Maths & the Evolution of Viruses

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1 / 24

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- 1 Motivation
- **2** Measuring escape
- **3** Epidemic models
 - Transient scenario
 - Endemic scenario
- 4 So what?
 - Results
 - Mathematical insights
- **5** Conclusion
- 6 Optional extra
- **7** Questions



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1 Motivation

- **2** Measuring escape
- **3** Epidemic models
- 4 So what?
- **5** Conclusion
- 6 Optional extra





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Motivation



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3 Epidemic models

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6 Optional extra





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Measuring escape: it's difficult!

Quantify the "immune/antigenic/vaccine escape pressure" exerted on the virus towards mutations with the ability to escape immunity (eg antibodies).

Can think of it as the probability that an escape variant appears during the epidemic:

• Escape at probability $p_U \ll 1$ per infection in the unvaccinated.

• Escape at probability $p_V \ll 1$ per infection in the vaccinated.

$$P = 1 - (1 - p_U)^{N_U} (1 - p_V)^{N_V} pprox p_U N_U + p_V N_V$$

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Measuring escape: the unknowns

Only care about **relative escape pressure** (relative to epidemic without vaccination). Define $\theta_E = p_V/p_U$, so

 $P \propto N_U + \theta_E N_V$

Is $\theta_E < 1$ or $\theta_E > 1$?

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Measuring escape: the unknowns

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 $P \propto N_U + \theta_E N_V$

Is $\theta_E < 1$ or $\theta_E > 1$? We have no idea, θ_E could be in any value in $[0,\infty)!!$

- Unvaccinated individuals have higher viral load
 more mutations
- Escape variants evade immunity in vaccinated individuals => more selection pressure



Figure 1: *Phylodynamics* (Grenfell et al. Science)

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Measuring escape: imperfect vaccines



Probability of infection reduced by a factor θ_S (susceptibility reduction)

Probability of transmission reduced by a factor θ_I (infectivity reduction)

Probability of escape mutation changed by a factor θ_E

Image: A image: A

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1 Motivation

2 Measuring escape

3 Epidemic models

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- Endemic scenario

4 So what?

6 Conclusion

6 Optional extra

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9 / 24

Background: the SIR model

Use proportions of the total population, start with everyone susceptible S(0) = 1, except an infinitesimal proportion of infected $I(0) = \epsilon$, $R_0 > 1$ (initial outbreak will grow)

Susceptible	$\dot{S} = -R_0 SI$
Infectious	$\dot{I} = R_0 S I - I$
Recovered	$\dot{R} = +I$

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Can eliminate R = 1 - S - I, and solve the 2D system:

$$\frac{dI}{dS} = \frac{\dot{I}}{\dot{S}} = \frac{R_0 SI - I}{-R_0 SI} = -1 + \frac{1}{R_0 S} \implies I = 1 - S + R_0^{-1} \log S$$

For final size, take I
ightarrow 0: $S_{\infty} = 1 - R_0 \log S_{\infty}^{-1}$ (implicit), total infections is

$$N = 1 - S_{\infty} = 1 + rac{W[-R_0e^{-R_0}]}{R_0}$$

where W is the Lambert W function.

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Aside: the Lambert W function

W(z) solves $We^W = z$ in the complex plane, has multiple branches!



For real numbers, only need two branches, W_0 and W_{-1} For the SIR model, only need principal branch $W_0(x) \in [-1,0)$ for $x \in [-e^{-1},0)$

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Transient scenario: setting up the system

Duplicated compartments for vaccinated or unvaccinated

Full permanent immunity from vaccines or infections

Vaccination before the outbreak, a proportion c of the population:

 $S_U(0) = (1 - c) \leftarrow$ susceptible unvaccinated, $S_V(0) = c\theta_S \leftarrow$ susceptible vaccinated

	Unvaccinated	Vaccinated
Susceptible	$\dot{S}_U = -R_0 S_U \lambda(t)$	$\dot{S}_V = -R_0 S_V \lambda(t)$
Infectious	$\dot{I}_U = R_0 S_U \lambda(t) - I_U$	$\dot{I}_V = R_0 S_V \lambda(t) - I_V$
Recovered	$\dot{R}_U = I_U$	$\dot{R}_V = I_V$

 $\lambda(t) = I_U + \theta_I I_V$ (effective number of infectious individuals)

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Transient scenario: solving the system 1/2

Claim:
$$(S_V, I_V, R_V) = \frac{c\theta_S}{1-c}(S_U, I_U, R_U)$$
 at all times.

