



Lesson 6

How to solve mathematical models

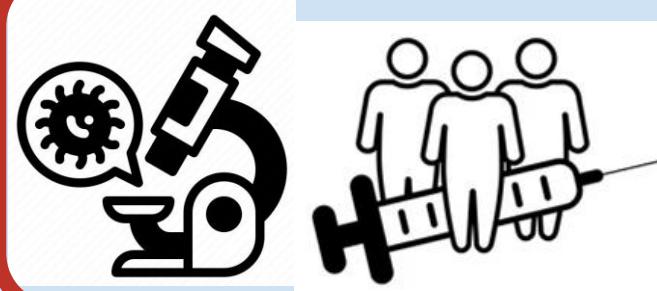
Cameline Orlando

Learning outcomes

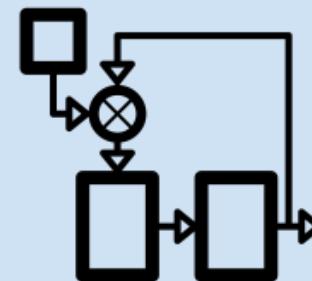
By the end of this lesson, you will be able to:

- **Construct** the basic SIS diagram; define compartments (S, I, S), transitions, and rates and build the differential equations step by step.
- **Define** equilibrium (disease-free and endemic).
- **Know how to calculate R₀, and how to define the threshold**

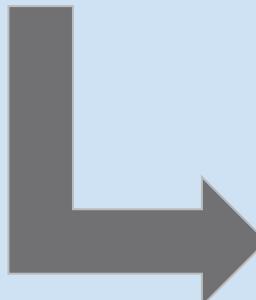
Recap: Modeling = a process where people from different worlds meet on a common ground



biologist/epidemiologist/entomologists:
*translate biological and population-level
observations into flow diagrams*

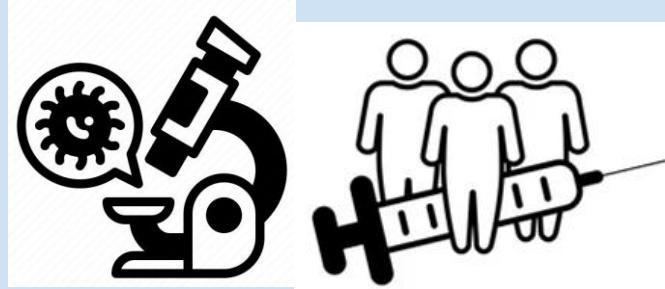


mathematician:
translates flow diagrams into equations

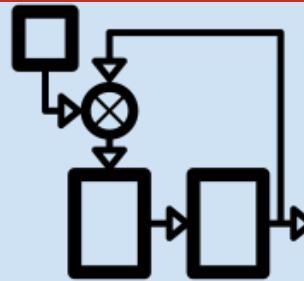
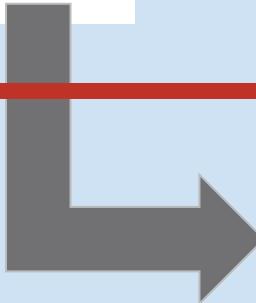


software engineer/computer science:
translates equations into computer code

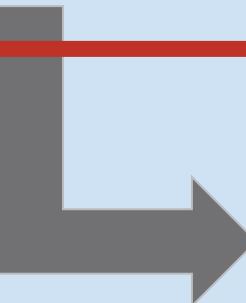
Modeling = a process where people from different worlds meet on a common ground



biologist/epidemiologist/entomologists:
translate biological and population-level observations into flow diagrams

