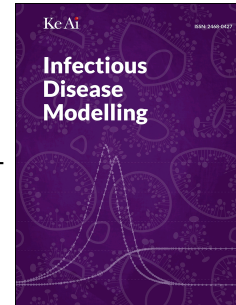


Journal Pre-proof

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Modelling policy combinations of vaccination and transmission suppression of SARS-CoV-2 in Rio de Janeiro, Brazil

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Abstract

COVID-19 vaccination in Brazil required a phased program, with priorities for age groups, health workers, and vulnerable people. Social distancing and isolation interventions have been essential to mitigate the advance of the pandemic in several countries. We developed a mathematical model capable of capturing the dynamics of the SARS-CoV-2 dissemination aligned with social distancing, isolation measures, and vaccination. Surveillance data from the city of Rio de Janeiro provided a case study to analyze possible scenarios, including non-pharmaceutical interventions and vaccination in the epidemic scenario. Our results demonstrate that the combination of vaccination and policies of transmission suppression potentially lowered the number of hospitalized cases by 380+ and 66+ thousand cases, respectively, compared to an absence of such policies. On top of transmission suppression-only policies, vaccination impacted more than 230+ thousand averted hospitalized cases and 43+ thousand averted deaths. Therefore, health surveillance activities should be maintained along with vaccination planning in scheduled groups until a large vaccinated coverage is reached. Furthermore, this analytical framework enables evaluation of such scenarios.

1 Introduction

2 Since the emergence of the SARS-CoV-2 virus, the COVID-19 pandemic has reached many coun-
3 tries causing millions of severe cases and deaths (Tangcharoensathien et al., 2021). The need for
4 interventions was necessary to mitigate the pandemic by reducing dissemination and in the next
5 phase by starting vaccination (Brauner et al., 2021; Perra, 2021). Even with a phased vaccination,
6 some measures remain essential such as social distancing and isolation of cases until extensive vaccine
7 coverage is achieved. Many authors in the literature have pointed out the importance of combining
8 non-pharmaceutical and pharmaceutical interventions to hinder the pandemic (Huang et al., 2021;
9 Borchering et al., 2021; Gumel et al., 2021; Patel et al., 2021). However, these interventions are
10 geographically determined, depending on diverse factors from each city to be applied, having different
11 outcomes in different regions and populations.

12 Several models studied the impact of social distancing (Matrajt and Leung, 2020). Models
13 range from understanding the epidemiological mechanisms behind SARS-CoV-2 and also to predict
14 the dynamics of the epidemic. Schematic and extensive reviews by Wynants et al. (2020) and
15 Padmanabhan et al. (2021) evaluates diverse models against their predictive capabilities. As SARS-
16 CoV-2 is a challenging disease in terms of modelling due to its diverse epidemiological mechanisms
17 that involve different comorbidities (Gude-Sampedro et al., 2021), heavy dependence on public health
18 capacity (Garcia et al., 2020), different impact by age groups (Wu and McGoogan, 2020), the role of
19 asymptomatic individuals (Day, 2020), and is heavily affected by several interventions (Zamir et al.,
20 2020; Lai et al., 2020). Models in the literature have been specializing in understanding not only the
21 general dynamics but also the impact of each factor to tackle each problem assertively. Regarding the
22 diverse epidemiological scenario, models have been developed to enlighten the question of which are
23 the high-risk population where public health authorities could focus resources (Gude-Sampedro et al.,
24 2021; Das et al., 2021b,a).

25 Several non-pharmaceutical interventions require prior careful analysis since they involve not only
26 the number of cases and deaths, but many also address psychological issues (Adeniyi et al., 2022;
27 Rahaman et al., 2020), the necessary resources for their application, and other health issues related
28 to them, due to the emergence of other diseases during the pandemic (Rana et al., 2021; Shimizu
29 et al., 2021; Torner, 2020). Nonetheless, these interventions have been of paramount necessity in
30 reducing the number of deaths and hospitalizations worldwide (Spinelli et al., 2021; Perra, 2021; Lai
31 et al., 2020; Flaxman et al., 2020; Zamir et al., 2020; Jorge et al., 2021).