Proof: For
$$\mathbf{u}^T = (S_U(t), I_U(t), R_U(t))$$
 and $\mathbf{v}^T = (S_V(t), I_V(t), R_V(t))$,
 $\dot{\mathbf{u}} = \mathbf{M}\mathbf{u}$ $\dot{\mathbf{v}} = \mathbf{M}\mathbf{v}$

with
$$\mathbf{M}(t) = \begin{pmatrix} -R_0\lambda(t) & 0 & 0\\ R_0\lambda(t) & -1 & 0\\ 0 & +1 & 0 \end{pmatrix}$$
 and $\mathbf{v}(0) = \frac{c\theta_s}{1-c}\mathbf{u}(0)$.
Both $\hat{\mathbf{u}}(t) := \frac{1-c}{c\theta_s}\mathbf{v}(t)$ and $\mathbf{u}(t)$ obey the same first order ODE and IC $\mathbf{u}(0) = \hat{\mathbf{u}}(0)$.
Hence (by uniqueness of solutions), $\hat{\mathbf{u}}(t) = \mathbf{u}(t)$. Thus $\mathbf{v}(t) = \frac{c\theta_s}{1-c}\mathbf{u}(t)$.

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Transient scenario: solving the system 2/2

This proportion reduces the system to a standard SIR model

$$\dot{S}_U = -S_U I_U R_0 \left(1 + rac{c heta_S heta_I}{1 - c}
ight) \qquad \dot{I}_U = +S_U I_U R_0 \left(1 + rac{c heta_S heta_I}{1 - c}
ight) - I_U
onumber \ S_U(0) = 1 - c \qquad I_U(0) = \epsilon \ll 1$$

with effective R-number $R_e = R_0(1 - c + c\theta_S\theta_I)$. $R_e = 1$ gives herd-immunity threshold $\tilde{c} = (1 - R_0^{-1})/(1 - \theta_S\theta_I)$. For $c < \tilde{c}$, get $N_U = (1 - c) (1 + W(-R_e e^{-R_e})/R_e) = (1 - c)(c\theta_S)^{-1}N_V$

$$P = N_U + \theta_E N_V = (1 - c + \theta_S \theta_E c) \left(1 + \frac{1}{R_e} W \left(-R_e e^{-R_e} \right) \right)$$

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 14 / 24
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Endemic scenario: setting up the system

	Unvaccinated	Vaccinated
Susceptible	$\dot{S}_U = \mu(1-c) + R_0 S_U \lambda - \mu S_U$	$\dot{S}_V = \mu c \theta_S - R_0 S_V \lambda - \mu S_V$
Infectious	$\dot{I}_U = R_0 S_U \lambda - I_U - \mu I_U$	$\dot{I}_V = R_0 S_V \lambda - I_V - \mu I_V$
Recovered	$\dot{R}_U = I_U - \mu R_U$	$\dot{R}_V = I_V - \mu R_V$

 μ is per capita birth/death rate, vaccinate proportion c of births



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Endemic scenario: solving the system

Claim: $(S_V, I_V, R_V) = \frac{c\theta_S}{1-c}(S_U, I_U, R_U)$ at all times. Proof: $\dot{\mathbf{u}} = \mathbf{x}_U + \mathbf{M}(t)\mathbf{u}, \ \dot{\mathbf{v}} = \mathbf{x}_V + \mathbf{M}(t)\mathbf{v}$, with $\mathbf{M}(t) = \begin{pmatrix} -R_0\lambda(t) - \mu & 0 & 0 \\ R_0\lambda(t) & -1 - \mu & 0 \\ 0 & +1 & -\mu \end{pmatrix}$ and $\mathbf{x}_U^T = (\mu(1-c), 0, 0) = \frac{1}{\alpha}\mathbf{x}_V^T, \ \mathbf{v}(0) = \alpha\mathbf{u}(0)$ for $\alpha = c\theta_S/(1-c)$. Both $\mathbf{u}(t), \ \hat{\mathbf{u}}(t) := \alpha^{-1}\mathbf{v}(t)$ solve the IVP: $\dot{\mathbf{y}}(t) = \mathbf{F}(t, \mathbf{y}(t)) = \mathbf{M}(t)\mathbf{y}(t) + \mathbf{x}_U, \ \mathbf{y}(0) = \mathbf{u}(0)$, so $\mathbf{u}(t) = \ \hat{\mathbf{u}}(t)$

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Endemic scenario: solving the system

