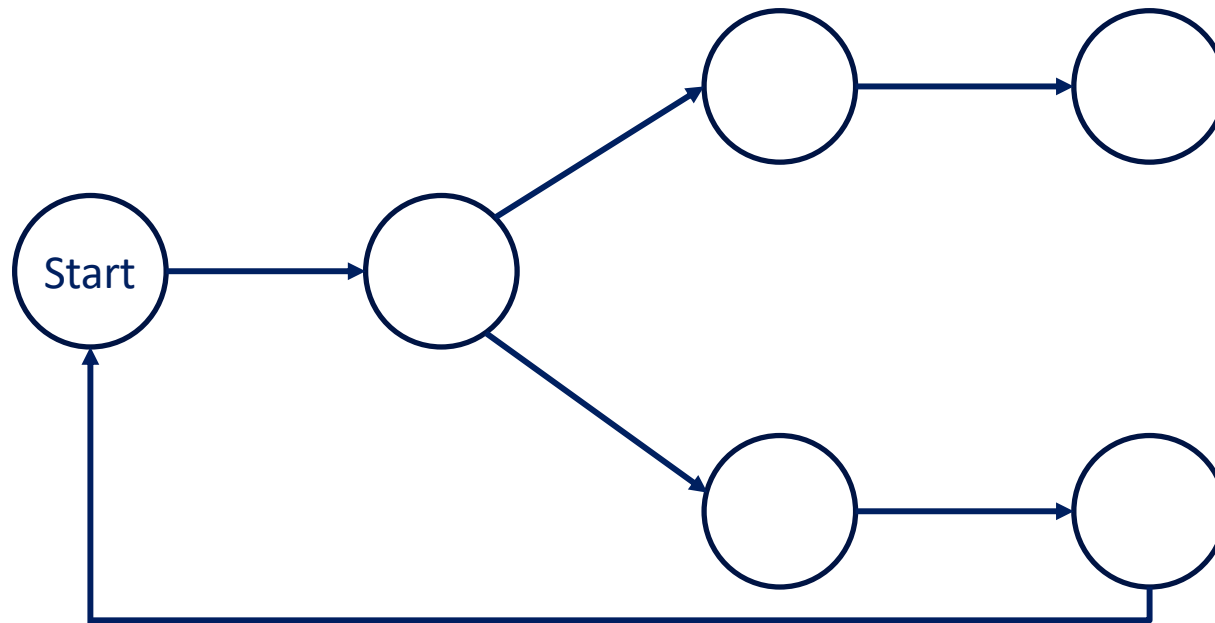
State Transition Matrix

	Next State		
State	A	B	C
A	0.0	0.5	0.5
B	0.1	0.8	0.1
C	0.0	0.0	1.0

To select next state:

1. Compute probabilities in row of current state
2. Select next state using the distribution

## FRED Transition Rules



Rules can be:

- Logic-driven
- Probability-driven
- Event-driven



## Factors that can affect an agent's state

### Social determinants:

- Demographics: age, race, sex, number of children
- Household: location (state, county or census tract), size, income
- Workplace or school: location, size
- Activity profile: pre-school, student, employed, retired, prisoner, military, nursing home resident, dorm resident

### Agent's own state

- Current health states: active conditions, immunity status, infection status, symptoms, where exposure occurred

### Other People's States

- Current epidemic levels of any user-defined condition in the population or in my mixing groups

