

# Modeling COVID-19 Incidence

Interim Presentation for Stat 222 (Spring 2023)

## Background and motivation

- ▶ By mid-summer, 2021, vaccination eligibility for COVID-19 was widespread and preventive public health measures were significantly loosened
- ▶ Return to normalcy in the presence of vaccination led to concerns of the emergence of a vaccine-resistant strain
- ▶ In July 2021, Rella et al. (2021) published simulations of outbreak trajectories under various emergence probabilities
  - ▶ Resistant strains never established during periods of preventive public health measures
- ▶ On November 30, 2021, the first case of the Omicron variant (B.1.1.529) in the US was confirmed (CDC 2021)

## Insights from Rella et al. (2021)

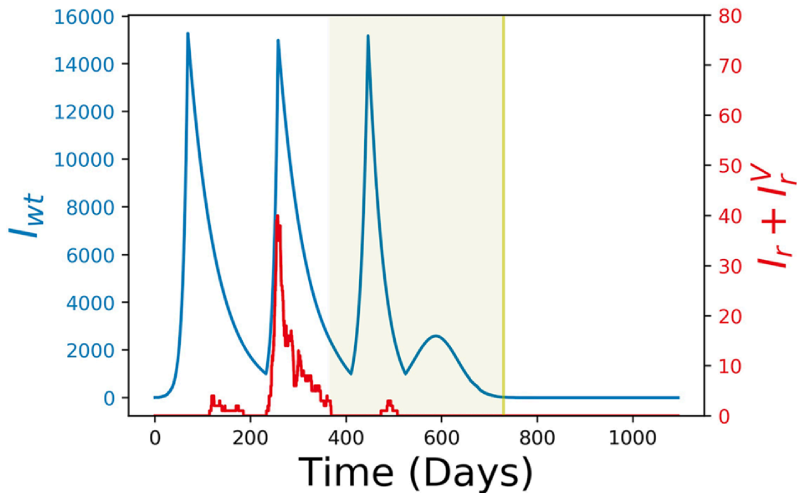


Figure 1: Low emergence probability

## Insights from Rella et al. (2021)

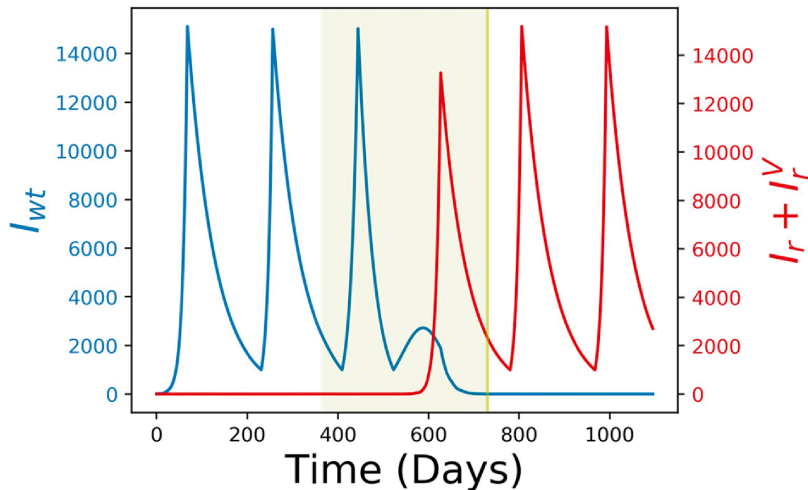
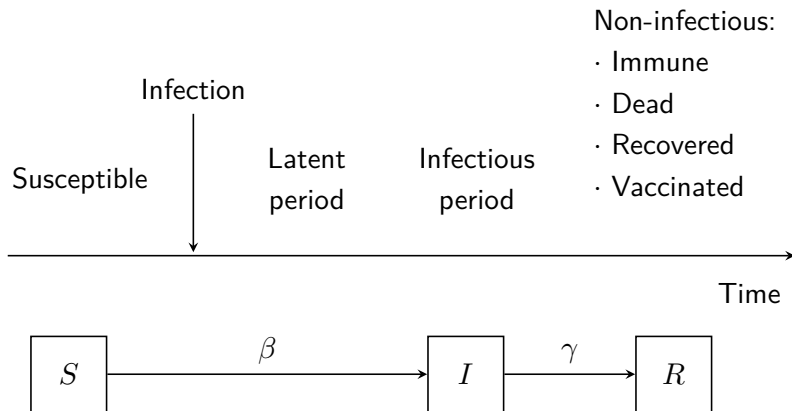


