

**DRP AM-7**

# Modelling the COVID-19 pandemic

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# Outline

## Introduction

- Background
- Model Framework
- Model Equations/Parameters

## Results

- Waning Immunity + Reinfection
- Vaccination Effects
- Multiple Variants

## Conclusion

- Strengths
- Limitations
- Future work



# Background and Objective

**Objective:** Studying the impact of vaccination, immunity, and different variants on a global pandemic.

## Background:

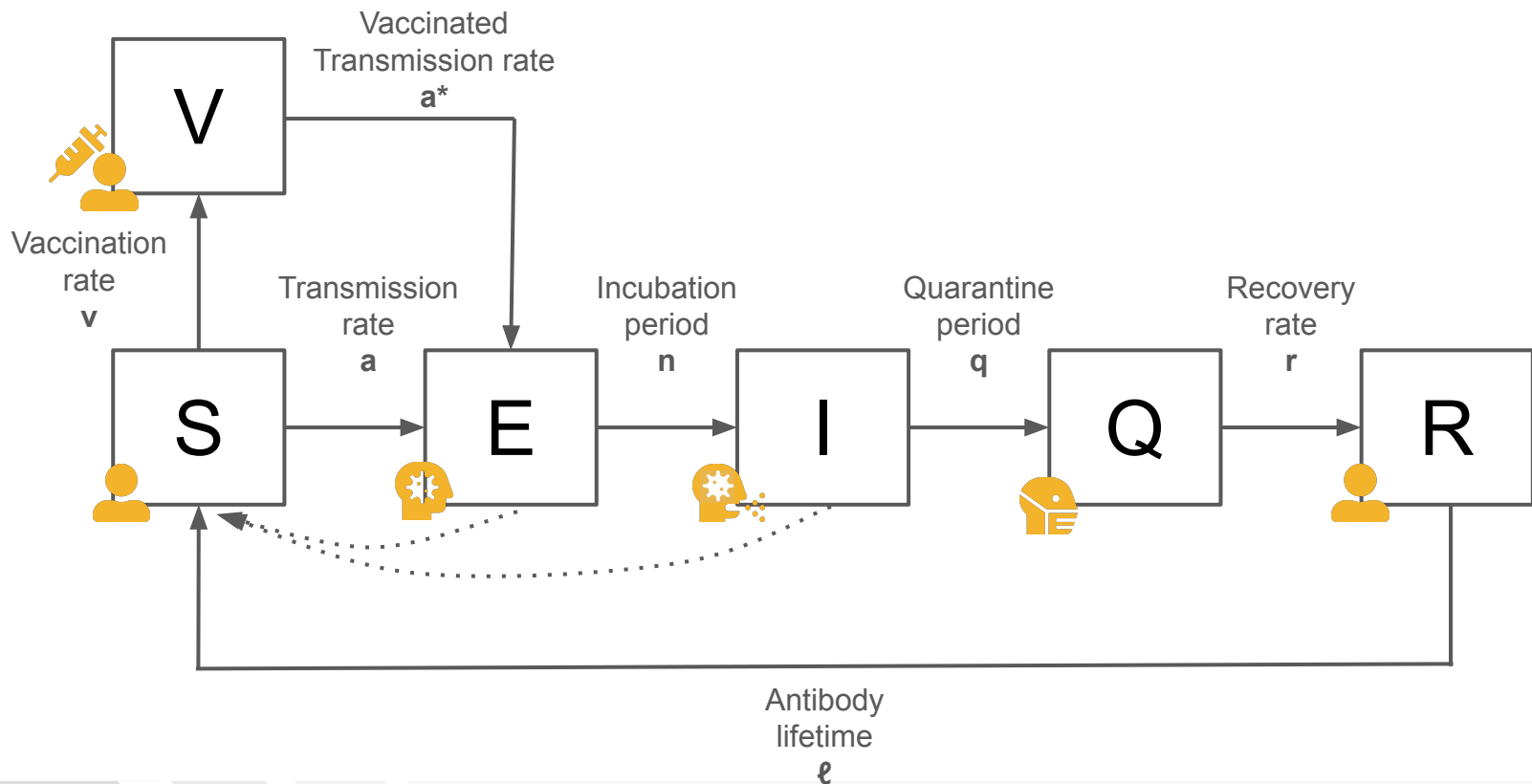
- COVID-19 created large-scale health and social disruption worldwide, and its spread is shaped by changing infection, recovery, vaccination, and immunity patterns
- Linear ODE modeling captures complex disease dynamics by tracking population flows between disease states

