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SIRS epidemics with individual heterogeneity of immunity waning

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Abstract

In the current paper we analyse an extended SIRS epidemic model in which immunity at the individual level wanes gradually at exponential rate, but where the waning rate may differ between individuals, for instance as an effect of differences in immune systems. The model also includes vaccination schemes aimed to reach and maintain herd immunity. We consider both the *informed* situation where the individual waning parameters are known, thus allowing selection of vaccinees being based on both time since last vaccination as well as on the individual waning rate, and the more likely *uninformed* situation where individual waning parameters are unobserved, thus only allowing vaccination schemes to depend on time since last vaccination. The optimal vaccination policies for both the informed and uninformed heterogeneous situation are derived and compared with the homogeneous waning model (meaning all individuals have the same immunity waning rate), as well as to the classic SIRS model where immunity at the individual level drops from complete immunity to complete susceptibility in one leap. It is shown that the classic SIRS model requires least vaccines, followed by the SIRS with homogeneous gradual waning, followed by the informed situation for the model with heterogeneous gradual waning. The situation requiring most vaccines for herd immunity is the most likely scenario, that immunity wanes gradually with unobserved individual heterogeneity. For parameter values chosen to mimic COVID-19 and assuming perfect initial immunity and cumulative immunity of 12 months, the classic homogeneous SIRS epidemic suggests that vaccinating individuals every 15 months is sufficient to reach and maintain herd immunity, whereas the uninformed case for exponential waning with rate heterogeneity corresponding to a coefficient of variation being 0.5, requires that individuals instead need to be vaccinated every 4.4 months.

Keywords: SIRS model, Immunity waning, Heterogeneity, Vaccination, Herd immunity.

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1 Introduction

Among other things, the COVID-19 pandemic showed that immunity waning as well as immunity escape for new virus strains play important roles when designing vaccination schemes to reduce and ultimately stop the spreading of an epidemic. In the current paper the focus lies on immunity waning for a fixed and specific strain and we thus study an epidemic model for an infectious disease where immunity, both from vaccination as well as natural infection, wanes gradually and monotonically following an exponential mode Wheatley et al. (2021).

The classic SIRS model Hethcote (1976, 1978) is the first model to consider immunity waning, and in this model population immunity decays gradually, but at the individual level each individual is either fully immune or fully susceptible. In the last few years this assumption has been relaxed (e.g. Reluga et al. (2008); Martcheva (2015); Forien et al. (2022); El Khalifi and Britton (2023)) thus allowing for gradual waning of immunity also at the individual level, resulting in individuals having different immunity levels, defined either discretely or continuously. These models still assume that immunity wanes in a similar fashion for all individuals, most often defined by a waning rate ω common for all individuals. For these models it has been shown that such gradual immunity waning requires more frequent vaccination to reach and maintain herd immunity as compared to the classic SIRS model which assumes one single jump from fully immune to fully susceptible (having the same average cumulative immunity) El Khalifi and Britton (2023).

Empirical measurements of antibodies however suggest large individual differences in antibody decay between individuals Fabiani et al. (2022); Shrotri et al. (2021); Widge et al. (2021); Pérez-Alós et al. (2022) thus suggesting different waning rates between individuals. In the present paper we therefore extend a model with homogeneous gradual immunity waning to a situation where the waning rate may differ between individuals. The general situation, where waning rates of individuals are drawn independently from some general random distribution is complicated to analyse, so here we focus on the situation where there are two types of waning rates ω_1 and ω_2 with population frequencies p and $1 - p$ respectively. We compare the heterogeneous situation with the homogeneous case having the same cumulative immunity ($= 1/\omega$) and we quantify the amount of heterogeneity by the coefficient of variation of the immunity distribution.

This paper is structured as follows. In the next section we present the SIRS models with heterogeneity under both situations: sudden loss and continuous waning of immunity. In Section 3 we formulate the SIR^(k)S model with heterogeneity. In Section 4, we introduce vaccination into the model by taking into account the effect of the available information on individuals immunity. To illustrate the results for our models, in Section 5, we compare the long term prevalence and the optimal vaccination schemes under parameter values mimicking the COVID-19 pandemic. We conclude the paper in Section 6 with a discussion and draw some perspectives.

2 Models

First, we define a model where immune individuals lose their immunity at once. Next we modify the model to allow for gradual (exponential) waning of individual immunity. For both models we divide the population into two immunity waning classes with waning rates ω_1 and ω_2 with fractions p and $1 - p$, allowing for a certain degree of heterogeneity. We compare the homogeneous case with cumulative immunity $1/\omega$ to the heterogeneous case with $1/\omega_1 = (1 - \alpha)(1/\omega)$ and $1/\omega_2 = (1 + \alpha \frac{p}{1-p})(1/\omega)$ for some α ($0 \leq \alpha \leq 1$) so that the

cumulative immunity is set to $1/\omega$. The coefficient of variation of the immunity distribution is given by $\sigma = \alpha\sqrt{p/(1-p)}$ which from now on is used as heterogeneity parameter rather than α . Hence, we have

$$\frac{1}{\omega_1} = \frac{1}{\omega} \left(1 - \sigma\sqrt{\frac{1-p}{p}}\right) \quad \text{and} \quad \frac{1}{\omega_2} = \frac{1}{\omega} \left(1 + \sigma\sqrt{\frac{p}{1-p}}\right).$$

2.1 The SIRS model with heterogeneity

The model that we consider in this section, also taking births and deaths into account, is defined as follows. Let $l \in \{1, 2\}$ be the index of the immunity waning classes and denote $s_l(t)$, $i_l(t)$, and $r_l(t)$ the community fractions of susceptible, infectious, and recovered l -individuals at time t , respectively. The model parameters are as defined in Table 1. Then,

Table 1: Model Parameters and interpretation.

Parameter	Description
μ	Birth and death rate
β	Effective infection rate
γ	Recovery rate
ω	Immunity waning rate
σ	Coefficient of variation of immunity distribution

the differential equations for the SIRS model with heterogeneity are given by

$$\begin{aligned} s_l'(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) - \mu s_l(t) + \omega_l r_l(t), \\ i_l'(t) &= \beta s_l(t) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \\ r_l'(t) &= \gamma i_l(t) - (\mu + \omega_l) r_l(t), \end{aligned} \tag{1}$$

with $l \in \{1, 2\}$ (so $s_l(t) + i_l(t) + r_l(t) = p_l$), $p_1 = p$ and $p_2 = 1 - p$. We define the basic reproduction number to be $R_0 = \frac{\beta}{\gamma + \mu}$ representing the average number of new infections generated by an infectious person in an entirely susceptible population. We have the following standard result for our model.

Proposition 2.1. *The solution to Eq. (1) has a unique endemic equilibrium if and only if $R_0 > 1$.*

When the endemic equilibrium exist, the endemic level is given by the sum of the constant fractions of infectives \hat{i}_1 and \hat{i}_2 in the type-1 and type-2 communities respectively. We also have the following result regarding the dependence of the endemic level on the population heterogeneity. Recall the p is the community fraction having lower immunity and hence higher waning rate ω_1 .

Proposition 2.2. *Assume that $R_0 > 1$ and $1/2 \leq p < 1$. Then, the endemic level is an increasing function of the coefficient of variation σ on $[0, \sqrt{p/(1-p)})$.*

The proofs of the Propositions 2.1 and 2.2 are given in the Appendix A. Although the monotonicity in Proposition 2.2 is only proved for a fraction p satisfying $1/2 \leq p < 1$, numerical simulations suggest that the endemic level is also increasing in σ on $[0, \sqrt{p/(1-p)})$ for any $0 < p < 1/2$.

2.2 The SIR^(∞)S model with heterogeneity

When immunity wanes continuously and following an exponential decay, the recovered equation could be modelled using a PDE evolving in calendar time t and time-since-recovery a , and the model equations become

$$\begin{aligned} s'_l(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) - \mu s_l(t), \\ i'_l(t) &= \beta \left(s_l(t) + \int_0^\infty (1 - e^{-\omega_l a}) r_l(t, a) da \right) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \end{aligned} \quad (2)$$

$$\frac{\partial r_l(t, a)}{\partial t} + \frac{\partial r_l(t, a)}{\partial a} = -\beta (1 - e^{-\omega_l a}) r_l(t, a) (i_1(t) + i_2(t)) - \mu r_l(t, a), \quad a > 0,$$

with the boundary condition $r_l(t, 0) = \gamma i_l(t)$, $l \in \{1, 2\}$ where $p_1 = p$ and $p_2 = 1 - p$.

We call the model (2) the SIR^(∞)S model with heterogeneity as it can be seen as the k -limit of the heterogeneous SIR^(k)S model where immunity drops in k steps, $1/k$ each time El Khalifi and Britton (2023). We refer to El Khalifi and Britton (2023) for more details on the construction (see also Section 3). We have the following expected result for the model (2).

Proposition 2.3.

- Assume that $R_0 \leq 1$. Then, the solution to Eq. (2) converges to the disease-free equilibrium.
- Assume that $R_0 > 1$. Then, Eq. (2) has a unique endemic equilibrium.

The proof of the Proposition 2.3 is given in Appendix A.1.3.