Figure 2: High emergence probability

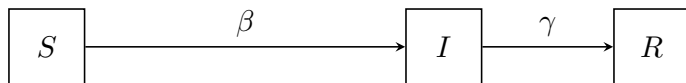
# Basic concepts in infectious disease epidemiology

- ▶ *Infectious agent*: biological causal locus of an infectious disease
- ▶ *Contact*: interaction between potential hosts and the infectious agent
- ▶ *Infection*: entry of the infectious agent into the host
- ▶ *Latent period*: time between infection and infectiousness
- ▶ *Infectious period*: the period of time during which contact with hosts means contact with the infectious agent
- ▶ These periods make up the *natural history timeline* for infectious disease

# Natural history timeline



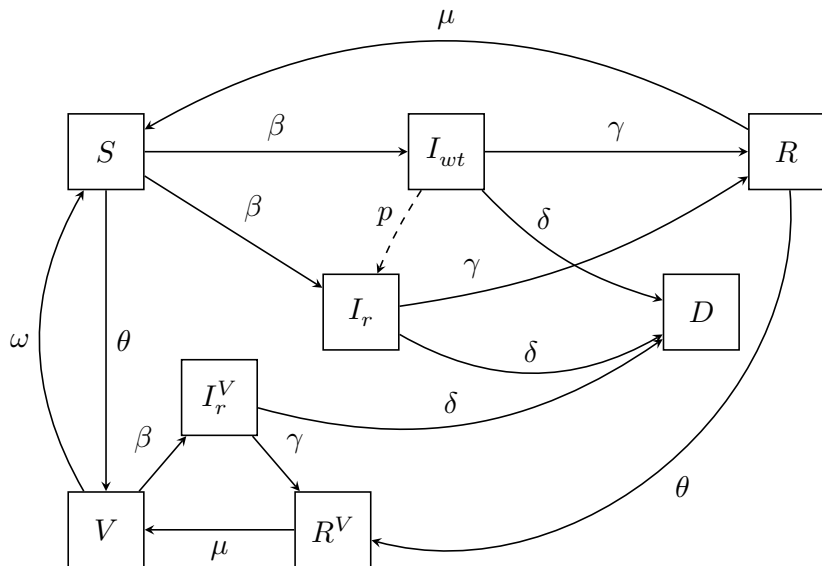
## Compartmental modeling: simple SIR case



$$\begin{cases} \frac{d}{dt}S = -\frac{\beta}{N}IS \\ \frac{d}{dt}I = \frac{\beta}{N}IS - \gamma I \\ \frac{d}{dt}R = \gamma I \end{cases}$$
$$S + I + R = N$$

See Harko, Lobo, and Mak (2014) for closed-form solutions.

# Compartmental modeling: theory of Rella et al. (2021)





## Modeling waves in epidemiologic trajectory

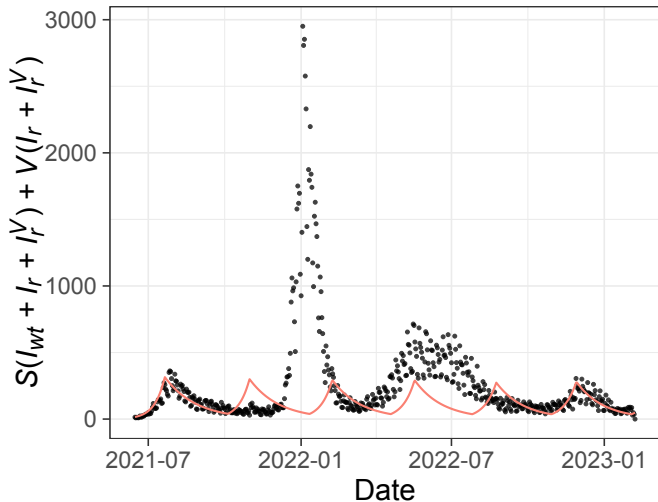
$$\beta(t) = R_0(t) \times \gamma \times N^{-1}$$