### Environmental States

- Time: year, month, day of week

## SOCIAL DETERMINANTS

age  
age\_squared  
log\_of\_age  
age\_is\_<a> = 0.0 for a in {0,110}  
age\_is\_<a>+ = 0.0 for a in {0,110}  
age\_is\_<a>-<b> for a in {0,5,...,110}  
and b in {a+4,a+9,...}  
is\_female  
is\_male  
race\_is\_white  
race\_is\_nonwhite  
race\_is\_african\_american  
race\_is\_american\_indian  
race\_is\_alaska\_native  
race\_is\_tribal  
race\_is\_asian  
race\_is\_hawaiian\_native  
race\_is\_other  
race\_is\_multiple  
has\_household\_income\_in\_first\_quartile  
has\_household\_income\_in\_second\_quartile  
has\_household\_income\_in\_third\_quartile  
has\_household\_income\_in\_fourth\_quartile  
has\_household\_income\_below\_median  
has\_household\_income\_above\_median  
household\_size\_is\_1  
household\_size\_is\_2  
household\_size\_is\_3  
household\_size\_is\_4  
household\_size\_is\_5  
household\_size\_is\_6  
household\_size\_is\_7  
household\_size\_is\_8  
household\_size\_is\_9  
household\_size\_is\_10+  
is\_resident\_in\_group\_quarters  
is\_resident\_in\_college\_dorm  
is\_resident\_in\_nursing\_home  
is\_resident\_in\_military\_barracks  
is\_resident\_in\_prison

## SOCIAL DETERMINANTS (cont'd)

is\_student  
is\_employed  
is\_unemployed  
is\_retired  
number\_of\_children\_is\_0  
number\_of\_children\_is\_1  
number\_of\_children\_is\_2  
number\_of\_children\_is\_3  
number\_of\_children\_is\_4  
number\_of\_children\_is\_5  
number\_of\_children\_is\_6+  
state\_fips\_is\_NN  
county\_fips\_is\_NNNNN  
census\_tract\_fips\_is\_NNNNNNNNNNN  
household\_threshold\_exceeded  
household\_threshold\_not\_exceeded  
neighborhood\_threshold\_exceeded  
neighborhood\_threshold\_not\_exceeded  
school\_threshold\_exceeded  
school\_threshold\_not\_exceeded  
workplace\_threshold\_exceeded  
workplace\_threshold\_not\_exceeded

## ENVIRONMENTAL STATES

year\_is\_2000  
...  
year\_is\_2100  
log\_current\_year  
month\_is\_january  
..  
month\_is\_december  
day\_of\_week\_is\_sunday  
...  
day\_of\_week\_is\_saturday  
is\_weekday  
is\_weekend

## PERSONAL STATES

is\_in\_state\_COND.STATE  
was\_ever\_exposed\_to\_COND  
has\_symptoms\_of\_COND  
is\_transmissible\_for\_COND  
exposed\_to\_COND\_in\_household  
exposed\_to\_COND\_in\_neighborhood  
exposed\_to\_COND\_in\_school  
exposed\_to\_COND\_in\_workplace

## OTHER PEOPLE'S STATES

any\_others\_in\_household\_in\_state\_COND.STATE  
any\_others\_in\_neighborhood\_in\_state\_COND.STATE  
any\_others\_in\_school\_in\_state\_COND.STATE  
any\_others\_in\_workplace\_in\_state\_COND.STATE  
log\_of\_number\_in\_household\_in\_state\_COND.STATE  
log\_of\_number\_in\_neighborhood\_in\_state\_COND.STATE  
log\_of\_number\_in\_school\_in\_state\_COND.STATE  
log\_of\_number\_in\_workplace\_in\_state\_COND.STATE  
incidence\_threshold\_exceeded\_for\_state\_COND.STATE  
prevalence\_threshold\_exceeded\_for\_state\_COND.STATE  
cumulative\_threshold\_exceeded\_for\_state\_COND.STATE  
log\_of\_incidence\_count\_for\_state\_COND.STATE  
log\_of\_prevalence\_count\_for\_state\_COND.STATE  
log\_of\_cumulative\_count\_for\_state\_COND.STATE

## Logic-driven State Transitions

- You can assign transition probabilities based on *any logical combinations of the factors*
  - FRED is *propositionally complete*, meaning that you can control state transitions based on any combination of IF-THEN-ELSE, AND, OR, and NOT statements
- This often involves transient (duration = 0) states as intermediary states