32 Since the beginning of the pandemic, vaccination and other pharmaceutical interventions have
33 been an object of study (Huang et al., 2021). However, only most recently have we reached more

34 thorough vaccination in diverse countries. Understanding the specific dynamics that separate the
 35 impacts of non-pharmaceutical and pharmaceutical interventions is still debated in the literature. It
 36 demands a modeling task that requires careful exploration of different classes of individuals through
 37 age groups, and their specificities (Wu and McGoogan, 2020).

38 In this work, we develop and evaluate how a model can capture the dynamics of the SARS-
 39 CoV-2 pandemic and compare scenarios with and without interventions to better deal with the
 40 ongoing SARS-CoV-2 pandemic and understand the real impact of these measures. Results consider
 41 specifically the dynamics of the pandemic in the city of Rio de Janeiro, Brazil, as a case study.
 42 However, implications of the results are general, such that they could be extended to other similar
 43 cities. Also, as non-pharmaceutical measures are essential to mitigate the effects of the pandemic, the
 44 perspective of controlling it comes with vaccination. However, its policies and methods for application
 45 need yet to be systematically addressed. Health surveillance should be maintained along with the
 46 planning for effective vaccination.

47 2 Methods

48 2.1 Model

49 We modeled different scenarios with an ODE-based compartmental model. In the model, susceptible
 50 individuals (S) can evolve to exposed (E) condition when in contact with infected individuals. The
 51 group of infected individuals is divided between asymptomatic cases (Y), symptomatic cases (C),
 52 which includes both mild and moderate cases (Cx), which can evolve to severe cases (H). This last
 53 group occurs from the evolution of the symptomatic group and, therefore, is considered to represent
 54 hospitalized individuals. All infected individuals can evolve to death (D) or recovered (R). We have
 55 also included the dynamics of vaccination for a single dose. Vaccination individuals (V) can evolve
 56 into Immunized (I) or non-immunized individuals (Im). Each of these model classes is stratified by
 57 age groups, from 0 to 100 years, in 5-year intervals, added by one last age group of higher than 100
 58 years, in a total of 21 groups of distinct age ranges for each compartment of the model.

59 Vaccination is included in the model as susceptible individuals are vaccinated at a coverage rate
 60 of η . These vaccinated individuals will take a pre-determined time τ_{immun} to develop immunity at
 61 a probability of ρ_I when they evolve to immunized status (I). Due to incomplete vaccine efficacy,
 62 we included the possibility of the vaccinated individual not developing the required immunization
 63 and still being susceptible (Im). Although some individuals are to be considered non-immunized,
 64 as reported by other authors (Hogan et al.), these individuals are less likely to be infected, develop

65 symptoms, be hospitalized, and die. Therefore, these individuals evolve to different but relatable
 66 classes of individuals, as shown in Fig. 1 with the classes that end with the letter "v". The reduced
 67 parameters related to these different degrees of severity were previously reported (Hogan et al.).

68 The infection rate between susceptible individuals and symptomatic is β , and with asymptomatic
 69 individuals is β_A . When they become exposed individuals, the time to evolve to infected is the
 70 incubation time τ_{inc} . At the end of this time, the individual has a probability ρ_S of developing
 71 symptoms.

72 The time required for an asymptomatic individual to evolve to death is α_A^{-1} , whereas for the
 73 symptomatic individuals is α^{-1} . It is expected that $\alpha > \alpha_A$ due to higher morbidity in the former
 74 case, besides the fact that asymptomatic individuals do not present themselves as clinical cases.
 75 Symptomatic individuals can evolve to a severe case with a risk probability of α_H . The symptomatic
 76 (C) and severe cases (H) individuals are modeled separately due to their different epidemiological
 77 mechanisms (Liu et al., 2020; Siordia Jr, 2020), and to allow the test of non-pharmaceutical
 78 methodologies that target these individuals separately. The separation between these individuals is
 79 mainly based on their symptoms, e.g., mild/symptomatic and symptoms requiring hospitalization.
 80 Severe cases exhibit clinical conditions for hospitalization, such as oxygen saturation lower than 93%,
 81 dyspnea, or multiple organ failure (Wu and McGoogan, 2020; Betti and Heffernan, 2021; Musa et al.,
 82 2021; Chevrier et al., 2021). Both can evolve to death (or the symptomatic case can evolve to the severe
 83 case) separately with different case-fatality ratios, as shown in the literature (Wu and McGoogan,
 84 2020). It is vital to understand whether isolating only the severe cases is an adequate measure to
 85 mitigate the pandemic or if we should apply a broader approach when applying non-pharmaceutical
 86 interventions.