3 SIR^(k)S model with heterogeneity

Similarly to the approach in El Khalifi and Britton (2023), we approximate the SIR^(∞)S model (2) by a system of ODEs allowing immunity to wane in k steps for some large value of k .

We now describe how this reduction of immunity in k small steps down to no immunity, each step having a high rate to drop to the next level. This can be done in several ways still reaching the same continuous limit as $k \rightarrow \infty$ and it is convenient to choose different choices for different constructions why we define the general construction. The most important thing is however that for large k immunity jumps in many small steps, each having a high jump rate. Let $\{r_{l,j}(t)\}_{j=1}^{k-1}$, $l \in \{1, 2\}$ be the fractions of recovered individuals, at time t , with the immunity levels $\{1 - f_{l,j}\}_{j=1}^{k-1}$ (or susceptibility levels $\{f_{l,j}\}_{j=1}^{k-1}$), $l \in \{1, 2\}$, and $\{c_{l,j}\}_{j=1}^k$, $l \in \{1, 2\}$ be the rates at which recovered individuals lose immunity portions through the k steps. Since k is fixed and typically large, we drop it from the notation. The resulting model equations are given by

$$\begin{aligned} s'_l(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) + c_{l,k} r_{l,k-1}(t) - \mu s_l(t), \\ i'_l(t) &= \beta \left(s_l(t) + \sum_{j=1}^{k-1} f_{l,j} r_{l,j}(t) \right) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \\ r'_{l,0}(t) &= \gamma i_l(t) - (c_{l,1} + \mu) r_{l,0}(t), \\ r'_{l,j}(t) &= c_{l,j} r_{l,j-1}(t) - \beta f_{l,j} r_{l,j}(t) (i_1(t) + i_2(t)) - (c_{l,j+1} + \mu) r_{l,j}(t), \end{aligned} \quad (3)$$

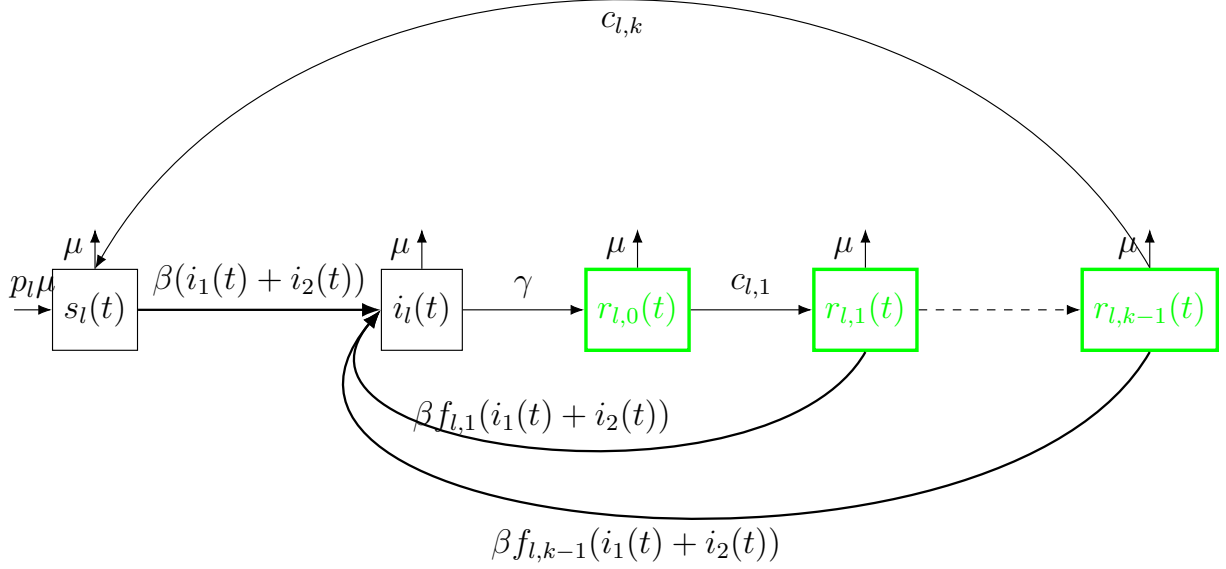


Figure 1: Diagram of the $\text{SIR}^{(k)}\text{S}$ epidemic model in the l -type individuals, $l = 1, 2$.

for $j = 1, \dots, k-1$ and $l \in \{1, 2\}$ where $p_1 = p$ and $p_2 = 1 - p$. We call this model the $\text{SIR}^{(k)}\text{S}$ model with heterogeneity. See Fig. 1 for a transition scheme and the Appendix A.2 for a derivation of the immunity jumps and the transition rates (see also El Khalifi and Britton (2023)).

4 $\text{SIR}^{(k)}\text{S}$ model with heterogeneity and vaccination

When immunity wanes over time, it is important to allow the vaccination strategies to depend on time since last vaccination and vaccines should not be uniformly distributed. Hence the vaccination rate in the j 'th susceptibility class of an l -type individuals might depend on both l and j . Let $\eta_{l,j}$ to denote this vaccination rate. A vaccination strategy is hence specified by these rates $\{\eta_{l,j}\}$, many rates often being 0 since strategies would often be defined by vaccinating once immunity drops to a certain level. Then, the $\text{SIR}^{(k)}\text{S}$ model with vaccination is given by the following equations

$$\begin{aligned}
 s'_l(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) + c_{l,k} r_{l,k-1}(t) - (\mu + \eta_{l,k}) s_l(t), \\
 i'_l(t) &= \beta \left(s_l(t) + \sum_{j=1}^{k-1} f_{l,j} r_{l,j}(t) \right) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \\
 r'_{l,0}(t) &= \eta_{l,k} s_l(t) + \sum_{j=1}^{k-1} \eta_{l,j} r_{l,j}(t) + \gamma i_l(t) - (c_{l,1} + \mu) r_{l,0}(t), \\
 r'_{l,j}(t) &= c_{l,j} r_{l,j-1}(t) - \beta f_{l,j} r_{l,j}(t) (i_1(t) + i_2(t)) - (c_{l,j+1} + \mu + \eta_{l,j}) r_{l,j}(t),
 \end{aligned} \tag{4}$$

for $j = 1, \dots, k-1$ and $l \in \{1, 2\}$ where $p_1 = p$ and $p_2 = 1 - p$.

What is the best, or optimal, vaccination strategy differs depending on amount of available information. In the situation where no information is available, a potential strategy could be to randomly vaccinate in all non-infectious classes, including individuals with partial immunity. Here we distinguish between two situations: both individual time since last vaccination and waning rates are known (informed situation), and only time since last vaccination is known (uninformed situation).

4.1 Informed situation

When the individual waning rate and time since last vaccination for all individuals are known, vaccines will be administered to those with faster waning at different frequency compared to individuals with slower waning. In this situation, we assume that the susceptibility levels $\{f_{l,j}\}_{j=1}^{k-1}$ are the same for both types and $f_{l,j} = \frac{j}{k}, j = 1, \dots, k-1$. Hence the transition rates $\{c_{l,j}\}_{j=1}^k, l \in \{1, 2\}$ (depend on l) are such that the average cumulative immunity equals $1/\omega_1$ and $1/\omega_2$ respectively (see Appendix A.2). For any $l \in \{1, 2\}$, we denote by $\eta_{l,j}$ the vaccination rate in the class $r_{l,j}$ for $j = 1, \dots, k-1$, and by $\eta_{l,k}$ the vaccination rate of fully susceptible individuals s_l . The resulting model equations are given by

$$\begin{aligned}
 s'_l(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) + c_{l,k} r_{l,k-1}(t) - (\mu + \eta_{l,k}) s_l(t), \\
 i'_l(t) &= \beta \left(s_l(t) + \sum_{j=1}^{k-1} \frac{j}{k} r_{l,j}(t) \right) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \\
 r'_{l,0}(t) &= \eta_{l,k} s_l(t) + \sum_{j=1}^{k-1} \eta_{l,j} r_{l,j}(t) + \gamma i_l(t) - (c_{l,1} + \mu) r_{l,0}(t), \\
 r'_{l,j}(t) &= c_{l,j} r_{l,j-1}(t) - \beta \frac{j}{k} r_{l,j}(t) (i_1(t) + i_2(t)) - (c_{l,j+1} + \mu + \eta_{l,j}) r_{l,j}(t),
 \end{aligned} \tag{5}$$

for $j = 1, \dots, k-1$, and $l \in \{1, 2\}$ where $p_1 = p$ and $p_2 = 1 - p$.

Here the immunity class of each individual is known. The best vaccination scheme is then to vaccinate 1-individuals once they have lost j_1 steps of immunity and 2-individuals once they have lost j_2 immunity steps, for some values of j_1 and j_2 (see Fig. 2a). This corresponds to vaccinating the two types of individuals at (possible different) fixed times, t_1 and t_2 respectively, since last vaccination (or infection). Clearly, the smaller j_1 and j_2 the more vaccines are needed, and the optimal relation between j_1 and j_2 will depend on the immunity waning rates ω_1 and ω_2 .