## Asthma prevalence

Age	White		Nonwhite	
	Male	Female	Male	Female
0-4	3.7	3.7	11.5	9.0
5-14	9.5	8.4	20.1	17.7
15-19	9.4	11.2	12.8	17.9
20-24	6.5	11.5	14.3	13.2
25-34	6.5	10.5	8.8	12.5
35-64	5.9	10.7	6.3	12.7
65+	5.5	7.5	7.8	7.9

Source: National Health Interview Survey, National Center for Health Statistics, CDC

To use this in FRED:

1. Write rules to separate the population into the given subgroups
2. Assign the condition risk according to the given prevalence

## Logic-driven State Transitions

```
conditions = GROUP ASTHMA
```

```
GROUP.states = Start Female Male WhiteF WhiteM NonwhiteF NonwhiteM WhiteF0-4 ...
```

```
GROUP.Start.duration = 0
```

```
GROUP.transition[Start][Female].is_female = 1
```

```
GROUP.transition[Start][Male].is_male = 1
```

```
GROUP.Female.duration = 0
```

```
GROUP.transition[Female][WhiteF].race_is_white = 1
```

```
GROUP.transition[Female][NonwhiteF].race_is_nonwhite = 1
```

```
GROUP.WhiteF.duration = 0
```

```
GROUP.transition[WhiteF][WhiteF0-4].age_is_0-4 = 1
```

```
GROUP.transition[WhiteF][WhiteF5-9].age_is_5-9 = 1
```

```
...
```

# Probability-driven State Transitions

## Traditional Transition Matrix

	Next State		
State	A	B	C
A	0.0	0.5	0.5
B	0.1	0.8	0.1
C	0.0	0.0	1.0

### Challenges:

- Make transitions more individualized
- Take into account dynamic aspects
- State space explosion

## Probability-driven State Transitions

```
conditions = GROUP ASTHMA
```

```
GROUP.states = Start Female Male WhiteF WhiteM NonwhiteF NonwhiteM WhiteF0-4 ...
```

```
GROUP.Start.duration = 0
```

```
GROUP.transition[Start][Female].is_female = 1
```

```
GROUP.transition[Start][Male].is_male = 1
```

```
GROUP.Female.duration = 0
```

```
GROUP.transition[Female][WhiteF].race_is_white = 1
```

```
GROUP.transition[Female][NonwhiteF].race_is_nonwhite = 1
```

```
GROUP.WhiteF.duration = 0
```

```
GROUP.transition[WhiteF][WhiteF0-4].age_is_0-4 = 1
```

```
GROUP.transition[WhiteF][WhiteF5-9].age_is_5-9 = 1
```

```
...
```

```
ASTHMA.states = Start AtRisk None
```

```
ASTHMA.Start.duration = 0
```

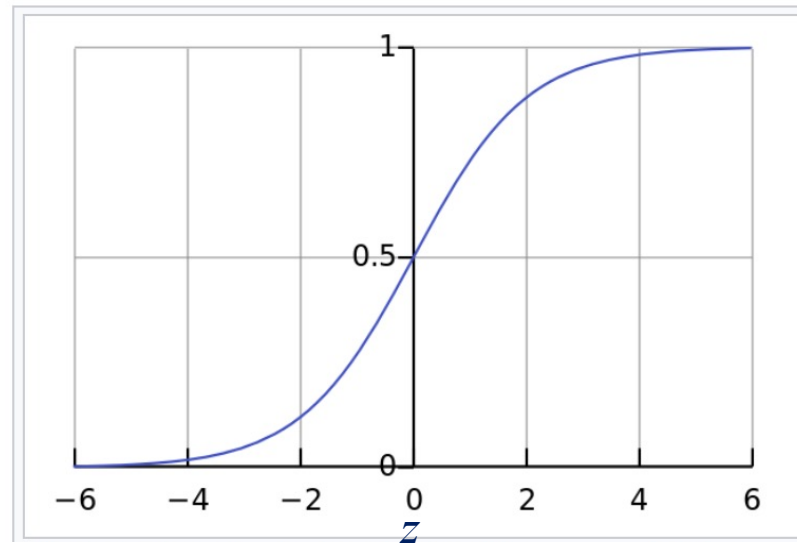
```
ASTHMA.transition[Start][AtRisk].is_in_state_GROUP.WhiteF0-4 = 0.037
```

```
ASTHMA.transition[Start][None].is_in_state_GROUP.WhiteF0-4 = 0.963
```

