



Swiss TPH



Lesson 6

How to solve mathematical models

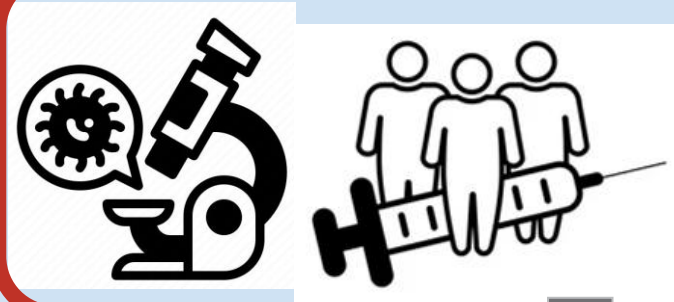
Cameline Orlando

Learning outcomes

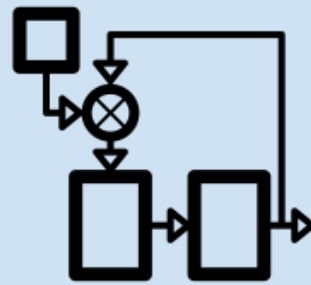
By the end of this lesson, you will be able to:

- **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** R_0 , and **how to define the threshold**

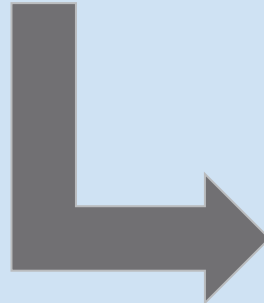
Recap: Modeling = a process where people from different worlds meet on a common ground