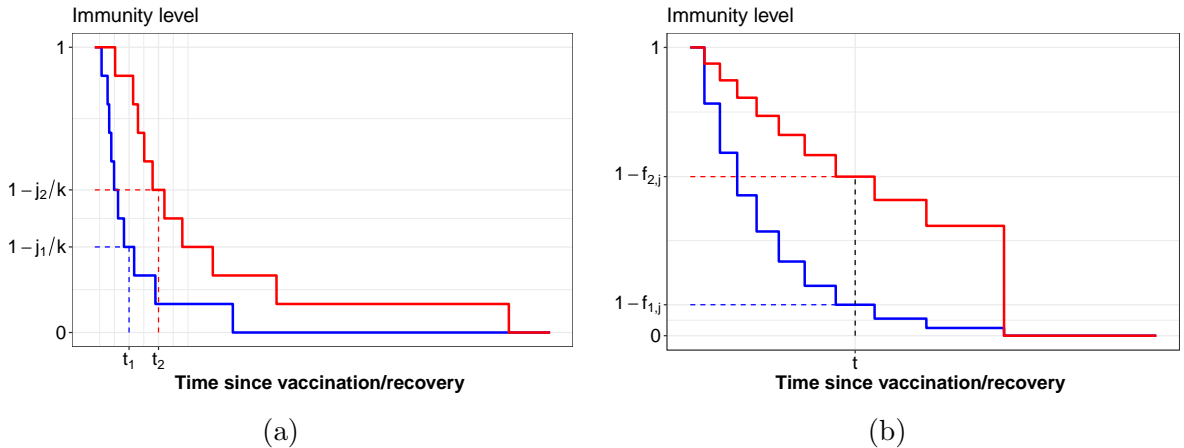


Figure 2: Examples showing the vaccination strategy when immunity wanes in $k = 10$ steps in (a) the informed situation (b) the uninformed situation.

4.2 Uninformed situation

Here we consider the more realistic situation where only time since last recovery/vaccination is known. In this situation, we let the transition rates $\{c_{l,j}\}_{j=1}^k, l \in \{1, 2\}$ to be independent on l , and then the susceptibilities $\{f_{l,j}\}_{j=1}^k$ are no longer the same for $l \in \{1, 2\}$ (see Appendix A.2). In addition, for both $l \in \{1, 2\}$ we let η_j to be the vaccination rate in the class $r_{l,j}$ for $j = 1, \dots, k-1$, and η_k to be the vaccination rate of fully susceptible individuals s_l

(the vaccination rate must be the same for both types in the uninformed situation). The resulting model equations are given by

$$\begin{aligned}
s'_l(t) &= p_l \mu - \beta s_l(t) (i_1(t) + i_2(t)) + c_k r_{l,k-1}(t) - (\mu + \eta_k) s_l(t), \\
i'_l(t) &= \beta \left(s_l(t) + \sum_{j=1}^{k-1} f_{l,j} r_{l,j}(t) \right) (i_1(t) + i_2(t)) - (\gamma + \mu) i_l(t), \\
r'_{l,0}(t) &= \eta_k s_l(t) + \sum_{j=1}^{k-1} \eta_j r_{l,j}(t) + \gamma i_l(t) - (c_1 + \mu) r_{l,0}(t), \\
r'_{l,j}(t) &= c_j r_{l,j-1}(t) - \beta f_{l,j} r_{l,j}(t) (i_1(t) + i_2(t)) - (c_{j+1} + \mu) r_{l,j}(t) - \eta_j r_{l,j}(t),
\end{aligned} \tag{6}$$

for $j = 1, \dots, k-1$, and $l \in \{1, 2\}$ where $p_1 = p$ and $p_2 = 1 - p$.

Here vaccination is the same for both types since they are unobserved, and all individuals are vaccinated (at the same time, t) once they have lost j steps of immunity (see Fig. 2b).

4.3 Extending to imperfect (leaky) vaccines

In the previous section, both infection and vaccination are assumed to initially confer perfect immunity. This could be relaxed by considering imperfect vaccines producing partial protection level to any vaccinated person. Although, these partial immunities could differ between the two subpopulations, we here consider a leaky vaccine conferring immunity e to all vaccinated individuals in both subpopulations.

4.4 Reproduction number and optimal vaccination

Recall that e is the protection level that vaccines are assumed to confer to any vaccinated individual. Each constant vaccination scheme gives rise to a disease free equilibrium $E_0 = (\hat{s}_1, \hat{s}_2, \hat{r}_{1,0}, \hat{r}_{2,0}, \dots, \hat{r}_{1,k-1}, \hat{r}_{2,k-1})$ (see Appendix A.3). The corresponding reproduction number R_v is given by

$$R_v = R_0 \sum_{l=1}^2 \left(\hat{s}_l + e \sum_{j=1}^{k-1} f_{l,j}^k \hat{r}_{l,j} \right). \tag{7}$$

In the expressions E_0 and R_v , we omit the dependence on the vaccination strategy (informed or uninformed) for the sake of convenience. Within a certain class of vaccination schemes, the optimal is the one solving the following optimization problem

$$\theta_c^k = \min_{\boldsymbol{\eta}} \theta^k(\boldsymbol{\eta}) \quad \text{subject to} \quad R_v \leq 1,$$

where $\theta^k(\boldsymbol{\eta})$ is the vaccine usage given by

$$\theta^k(\boldsymbol{\eta}) = \sum_{l=1}^2 \eta_{l,k} \hat{s}_l + \sum_{j=1}^{k-1} \eta_{l,j} \hat{r}_{l,j}, \tag{8}$$

for a $(2 \times k)$ -matrix of vaccination rates $\boldsymbol{\eta}$ within the class of possible vaccination schemes.

Informed optimal vaccination strategy

Within each type it is always better to vaccinate less immune individuals compared to more immune individuals. The optimal vaccination strategy in the informed situation is hence to vaccinate 1-individuals and 2-individuals as soon as their immunities drop below some levels $\iota_1 = 1 - j_1/k$ and $\iota_2 = 1 - j_2/k$, respectively, for some j_1 and j_2 . For finite

k and by referring to model (5), this is equivalent to not vaccinate type-1 individuals in states $(r_{1,0}, r_{1,1}, \dots, r_{1,j_1-1})$ (and $(r_{2,0}, r_{2,1}, \dots, r_{2,j_2-1})$ for type-2 individuals) up to some $j_1, j_2 \in \{1, \dots, k\}$, to vaccinate in r_{1,j_1} (resp. r_{1,j_2}) at some rate η_{1,j_1}^* (resp. η_{1,j_2}^*), and to immediately vaccinate individuals leaving the state r_{1,j_1} (resp. r_{1,j_2}). The states (j_1, j_2) and the rates $(\eta_{1,j_1}^*, \eta_{1,j_2}^*)$ correspond to the minimal immunity levels and the minimal vaccination rates respectively, satisfying $R_v \leq 1$. For large k this means we vaccinate type-1 and type-2 individuals once their immunities have dropped to $1 - j_1/k$ and $1 - j_2/k$ respectively. What are the optimal values of j_1 and j_2 for a given overall vaccination rate θ we solve numerically.

Uninformed optimal vaccination strategy

As only individual time since vaccination is known, the optimal vaccination strategy consists of vaccinating all individuals, irrespective of type, as soon as they reach some time t since their last vaccination. The shorter t , the bigger the vaccine coverage θ_c^k . For finite k and by referring to model (6), this is to not vaccinate up to some $j \in \{1, \dots, k\}$, to vaccinate in both $r_{1,j}$ and $r_{2,j}$ at some rate η_j^* , and immediately vaccinate individuals leaving the states $r_{1,j}$ and $r_{2,j}$. The state j and the rate η_j^* correspond to the minimal immunity levels and the minimal vaccination rate respectively, satisfying $R_v \leq 1$ and this we also solve numerically. At time t , individuals immunities are different and equal to $\iota_{u,1} = 1 - f_{1,j}$ and $\iota_{u,2} = 1 - f_{2,j}$ for type-1 individuals and type-2 individuals respectively.

Remark: While both informed and uninformed vaccination strategies will be considered when immunity wanes gradually, only the informed situation is considered when immunity wanes in one jump. Indeed, introducing an uninformed vaccination in the SIRS model necessitates the change of the distribution of immunity duration and probably using a PDE model for the dynamics of vaccinated/recovered individuals, something which we do not consider in this paper.

5 Results

We now illustrate our results numerically, studying the effect of heterogeneity of the gradual waning, for parameter values consistent with Covid-19 (of course lacking many other features of reality). Our primary focus is to study the effect of heterogeneity measured by its coefficient of variation σ , but also to compare the informed situation, which assumes the individual heterogeneities to be known, to the uninformed case. We also compare our model to the classical homogeneous SIRS epidemic model as well as to the heterogeneous SIRS (loosing all immunity at once).

To illustrate how various waning assumptions affect disease prevalence and the vaccination frequency needed to avoid an outbreak to occur, we use the following parameter values. The life expectancy is set to $\mu^{-1} = 80$ years, the mean infectious period is set to $\gamma^{-1} = 0.02$ years (one week) and the average cumulative immunity is to $\omega^{-1} = 1$ year. These parameter values are reasonable for several infectious diseases including COVID-19, influenza, common cold, etc Byrne et al. (2020); Davies et al. (2020); Hall et al. (2022); CDC (2022). Although the waning rate ω could be estimated using for instance the antibody decay data Goldberg et al. (2021); Bobrovitz et al. (2023), we do not attempt to do so here. The amount of waning heterogeneity is measured by the coefficient of variation σ of immunity heterogeneity, which is always smaller than $\sqrt{p/(1-p)}$. We vary β (or equivalently R_0) and σ (often with $p = 50\%$ fixed but sometimes also varying p). Using the different models we compare the

endemic prevalence levels without vaccination, and the required amount of vaccines to reach a sustainable herd immunity.