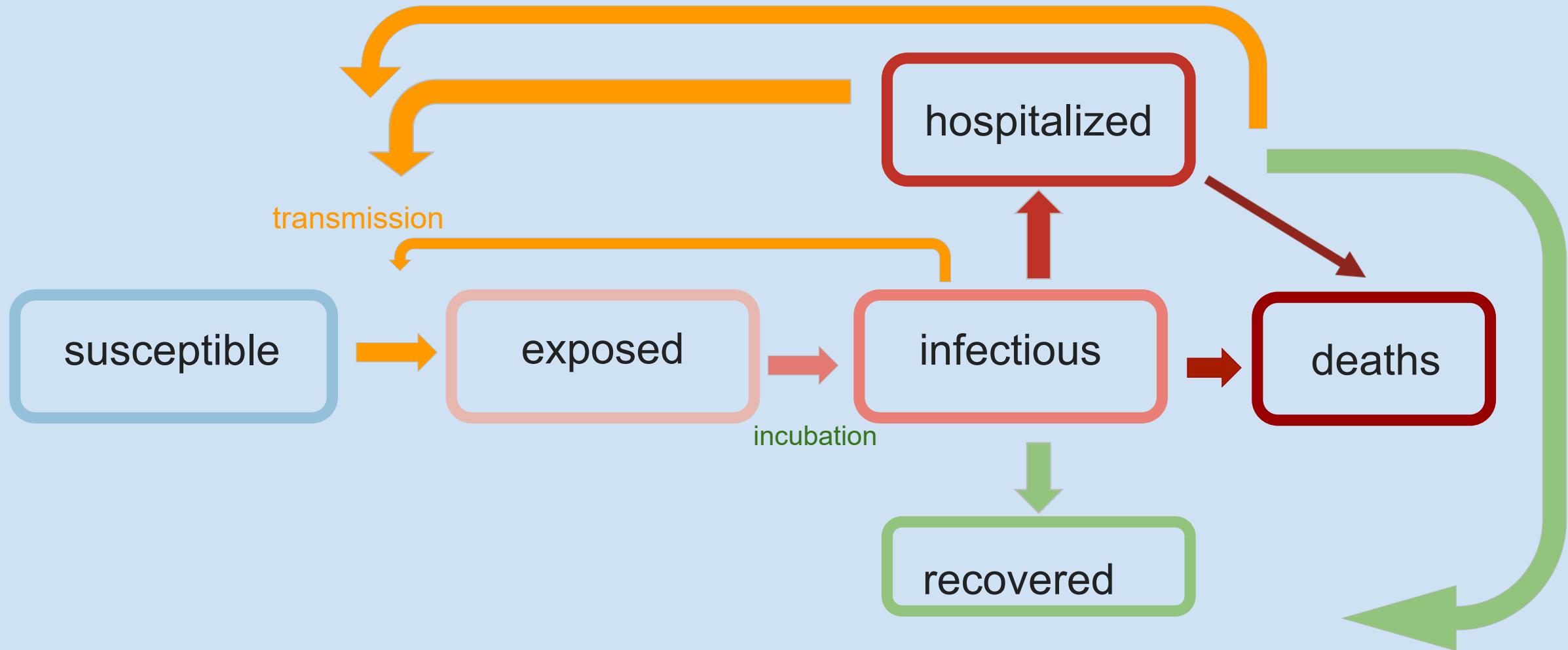


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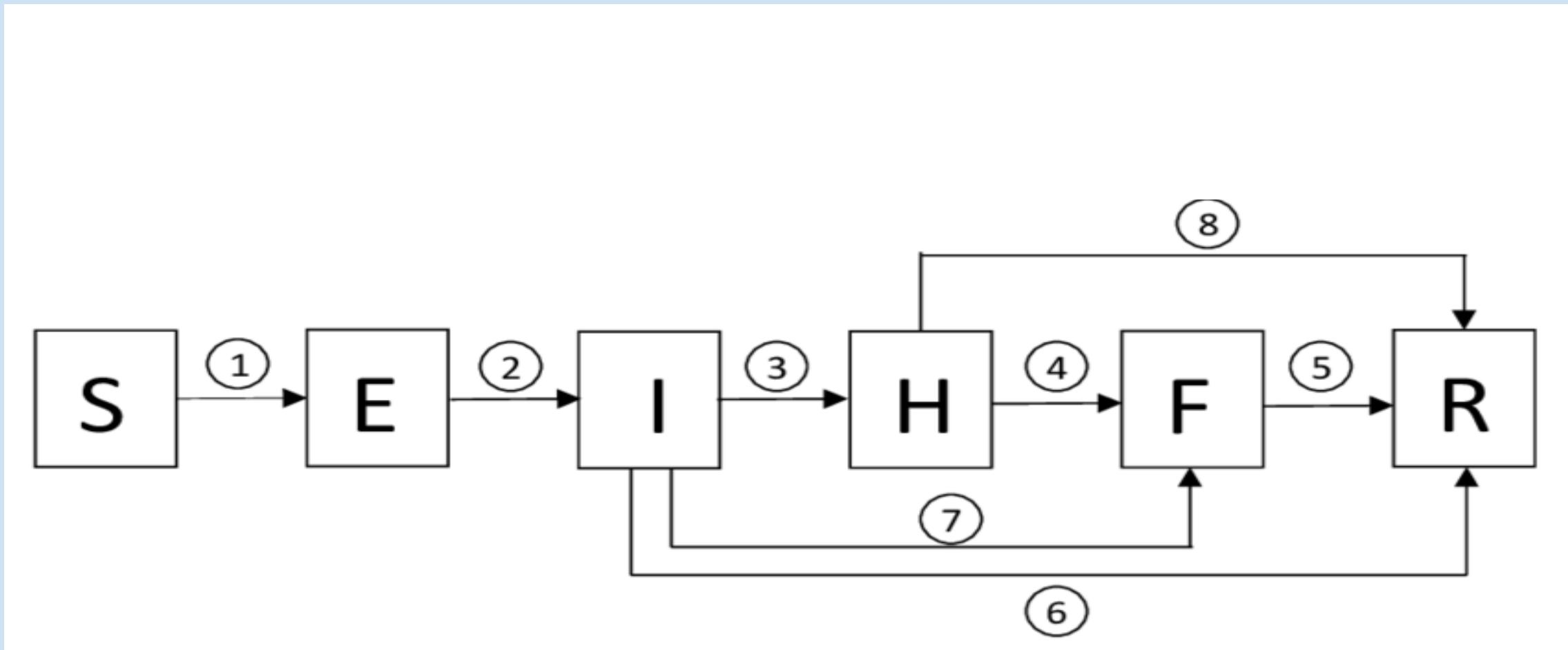