87 The parameters related to asymptomatic individuals, such as β_A and α_A are calculated through
 88 a product between a reducing factor (Byambasuren et al., 2020) f_A and the original parameter for
 89 symptomatic individuals β and α respectively. Regarding the individuals that were vaccinated but
 90 are not immunized, another reducing factor is considered (Palacios et al., 2021), f_v . This factor
 91 applies to reduce the infection rate β with the product $\beta \cdot f_v$. These individuals also have a reducing
 92 factor applied to their hospitalization risk (Palacios et al., 2021), $f_{v,H}$.

93 The recovery of infected individuals (symptomatic and severe) is controlled by the recovery
 94 rate (Zhou et al., 2020) γ , being modified to γ_H in the case of severe cases. Severe cases are
 95 hospitalized and thus receiving proper assistance confronting the sickness. The hospitalized individual
 96 can recover after a determined period, controlled by the discharge time τ_{disc} and dyspnea time τ_{dysp} .
 97 The discharge time is a median time where individuals who present a clinical recovery are therefore
 98 termed as recovered individuals and are no longer hospitalized. In contrast, the dyspnea time is the

99 time from illness onset to dyspnea clinical condition (Zhou et al., 2020). We calculated the recovery
 100 rate of individuals ($\gamma_H = \frac{1}{\tau_{disc} - \tau_{dysp}}$) using both the discharge and dyspnea time, as we considered a
 101 stochastic implementation of our model.

The ODE system which resumes this model is:

$$\frac{dS}{dt} = -\beta(C + H)S - \beta_A YS - \eta S \quad (1)$$

$$\frac{dE}{dt} = \beta(C + H)S + \beta_A YS + \beta f_v(C + H)Im + \beta_{Im,A} YS - \frac{E}{\tau_{inc}} \quad (2)$$

$$\frac{dC}{dt} = \rho_S \frac{E}{\tau_{inc}} - \frac{\alpha \gamma C}{1 - \alpha(1 - \gamma)} - \frac{\alpha_H \gamma C}{1 - \alpha_H(1 - \gamma)} - \gamma C \quad (3)$$

$$\frac{dY}{dt} = (1 - \rho_S) \frac{E}{\tau_{inc}} - \frac{\alpha_A \gamma C}{1 - \alpha_A(1 - \gamma)} - \gamma Y \quad (4)$$

$$\frac{dH}{dt} = \frac{\alpha_H \gamma C}{1 - \alpha_H(1 - \gamma)} - \frac{H}{\tau_{disc} - \tau_{dysp}} - \frac{\alpha \gamma_H H}{1 - \alpha(1 - \gamma_H)} \quad (5)$$

$$\frac{dD}{dt} = \frac{\alpha \gamma C}{1 - \alpha(1 - \gamma)} + \frac{\alpha_A \gamma C}{1 - \alpha_A(1 - \gamma)} + \frac{\alpha \gamma_H H}{1 - \alpha(1 - \gamma_H)} \quad (6)$$

$$\frac{dR}{dt} = \gamma C + \gamma Y + \frac{H}{\tau_{disc} - \tau_{dysp}} \quad (7)$$

$$\frac{dV}{dt} = \eta S - \frac{1}{\tau_I} V \quad (8)$$

$$\frac{dI}{dt} = \frac{\gamma_I}{\tau_I} V \quad (9)$$

$$\frac{dIm}{dt} = \frac{1 - \gamma_I}{\tau_I} V - \beta f_v(C + H)Im \quad (10)$$

$$(11)$$

102 2.1.1 Social distancing interventions

103 The model enables the application of intervention measures with the social distancing of specific age
 104 groups. Social distancing affects people in reducing the probability of encounters between infected
 105 and susceptible individuals. Thus, we simulate this condition by reducing the infection rates β , β_A ,
 106 β_I and β_{Im} for the specific age groups. Due to imperfect application of social distancing intervention,
 107 each intervention is controlled by a success rate.