5.1 Endemic prevalence

Fig. 3 shows heatmaps of the endemic level for the heterogeneous SIRS model (1) and our new heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model (2), as functions of R_0 and σ (without vaccination). It can be seen from Figs. 3a-3b that the long term prevalence is increasing in R_0 (as expected) but also in population heterogeneity σ . Moreover, Fig. A.1 in the Appendix A shows that the difference between homogeneous and heterogeneous populations increases with p (the community fraction having the lower immunity, i.e. higher waning rate). The homogeneous models ($\sigma = 0$) have the lowest endemic levels irrespective of the immunity waning mode (sudden or gradual loss).

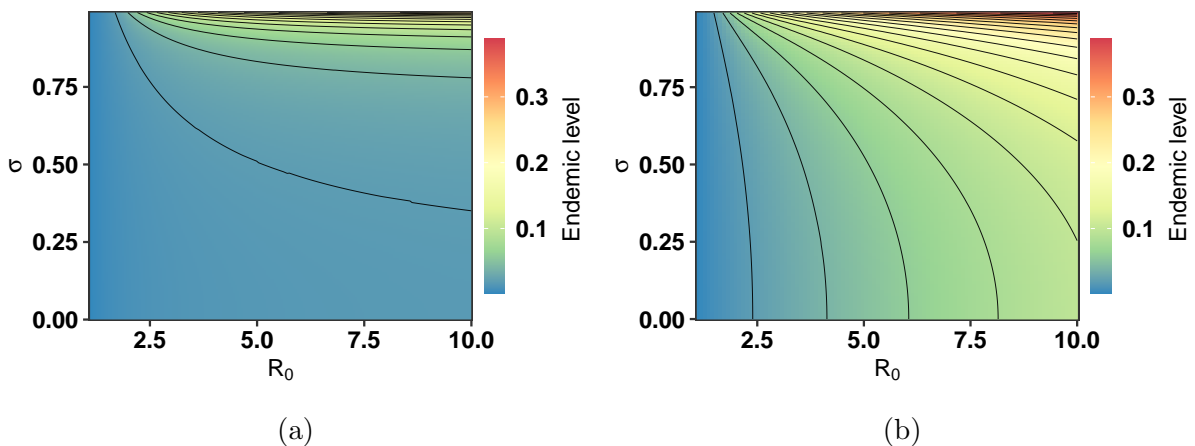
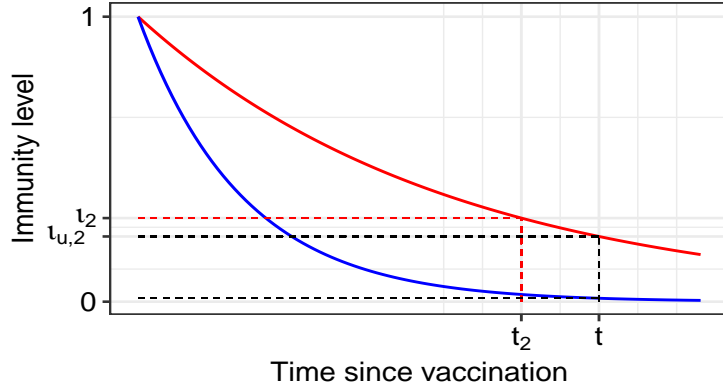


Figure 3: Heatmaps of the endemic level for different values of R_0 and σ ($p = 50\%$) in (a) the heterogeneous SIRS model and (b) the heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model. The value at the origin is 0 and the contour interval is 2% of the population.

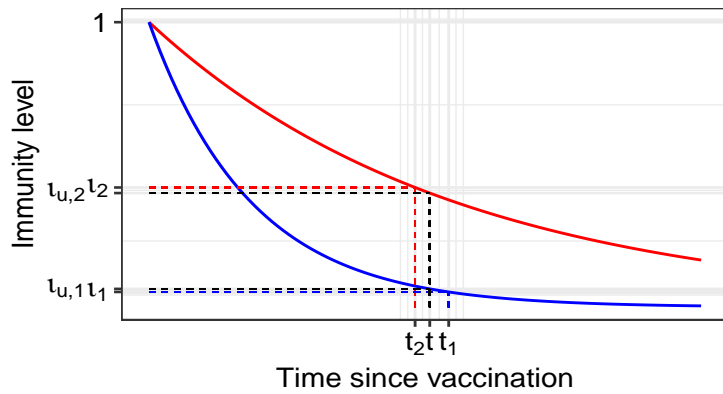
5.2 Optimal vaccination: Perfect vaccine

5.2.1 Possible vaccination times

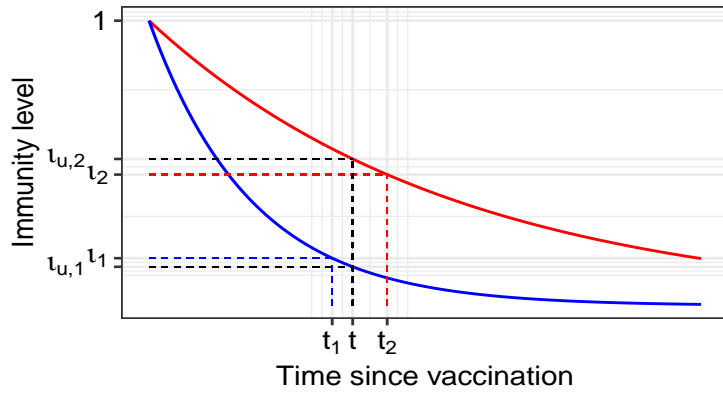
Fig. 4 shows the best vaccination strategies for the informed and uninformed situations. The optimal strategy always vaccinate type 1 (with higher waning rate) at a lower immunity level compared to type 2 individuals (both informed and uninformed). In the informed situation it may even be optimal to only vaccinate type 2 individuals, e.g. when R_0 is small enough (Fig. 4a) or if ω_1 is very large so these individuals lose their immunity very quickly implying that there is not much gain in vaccinating them. However, the time (since vaccination) at which we vaccinate type 1 could be bigger or smaller than the time for type 2. Plot 4a shows that only type-2 need to be vaccinated, at time t_2 , when $R_0 = 1.6$, but later on also type-1 have to be vaccinated, at time t_1 , when R_0 increases to 2 as shown in Fig. 4b. When $R_0 = 2.4$, the vaccination time t_1 of type-1 becomes smaller compared to t_2 for type-2 as seen in Fig. 4c. In the uninformed situation all individuals need to be vaccinated after the same time t since their last vaccination.



(a) $R_0 = 1.6$.



(b) $R_0 = 2$.



(c) $R_0 = 2.4$.

Figure 4: Informed and the uninformed optimal vaccination times for different values of R_0 with $\sigma = 0.5$ ($p = 50\%$). Blue and Red solid curves are immunity waning functions of type-1 and type-2 individuals respectively. The best informed vaccination strategy is to vaccinate 1-individuals at time t_1 (with the immunity level l_1) and 2-individuals at time t_2 (with the immunity level l_2). The best uninformed vaccination strategy is to vaccinate everyone at time t , that is, 1-individuals and 2-individuals at the immunity levels $l_{u,1}$ and $l_{u,2}$ respectively.

5.2.2 Optimal vaccination scheme

Fig. 5 shows the minimum number of vaccine doses per person per year to achieve and maintain herd immunity according to the heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model. It is evident from the plots that the critical amount of vaccine supply in the continuous waning situation (for fixed p) is increasing in the coefficient of variation of population heterogeneity σ . Moreover, the bigger the fraction p (of immune-weak type 1 individuals), the bigger the critical amount of vaccine supply. This indicates that heterogeneity in population immunity requires more frequent vaccination. It is worth mentioning that the optimal vaccine supply is not always increasing in heterogeneity when immunity wanes in one sudden leap as illustrated in Fig. A.2 in the Appendix.