where

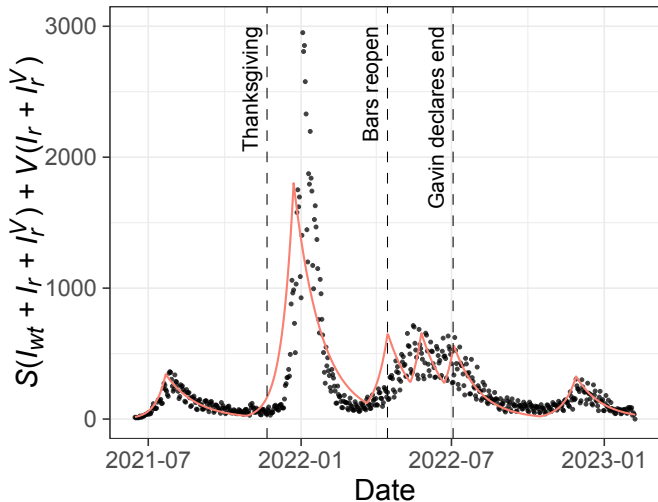
$$R_0(t) = \begin{cases} R_{0,\text{low}} & \text{if } I > N \times p_h \quad \text{and} \quad R_0(t - \Delta t) = R_{0,\text{high}} \\ R_{0,\text{high}} & \text{if } I \times p_h \times p_l \quad \text{and} \quad R_0(t - \Delta t) = R_{0,\text{low}} \end{cases}$$

where  $I = I_{wt} + I_r + I_r^V$

## Preliminary fit with time-invariant $p_h$ and $p_l$



## Preliminary fit with time-varying $p_h$ and $p_l$



## Parameter specification (time-permitting)

Parameter	Value	Interpretation
$\theta$	250.6335	Vaccination rate (vaccines per day)
$\delta$	$7.3 \times 10,000$	Death rate (deaths per day)
$R_{0h}$	2.2	$R_0$ during low caution
$R_{0l}$	0.65	$R_0$ during high caution
$\gamma$	1/14	Inv. disease duration (days)
$\mu$	1/(30 $\times$ 12)	Inv. duration of natural immunity
$\omega$	1/(30 $\times$ 12)	Inv. duration of vaccine protection

## Parameter specification (time-permitting)

Parameter	Value	Interpretation
$p_{h,1}$	1/350	Maximum prevalence before preventive measures in 1st wave
$p_{h,2}$	1/63	Maximum prevalence (2nd)
$p_{h,3}$	1/165	Maximum prevalence (3rd)
$p_{h,4}$	1/325	Maximum prevalence (4th)
$p_{l,1}$	$1/(350 \times 15)$	Allowable prevalence before loosening measures (1st)
$p_{l,2}$	$1/(63 \times 15)$	Allowable prevalence (2nd)
$p_{l,3}$	$1/(165 \times 2.5)$	Allowable prevalence (3rd)
$p_{l,4}$	$1/(325 \times 17)$	Allowable prevalence (4th)

## References

- CDC. 2021. "First Confirmed Case of Omicron Variant Detected in the United States." *First Confirmed Case of Omicron Variant Detected in the United States*.  
<https://www.cdc.gov/media/releases/2021/s1201-omicron-variant.html>.
- Harko, Tiberiu, Francisco SN Lobo, and MK3197716 Mak. 2014. "Exact Analytical Solutions of the Susceptible-Infected-Recovered (SIR) Epidemic Model and of the SIR Model with Equal Death and Birth Rates." *Applied Mathematics and Computation* 236: 184–94.
- Rella, Simon A, Yuliya A Kulikova, Emmanouil T Dermitzakis, and Fyodor A Kondrashov. 2021. "Rates of SARS-CoV-2 Transmission and Vaccination Impact the Fate of Vaccine-Resistant Strains." *Scientific Reports* 11 (1): 15729.