108 The fact that the model is stratified by age groups opens a new range of different scenarios, e.g.
 109 when applying the social distancing intervention to younger age groups, we can simulate limitation
 110 of school activities. The reduction is applied to the R_0 value, from which the infection rates are
 111 calculated, by multiplying it with the reduction factor $\kappa = 0.65$. The social distancing applied to the
 112 0-20 years old age groups is labeled SD-Y, when applied to the age groups higher than 60 years old is
 113 labeled SD-E, and when we apply the reduction to all age groups, we label this condition as SD-A.

114 2.1.2 Isolation interventions

115 The application of isolation interventions is made by reducing the encounter probability between
 116 susceptible and infected individuals. Different scenarios are tested in this work. In the lockdown
 117 scenario (L), we alter the susceptible flow equation to

$$\frac{dS}{dt} = -\beta(1 - \sigma_L)(C + H)S - \beta_A(1 - \sigma_L)YS - \eta S \quad (12)$$

118 Another intervention possibility is when tests are applied to the individuals, and a quarantine
 119 is applied where symptomatic cases are isolated with a probability σ and asymptomatic with a
 120 probability σ_A , this condition is labeled as TQ-C. In this scenario, we modify the susceptible flow
 121 equation to

$$\frac{dS}{dt} = -\beta(1 - \sigma)(C + H)S - \beta_A(1 - \sigma_A)YS - \eta S \quad (13)$$

122 If we only isolate the symptomatic cases (scenario TQ), we change the susceptible individuals
 123 flow equation to

$$\frac{dS}{dt} = -\beta(1 - \sigma)(C + H)S - \beta_A YS - \eta S \quad (14)$$

124 The scenario where we only isolate the severe cases is termed TQ-S, and we modify the susceptible
 125 flow equation to

$$\frac{dS}{dt} = -\beta(C + \sigma H)S - \beta_A YS - \eta S \quad (15)$$

126 The exposed, vaccinated, and partially immunized compartments are also changed as the susceptible
 127 flow, depending on the applied scenario. Table 1 summarizes the parameters used in the model
 128 with their respective values and references. Only four parameters were fitted to represent the SARI
 129 notification data for the city of Rio de Janeiro: the basic transmission rate (β) via R_0 , and the three
 130 probabilities of isolation (for symptomatic cases (σ), asymptomatic cases (σ_A) and lockdown scenario
 131 (σ_L)). The other parameters are recovered from the literature (Table 1).

132 The parameter β is calculated from the previous definition of R_0 value, the asymptomatic

Table 1: Description of parameters in the model and values used in simulations with references.

Parameter	Description	Value
β	Infection rate	Calculated using R_0
f_A	Asymptomatic factor	0.42 (Byambasuren et al., 2020)
β_A	Asymptomatic infection rate	$f_A \cdot \beta$
σ	Probability of successful isolation of symptomatic individuals	0.60
σ_A	Probability of successful isolation of asymptomatic individuals	0.20
σ_L	Probability of successful isolation during lockdown	0.75
ρ_S	Probability of developing symptoms	0.83 (Byambasuren et al., 2020)
α	Death risk	Depends on age group (Wu and McGoogan, 2020)
α_H	Hospitalization risk	Depends on age group (Stokes et al., 2020)
α_A	Death risk of asymptomatic individuals	$f_A \cdot \alpha$
τ_{dysp}	Time for dyspnea	7 days (Zhou et al., 2020)
τ_{disc}	Discharge time	22 days (Zhou et al., 2020)
τ_{inc}	Incubation time	5.1 days (Lauer et al., 2020)
γ	Recovery rate	1/6.5 (Zhou et al., 2020)
γ_H	Recovery rate for hospitalized individuals	Calculated using τ_{disc} and τ_{dysp}
γ_I	Immunization probability	0.493 (Palacios et al., 2021)
τ_I	Time to immunization	14 days (Palacios et al., 2021)
f_v	β reducing factor for I_m individuals	0.163 (Palacios et al., 2021)
$f_{v,H}$	α_H reducing factor for I_m individuals	0.163 (Palacios et al., 2021)