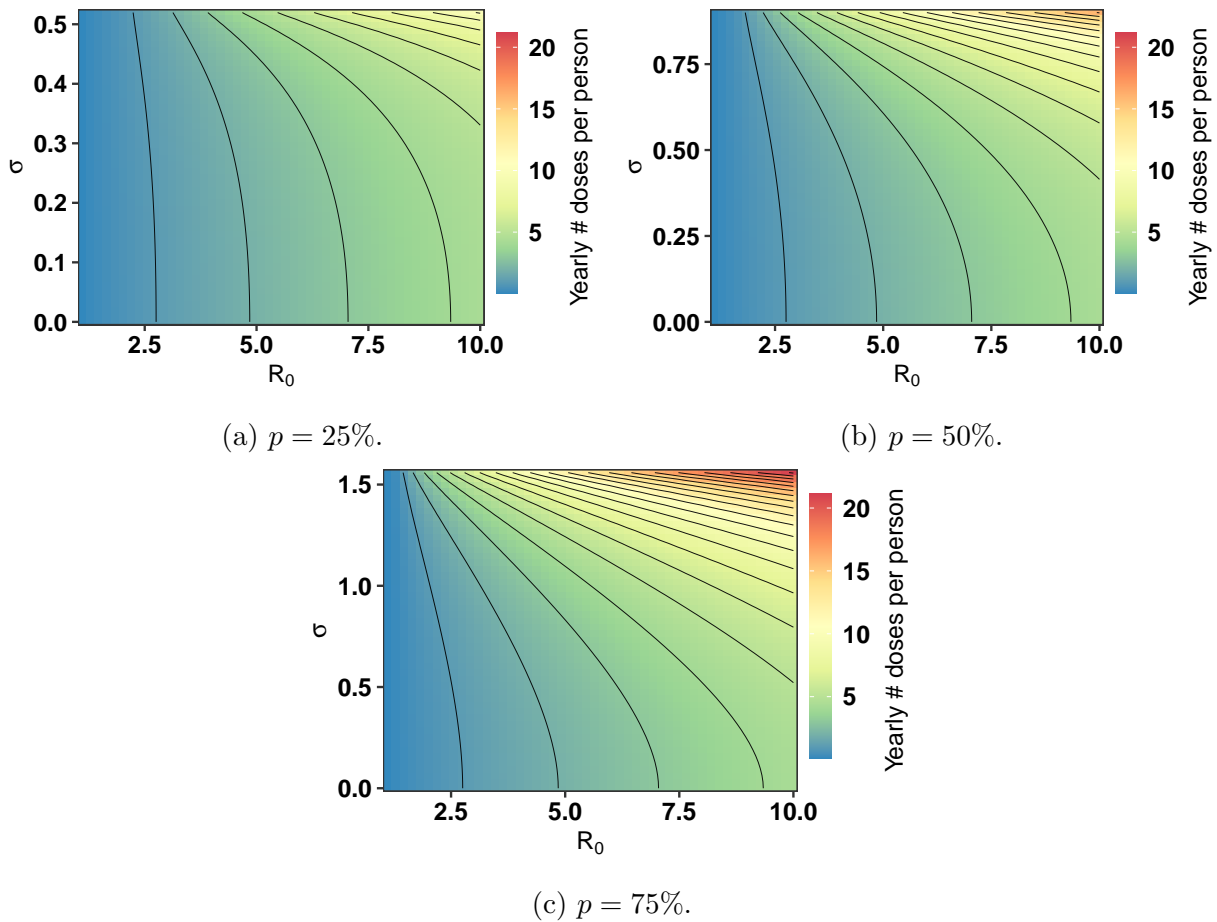


Figure 5: Heatmaps of the critical vaccine supply for different values of p in the uninformed scenario of the $\text{SIR}^{(\infty)}\text{S}$ model. The value at the origin is 0 and the contour interval is 1 yearly dose per person.

Table 2 compares the critical vaccination frequency for different models when $R_0 = 5$ (and $\sigma = 0.5$ for the heterogeneous models). While the simple SIRS model suggests to vaccinate individuals every 15 months (0.81 doses per year) to maintain herd immunity, the heterogeneous exponentially waning immunity model increases this vaccination frequency to every ≈ 4.6 months in the informed situation (≈ 2.62 doses per person per year), and to ≈ 4.4 months in the more realistic uninformed situation (≈ 2.76 doses per person per year). Table 3 compares the value of R_v for a given vaccine supply per year and shows that knowing individuals immunity status reduce the effective reproduction number compared to the uninformed situation, the difference is however moderate.

Table 2: Critical vaccination schemes for $R_0 = 5$ for the different models. The heterogeneous models are computed with $\sigma = 0.5$ and $p = 50\%$.

	Vaccination frequency (in months)	Yearly doses per person	# per
Hom. SIRS	15	0.81	
Het. SIRS: informed	12.8 (10 / 17.7) ¹	0.94	
Hom. exp. waning	5.5	2.15	
Het. exp. waning: informed	4.6 (3.8 / 5.7)	2.62	
Het. exp. waning: uninformed	4.4	2.76	

¹12.8 (10 / 17.7) means that vaccines are given to type-1 individuals every 10 months and type-2 every 17.7 months, resulting in vaccinating everyone every 12.8 months on average.

Table 3: Reproduction number for different values of individual vaccine supply per year given $R_0 = 5$ and $\sigma = 0.5$ ($p = 50\%$) under exponential waning of immunity.

Yearly # doses per person	$\theta = 0$	$\theta = 1/2$	$\theta = 1$	$\theta = 2$	$\theta = 3$
Informed	5	2.99	2.08	1.26	0.90
Uninformed	5	3.01	2.09	1.29	0.93

5.3 Optimal vaccination: Leaky vaccine

Table 4 compares the optimal vaccination frequency in case of a leaky vaccine for the considered models. It is clear from the table that the optimal vaccination frequency increases as the protection e becomes smaller. While herd immunity could be achieved with imperfect vaccines with relatively high efficacy when immunity wanes at once (e.g. by approximately administering 80%-effective vaccines every 8.9 months on average – Table 4), herd immunity under continuous waning would require very high vaccine efficacy and that in both homogeneous and heterogeneous situations.

5.4 Two extreme models comparison

In El Khalifi and Britton (2023), the standard SIRS model and the homogeneous $\text{SIR}^{(\infty)}\text{S}$ model were compared and found that the latter has the larger endemic level and the higher critical vaccine supply. Fig. 6 added a comparison with the heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model, with the critical vaccine supply plotted under the uninformed situation. It is clear that the biggest effect comes from the continuous waning of immunity compared to the sudden loss assumption. Still, heterogeneity makes the situation worse as it increases long-term prevalence and the critical vaccine coverage, in particular when heterogeneity is substantial.

Table 4: Critical vaccination frequency (in months) for the different models when $R_0 = 5$ ($\sigma = 0.5$ and $p = 50\%$ in heterogeneous settings).

Vaccine efficacy	$e = 100\%$ (perfect)	$e = 95\%$	$e = 90\%$	$e = 80\%$
SIRS	15	14	13.2	11.8
Het. SIRS: informed	12.8 (10 / 17.7) ¹	11.7 (8.7 / 17.7)	10.7 (7.7 / 17.7)	8.9 (6 / 17.7)
Hom. exp. waning	5.5	4.5	3.3	–
Het. exp. waning: informed	4.6 (3.8 / 5.7)	3.7 (2.7 / 4.8)	2.6 (1.8 / 3.4)	–
Het. exp. waning: uninformed	4.4	3.5	2.5	–

¹12.8 (10 / 17.7) means that vaccines are given to type-1 individuals every 10 months and type-2 every 17.7 months, resulting in vaccinating individuals every 12.8 months on average.

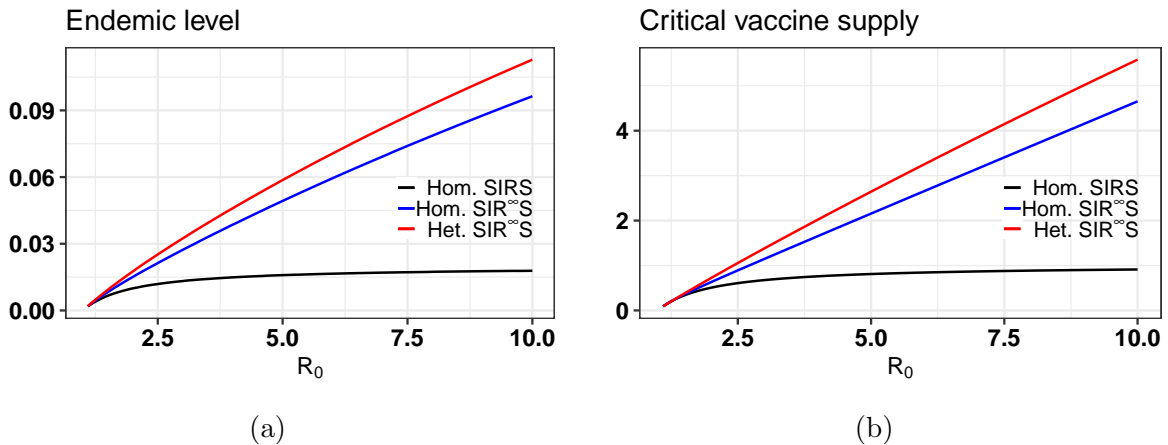


Figure 6: Comparison of the standard SIRS model, the homogeneous $\text{SIR}^{(\infty)}\text{S}$ model, and the heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model with $\sigma = 0.5$ ($p = 50\%$). (a) endemic levels and (b) the yearly number of vaccine doses per person required for herd immunity.

6 Discussion

In the current paper we have shown that if immunity wanes gradually but at different rates for different individuals, the effect of such heterogeneity is that endemic prevalence becomes higher, and when introducing vaccinations, more vaccines are required to reach and sustain herd immunity. This effect is shown to be substantial even when heterogeneity of immunity waning is moderate (e.g. coefficient of variation 0.5). An additional feature treated in our analysis is to distinguish between the informed situation where the waning heterogeneity is known and taken into account when designing vaccination policies, and the more likely uninformed scenario where such heterogeneities are unobserved. It is shown that the informed and uninformed situations differ in vaccination policies, but the required amount of vaccines for maintaining herd immunity is only moderately higher for the more likely uninformed situation.

