

1070 **Supplementary information**

1071 **SIR+ models: Accounting for interaction-dependent disease suscepti-**
1072 **bility in the planning of public health interventions**

1073 *Maria M. Martignoni, Aura Raulo, Omer Linkovski, Oren Kolodny*

1074 **Contents**

1075	1 Introduction	3
1076	2 Model and Methods	4
1077	2.1 The SIR+ model	4
1078	2.2 Processes driving changes interaction-dependent health benefits	5
1079	2.2.1 Acquisition of interaction-dependent health benefits	5
1080	2.2.2 Loss of interaction-dependent health benefits	7
1081	2.3 Processes determining infection transmission and disease severity	8
1082	2.3.1 Infection resistance and the $\alpha(H)$ function	8
1083	2.3.2 Disease tolerance and the $p(H)$ function	9
1084	2.4 The role of the contact rate C and scenarios of interest	10
1085	3 Results	11
1086	4 Discussion	13
1087	5 Concluding remarks	17
1088	A Default parameters	32
1089	B Function $p(H)$	33
1090	C Acquisition and loss of interaction-dependent health benefits over time	33
1091	D Results obtained for linear $\alpha(H)$ and $p(H)$, and for $\alpha_{min}, p_{min} = 0$	36

A Default parameters

Table A.1: Variables and default parameter values chosen for the simulations.

Symbol	Description	Unit	Default value
S	Proportion of susceptible population	–	–
I	Proportion of infected population	–	–
R	Proportion of recovered population	–	–
H	Average interaction-dependent health benefits of a population	Health benefits	–
C	Contact rate	Contact/time	0-30
k	Fractional contribution of each contact to an increase in H	–	0.25, 0.5, 0.75, 1
h	Maximal interaction-dependent health benefits acquired from each contact	Health benefits/contact	0.1
γ	Recovery rate	1/time	1/7
η	Recovery rate (severe cases)	1/time	1/10
δ	Rate of loss of interaction-dependent benefit	1/time	0.1
α_{max}	Probability of infection transmission given a contact (low H)	1/contact	0.04
α_{min}	Probability of infection transmission given a contact (high H)	1/contact	0.02
p_{max}	Probability of severe illness from infection (low H)	–	blue 0.05
p_{min}	Probability of severe illness from infection (high H)	–	0.03
p_I	Adjusted contribution of infected individuals to the acquisition of interaction-dependent health benefits	–	0.6
p_R	Adjusted contribution of recovered individuals to the acquisition of interaction-dependent health benefits	–	0.8
H_{max}	Critical value of H above which interaction-dependent health benefits can no longer be acquired through contact	Health benefits	10
H_i^α	Critical value of H determining a change in infection resistance	Health benefits	5
H_i^p	Critical value of H determining a change of disease tolerance	Health benefits	5

1093 **B Function $p(H)$**

1094 Function $p(H)$ represents the dependency on H of the probability p of experiencing severe
 1095 illness from infection, and can be modelled equivalently to the probability of infection trans-
 1096 mission $\alpha(H)$ (see Eqs. 3 and Eqs. 4, in the main manuscript). When considering that p
 1097 assumes a sigmoid form, we write

$$1098 \quad p(H) = p_{min} + \frac{p_{max} - p_{min}}{2} [1 - \tanh((H - H_i^p))]. \quad (6)$$

1099 Alternatively, the function $p(H)$ may assume a linear form, as

$$1100 \quad p(H) = \begin{cases} -\frac{(p_{max} - p_{min})}{2 H_i^p} H + p_{max} & \text{for } H \leq 2 H_i^p, \\ p_{min} & \text{for } H > 2 H_i^p. \end{cases} \quad (7)$$

1101 The graphical representation of Eqs. (6) and (7) is provided in Fig. 2(b).

1102 **C Acquisition and loss of interaction-dependent health** 1103 **benefits over time**

1104 Consider changes in the average interaction-dependent health benefits H experienced by an
 1105 initially susceptible population over time. From Eqs. (1) and (2), we know that:

$$1106 \quad \frac{dH}{dt} = \begin{cases} C k h - \delta H & \text{for } H \leq H_{max}, \\ -\delta H & \text{for } H > H_{max}. \end{cases} \quad (8)$$

1107 Note that, for constant contact rate C and in a fully susceptible population (i.e., when
 1108 $S = 1$ and $I, R = 0$), H reaches an equilibrium H^* which corresponds to

$$1109 \quad \lim_{t \rightarrow \infty} H(t) = H^* = \begin{cases} C k \frac{h}{\delta} & \text{for } C k \frac{h}{\delta} \leq H_{max}, \\ H_{max} & \text{for } C k \frac{h}{\delta} > H_{max}. \end{cases} \quad (9)$$