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Recap: Ebola paper



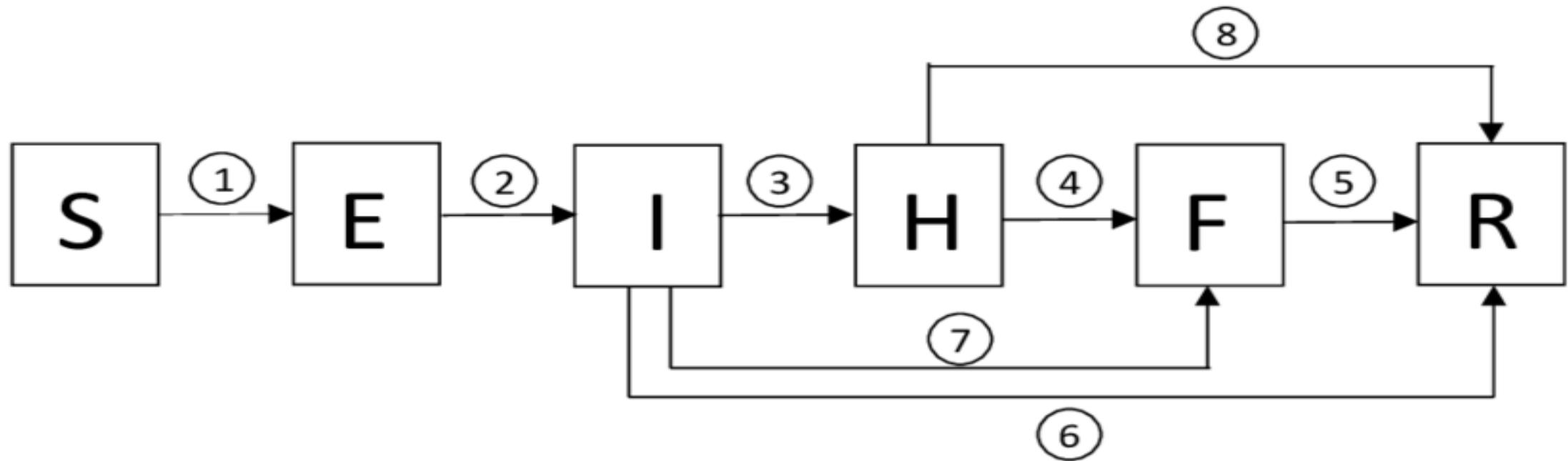
Recap: Ebola flow diagram



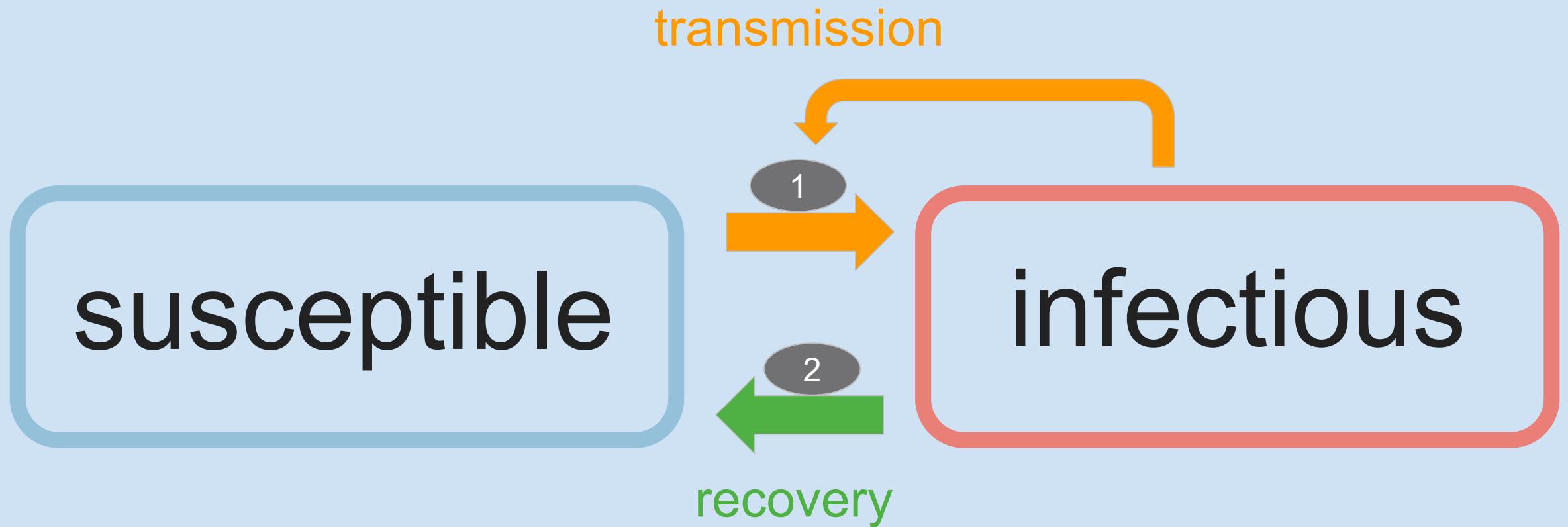
Recap: Ebola flow diagram

$$\begin{array}{ll} 1 & (\beta_I SI + \beta_H SH + \beta_F SF)/N \\ 2 & \alpha E \\ 3 & \gamma_I \theta_1 I \\ 4 & \gamma_H \delta_2 H \end{array}$$

$$\begin{array}{ll} 5 & \gamma_F F \\ 6 & \gamma_I (1-\theta_1)(1-\delta_1)I \\ 7 & \delta_1 (1-\theta_1) \gamma_I I \\ 8 & \gamma_H (1-\delta_2)H \end{array}$$



Recap: Flow diagrams for disease transmission



SIS model

Discussion



What do we need to know
to estimate how many
people are in the S
compartment on day 0?

transmission



In our class of 35
Zenabu is infectious
on day 0c

Discussion



What do we need to know
to estimate how many
people are in the S
compartment on day 0?