This comparison between the homogeneous and heterogeneous situations, calibrated by

assuming the same population average of cumulative immunity, hence has the heterogeneous situation as the worse case. As a consequence, models neglecting waning heterogeneities can estimate too low vaccination rates. This result is in contrast with many other comparisons in epidemic models in which the homogeneous situation is often the worst case scenario. Two such examples are Ball (1985) who considers variable susceptibility to the homogeneous situation where all individuals have the same susceptibility (see also Elbasha and Gumel (2021)), and the second example is epidemics on networks where the final size is maximized when all individuals have equal degree (if the transmission rate is large enough) Britton and Trapman (2012).

Our new epidemic model with gradual waning rate with individual heterogeneity neglects many other factors affecting diseases dynamics. Such factors may for example include demographic structure, behaviour change of the population, elements of chance, individual heterogeneity also with respect to infectivity and susceptibility, social population structures, and so on. Here we neglect such aspects and focus on heterogeneity of immunity waning. It would be of interest to study the effect of waning heterogeneity also when including other realistic model features. It is our belief that the same qualitative observation remains: heterogeneity in immunity waning makes the situation worse, but clearly this needs to be shown.

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Ethics declarations

The authors declare no competing interests.

Data availability

All data generated or analysed during this study are included in this published article and its supplementary information files.

References

- Ball, F. (1985). Deterministic and stochastic epidemics with several kinds of susceptibles. *Advances in applied probability*, 17(1):1–22.
- Bobrovitz, N., Ware, H., Ma, X., Li, Z., Hosseini, R., Cao, C., Selemon, A., Whelan, M., Premji, Z., Issa, H., et al. (2023). Protective effectiveness of previous sars-cov-2 infection and hybrid immunity against the omicron variant and severe disease: a systematic review and meta-regression. *The Lancet Infectious Diseases*.

- Britton, T. and Trapman, P. (2012). Maximizing the size of the giant. *Journal of Applied Probability*, 49(4):1156–1165.
- Byrne, A. W., McEvoy, D., Collins, A. B., Hunt, K., Casey, M., Barber, A., Butler, F., Griffin, J., Lane, E. A., McAloon, C., et al. (2020). Inferred duration of infectious period of SARS-CoV-2: rapid scoping review and analysis of available evidence for asymptomatic and symptomatic COVID-19 cases. *BMJ open*, 10(8):e039856.
- CDC (2022). How flu spreads.
- Davies, N. G., Klepac, P., Liu, Y., Prem, K., Jit, M., and Eggo, R. M. (2020). Age-dependent effects in the transmission and control of COVID-19 epidemics. *Nature medicine*, 26(8):1205–1211.
- El Khalifi, M. and Britton, T. (2023). Extending sirs epidemics to allow for gradual waning of immunity. *Journal of the Royal Society Interface*, 20(206).
- Elbasha, E. H. and Gumel, A. B. (2021). Vaccination and herd immunity thresholds in heterogeneous populations. *Journal of mathematical biology*, 83(6-7):73.
- Fabiani, M., Puopolo, M., Morciano, C., Spuri, M., Alegiani, S. S., Filia, A., DâAncona, F., Del Manso, M., Riccardo, F., Tallon, M., et al. (2022). Effectiveness of mRNA vaccines and waning of protection against sars-cov-2 infection and severe covid-19 during predominant circulation of the delta variant in italy: retrospective cohort study. *bmj*, 376.
- Forien, R., Pang, G., Pardoux, É., et al. (2022). Stochastic epidemic models with varying infectivity and susceptibility. *arXiv preprint arXiv:2210.04667*.
- Goldberg, Y., Mandel, M., Bar-On, Y. M., Bodenheimer, O., Freedman, L., Haas, E. J., Milo, R., Alroy-Preis, S., Ash, N., and Huppert, A. (2021). Waning immunity after the BNT162b2 vaccine in Israel. *New England Journal of Medicine*, 385(24):e85.
- Hall, V., Foulkes, S., Insalata, F., Kirwan, P., Saei, A., Atti, A., Wellington, E., Khawam, J., Munro, K., Cole, M., et al. (2022). Protection against SARS-CoV-2 after Covid-19 vaccination and previous infection. *New England Journal of Medicine*, 386(13):1207–1220.
- Hethcote, H. W. (1976). Qualitative analyses of communicable disease models. *Mathematical biosciences*, 28(3-4):335–356.
- Hethcote, H. W. (1978). An immunization model for a heterogeneous population. *Theoretical population biology*, 14(3):338–349.
- Martcheva, M. (2015). *An introduction to mathematical epidemiology*, volume 61. Springer.
- Pérez-Alós, L., Armenteros, J. J. A., Madsen, J. R., Hansen, C. B., Jarlhelt, I., Hamm, S. R., Heftdal, L. D., Pries-Heje, M. M., Møller, D. L., Fogh, K., et al. (2022). Modeling of waning immunity after sars-cov-2 vaccination and influencing factors. *Nature communications*, 13(1):1614.
- Reluga, T. C., Medlock, J., and Perelson, A. S. (2008). Backward bifurcations and multiple equilibria in epidemic models with structured immunity. *Journal of theoretical biology*, 252(1):155–165.
- Shrotri, M., Navaratnam, A. M., Nguyen, V., Byrne, T., Geismar, C., Fragaszy, E., Beale, S., Fong, W. L. E., Patel, P., Kovar, J., et al. (2021). Spike-antibody waning after second dose of bnt162b2 or chadox1. *The Lancet*, 398(10298):385–387.
- Wheatley, A. K., Juno, J. A., Wang, J. J., Selva, K. J., Reynaldi, A., Tan, H.-X., Lee, W. S., Wragg, K. M., Kelly, H. G., Esterbauer, R., et al. (2021). Evolution of immune responses to sars-cov-2 in mild-moderate covid-19. *Nature communications*, 12(1):1162.
- Widge, A. T., Roupshael, N. G., Jackson, L. A., Anderson, E. J., Roberts, P. C., Makhene, M., Chappell, J. D., Denison, M. R., Stevens, L. J., Pruijssers, A. J., et al. (2021). Durability of responses after sars-cov-2 mRNA-1273 vaccination. *New England Journal of Medicine*, 384(1):80–82.

A Appendix

A.1 Proof

A.1.1 Proposition 2.1

First, it is easy to see that only the disease free equilibrium exists when $R_0 < 1$. Next, we assume that $R_0 > 1$ and let $\hat{i} = \hat{i}_1 + \hat{i}_2$ to denote the endemic level. By equating the right hand side equations of (1) to 0 and after some simplifications, we get into

$$\hat{i} = \frac{pR_0(\mu + \gamma)(\mu + \omega_1)\hat{i}}{(R_0(\mu + \gamma)\hat{i} + \mu)(\mu + \gamma + \omega_1) + \omega_1\gamma} + \frac{(1-p)R_0(\mu + \gamma)(\mu + \omega_2)\hat{i}}{(R_0(\mu + \gamma)\hat{i} + \mu)(\mu + \gamma + \omega_2) + \omega_2\gamma}, \quad (9)$$

where we replaced β by $R_0(\mu + \gamma)$. That is, \hat{i} is the (positive) fixed point of the function ψ defined by

$$\psi(x) = \frac{pR_0(\mu + \gamma)(\mu + \omega_1)x}{(R_0(\mu + \gamma)x + \mu)(\mu + \gamma + \omega_1) + \omega_1\gamma} + \frac{(1-p)R_0(\mu + \gamma)(\mu + \omega_2)x}{(R_0(\mu + \gamma)x + \mu)(\mu + \gamma + \omega_2) + \omega_2\gamma},$$

which is increasing on the positive real half line and verifies $\lim_{x \rightarrow \infty} \psi(x) < 1$. Moreover, it can be shown that its derivative at $x = 0$ satisfies $\psi'(0) > 1$ as long as $R_0 > 1$ (and equals to 1 when $R_0 = 1$). Hence, ψ has a unique positive fixed point \hat{i} , the endemic level, provided that $R_0 > 1$. Consequently, the equation (1) has a unique endemic equilibrium if and only if $R_0 > 1$.

A.1.2 Proposition 2.2

Now, we recall that $\omega_1 = \omega / (1 - \sigma\sqrt{(1-p)/p})$ and $\omega_2 = \omega / (1 + \sigma\sqrt{p/(1-p)})$ with $0 \leq \sigma < \sqrt{p/(1-p)}$. To prove that the endemic level is increasing in σ , it is enough to show that the right hand side Eq. (9) is increasing in σ . This function could be defined by

$$f(\sigma) = \frac{p(\mu + \omega) - \sigma\mu\sqrt{p(1-p)}}{(\beta i + \mu)(\mu + \gamma) + \omega(\beta i + \mu + \gamma) - \sigma\sqrt{(1-p)/p}(\beta i + \mu)(\mu + \gamma)} + \frac{(1-p)(\mu + \omega) + \sigma\mu\sqrt{p(1-p)}}{(\beta i + \mu)(\mu + \gamma) + \omega(\beta i + \mu + \gamma) + \sigma\sqrt{p/(1-p)}(\beta i + \mu)(\mu + \gamma)},$$

Then, by differentiating, we obtain that the sign of $f'(\sigma)$ is the same as the sign of

$$2((\beta i + \mu)(\mu + \gamma) + \omega(\beta i + \mu + \gamma)) + \sigma(\beta i + \mu)(\mu + \gamma) \left(\sqrt{\frac{p}{1-p}} - \sqrt{\frac{1-p}{p}} \right),$$

which is positive provided that $p \geq 1/2$. Hence, f is increasing in σ , and so is the endemic level.