1110 As Eq. (8) is not continuous, in order to solve it numerically, we use a smooth version of
 1111 the function dH/dt , for which $\zeta(H)$ is given by

$$1112 \quad \zeta(H) = C k h \frac{1}{2} [1 - \tanh(3(H - H_{max}))] \quad (10)$$

1113 and the differential equation dH/dt is given by

$$1114 \quad \frac{dH}{dt} = C k h \frac{1}{2} [1 - \tanh(3(H - H_{max}))] - \delta H. \quad (11)$$

1115 A graphical representation of Eq. (10) is provided in Fig. C.1.

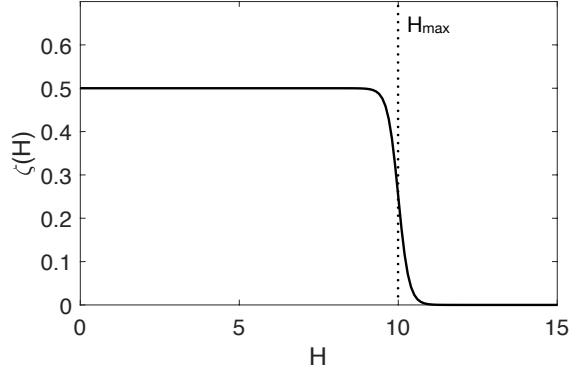


Fig. C.1: Function $\zeta(H)$ of Eq. (10), with $H_{max} = 10$, $C = 10$, $h = 0.1$ and $k = 0.5$.

1116 In Fig. C.2, the solution $H(t)$ of Eq. 11 is plot as a function of time for different contact
 1117 rates (parameter C) and for different fractional contribution of each contact to H (parameter
 1118 k). As given in Eq. (9), the solution $H(t)$ tends to the equilibrium value $H^* = Ckh/\delta$
 1119 for $H \leq H_{max}$. When $H > H_c^m$, the value of $H(t)$ would decrease and stabilize around
 1120 $H^* = H_{max}$. Note that, for constant ratio h/δ , the equilibrium value H^* is reached much
 1121 faster for a higher contact rates C and when a higher fraction of contacts contributes to
 1122 the acquisition of health benefits (parameter k). Increasing the ratio h/δ will also cause the
 1123 equilibrium value to be reached faster.

1124 In Figs. C.3 and C.4 we show how interaction-dependent health benefits (H), the prob-
 1125 ability of infection transmission given a contact ($\alpha(H)$) and the probability of experiencing
 1126 severe disease ($p(H)$) vary as a function of the contact rate C , in an initially fully susceptible
 1127 population.

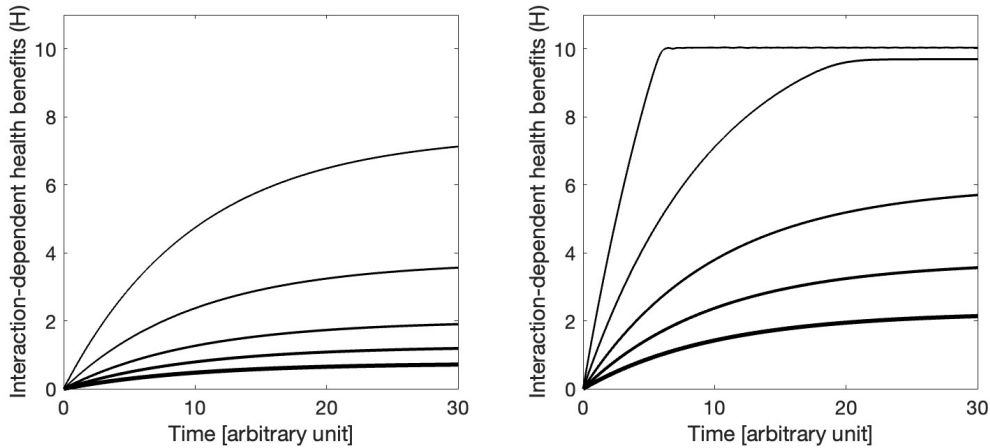


Fig. C.2: The average interaction-dependent health benefits H experienced by a population as a function of time, for different contact rates (from the thicker to thinner lines: $C = 3, 5, 8, 15, 30$, for (a) $k = 0.25$, and (b) $k = 0.75$). For the simulations, dH/dt is modeled according to Eq. 11, with $H_{max} = 10$

. Other default parameter values are provided in Table A.1.