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

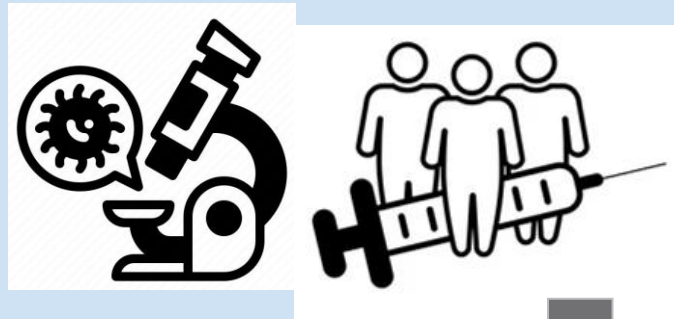


mathematician:
translates flow diagrams into equations

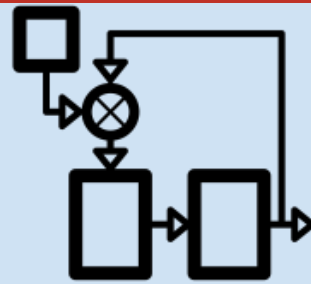


software engineer/computer science:
translates equations into computer code

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

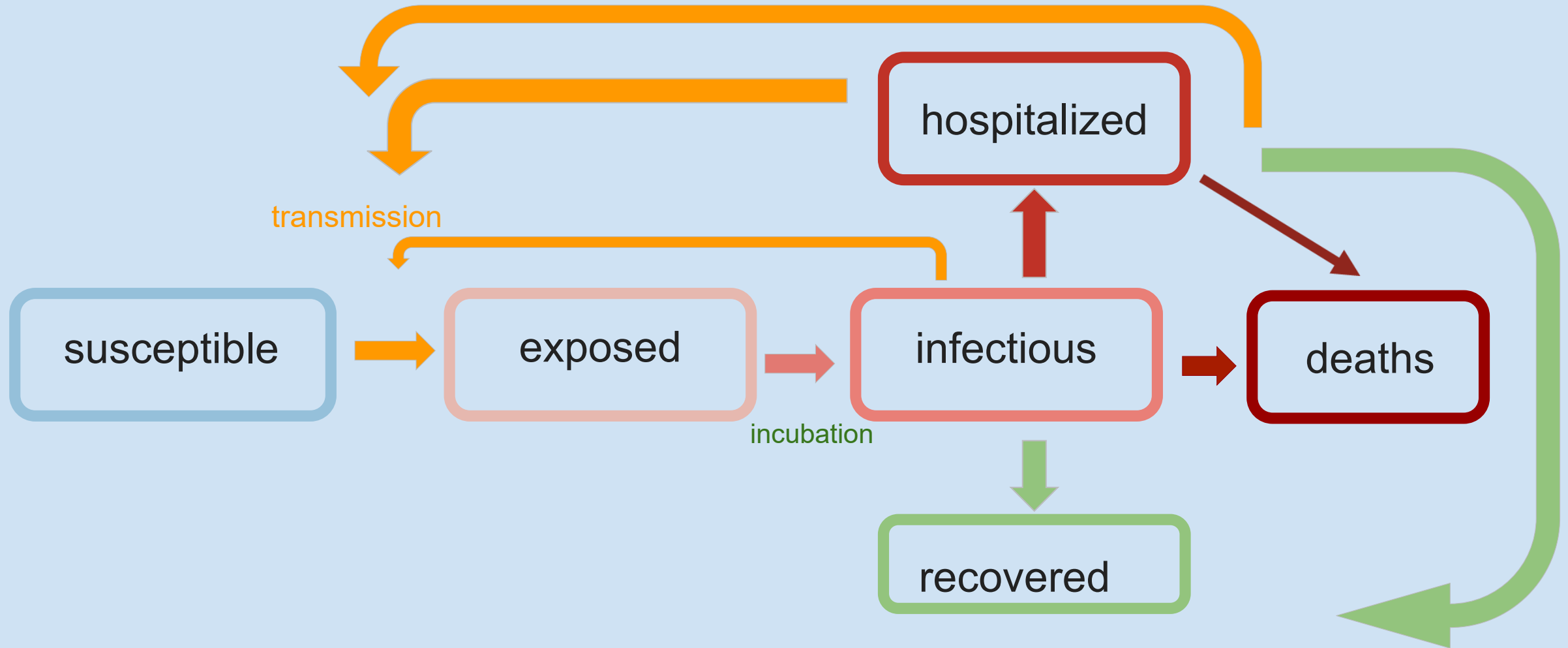


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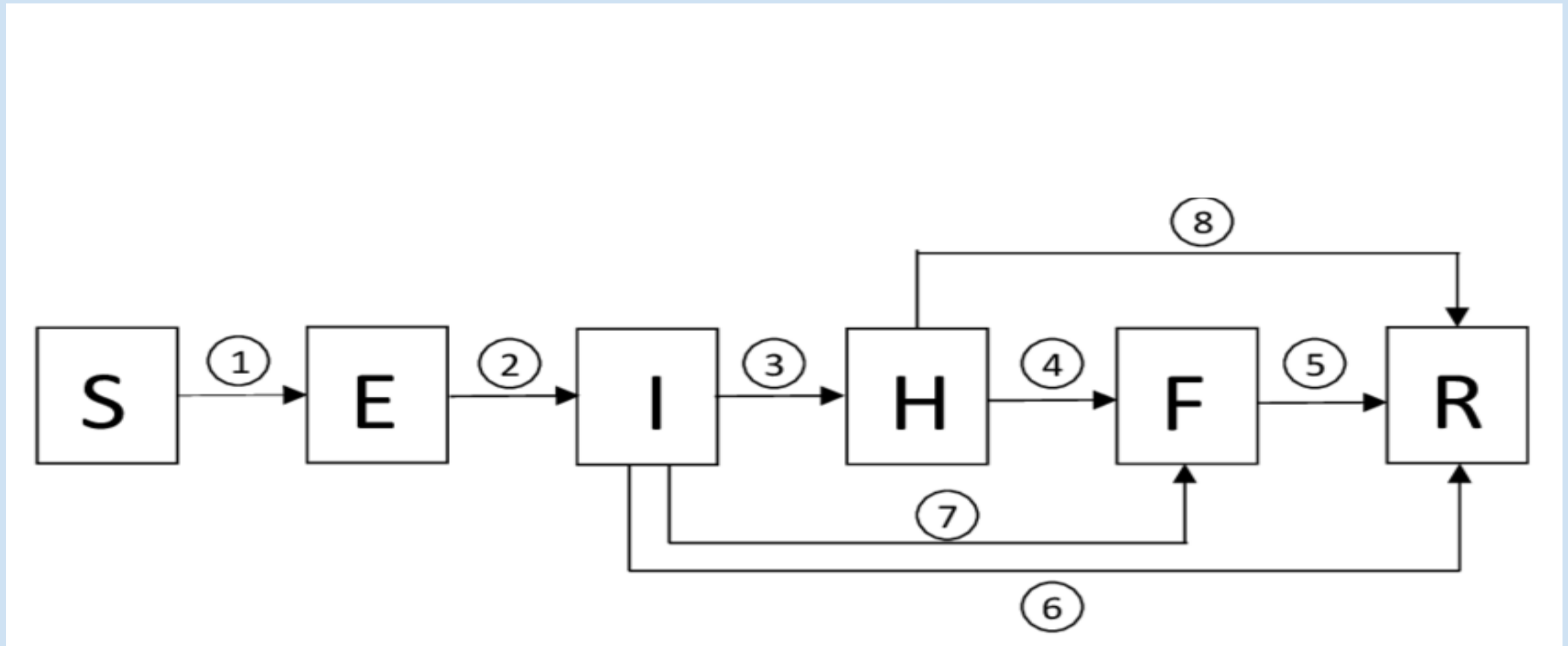


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translates equations into computer code

Recap: Ebola paper



Recap: Ebola flow diagram

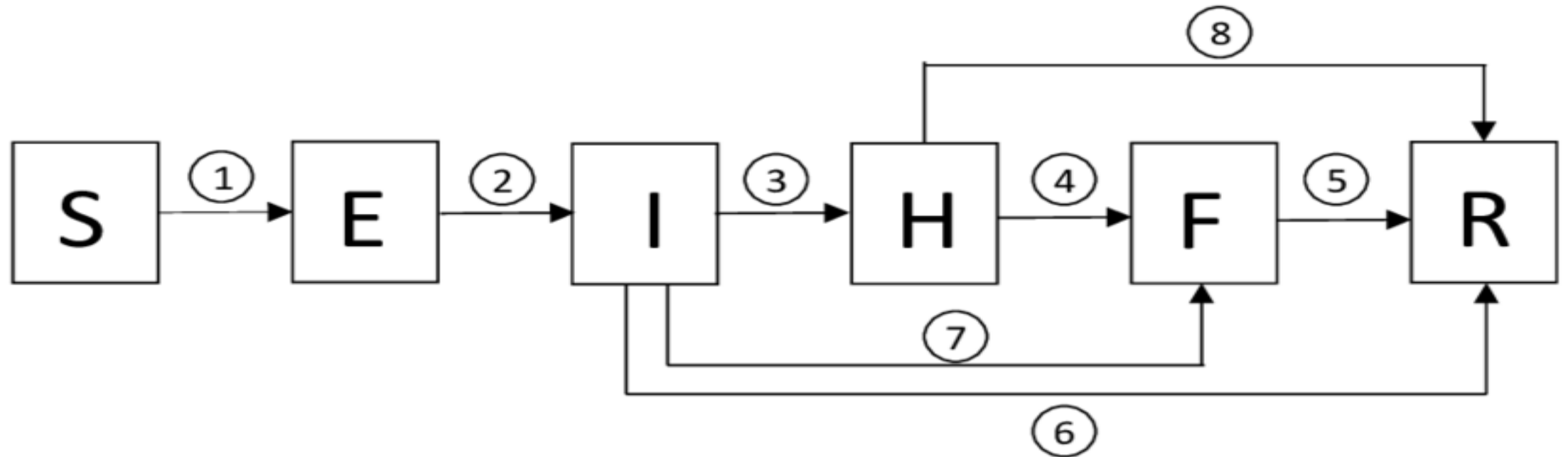


Hernandez-Suarez, C., & Lopez, O. M. (2022).
<https://doi.org/10.48550/arXiv.2208.11509>