A.1.3 Proposition 2.3

From the infective equations in Eq. (2), and using the fact that

$$s_1(t) + s_2(t) + \int_0^t r_1(\tau) d\tau + \int_0^t r_2(\tau) d\tau = 1 - (i_1(t) + i_2(t)),$$

the differential equation of the total infective fraction $i = i_1 + i_2$ verifies

$$i'(t) \leq \beta i(t) (-i(t) + 1 - 1/R_0). \quad (10)$$

For any positive initial point z_0 , the solution to the ODE $z'(t) = \beta z(t) (-z(t) + 1 - 1/R_0)$ converges to 0 when $R_0 \leq 1$. Then from the Ineq. (10), we obtain that $i(t) \rightarrow 0$ as $t \rightarrow \infty$ when $R_0 \leq 1$. Hence, the infective and recovered fractions vanish. Moreover, the total susceptible fraction $s_1 + s_2$ converges to 1. This proves the first assertion of the Proposition 2.3.

Now, we proceed to prove the second assertion of Proposition 2.3. Solving the endemic equilibrium of system (2) allows to write

$$\frac{\mu}{2} - \beta s_l(i_1 + i_2) - \mu s_l = 0, \quad (11)$$

$$\frac{\partial r_l(a)}{\partial a} = -\beta (1 - e^{-\omega a}) r_l(a)(i_1 + i_2) - \mu r_l(a), \quad r_l(0) = \gamma i_l, \quad (12)$$

for $l \in \{1, 2\}$, coupled with

$$i_1 = p - s_1 - \int_0^\infty r_1(\tau) d\tau, \quad \text{and} \quad i_2 = 1 - p - s_2 - \int_0^\infty r_2(\tau) d\tau. \quad (13)$$

It is easy to see that an endemic equilibrium verifies both $i_1 \neq 0$ and $i_2 \neq 0$. Set $i = i_1 + i_2$, then solving the ordinary differential equation for the recovered equations, we obtain

$$r_l(\tau) = \gamma i_l \phi_l(i), \quad l \in \{1, 2\}, \quad (14)$$

where $\phi_l, l = 1, 2$, are the functions defined by

$$\phi_l(x) = \int_0^\infty \exp\left(-\mu\tau - \beta x \int_0^\tau (1 - e^{-\omega a}) da\right) d\tau. \quad (15)$$

Then, we arrived to

$$\begin{aligned} i_1 &= p \left(1 - \frac{\mu}{\beta i + \mu} - \gamma i_1 \phi_1(i)\right), \\ i_2 &= (1 - p) \left(1 - \frac{\mu}{\beta i + \mu} - \gamma i_2 \phi_2(i)\right). \end{aligned} \quad (16)$$

Re-arranging both equations allows to write i_1 and i_2 in terms of i as

$$\begin{aligned} i_1 &= \frac{\beta i}{\beta i + \mu} \frac{p}{1 + \gamma \phi_1(i)}, \\ i_2 &= \frac{\beta i}{\beta i + \mu} \frac{1 - p}{1 + \gamma \phi_2(i)}. \end{aligned} \quad (17)$$

Taking the sum, it yields that

$$i = \frac{\beta i}{\beta i + \mu} \left(\frac{p}{1 + \gamma \phi_1(i)} + \frac{1 - p}{1 + \gamma \phi_2(i)} \right). \quad (18)$$

As $i \neq 0$, we cancel one i and get to the following equation

$$i = \left(\frac{p}{1 + \gamma \phi_1(i)} + \frac{1 - p}{1 + \gamma \phi_2(i)} \right) - \frac{\mu}{\beta}. \quad (19)$$

The right-hand side of (19) is increasing in i and smaller than $1 - \mu/\beta$. Moreover, it converges to $\mu(R_0 - 1)/\beta$ as $i \rightarrow 0$. On the other hand, as $\phi_l, l = 1, 2$, are convex functions, the right-hand side of (19) is a concave down function. That is, Eq. (19) has a unique positive solution when $R_0 > 1$ and no positive solution when $R_0 \leq 1$. Since the right-hand side functions of (17) are increasing in i , the steady points i_1 and i_2 are defined from i uniquely. This completes the proof.

A.2 Formulation of the SIR^(k)S model with heterogeneity

Here we present the details of the determination of the immunity levels $\{1 - f_{l,j}\}_{j=1}^{k-1}$ and the immunity jumps rates $\{c_{l,j}\}_{j=1}^k$ in the model (3) in the main text for fixed $l \in \{1, 2\}$. Let $\{f_{l,j}\}_{j=1}^{k-1}$ be an increasing sequence of elements of $(0, 1)$. An l -individual recently recovered stays perfectly immune for an exponentially time with mean duration $1/c_{l,1}$, after that immunity drops to $1 - f_{l,1}$. Each $1 - f_{l,j}$ immunity level lasts for an exponentially time with mean $1/c_{l,j+1}$ for $j = 1, \dots, k-1$. The susceptibility levels and the rates are chosen to fit the exponential waning with rate w_l and to satisfy the constant average cumulative immunity equation

$$\frac{1}{c_{l,1}} + \sum_{j=1}^{k-1} (1 - f_{l,j}) \frac{1}{c_{l,j+1}} = \frac{1}{w_l}. \quad (20)$$

We mention that there is no unique way to define the immunity jumps and the rates above, yet their choice would not affect the results for typically large k as long as all jumps become small and rates large. In the informed situation, we choose $f_{l,j} = j/k$ so the immunity jumps by $1/k$ each step. We then define the rates by

$$\frac{1}{c_{l,1}} = -\frac{1}{w_l} \log \left(\frac{k-1}{k} + \frac{x}{k} \right), \quad (21)$$

$$\frac{1}{c_{l,j}} = \frac{1}{w_l} \left(-\log \left(\frac{k-j}{k} + \frac{x}{k} \right) + \log \left(\frac{k-j+1}{k} + \frac{x}{k} \right) \right), \quad j = 2, \dots, k, \quad (22)$$

where $x \in (0, 1)$ solves the cumulative immunity equation (20), that is, the equation

$$-\log \left(\frac{k-1}{k} + \frac{x}{k} \right) + \sum_{j=1}^{k-1} \frac{k-j}{k} \left(-\log \left(\frac{k-j}{k} + \frac{x}{k} \right) + \log \left(\frac{k-j+1}{k} + \frac{x}{k} \right) \right) = 1. \quad (23)$$

In the uninformed situation we assume fixed (in l) immunity jump rates, that is $c_{1,j} = c_{2,j} = c_j$ and set $c_j = k - j + 1$. We define the susceptibility levels by

$$f_{l,j} = 1 - \exp \left(-\omega_l (1 + x_l) \sum_{n=1}^j \frac{1}{c_n} \right), \quad (24)$$

with $x_l \in (0, 1)$ is the solution to the the cumulative immunity equation (20), that is, the equation

$$\sum_{j=0}^{k-1} \frac{1}{k-j} \exp \left(-\omega_l (1 + x_l) \sum_{n=1}^j \frac{1}{c_n} \right) = \frac{1}{\omega_l}. \quad (25)$$

A.3 Disease-free equilibria

Informed situation: The disease-free equilibrium $E_0^{inf} = (\hat{s}_1, \hat{s}_2, \hat{r}_{1,0}, \hat{r}_{2,0}, \dots, \hat{r}_{1,k-1}, \hat{r}_{2,k-1})$ of the models (5) is given by

$$\hat{s}_l = \frac{p_l \mu}{\mu + \eta_{l,k} \left(1 - c_{l,k}^k A_k^l B_{k-1}^l \right)}, \quad l = 1, 2, \quad (26)$$

$$\hat{r}_{l,j} = \eta_{l,k} A_k^l B_j^l \hat{s}_l, \quad j = 1, \dots, k-1, \quad l = 1, 2, \quad (27)$$

$$\hat{r}_{l,0} = \frac{1}{2} - \hat{s}_l - \sum_{j=1}^{k-1} \hat{r}_{l,j}, \quad l = 1, 2, \quad (28)$$

where $A_k^l = \left(\mu + c_{l,1} - \sum_{j=1}^{k-1} \eta_{l,j} B_j^l \right)^{-1}$ and $B_j^l = \prod_{n=1}^j \frac{c_{l,n}}{\mu + c_{l,n+1} + \eta_{l,n}}$, for $j = 1, \dots, k-1$ and $l = 1, 2$.