# Model Framework





# Model Framework

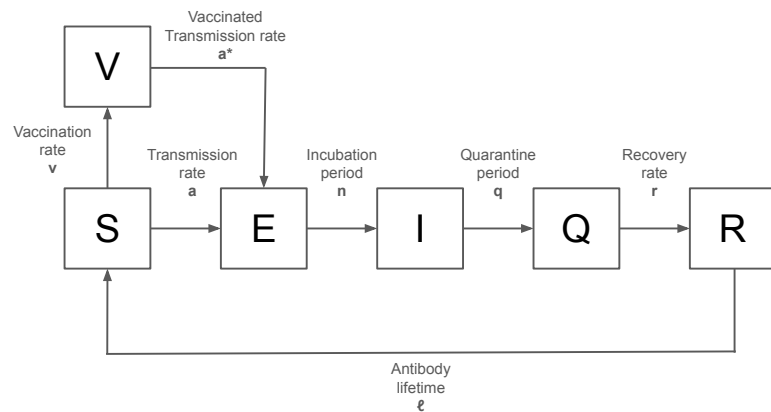




# Parameters

Parameter	Definition	Value
$a$	transmission rate ( $a_v$ for vaccinated)	$\frac{R_0 r}{S_0}$
$v$	vaccination rate	Varies
$n$	incubation rate	$\frac{1}{7}$
$q$	quarantine rate	$\frac{1}{14}$
$r$	recovery rate	$\frac{1}{14}$
$l$	antibody deterioration rate	$\frac{1}{90}$

Variant	Alpha	Beta	Delta	Gamma	Omicron
$R_0$	1.143	1.101	1.177	1.115	1.559





# Equations

Susceptible  $S$

$$\frac{dS}{dt} = \underbrace{-aS(E + I)}_{\text{become exposed}} + \underbrace{lR}_{\text{return from recovered}} + \underbrace{(-vS)}_{\text{vaccination loss}}$$

Vaccinated  $V$

$$\frac{dV}{dt} = \underbrace{vS}_{\text{vaccination gain}} - \underbrace{a_v(aVE + aVI)}_{\text{exposed despite vaccination}}$$

Exposed  $E$

$$\frac{dE}{dt} = \underbrace{aS(I + E)}_{\text{exposure from susceptibles}} + \underbrace{(-nE)}_{\text{progression to infected}} + \underbrace{a_v(aV(E + I))}_{\text{exposure among vaccinated}}$$

Infected  $I$

$$\frac{dI}{dt} = \underbrace{nE}_{\text{becoming infected}} + \underbrace{(-qI)}_{\text{to quarantine}}$$

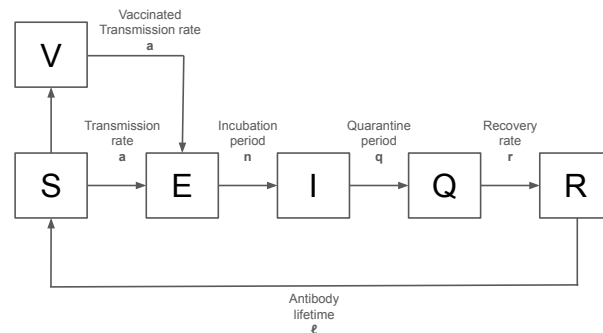
Quarantined  $Q$

$$\frac{dQ}{dt} = \underbrace{qI}_{\text{enter quarantine}} + \underbrace{(-rQ)}_{\text{recovery from quarantine}}$$

Recovered  $R$

$$\frac{dR}{dt} = \underbrace{rQ}_{\text{recovery from quarantine}} + \underbrace{-lR}_{\text{loss of immunity}}$$