133 value (Byambasuren et al., 2020) f_A , the probability of developing symptoms (Byambasuren et al.,
 134 2020) ρ_S , and the incubation time (Lauer et al., 2020) τ_{inc} with

$$\beta = \frac{R_0}{\tau_{inc}(\rho_S + (1 - \rho_S)f_A)} \quad (16)$$

135 Fig. 1 depicts a schematic diagram showing the model compartments.

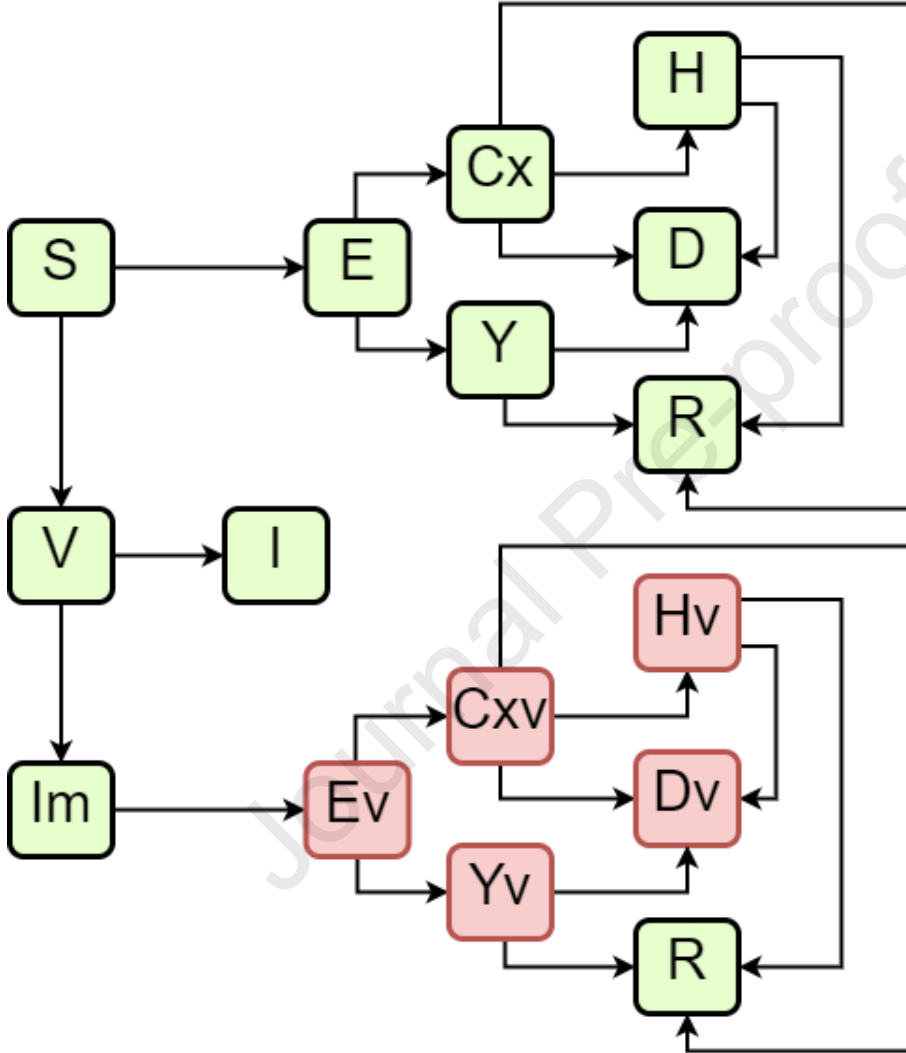


Figure 1: Schematic diagram of the model compartments.

136 2.1.3 Stochastic implementation

137 The model is implemented in a discrete-time fashion. Days were applied as the time units since cases
 138 are reported on a daily basis and the change of interventions could be simulated on specific dates.
 139 However, the algorithm could use other time units such as weeks, if applied adjusted parameters.
 140 The algorithm requires initial values for the variables used in the model. The transitions given in the

141 equations in the ODE system are used to obtain the transition probabilities (Allen, 2017). Typically,
142 for each time step, all transitions are evaluated as probabilities and the number transitioning from
143 a compartment to the other linked compartments, including keeping the state, are drawn from a
144 multinomial distribution. Multiple simulations generate multiple sample paths, which are evaluated
145 allowing to obtain mean values and intervals. Since the transitions follow distributions with the
146 parameters used in the model, after multiple simulations the mean values are expected very close to
147 the deterministic realization. A stochastic simulation code was implemented using Rstudio (Racine,
148 2012) Version 1.2.5042 with R software version 4.0.0 (<http://www.r-project.org>) was used for all
149 calculations, data importation, and curation.