Uninformed situation: The disease-free equilibrium $E_0^{uni} = (\hat{s}_1, \hat{s}_2, \hat{r}_{1,0}, \hat{r}_{2,0}, \dots, \hat{r}_{1,k-1}, \hat{r}_{2,k-1})$ of the models (6) is given by

$$\hat{s}_l = \frac{p_l \mu}{\mu + \eta_k \left(1 - c_k^k A_k B_{k-1} \right)}, \quad l = 1, 2, \quad (29)$$

$$\hat{r}_{l,j} = \eta_k A_k B_j \hat{s}_l, \quad j = 1, \dots, k-1, \quad l = 1, 2, \quad (30)$$

$$\hat{r}_{l,0} = \frac{1}{2} - \hat{s}_l - \sum_{j=1}^{k-1} \hat{r}_{l,j}, \quad l = 1, 2, \quad (31)$$

where $A_k = \left(\mu + c_1 - \sum_{j=1}^{k-1} \eta_j B_j \right)^{-1}$ and $B_j = \prod_{n=1}^j \frac{c_n}{\mu + c_{n+1} + \eta_n}$, for $j = 1, \dots, k-1$.

A.4 Endemic level: varying p

Fig. A.1 shows how the endemic level varies with R_0 and σ for different values of p for the heterogeneous SIRS model (sudden loss of immunity) and the heterogeneous $\text{SIR}^{(\infty)}$ S model (continuous waning).

A.5 Critical vaccine supply when immunity wanes in one sudden leap

Fig. A.2 plots the critical vaccine supply for different values of p for the informed case of the heterogeneous SIRS model with sudden loss of immunity.

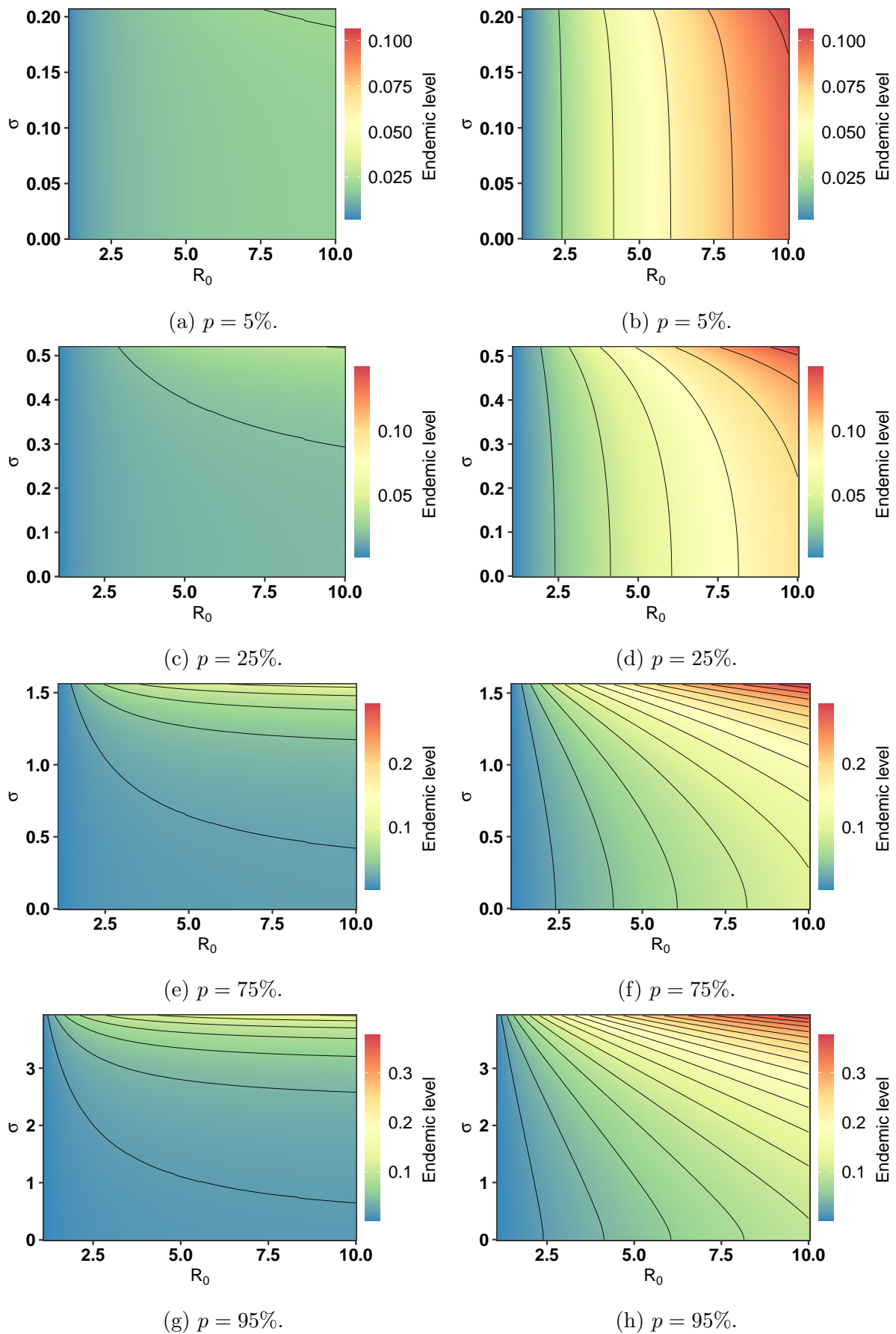


Figure A.1: Heatmaps of the endemic level for different values of R_0 and σ . Left hand panel: Heterogeneous SIRS model. Right hand panel: Heterogeneous $\text{SIR}^{(\infty)}\text{S}$ model. The value at the origin is 0 and the contour interval is 2% of the population.

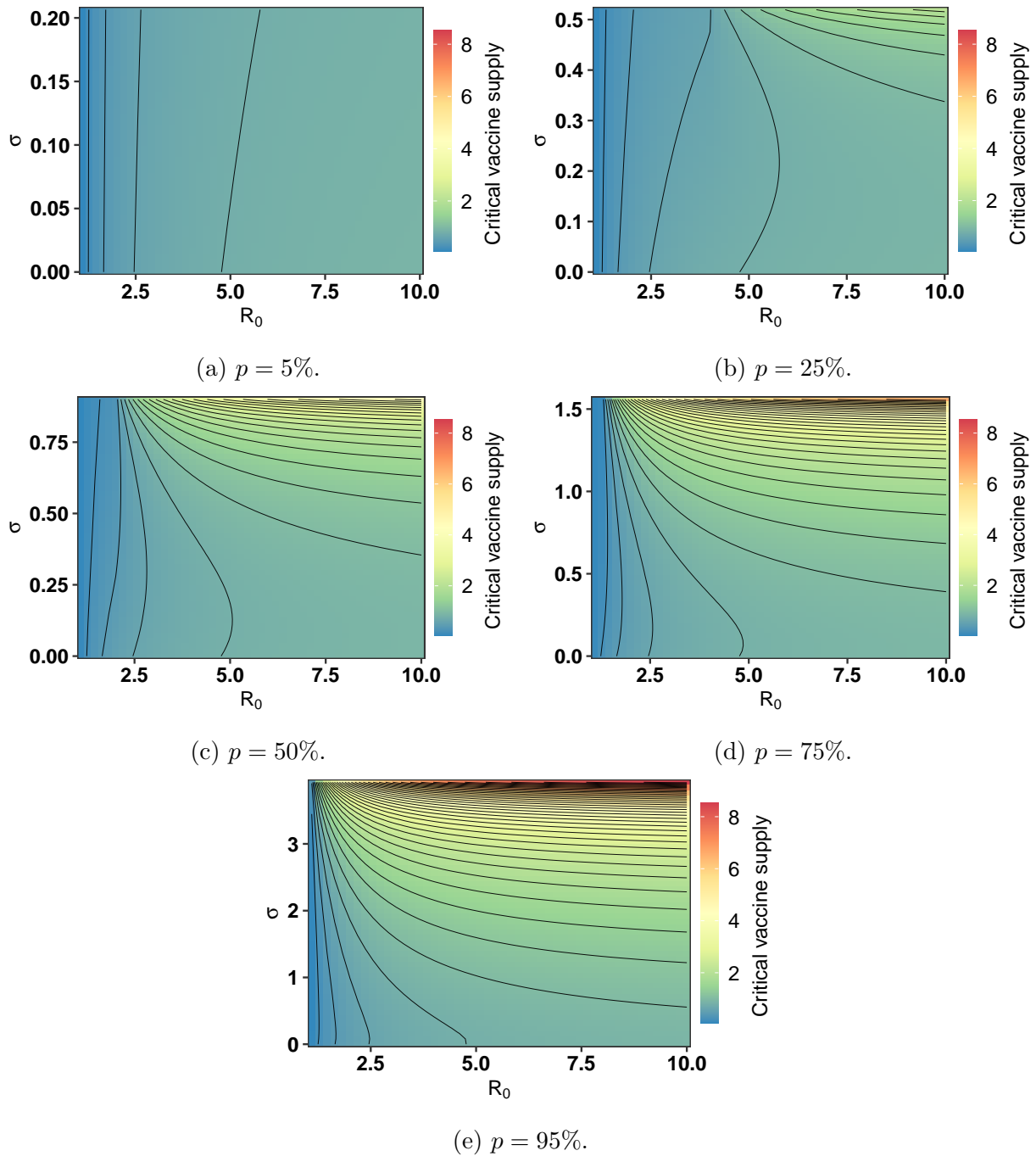


Figure A.2: Heatmaps of the critical vaccine supply for different values of p in the informed SIRS model with sudden loss of immunity. The value at the origin is 0 and the contour interval is 1 yearly dose per person.