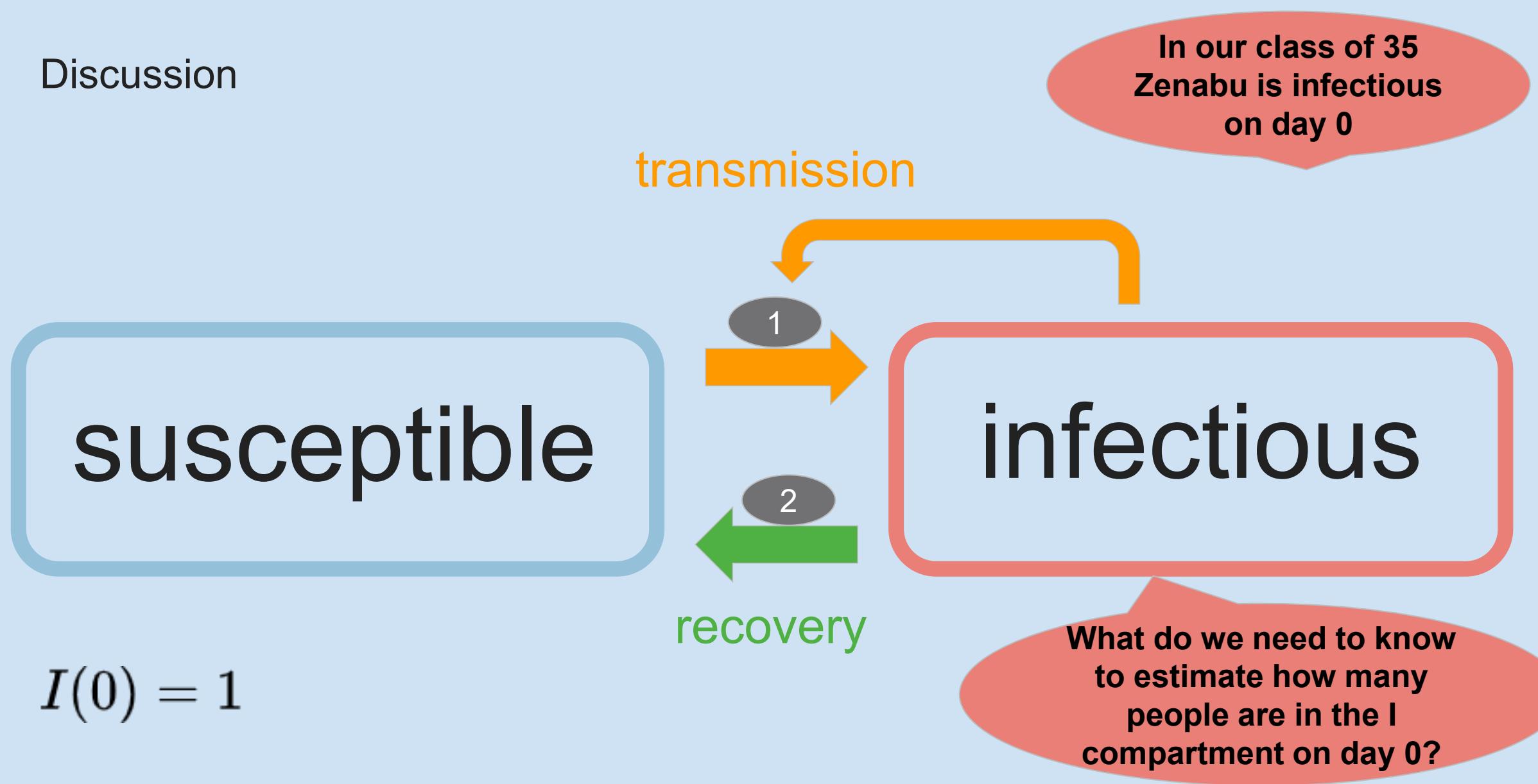
transmission



In our class of 35
Zenabu is infectious
on day 0

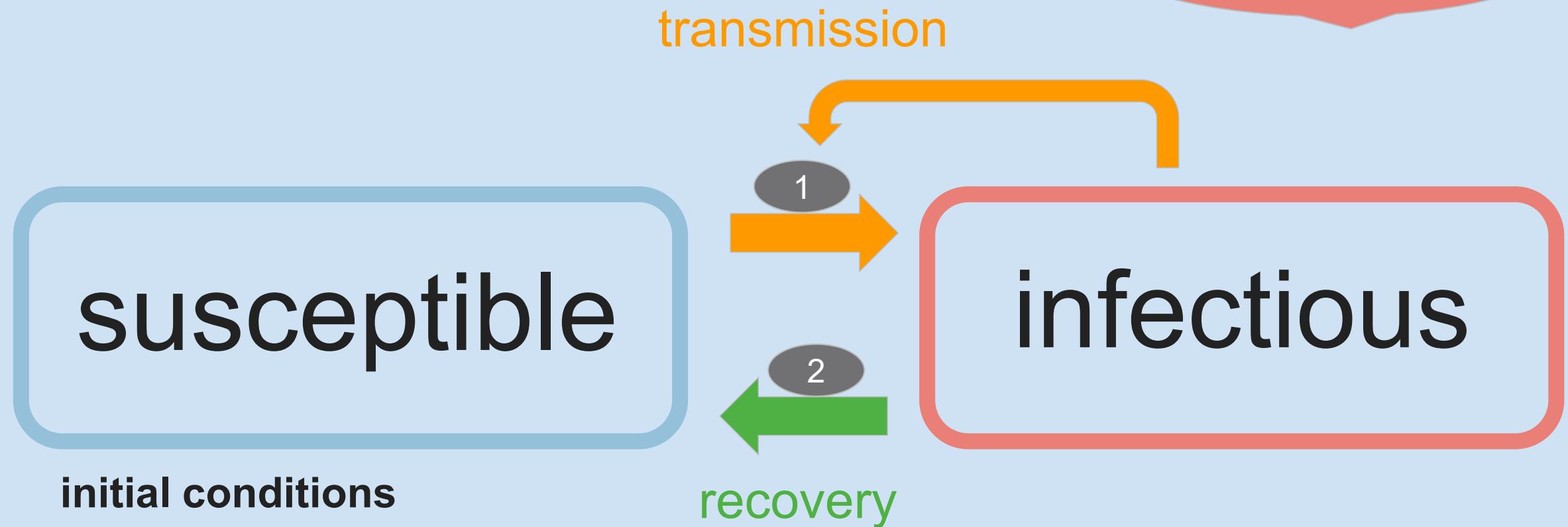
$$S(0) = 34$$

Discussion



Discussion

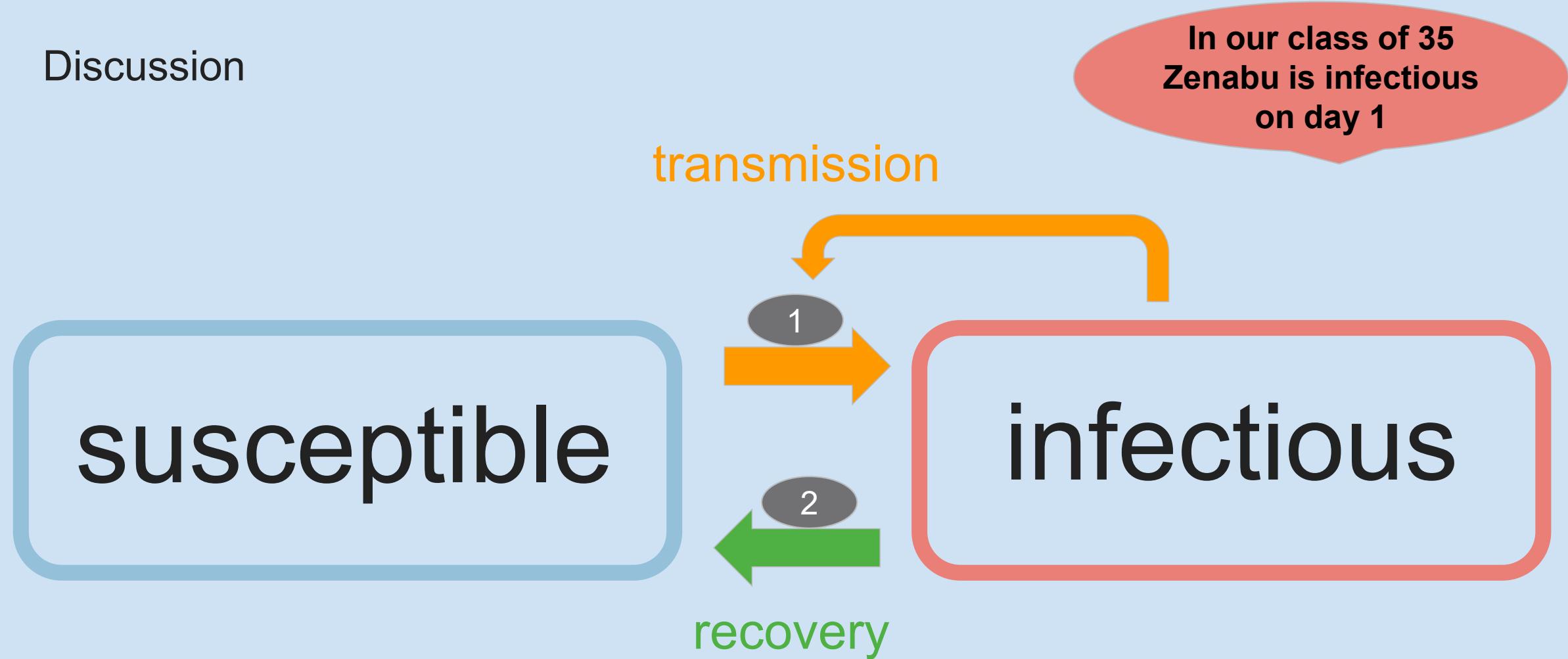
In our class of 35
Zenabu is infectious
on day 0



initial conditions

$$S(0) = 34, \\ I(0) = 1$$

Discussion



In our class of 35
Zenabu is infectious
on day 1

infectious

What do we need to know to estimate how many people are in the S and I compartments on day 1?

Discussion summary

In our class of 35
Zenabu is infectious
on day 1

1. What do we need to know to estimate how many people are infectious on day 1?

How many new infections occur and how many recover.

1. Who can become newly infected? *Only those in the susceptible group.*
2. How does the number of infectious students from the previous day affect the risk for susceptibles today?

The more proportion of the infectious population there are, the higher the risk.

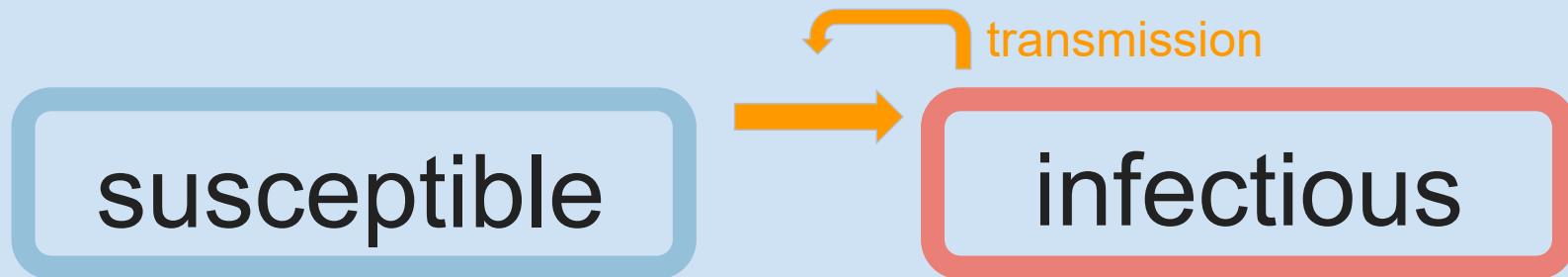
1. If a susceptible meets an infectious, do they always get infected?

(chance of transmission per contact).

1. And what about the students already infectious — do they all stay infectious tomorrow, or do some recover?

(chance of recovery each day)

Flow diagrams as differential equations



outflow **from** susceptible

$$S(t+1) = S(t) - \beta \frac{I(t)}{N} S(t)$$

inflow **into** infectious

$$I(t+1) = I(t) + \beta \frac{I(t)}{N} S(t)$$

assumption: all individuals are well mixed

: the population is closed,
 $S(t) + I(t) = N$

Flow diagrams as differential equations

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: the population is $S(t) + I(t) = N$

closed,

outflow from susceptible

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Transmission rate β

The rate at which a susceptible individual becomes infected per unit time, given contact with infectious individuals.

Flow diagrams as differential equations

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closed,

outflow **from** susceptible

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inflow **into** infectious

$$I(t+1) = I(t) + \beta \frac{I(t)}{N} S(t)$$

Proportion infectious $\frac{I(t)}{N}$

The fraction of the population that is infectious
at time t .

captures the “risk of exposure” for any
susceptible individual.