- Negatives imply people leaving the group
- Positives imply entering the group



# Results



# Waning Immunity ( $R \rightarrow S$ )

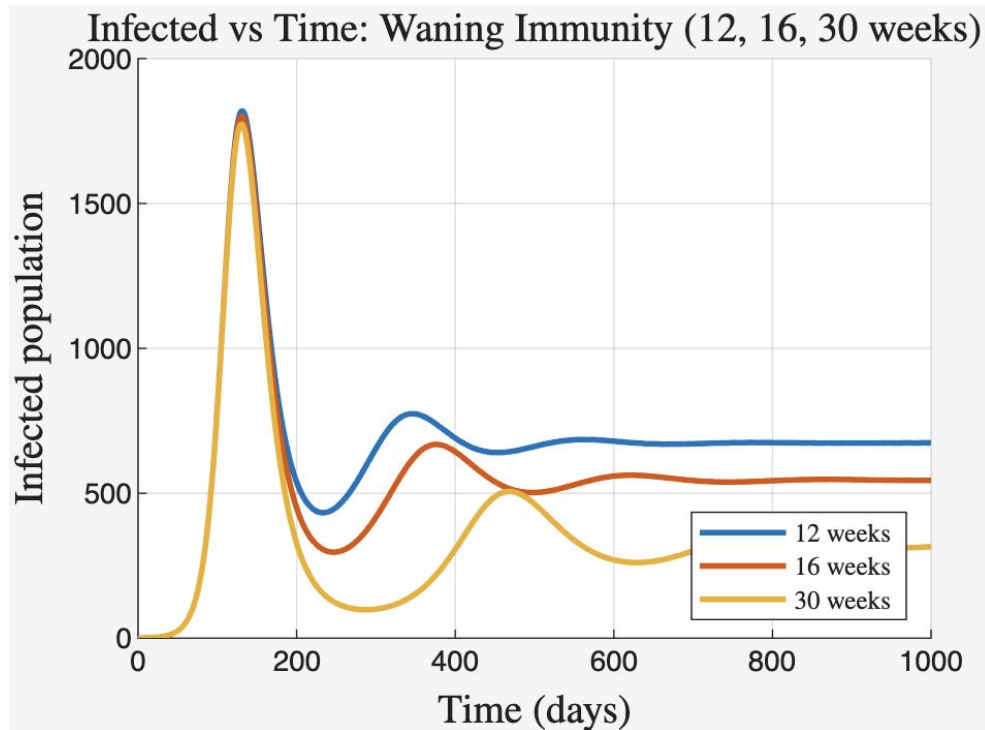
$\ell$  = antibody deterioration rate ( $R \rightarrow S$ )

Tested immunity durations:

- $\ell = 1/84 \approx 12$  weeks immunity
- $\ell = 1/112 \approx 16$  weeks immunity
- $\ell = 1/210 \approx 30$  weeks immunity



# Infection $I(t)$



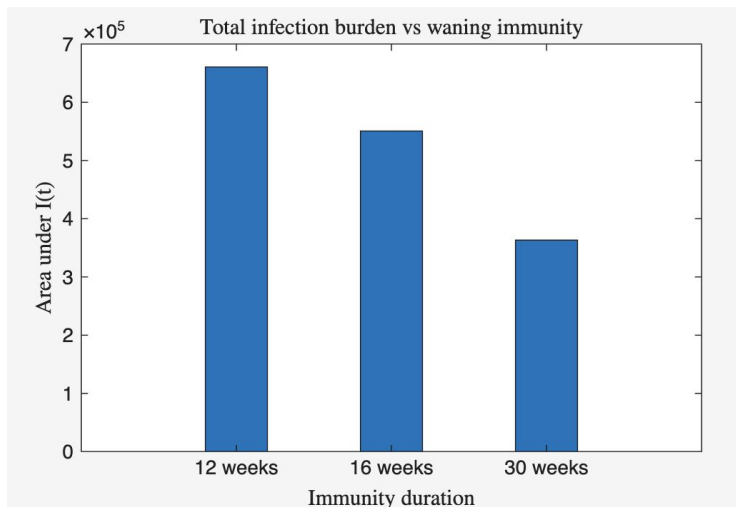
## Key observations:

- 30 weeks (yellow): Sharp single wave
- 16 weeks (red): Broader, higher baseline
- 12 weeks (blue): Largest peak (1820), sustained transmission



# Total Epidemic Size

Immunity duration	I value	Peak I	Peak time	AUC (total infections)
12 weeks	0.01190	1820.3	131 days	660,614
16 weeks	0.00893	1801.7	131 days	550,618
30 weeks	0.00476	1774.2	130 days	363,242