Recap: Ebola flow diagram

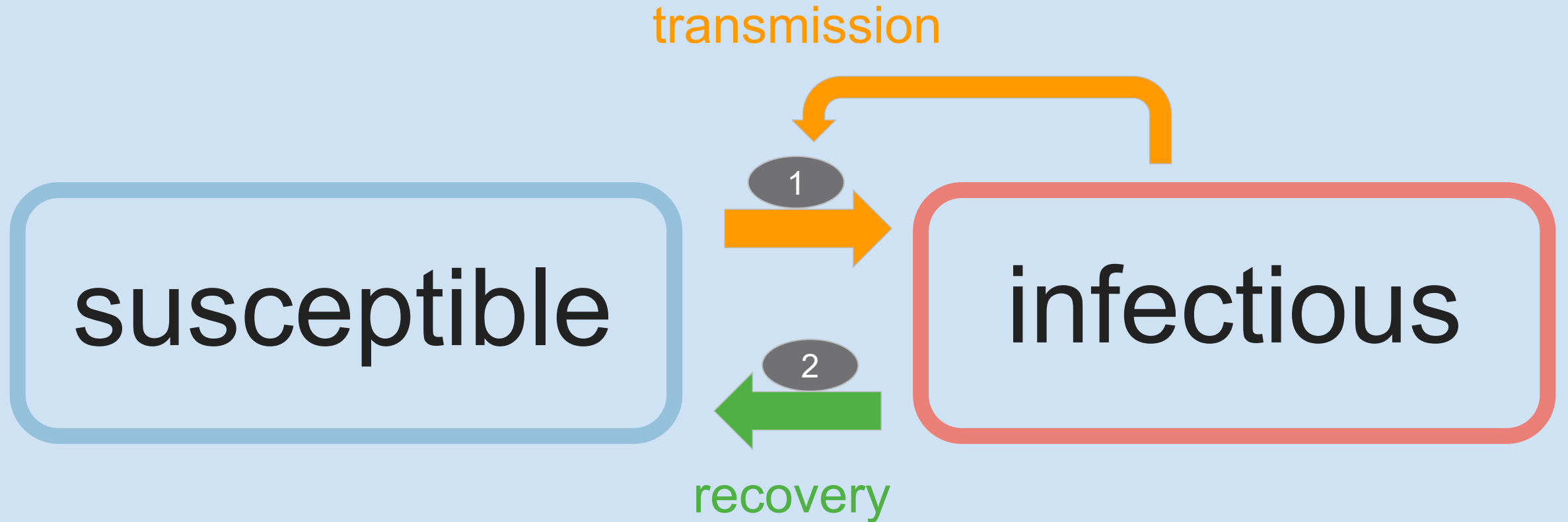
- 1 $(\beta_I SI + \beta_H SH + \beta_F SF)/N$
- 2 αE
- 3 $\gamma_I \theta_1 I$
- 4 $\gamma_H \delta_2 H$

- 5 $\gamma_F F$
- 6 $\gamma_I(1-\theta_1)(1-\delta_1)I$
- 7 $\delta_1(1-\theta_1)\gamma_I I$
- 8 $\gamma_H(1-\delta_2)H$



Hernandez-Suarez, C., & Lopez, O. M. (2022).
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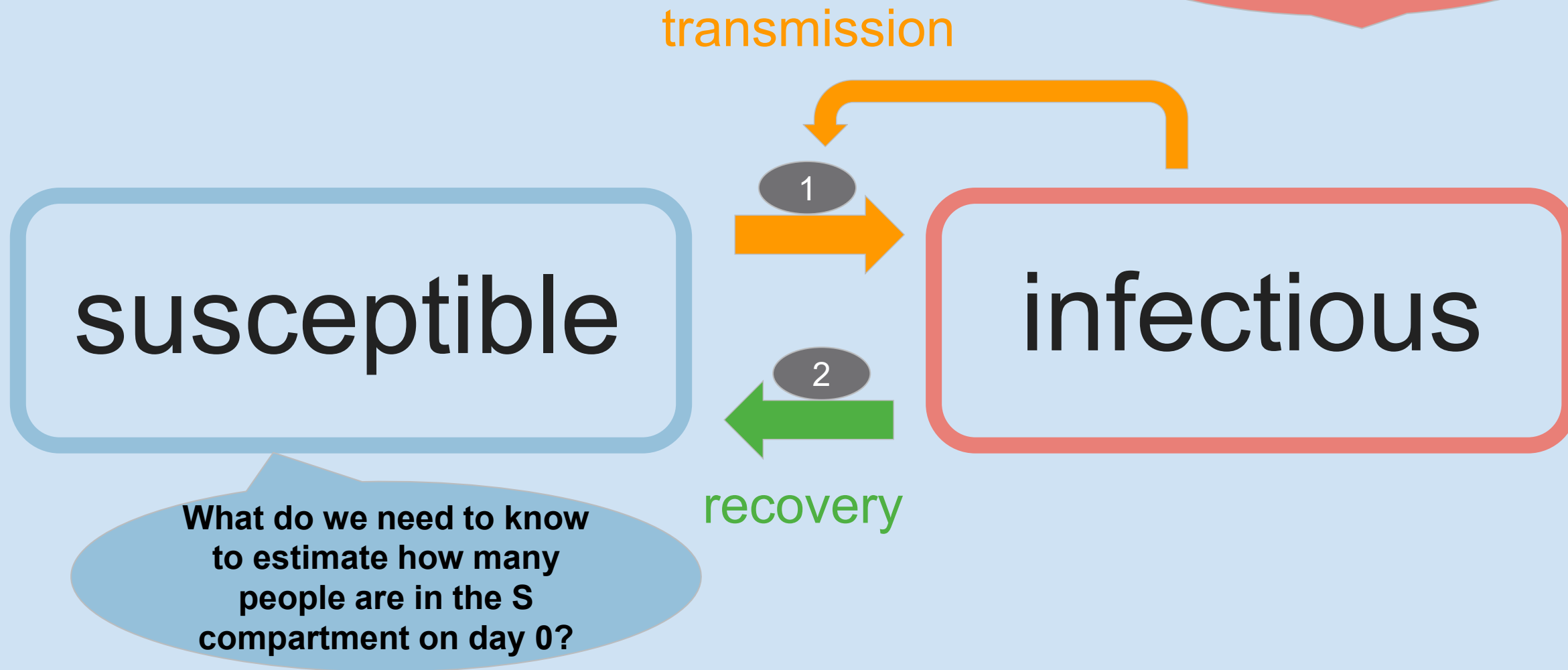
Recap: Flow diagrams for disease transmission