Flow diagrams as differential equations

assumption: all individuals are well mixed

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outflow **from** susceptible

$$S(t+1) = S(t) - \beta \frac{I(t)}{N} S(t)$$

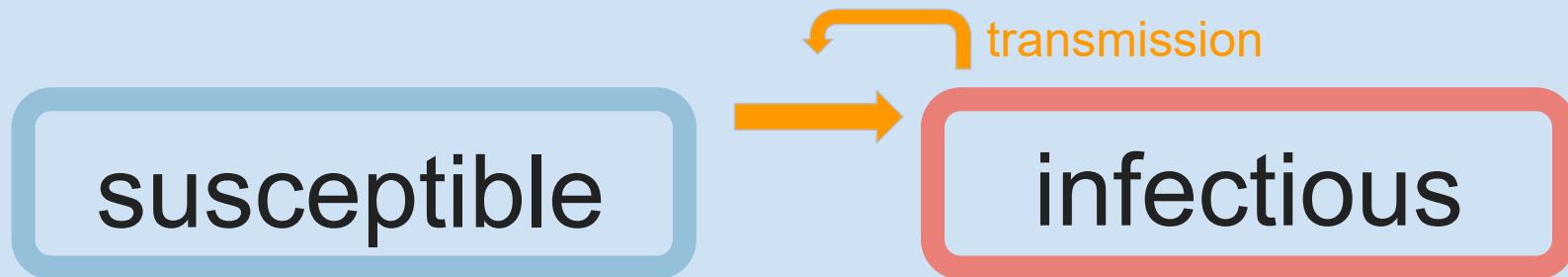
inflow **into** infectious

$$I(t+1) = I(t) + \beta \frac{I(t)}{N} S(t)$$

Susceptible population $S(t)$

How many susceptible people are present at time t .

Flow diagrams as differential equations



outflow **from** susceptible

$$S(t+1) = S(t) - \underbrace{\beta \frac{I(t)}{N} S(t)}_{\text{outflow from susceptible}}$$

inflow **into** infectious

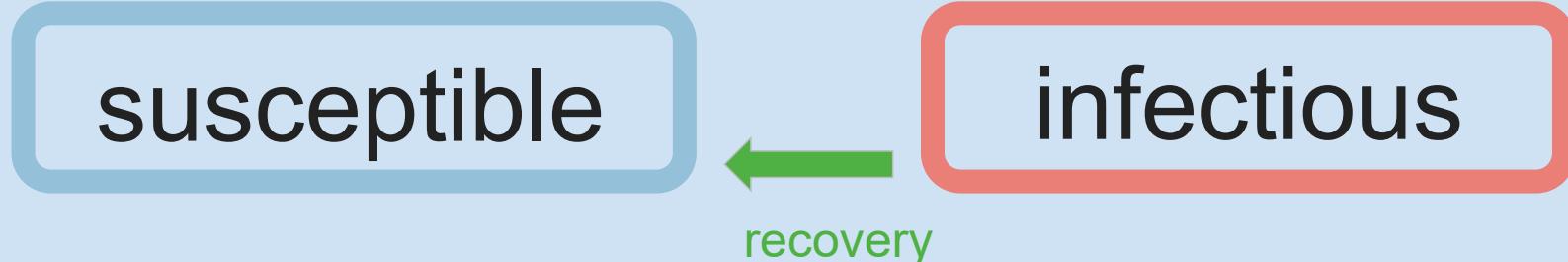
$$I(t+1) = I(t) + \beta \frac{I(t)}{N} S(t)$$

assumption: all individuals are well mixed
: the population is closed,

$$S(t) + I(t) = N$$



Flow diagrams as differential equations



inflow **into** susceptible

$$S(t + 1) = S(t) + \gamma I(t)$$

infectious

outflow **from** infectious

$$I(t + 1) = I(t) - \gamma I(t)$$

assumption: all individuals are well mixed

: the population is closed,
 $S(t) + I(t) = N$

Flow diagrams as differential equations

assumption: all individuals are well mixed

: the population is $S(t) + I(t) = N$

closed,

inflow **into** susceptible

outflow **from** infectious

$$S(t+1) = S(t) + \gamma I(t)$$

$$I(t+1) = I(t) - \gamma I(t)$$

Recovery parameter γ

The rate at which infectious individuals recover per unit time.

Average infectious period: $\frac{1}{\gamma}$

Flow diagrams as differential equations

assumption: all individuals are well mixed

: the population is $S(t) + I(t) = N$

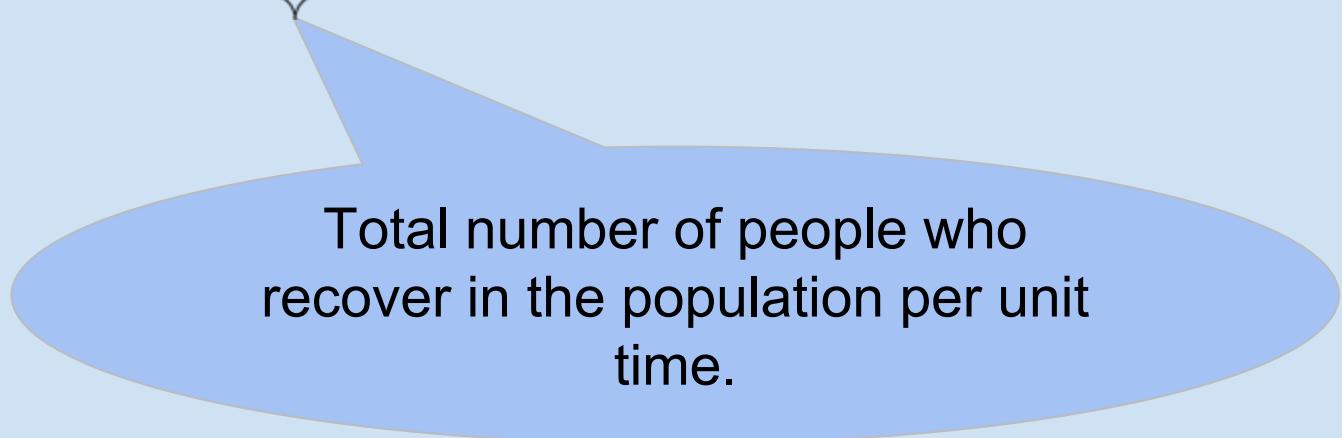
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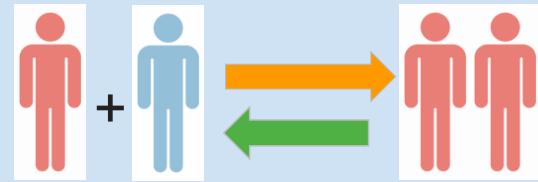
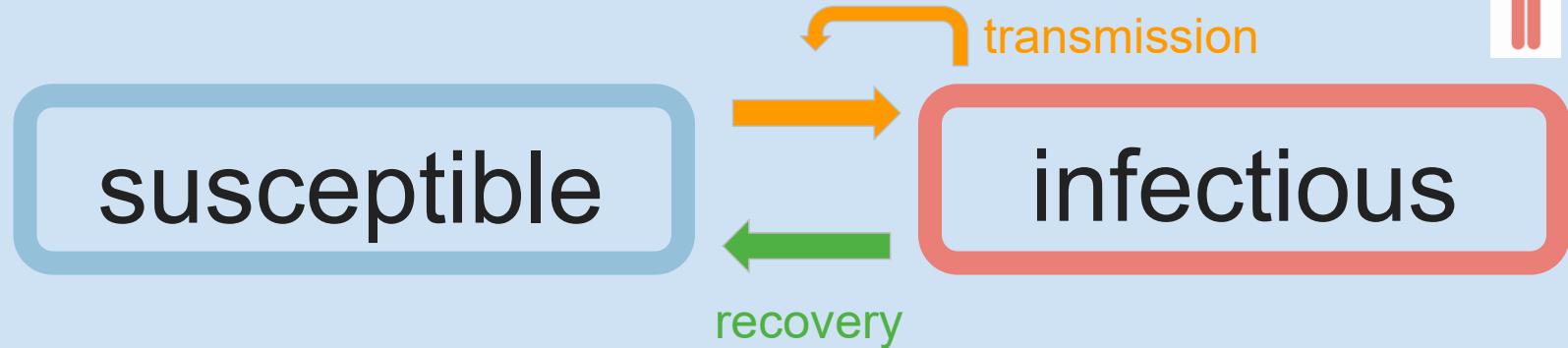
$$S(t+1) = S(t) + \gamma I(t)$$