## Logistic Regression

$$\sigma(z) = 1 / (1 + \exp(-z)) \quad \text{and} \quad z = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_n x_n$$

coefficient      risk factor



Wikipedia

Standard logistic regression function  $\sigma(z)$ .  
Note that  $\sigma(z)$  returns a value (0, 1) for all  $z$ .



# Probability-driven State Transitions

## FRED Transition Matrix

	Next State		
State	A	B	C
A	$\alpha(z_{AA})$	$\alpha(z_{AB})$	$\alpha(z_{AC})$
B	$\alpha(z_{BA})$	$\alpha(z_{BB})$	$\alpha(z_{BC})$
C	$\alpha(z_{CA})$	$\alpha(z_{CB})$	$\alpha(z_{CC})$

Each entry has its own set of coefficients.

To select next state:

1. Compute probabilities in row of current state
2. Normalize
3. Select next state using the distribution

## Probability-driven State Transitions

(with thanks to Hawre)

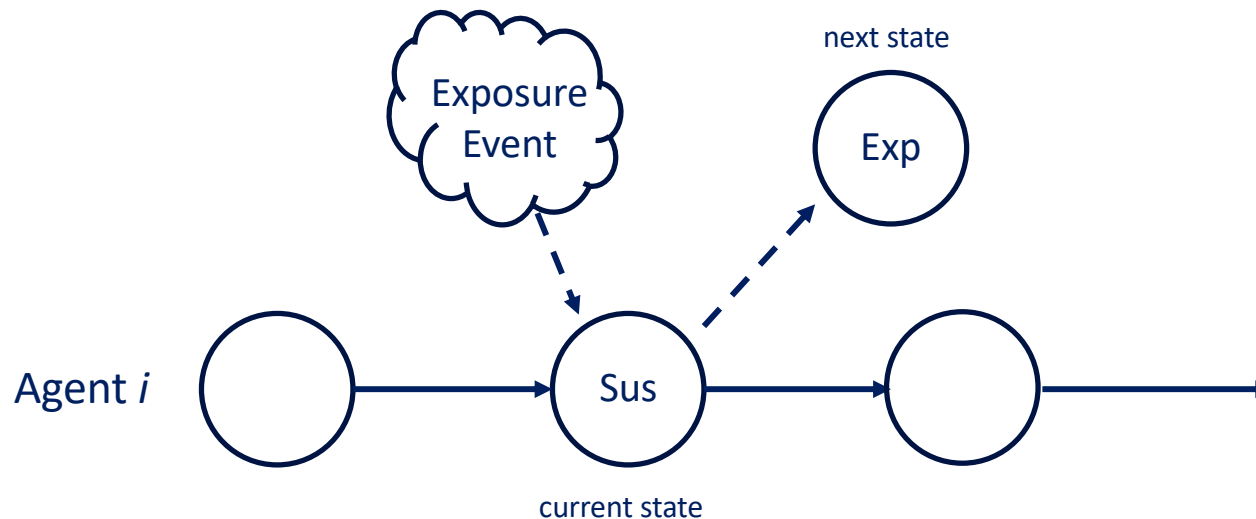
The probability of heroin use onset is described by:

$$\text{Prob(Heroin)} = 1 / (1 + \exp(-z))$$

where  $z = -214.8 + 0.112 * \text{age} - 0.00151 * \text{age}^2 - 0.18 * \text{female} + 1.89 * \text{po} + 2.447 * \text{cocaine} + 0.979 * \text{meth} + 1.256 * \text{mj}$