150 2.2 Case study

151 Parameters of the model were adjusted to the number of cases and the dynamics observed in the
152 municipality of Rio de Janeiro. Data from Severe Acute Respiratory Illness (SARI) are compared
153 to the results of new daily hospitalizations. In contrast, data from Acute Respiratory Illness (ARI)
154 notified cases are compared to the results of new daily cases. All notified data is retrieved from the
155 public database OpenDataSus (available at <https://opendatasus.saude.gov.br/dataset>).

156 Throughout the pandemic, the scenario was altered several times due to governmental decisions
157 of applying the interventions or making them more flexible and the incomplete adherence of the
158 population. In this section, we evaluate how the model behaves when we use the same quarantine
159 severity as applied by the government for each period while comparing the results to real-time data.
160 Our approach is based on the Rio de Janeiro municipality and state real pandemic decrees, with slight
161 adjustments, as the accordance of the population to governmental decisions is not straightforward.
162 We consider no intervention done between 01 January 2020 and 15 March 2020 (day 1 to day 74).
163 Starting from 16 March 2020 until 27 March 2020 (day 75 to day 86), we consider that this is the
164 beginning of the pandemic, where the government started to apply some intervention measures. The
165 population's adherence to non-pharmaceutical interventions of the government in Brazil was not
166 strictly followed (de Moura Villela et al., 2021; Jorge et al., 2021; Szwarcwald et al., 2020), decreasing
167 with the temporal advance of the pandemic. Therefore, to model the notification data, we have
168 considered the non-pharmaceutical interventions during 2021 after the vaccination as adjustable when
169 necessary. The different isolation and social distancing scenarios are summarized in Table 2.

Table 2: Isolation and social distancing scenarios for the different data ranges throughout the years of 2020 and 2021.

Data range (DD.MM.YY)	Isolation	Social Distancing
16.03.2020 - 27.03.2020	TQ-S	SD-Y+SD-E
28.03.2020 - 03.04.2020	TQ-S	SD-A
05.04.2020 - 14.05.2020	TQ-C	SD-A
15.05.2020 - 29.05.2020	L	SD-A
30.05.2020 - 02.06.2020	TQ	SD-A
03.06.2020 - 12.07.2020	TQ-C	SD-A
13.07.2020 - 02.09.2020	TQ-C	SD-Y + SD-E
03.09.2020 - 22.09.2020	TQ-C	SD-A
23.09.2020 - 31.10.2020	TQ-C	SD-Y + SD-E
01.11.2020 - 16.11.2020	TQ-S	SD-A
17.11.2020 - 21.11.2020	TQ-C	SD-A
22.11.2020 - 01.12.2020	TQ	SD-A
02.12.2020 - 30.01.2020	TQ-C	SD-A
31.01.2021 - 07.03.2021	TQ-S	SD-A
08.03.2021 - 18.03.2021	TQ-S	SD-Y+SD-E
19.03.2021 - 02.04.2021	TQ	SD-A
03.04.2021 - 06.04.2021	-	SD-A
07.04.2021 - 18.04.2021	TQ-S	SD-A
19.04.2021 - 22.04.2021	TQ-C	SD-A
23.04.2021 - 30.04.2021	TQ-S	SD-A
01.05.2021 - 04.05.2021	TQ-C	SD-A
04.05.2021 - 14.05.2021	TQ-S	SD-Y + SD-E
15.05.2021 - 19.05.2021	TQ-C	SD-Y + SD-E
20.05.2021 - 30.06.2021	TQ-S	SD-A

170 To better fit the model to the real notification data, we estimated initially $R_0 = 2.6$, the reduction
171 factor of the social distancing during lockdown to be 0.75, the success in isolating symptomatic
172 cases to be 0.60, while 0.20 for the asymptomatic cases. Also, we considered that the first cases
173 were imported on 11 February 2020. Reporting rate of severe cases (SARI) is 96% of the real cases,
174 accounting for small under-reporting, whereas under-reporting of notified ARI disease cases is 20% of
175 the actual number of ARI cases. The number of SARI cases notified in the city of Rio de Janeiro,

176 daily aggregated, is evaluated from January to the end of June of 2021. This data range is considered
 177 an acceptable range to avoid the effect of dramatic sub notification due to notification delay.