SIS model

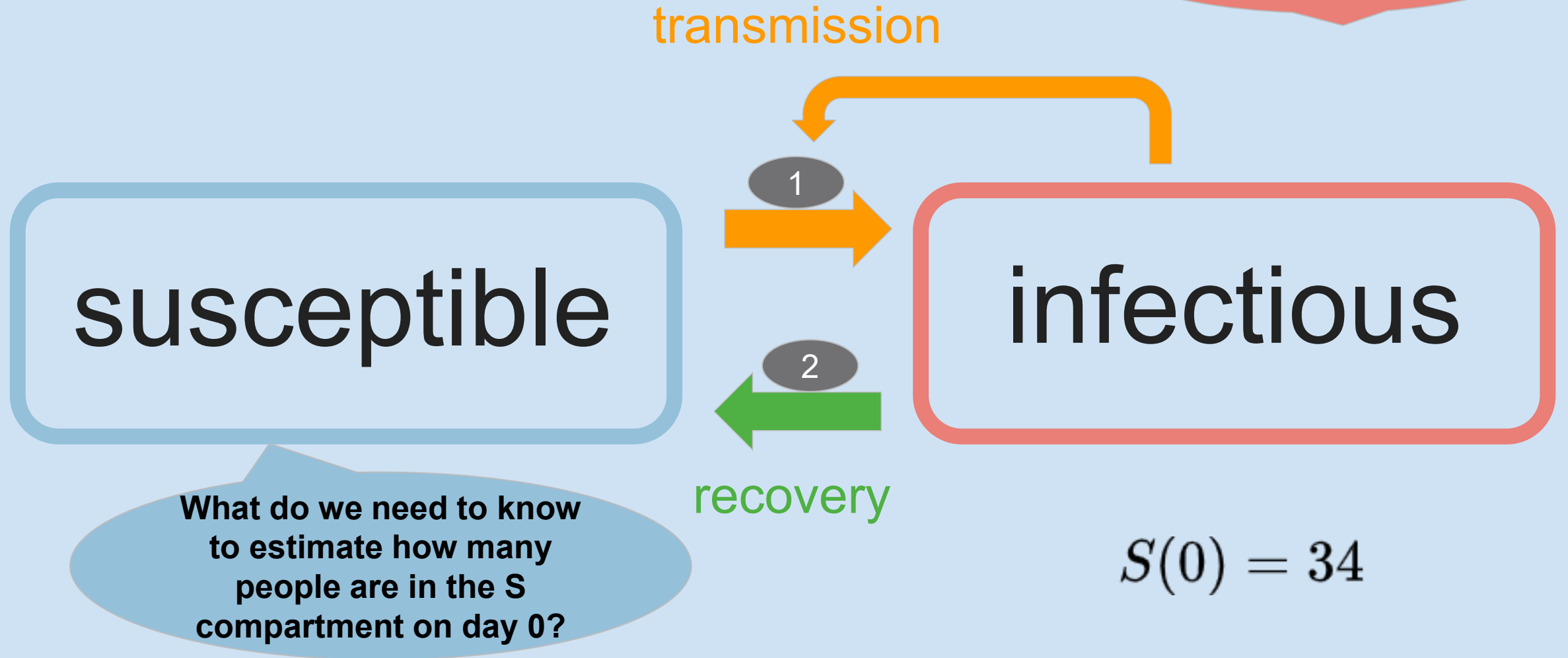
Discussion

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



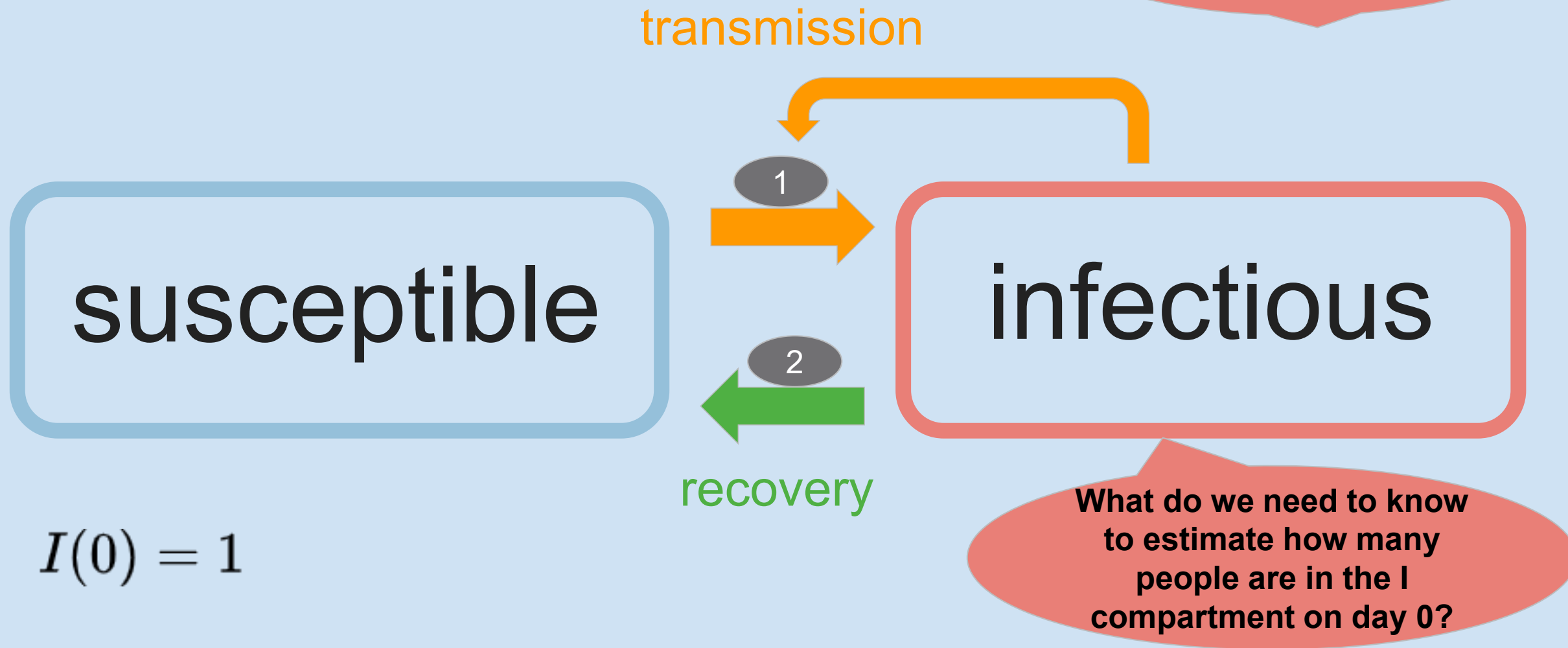
Discussion

In our class of 35
Zenabu is infectious
on day 0



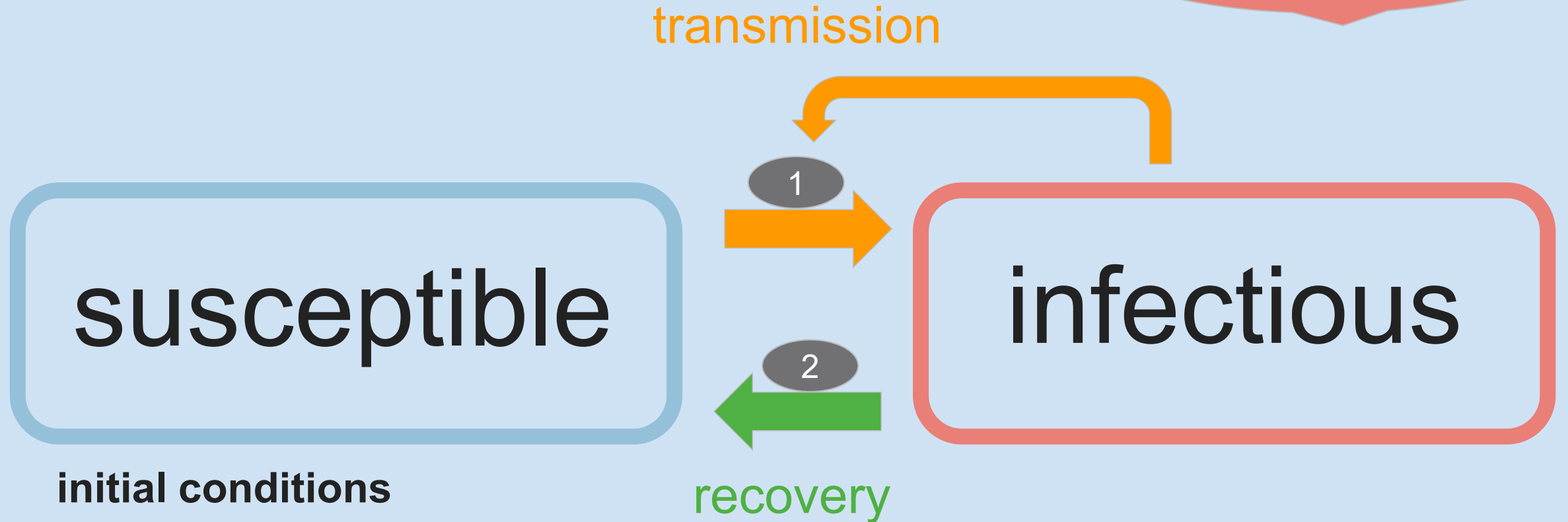
Discussion

In our class of 35
Zenabu is infectious