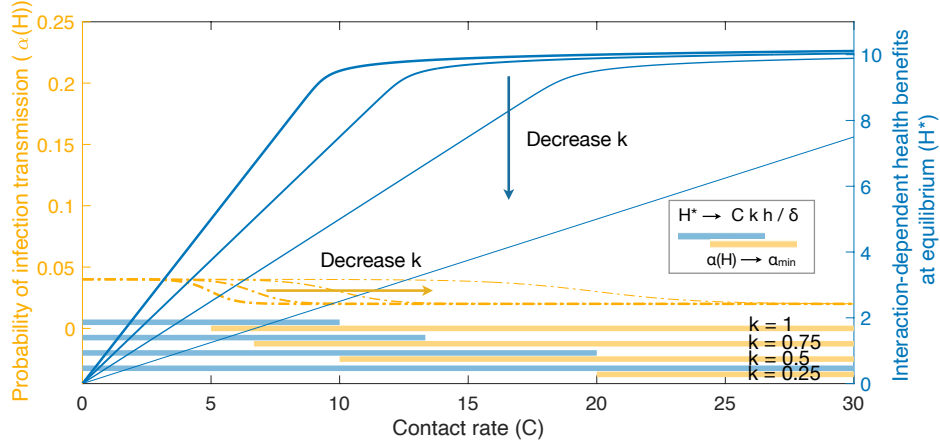


Fig. C.3: Left vertical axis: Probability of infection transmission given a contact (i.e., function $\alpha(H)$, Eq. (4)), as a function of the contact rate C . An equivalent plot can be obtained when considering disease tolerance, and the probability of experiencing severe illness from infection (i.e., function $p(H)$, see Fig. C.4). Right vertical axis: Average interaction-dependent health benefits H experienced by a population at equilibrium (H^* , Eq. (9)), as a function of the contact rate C . Both functions are represented for $k = 0.25, 0.5, 0.75, 1$, with $k = 1$ indicating that all contacts contribute to an increase in H , and $k = 0.25$ indicating that only 1/4 of the contacts contribute to an increase in H . Note that $H^* = Ckh/\delta$ for $Ckh/\delta \leq H_{max}$, and $H^* = H_{max}$ for $Ckh/\delta > H_{max}$ (see Eq. 9). The parameter range of C for which $H^* \rightarrow Ckh/\delta$ (i.e., $C = [0, H_{max}\delta/(hk)]$) for different values of k is indicated in blue at the bottom of the figure. The probability of infection given a contact $\alpha(H)$ decreases with increasing H^* , and thus with increasing C , where $\alpha(H)$ tends to its minimal value α_{min} when $H > H_i^\alpha$. The range $C > H_i^\alpha\delta/k$ for which $\alpha(H) \rightarrow \alpha_{min}$ is indicated for different values of k in orange at the bottom of the figure. Default parameter values chosen for the simulations are provided in Table A.1.

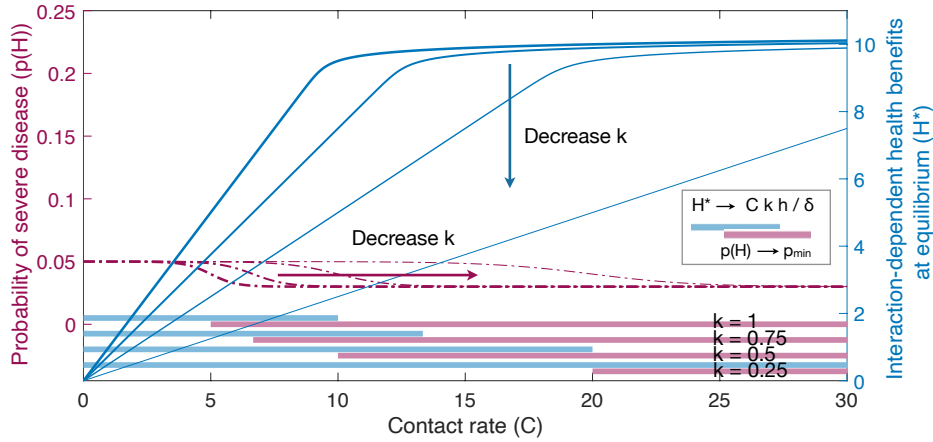


Fig. C.4: Left vertical axis: Plot of disease tolerance, expressed as the probability of experiencing severe illness from infection (i.e., function $p(H)$, Eq. 6), as a function of the contact rate C . Right vertical axis: Average interaction-dependent benefits H experienced by a population at equilibrium (i.e., H^* , Eq. (9)), as a function of the contact rate C . Both functions are represented for $k = 0.25, 0.5, 0.75, 1$. See Fig. C.3 for a more detailed explanation of the figure.