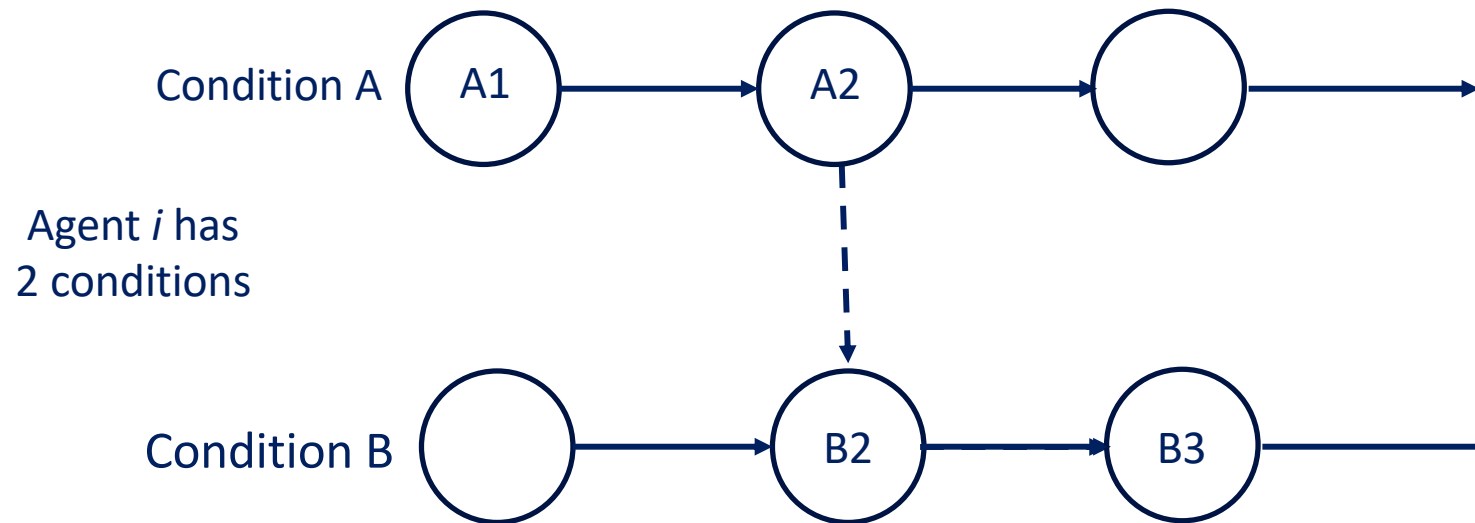
```
conditions = HEROIN PO COCAINE METH MJ
HEROIN.states = NonUse Use ProblemUse
HEROIN.duration = 30
HEROIN.transition[NonUse][Use].coeff = -214.8
HEROIN.transition[NonUse][Use].age = 0.112
HEROIN.transition[NonUse][Use].age_squared = -0.00151
HEROIN.transition[NonUse][Use].is_female = -0.18
HEROIN.transition[NonUse][Use].is_in_state_PO.Use = 1.89
HEROIN.transition[NonUse][Use].is_in_state_COCAINE.Use = 0.979
HEROIN.transition[NonUse][Use].is_in_state_MJ.Use = 1.256
```