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$$(S_V, I_V, R_V) = \frac{c\theta_S}{1-c}(S_U, I_U, R_U)$$
 at all times.
Proof: $\dot{\mathbf{u}} = \mathbf{x}_U + \mathbf{M}(t)\mathbf{u}, \ \dot{\mathbf{v}} = \mathbf{x}_V + \mathbf{M}(t)\mathbf{v}$, with $\mathbf{M}(t) = \begin{pmatrix} -R_0\lambda(t) - \mu & 0 & 0\\ R_0\lambda(t) & -1 - \mu & 0\\ 0 & +1 & -\mu \end{pmatrix}$
and $\mathbf{x}_U^T = (\mu(1-c), 0, 0) = \frac{1}{\alpha}\mathbf{x}_V^T$, $\mathbf{v}(0) = \alpha\mathbf{u}(0)$ for $\alpha = c\theta_S/(1-c)$.
Both $\mathbf{u}(t), \ \hat{\mathbf{u}}(t) := \alpha^{-1}\mathbf{v}(t)$ solve the IVP:
 $\dot{\mathbf{y}}(t) = \mathbf{F}(t, \mathbf{y}(t)) = \mathbf{M}(t)\mathbf{y}(t) + \mathbf{x}_U, \mathbf{y}(0) = \mathbf{u}(0)$, so $\mathbf{u}(t) = \ \hat{\mathbf{u}}(t)$

Endemic equilibrium: Set derivatives to zero, look for $(S_U^*, I_U^*, R_U^*) = \frac{1-c}{c\theta_S}(S_V^*, I_V^*, R_V^*)$ For $R_e > 1 + \mu$, $I_U^* = \mu(1-c) [1/(1+\mu) - 1/R_e] = (1-c)I_V^*/c\theta_S$. Hence $P^* = I_S^* + \theta_E I_U^* = (1 - c(1 - \theta_E \theta_S))\mu [1/(1+\mu) - 1/R_e]$. Take limit $\mu \ll 1$

$${{\it P}^{st}} \propto \left(1-{\it c}(1- heta_{{\it E}} heta_{{\it S}})
ight) \left(1-{\it R}_{{\it e}}^{-1}
ight)$$

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Results: escape pressure as a function of c

- ▶ Peaks at intermediate vaccination c for θ_E above a threshold $\hat{\theta}_E$: this is when vaccinated individuals contribute a lot to escape
- Monotonically decreasing with *c* for $\theta_E < \hat{\theta}_E$





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18 / 24

Finding the maximiser and the bifurcation point

$$P^* = (1 + c(\theta_S \theta_E - 1))(1 - R_e^{-1}), R_e = R_0(1 - c(1 - \theta_S \theta_I))$$

Differentiate to find gradient and evaluate at c = 0:

$$\frac{dP}{dc} = (\theta_{S}\theta_{E} - 1)(1 - R_{e}^{-1}) - (1 + c(\theta_{S}\theta_{E} - 1))R_{e}^{-2}R_{0}(1 - \theta_{S}\theta_{I})$$

$$\left. \frac{dP}{dc} \right|_{c=0} = (\theta_{S}\theta_{E} - 1)(1 - R_{0}^{-1}) - R_{0}^{-1}(1 - \theta_{S}\theta_{I})$$

This is negative $(P(c) \text{ decreasing } \forall c)$ iff $(\theta_E \theta_S - 1) < (1 - \theta_S \theta_I)/(R_0 - 1)$

Else maximiser is
$$c_m = (1 - \theta_S \theta_I)^{-1} \left[1 - \sqrt{1 + (1 - \theta_S \theta_I)/(\theta_S \theta_E - 1)} / \sqrt{R_0} \right]$$

1 Motivation

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Conclusion and takeaways

- θ_E is key in regulating how vaccination affects escape
 - Intermediate levels of vaccination can be the most dangerous
 - But also possible that any vaccination level decreases escape
- Other evo-epi models should not neglect $\theta_E!$
- Would be very helpful to empirically estimate θ_E
- > At some point, vaccines could also be evaluated based on their θ_E
- ▶ While θ_E remains unknown, keep vaccinating to reduce escape pressure
- Perhaps escape variants more likely to come from countries with lower vaccination rates (so donate vaccines!)
- Maths is super useful in this context

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Where's my model going wrong or what is missing?

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Where's my model going wrong or what is missing?

- Transmission of escape variants
- Reinfections
- Waning immunity
- Different individuals likely contribute differently to escape
- Multiple vaccine doses
- Multiple vaccine types
- Multiple variants in circulation

All models are wrong, but some are useful...(George Box)

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Any questions?

Thank you for listening and thanks to my PhD supervisor, Prof. Julia Gog



https://mariaalegriagutierrez.wordpress.com