D Results obtained for linear $\alpha(H)$ and $p(H)$, and for

$$\alpha_{min}, p_{min} = 0$$

We would like to explore how the scenarios described in Fig. 3 may change when considering linear $\alpha(H)$ and $p(H)$ functions (Eqs. (3) and (7)), instead of a sigmoid function (Eqs. (4) and (6)). Additionally, we investigate the impact of setting parameters α_{min} and p_{min} to zero on the dynamics. Results are shown in Figs. D.1 and D.2.

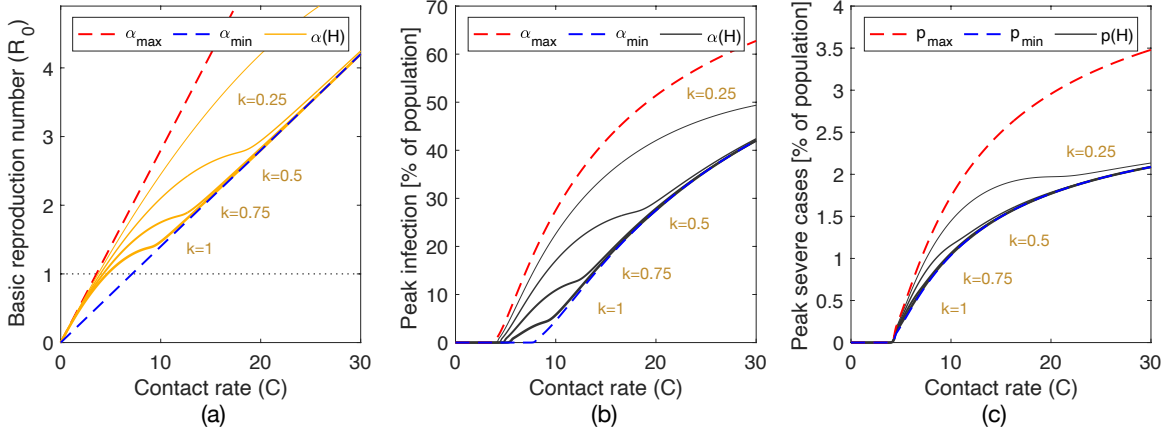


Fig. D.1: Plot equivalent to Fig. 3, but considering that (a,b) the probability of infection transmission given a contact ($\alpha(H)$), and (c) the probability of experiencing severe illness ($p(H)$), decrease linearly with increasing interaction-dependent health benefits (see Eqs. (3) and (7)).

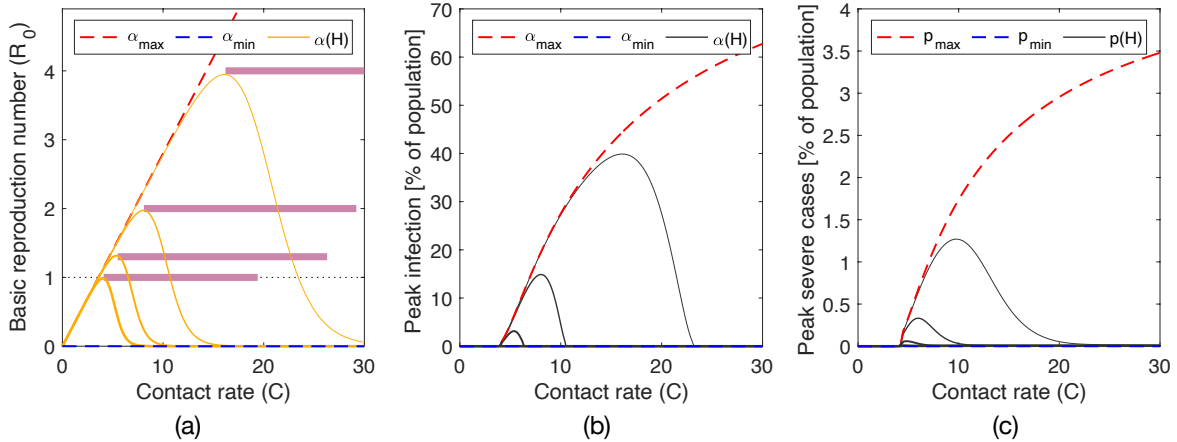


Fig. D.2: Plot equivalent to Fig. 3, but considering that (a,b) the probability of infection transmission given a contact α_{min} and (c) the probability of experiencing severe illness p_{min} reduce to zero when interaction-dependent health benefits are high. Note that in this case (b) an epidemic outbreak, or (c) severe disease occurs only if physical distancing is in place.