on day 0



Discussion

In our class of 35
Zenabu is infectious
on day 0

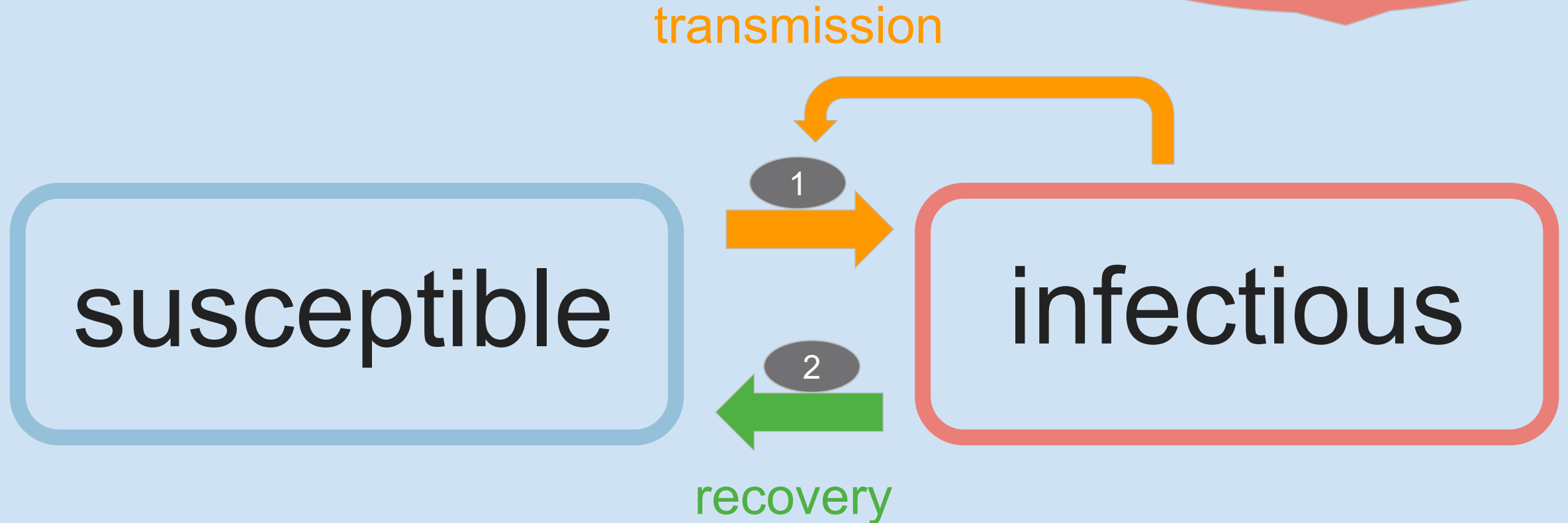


initial conditions

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

Discussion

In our class of 35
Zenabu is infectious
on day 1



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.