178 In order to evaluate the vaccination program, we used real vaccination data notification from the
 179 city of the Rio de Janeiro applied to each group at the specific dates on which they were applied.
 180 Figures containing the reported vaccination data are available in the supplementary material of this
 181 work. As our model accounts for only one dose of vaccination, we applied to the simulations the
 182 dates of first dose to reach the different scenarios, using data from all applied vaccines. In Brazil, the
 183 vaccination program covers both two-dose and single dose vaccines (Hung and Poland, 2021; Ranzani
 184 et al., 2021; Villela et al., 2021). To capture the general mechanism provided by the pharmaceutical
 185 interventions, our approach has only the application of a single dose program that also includes
 186 the infection-rate reduction (Hogan et al.) and hospitalization risk reduction (Palacios et al., 2021).
 187 Furthermore, the protection provided by the vaccination starts after the first dose (Iacobucci and
 188 Mahase; Tuite et al., 2021), although not full nor long-lasting, as these mechanisms help to represent
 189 the notification data using only a single dose program simulation. Also, we analyzed the prevention
 190 of deaths and hospitalizations for different scenarios considering the cumulative curves of each case
 191 using the equation for number of prevented (deaths or hospitalizations) $\lambda(t)$,

$$\lambda(t) = \lambda_{specific}(t) - \lambda_{non}(t) \quad (17)$$

192 where $\lambda(t)$ represents cumulative deaths or hospitalization at time t , *specific* refers to the specific
 193 scenario studied scenario, and *non* represents the scenario without vaccination and restrictions.

194 3 Results

195 The model captured the dynamics of the epidemics in Rio de Janeiro successfully regarding the
 196 hospitalizations compared to SARI notified cases (Fig. 2). As the model does not account for all the
 197 influenza-like illness, but it is limited to the SARS-CoV-2 cases, there should be a difference between
 198 the notification data and the SARI notified cases, also due some natural errors within notification
 199 systems. To cover this problem, we considered a reporting parameter of 0.95 to data.

200 The model also presented a good fit to notification data when using the vaccination data. As
 201 expected, the combination of vaccination, social distancing and isolation measures was responsible to
 202 significantly lower the number of SARI notified cases throughout the years of 2020 and 2021 in Rio de
 203 Janeiro. However, if more restrictive measures were applied, the resulting effect was clearly stronger.