## Event-driven State Transitions (Part 1)



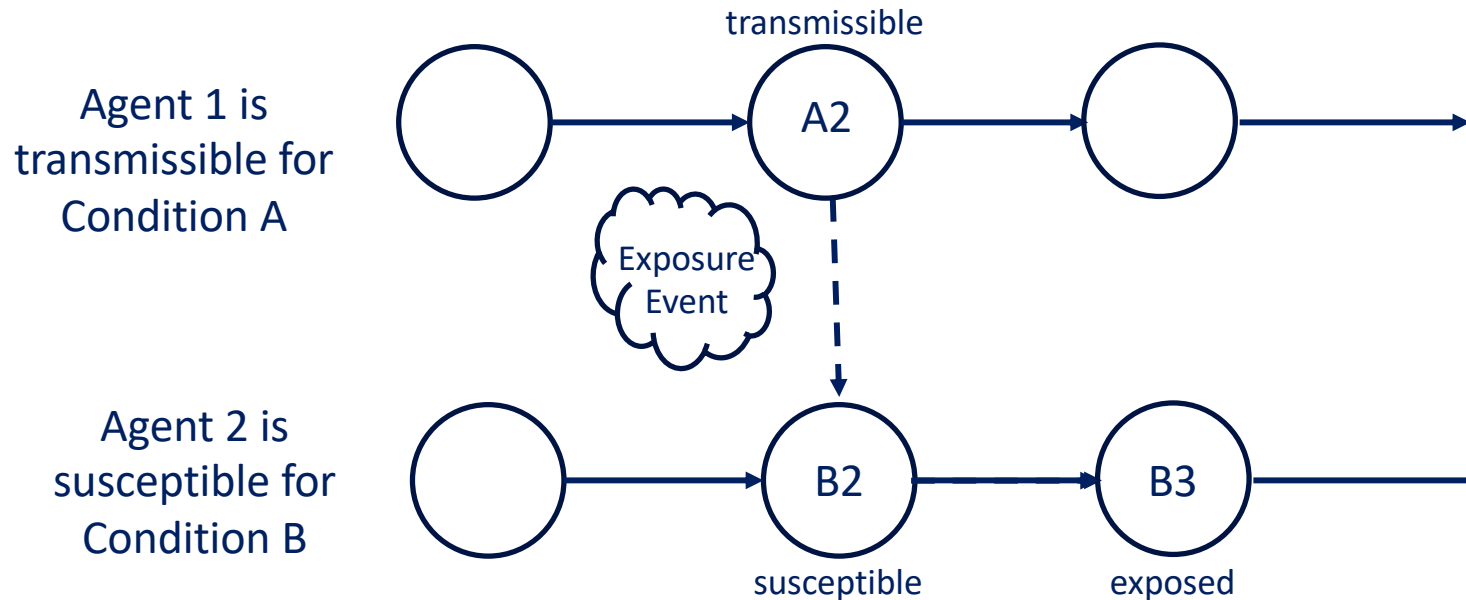
- Agents can be exposed to any condition through simulation events
  - *interaction with another agent who is transmissible for this condition*
  - *imported cases caused by an unidentified source*
- User defines the schedule of imported cases: day, location, quantity
- Agents must be in a susceptible state to be exposed

## Event-driven State Transitions (Part 2)



- When an agent enters a state, it can have a *side effect* on a state in another condition, causing the agent to change to a different state
- Example: when a person in a VACCINATION condition changes from state *Received* into the *Success* state, that person changes its state in the INFLUENZA condition from *Susceptible* to *Immune*

## Event-driven State Transitions (Part 3)



- We can tell FRED that Condition A causes exposure to Condition B
- If agent 1 "transmits" to agent 2, then agent 2 is exposed to condition B
- Examples:
  - drug dealer sells to drug user
  - perpetrator inflicts violence on victim
  - offspring inherits wealth from parent