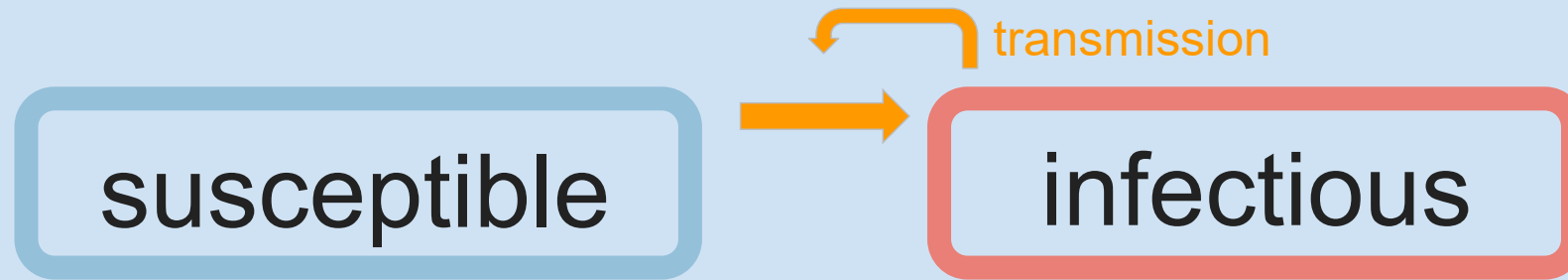
1. If a susceptible meets an infectious, do they always get infected?

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

closed,

Flow diagrams as differential equations

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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

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

inflow **into** infectious

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

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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closed,

outflow **from** susceptible

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

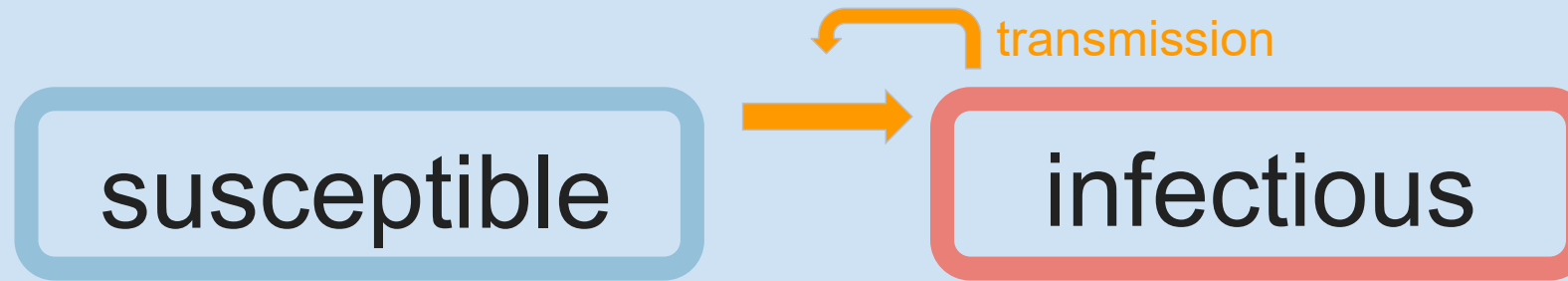
inflow **into** infectious

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

Susceptible population $S(t)$

How many susceptible people are present at time t .

Flow diagrams as differential equations



outflow **from** susceptible

$$S(t + 1) = S(t) - \underbrace{\beta \frac{I(t)}{N} S(t)}_{\text{total new infections per unit time}}$$

inflow **into** infectious

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

assumption: all individuals are well mixed

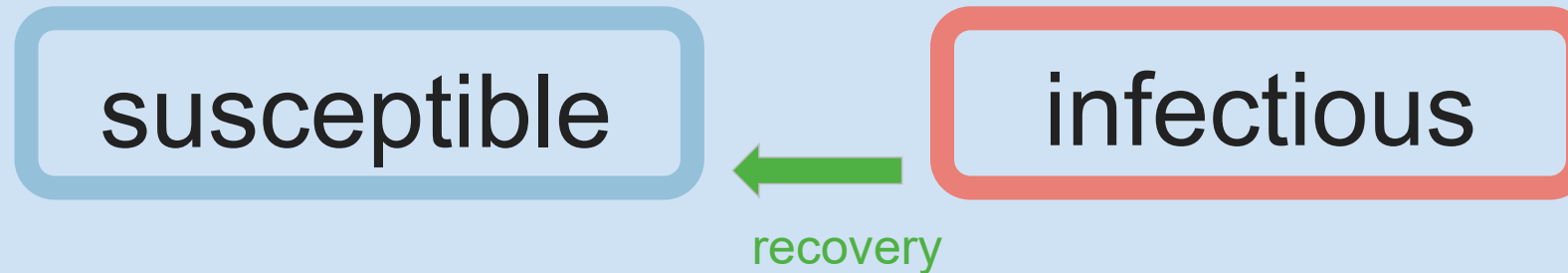
: the

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

population is closed,

total new infections per unit
time

Flow diagrams as differential equations



inflow **into** susceptible

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

outflow **from** infectious

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

assumption: all individuals are well mixed

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

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Flow diagrams as differential equations

assumption: all individuals are well mixed

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

closed,

inflow **into** susceptible

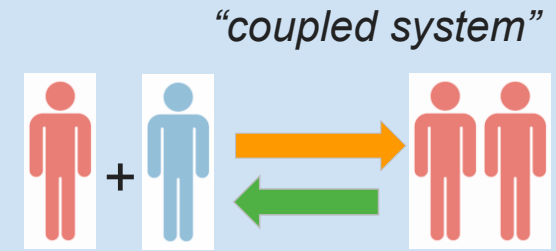
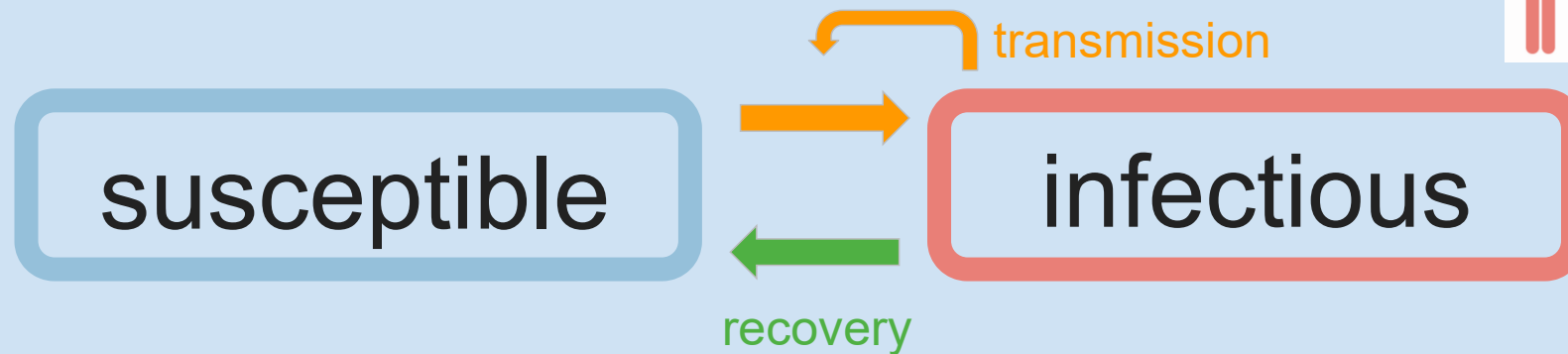
outflow **from** infectious