$$I(t+1) = I(t) - \gamma I(t)$$



Total number of people who recover in the population per unit time.

Flow diagrams as differential equations



$$\frac{dS(t)}{dt} = -\beta \frac{I(t)}{N} S(t) + \gamma I$$

$$S(0) = S_0$$

$$I(0) = I_0$$

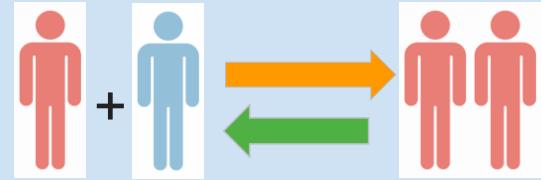
$$\frac{dI(t)}{dt} = \beta \frac{I(t)}{N} S(t) - \gamma I$$

$$S(t) + I(t) = N$$

$$\Delta \rightarrow 0$$

"coupled system"

Flow diagrams as differential equations



$$\frac{dS(t)}{dt} = -\beta \frac{I(t)}{N} S(t) + \gamma I$$

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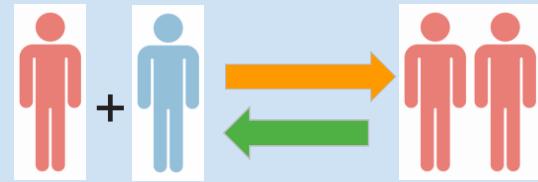
$$S(t) + I(t) = N$$

$$\Delta \rightarrow 0$$

Rate of change of susceptible
individuals over time

"coupled system"

Flow diagrams as differential equations



$$\frac{dS(t)}{dt} = -\beta \frac{I(t)}{N} S(t) + \gamma I$$

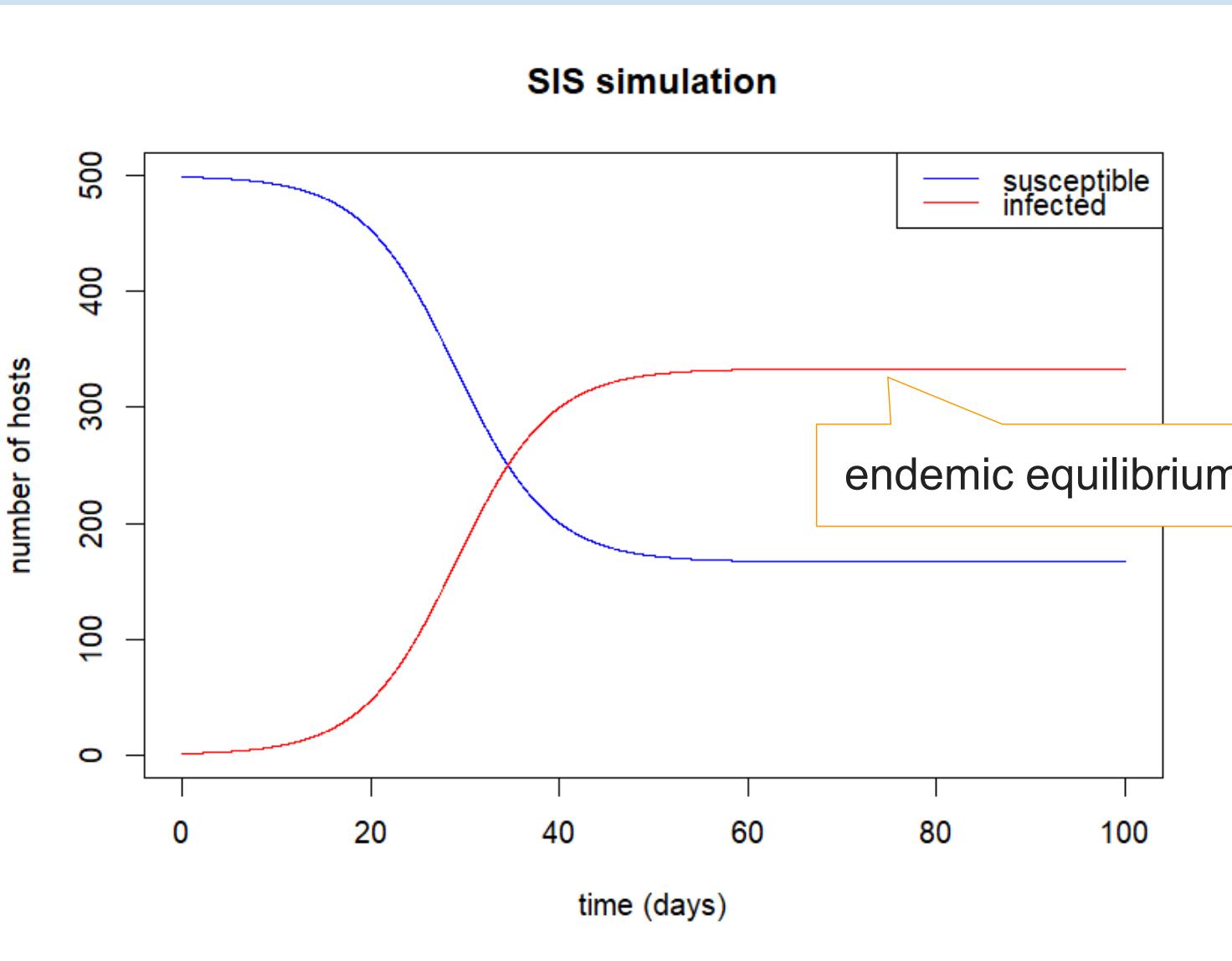
$$\frac{dI(t)}{dt} = \beta \frac{I(t)}{N} S(t) - \gamma I$$