Key observation: 12-week immunity creates 1.8x larger epidemics

# SEIQRV

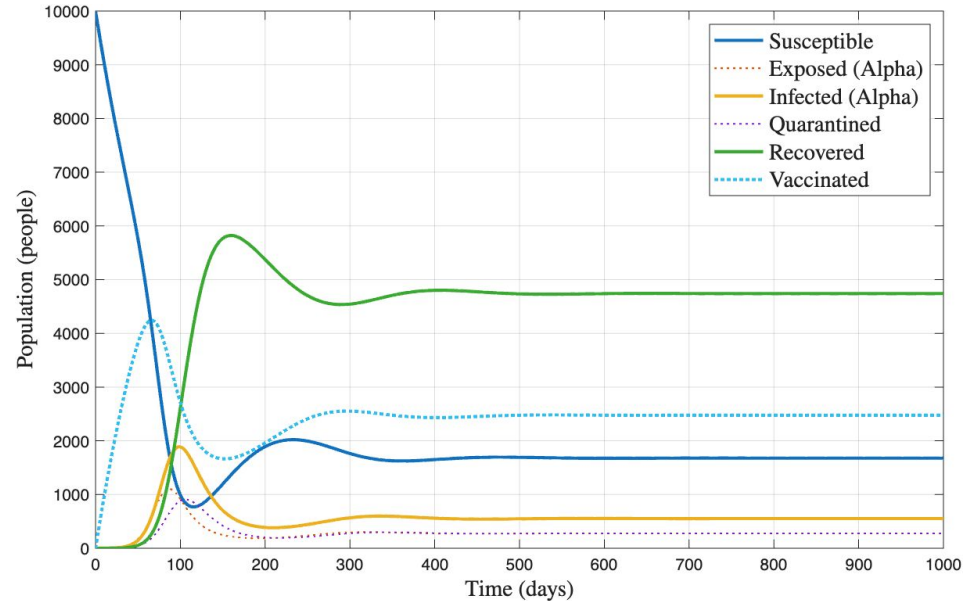
Susceptible Exposed Infected Quarantined Recovered Vaccinated



# What are we looking at?

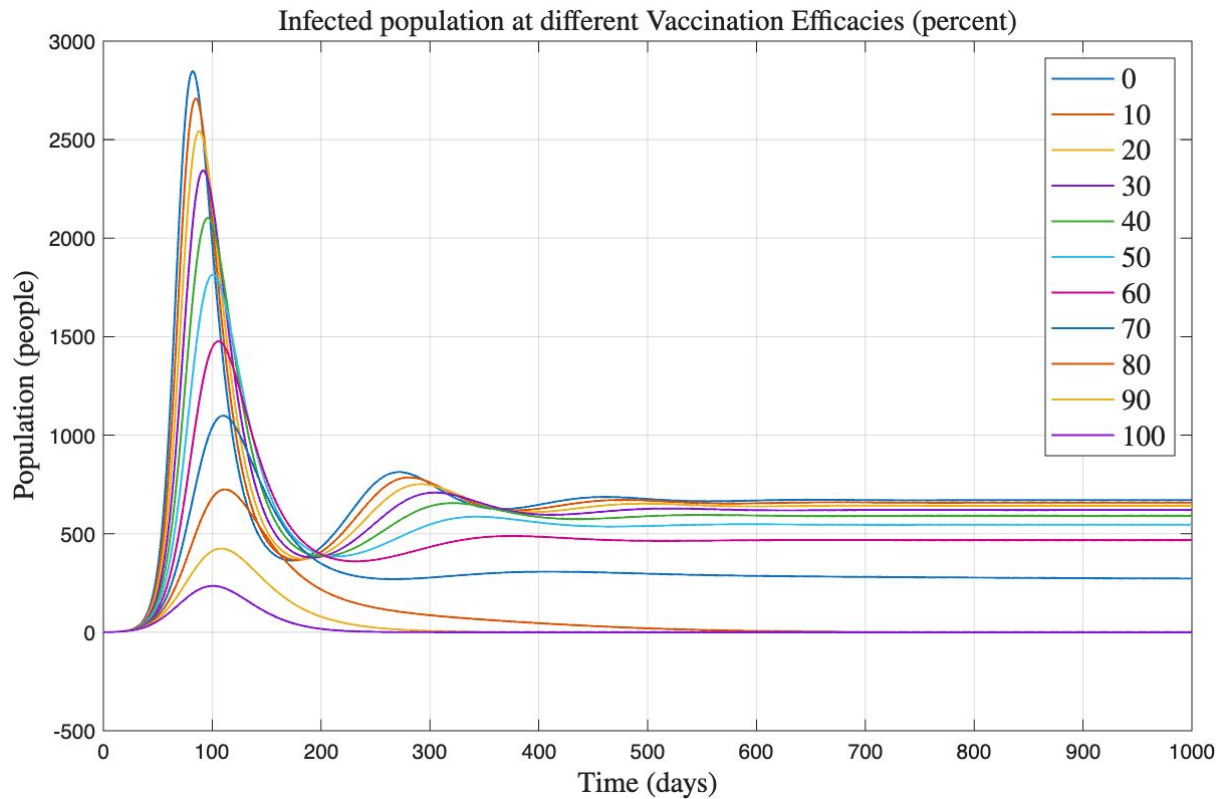
- **Vaccine efficacy:** reduced transmission
  - Different efficacy rates (how much transmission is reduced)
- **Vaccination rate:**
  - Different vaccination rates