$$S(t + 1) = S(t) + \underbrace{\gamma I(t)}$$

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

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

Flow diagrams as differential equations



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

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

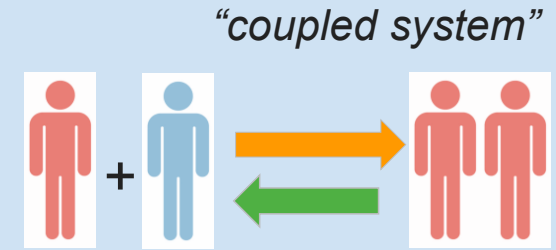
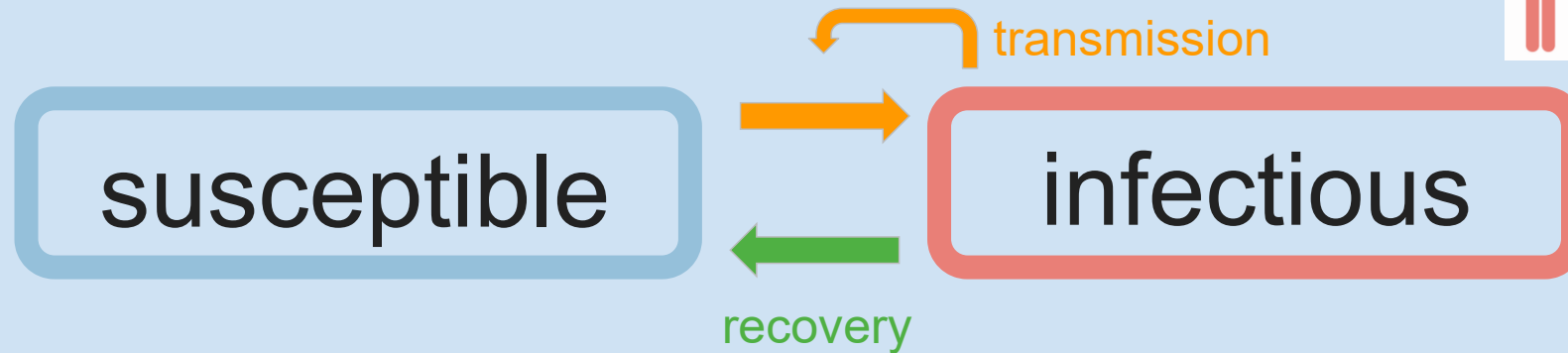
$$S(0) = S_0$$

$$I(0) = I_0$$

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

$$\Delta \rightarrow 0$$

Flow diagrams as differential equations



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

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

$$S(0) = S_0$$

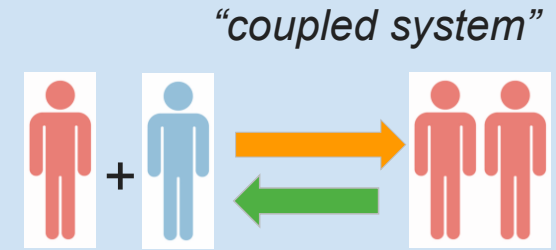
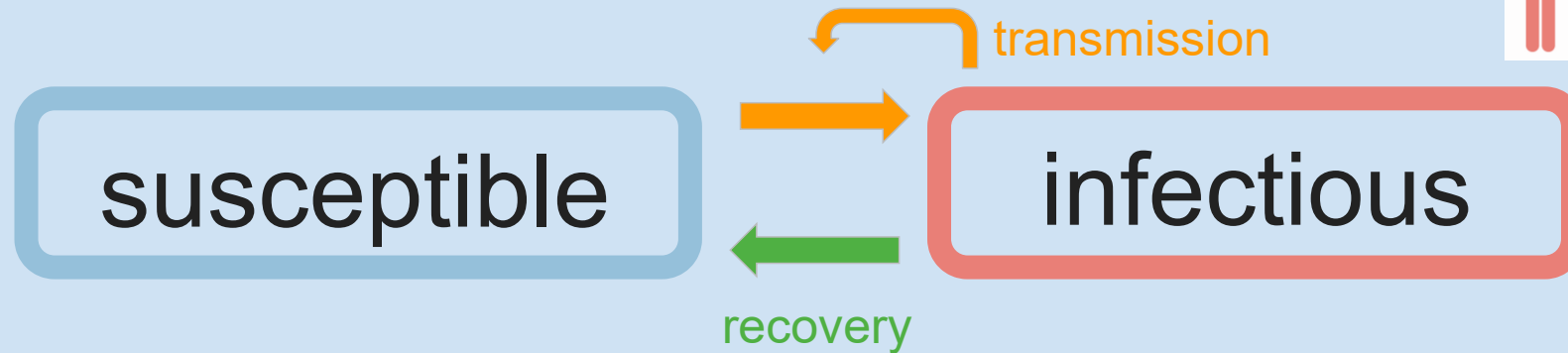
$$I(0) = I_0$$