$$S(0) = S_0$$
$$I(0) = I_0$$

$$S(t) + I(t) = N$$
$$\Delta \rightarrow 0$$

Rate of change of infectious individuals
over time

Endemic and disease-free equilibrium for SIS system



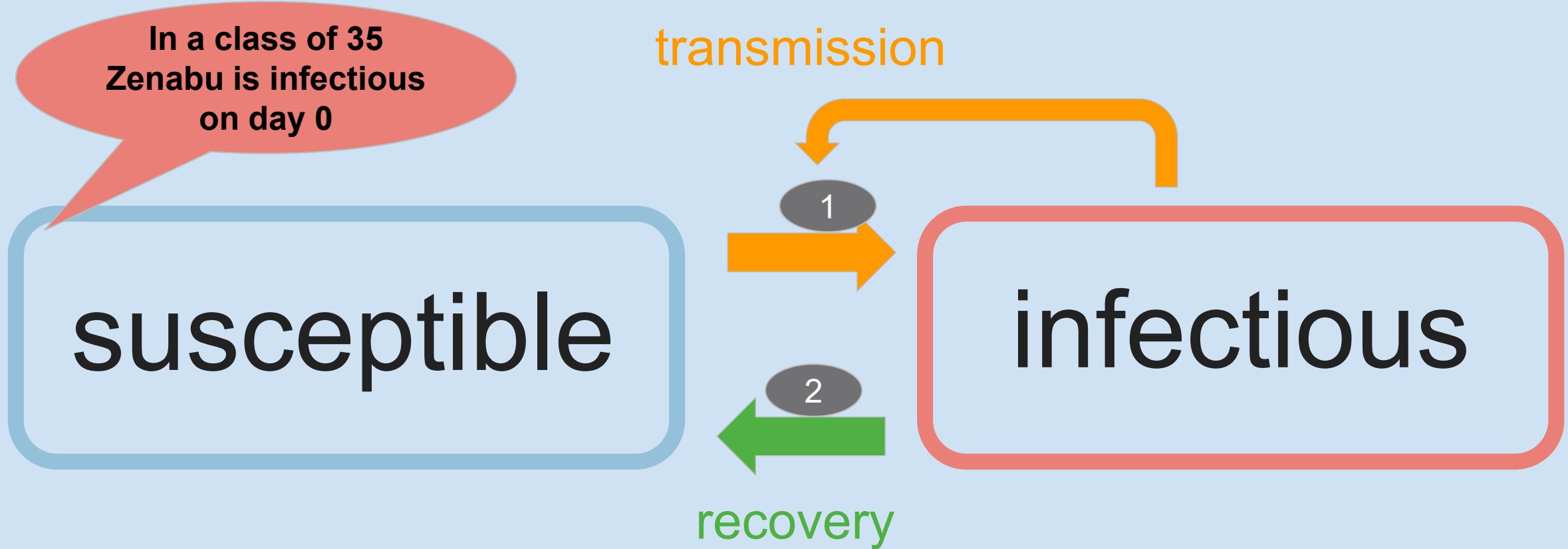
$$0 = -\beta \frac{I(t)}{N} S(t) + \gamma I$$

$$0 = \beta \frac{I(t)}{N} S(t) - \gamma I$$

$$\Leftrightarrow S^*(\infty) = \frac{\gamma}{\beta} N$$

$$I^*(\infty) = N - S^*(\infty) = N \left(1 - \frac{1}{\mathcal{R}_0}\right)$$

Practical: Infection OR No Infection



SIS model

Basic reproduction number

Introducing a single infection into a entirely susceptible population

$$\frac{dI}{dt} = \beta S \frac{I}{N} - \gamma I \quad S \sim N, I(0) = I_0$$

$$\beta - \gamma > 0 \text{ exponential growth} \Leftrightarrow \beta > \gamma \Leftrightarrow \frac{\beta}{\gamma} > 1$$

$$\beta - \gamma < 0 \text{ exponential growth} \Leftrightarrow \beta < \gamma \Leftrightarrow \frac{\beta}{\gamma} < 1$$

“basic reproduction number”

$$\mathcal{R}_0 = \frac{\beta}{\gamma} \quad I(\infty) = N \left(1 - \frac{1}{\mathcal{R}_0} \right)$$

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- Geoffrey Githinji

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