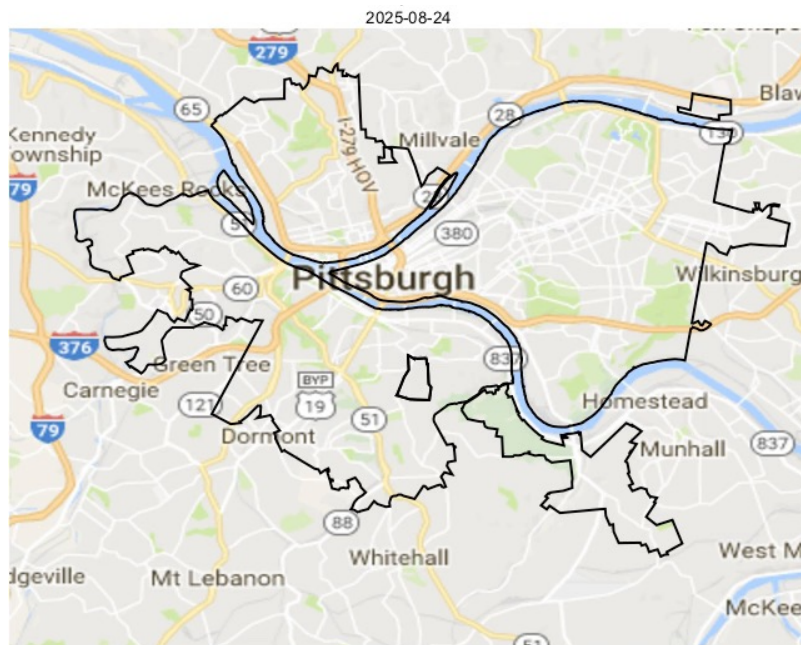
## Event-driven State Transitions (Part 3)

```
conditions = PERPETRATOR VICTIM
PERPETRATOR.states = Start Inactive Active Arrested
VICTIM.states = Safe AtRisk Injured Recovered Dead
...
# Condition Properties
PERPETRATOR.transmission_mode = proximity
PERPETRATOR.transmissibility = 0.01
...
VICTIM.exposed_state = Injured
...
# State Properties
PERPETRATOR.Active.transmissibility = 1.0
PERPETRATOR.Active.condition_to_transmit = VICTIM
...
VICTIM.AtRisk.susceptibility = 1
```

## Plots and Visualization

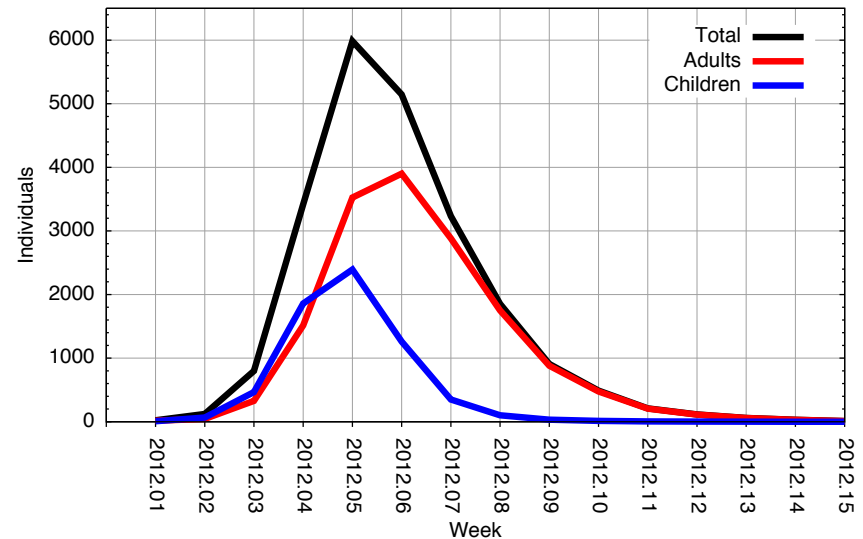
- FRED can produce time plots and movies for *any condition and state*

FRED Simulation of EMS Requests  
Severity Level 5



Blue = Asthma Attacks    Red = Heat Stroke    Black = All EMS Calls

Basic Influenza Model



## Environmental Resources

- You can define any number of *environmental resources*
  - e.g., pharmacies, clinics, vaccine doses, treatment beds, bags of heroin, cooling centers
- You declare the resources properties:
  - timing (days available)
  - location (lat-lon)
  - quantity (count or per capita)
- You can declare that an agent must be able to access a resource to enter a given state
  - how far can the agent travel
  - how many resources are needed?
  - does the agent return the resources when entering another state?

```
resources = BAGS  
BAGS.import_file = delivery_schedule.txt
```

```
HEROIN.Use.resource = BAGS  
HEROIN.Use.resource_required = 1  
HEROIN.Use.resource_returned = 0  
HEROIN.Use.resource_distance = 10.0
```



## Near-term Plans

- FRED Web interface
  - wizards to help you define new conditions
  - setup experiments
  - run FRED experiments
  - create reports (csv files), plots and movies
  - save your work
- FRED community
  - shareable condition definitions
  - shareable models on the web site
  - wiki
- FRED training and documentation
  - Using FRED Web
  - Plotting and Visualization Tools
  - Library of simple models
  - Calibration Methods
  - Sensitivity and Uncertainty Analysis
  - Book?