Epidemic progression

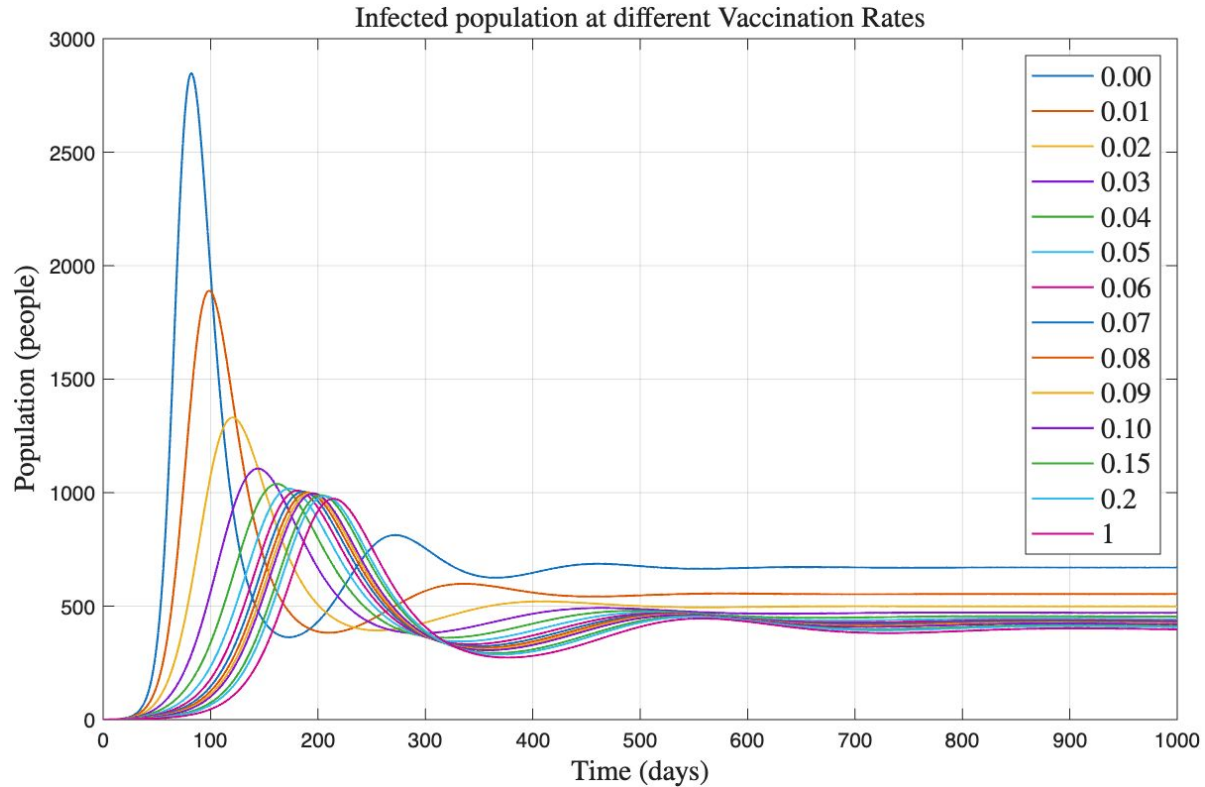




# Vaccination Efficacy



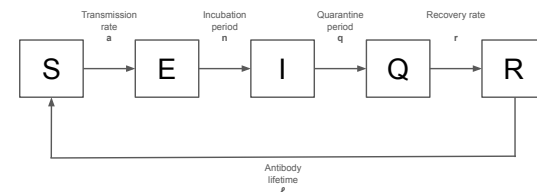
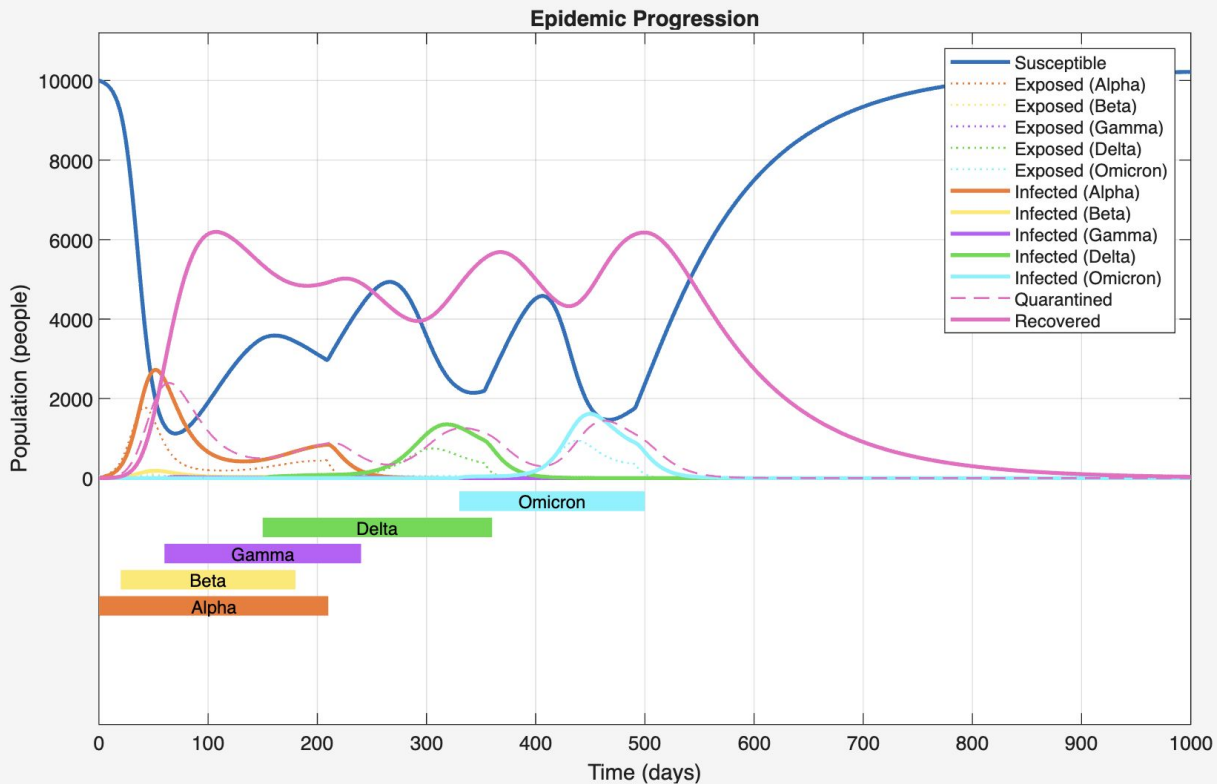
# Vaccination Rate



# SEIQR (+ Variants)

Susceptible Exposed Infected Quarantined Recovered

# Results - COVID Variants



- Alpha, Beta, Gamma, Delta, Omicron Variants
- Drops in susceptible population  
→ peaks in infected groups

# Conclusion



# Strengths

- Projection of disease progression
- Can see how different interventions affect epidemic in a “controlled” environment
  - Tells us what to prioritize
  - e.g. Vaccination: prioritize efficacy over rate



# Limitations

## Model assumptions

- Fixed population:  $N = 10,000$  (no deaths/births) but COVID mortality + natural demographics change susceptibility
- Everyone follows  $S \rightarrow E \rightarrow I \rightarrow Q \rightarrow R \rightarrow S$  exactly, but there are asymptomatic cases, misdiagnosis, treatment failures

## Linear ODE assumptions

- No travel, superspreaders, regional outbreaks
- No behavioural changes (lockdowns, testing, masks)
- Single homogeneous population "box"



# Future Work

Opinion dynamics



Human behaviour



Closure



Thank you!

Questions?

# References

Manathunga, S. S., Abeyagunawardena, I. A., & Dharmaratne, S. D. (2023). A comparison of transmissibility of SARS-CoV-2 variants of concern. *Virology journal*, 20(1), 59. <https://doi.org/10.1186/s12985-023-02018-x>  
*SARS-COV-2 variants in analyzed sequences*. Our World in Data. (n.d.). <https://ourworldindata.org/grapher/covid-variants-area>