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

$$\Delta \rightarrow 0$$

Rate of change of susceptible individuals over time

Flow diagrams as differential equations



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

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

$$S(0) = S_0$$

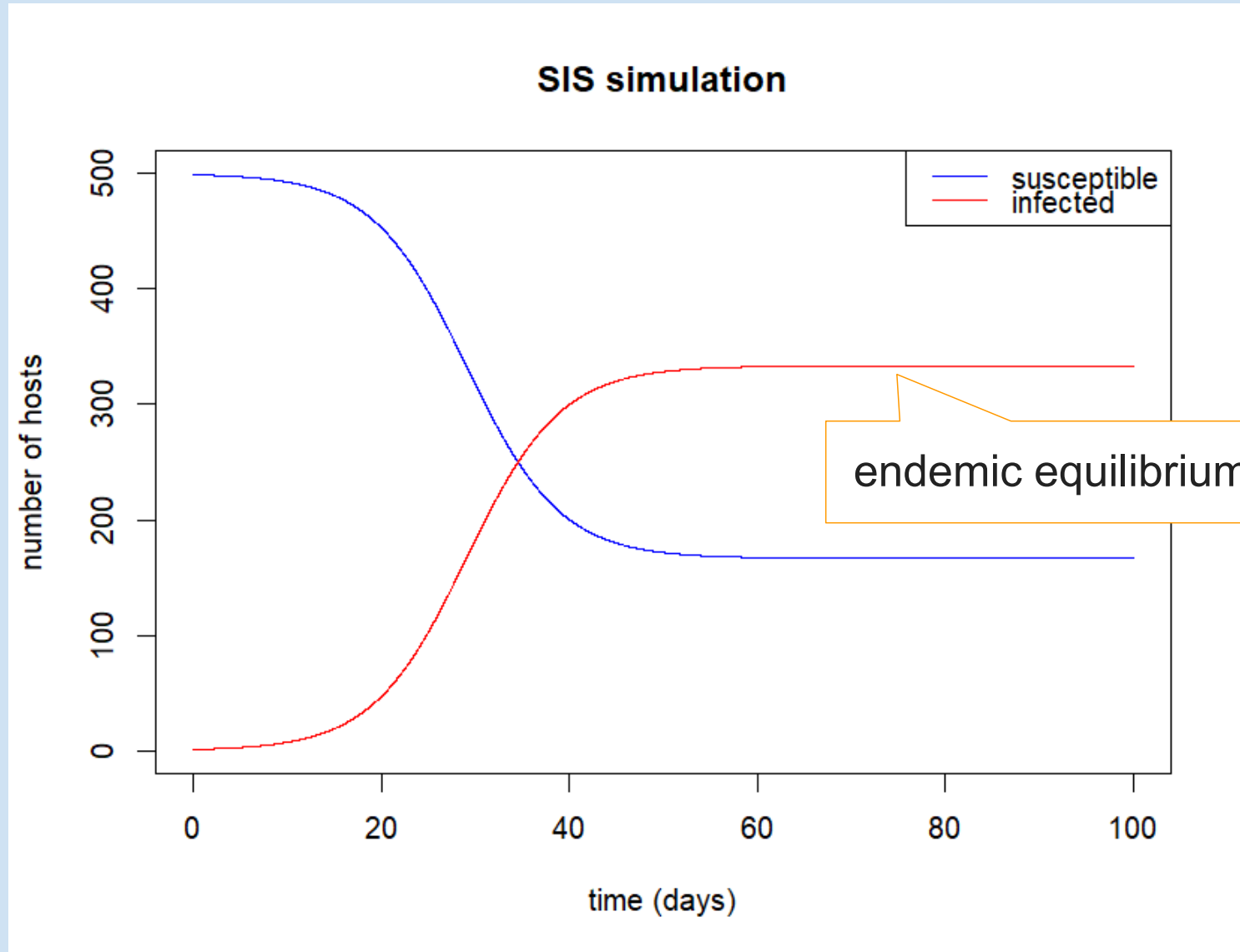
$$I(0) = I_0$$

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

$$\Delta \rightarrow 0$$

Rate of change of infectious individuals
over time

Endemic and disease-free equilibrium for SIS system



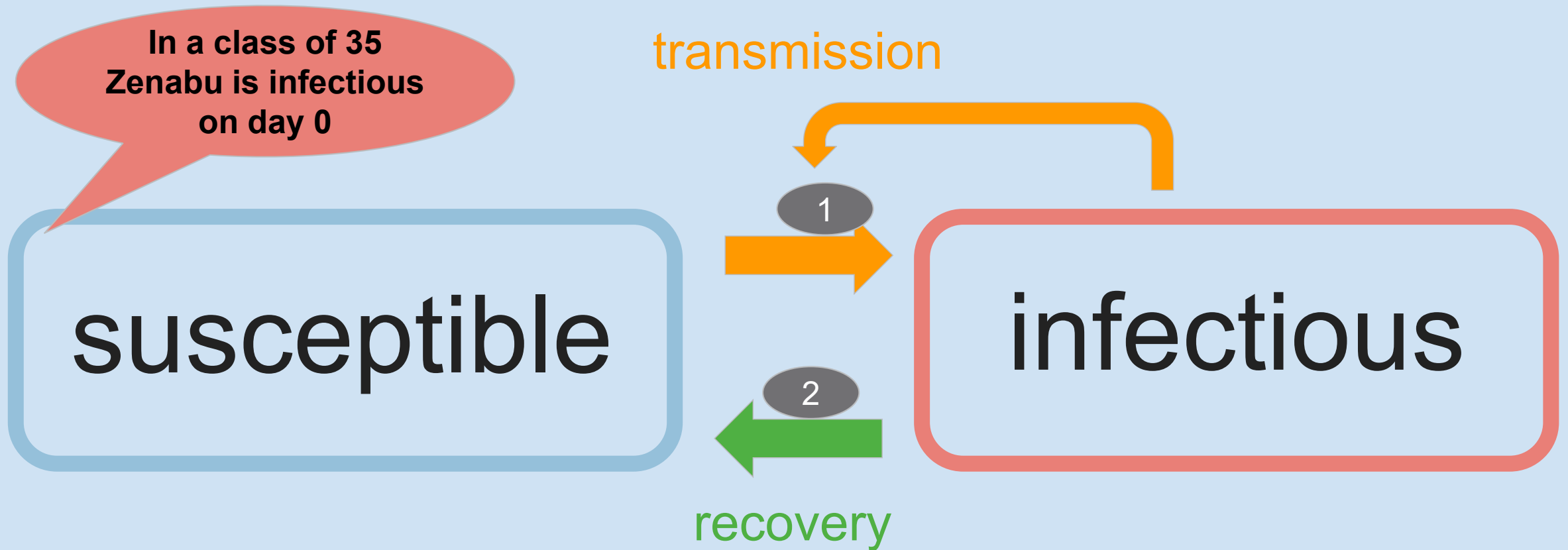
$$0 = -\beta \frac{I(t)}{N} S(t) + \gamma I$$

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

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

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

Practical: Infection OR No Infection



SIS model

Basic reproduction number

Introducing a single infection into a entirely susceptible population

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

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

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

“basic reproduction number”

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

Acknowledgements



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- Emilie Pothin
- Billy Bauzille
- Zenabu Suboi
- Geoffrey Githinji

Asanteni sana

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