## Condition Properties

```
# State definitions
COND.states = state_1 state_2 ... state_n
COND.maternity_state = none | <state_name>
COND.fatal_state = none | <state_name>

# Transmission
COND.transmission_mode = none | proximity | environmental |
                          respiratory | sexual | vector
COND.transmissibility = 0.0
COND.exposed_state = none | <state_name>
COND.import_file = none

# Transitions
COND.transition[state_i][state_j] = 0.0
COND.transition[state_i][state_j].coeff = 0.0
COND.transition[state_i][state_j].RISK_FACTOR = 0.0
COND.transition[state_i][state_j].RISK_FACTOR.coeff = 0.0
...
# Reporting
COND.enable_health_records = 0
```

## State Properties (1)

```
# Duration
COND.STATE.duration_distribution = normal | lognormal |
                                exponential | uniform

COND.STATE.duration_mean = -1.0
COND.STATE.duration_stdev = 0.0
COND.STATE.duration_median = -1.0
COND.STATE.duration_dispersion = 1.0
COND.STATE.duration_mean = -1.0
COND.STATE.duration_lower_bound = 0.0
COND.STATE.duration_upper_bound = 99999.0
COND.STATE.duration_offset = 0.0

# Transition
COND.STATE.is_dormant = 0
COND.STATE.default_next_state = <current state>

# Resources
COND.STATE.resource = none
COND.STATE.resources_required = 0
COND.STATE.resource_distance = -1

# Visualization
COND.STATE.visualize = 0
```

## State Properties (2)

### # Health side effects

COND.STATE.susceptibility = 0.0

COND.STATE.transmissibility = 0.0

COND.STATE.condition\_to\_transmit = COND2

COND.STATE.symptoms = none | mild | moderate | severe

### # Contact side effects

COND.STATE.probability\_of\_household\_confinement = 0.0

COND.STATE.decide\_household\_confinement\_daily = 0

COND.STATE.probability\_absent\_from\_household = 0.0

COND.STATE.probability\_absent\_from\_neighborhood = 0.0

COND.STATE.probability\_absent\_from\_work = 0.0

COND.STATE.probability\_absent\_from\_school = 0.0

COND.STATE.probability\_of\_isolation = 0.0

COND.STATE.probability\_of\_hospitalization = 0.0

COND.STATE.probability\_of\_visiting\_outpatient\_clinic = 0.0

### # Side effects on other Conditions

COND.STATE.multiply\_susceptibility\_to\_COND2\_by = 1.0

COND.STATE.multiply\_transmissibility\_of\_COND2\_by = 1.0

COND.STATE.multiply\_symptoms\_of\_COND2\_by = 1.0

COND.STATE.change\_state\_from\_COND2.STATE2\_to = STATE3

## Summary

- FRED is a unique tool for social science modeling
- Agents in FRED can have any number of conditions and can change states based on a variety of interactions with other agents and with the environment
- FRED requires no computer programming skills to define complex agent based models
- FRED requires a systems thinking approach to identify conditions of interest, their states, and the rules for changing states

*Enjoy!*

## FRED Resources

- FRED Web:

<http://fred.publichealth.pitt.edu>

- FRED Github:

<https://github.com/PublicHealthDynamicsLab/FRED>

- FRED Team:

John Grefenstette

David Galloway

Mary Krauland

Bob Frankeny

Mike Lann

David Sinclair