204 After the beginning of the vaccination program, the downfall of the pandemic is advanced and

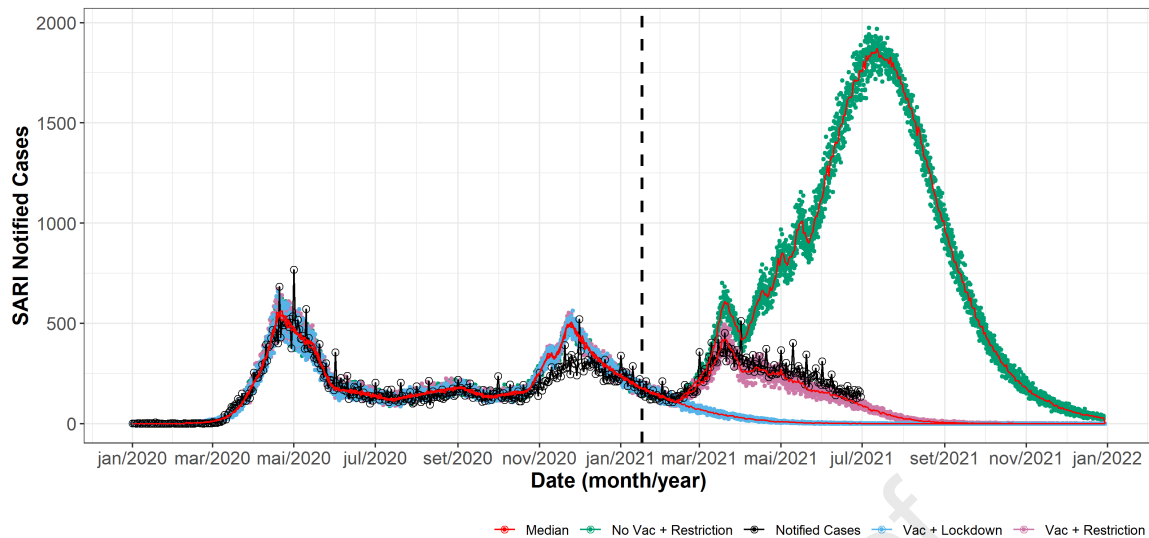


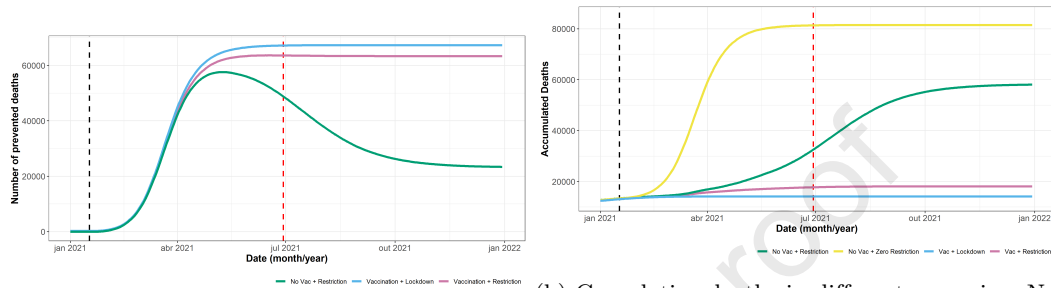
Figure 2: Model results for new daily hospitalizations and cases of SARI in Rio de Janeiro. Notified cases of SARI in Rio de Janeiro are represented by black lines, other colors represent the different simulated vaccination scenarios: vaccination with the applied restrictions (pink), no vaccination but applying the same restrictions as the pink case (green), and vaccination with lockdown scenario (blue). Red lines represents the median values in each scenario.

205 accelerated, which is evidenced by the observed inflection point. Abandoning social distancing,
 206 however, generates an increase in the number of expected SARI cases as shown. As shown by a
 207 last peak of simulation data, the advancement of vaccination dates is responsible to reduce the
 208 number of cases in a downward direction in conditions where a new peak would rise. If there were
 209 no flexibilization during vaccination, no peaks would be observed. The number of cumulative and
 210 prevented deaths and hospitalizations, are shown in Fig. 3.

211 As shown in Fig. 3, the vaccination had a major role in reducing the number of hospitalizations
 212 and deaths due to SARI. The reduction in number of cases after vaccination and suppression policies,
 213 compared to a no-policy scenario, was 380+ thousand hospitalized cases and 66+ thousand cases,
 214 considering until June 2021. Vaccination is expected in this case to avert more than 230+ thousand
 215 hospitalized cases and 43+ thousand deaths.

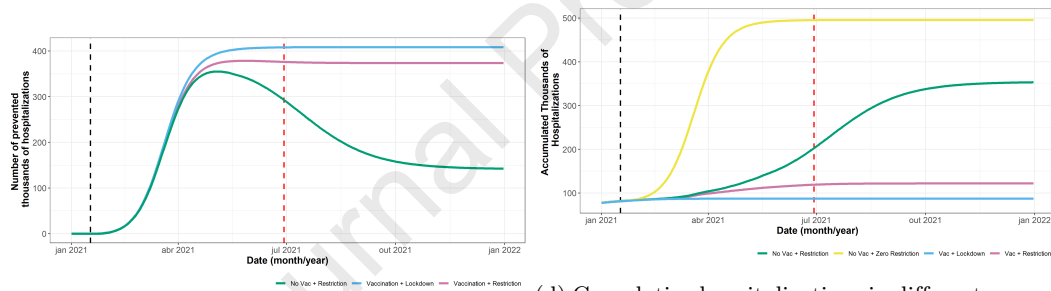
216 The prevented deaths and hospitalizations are only observed to remain high when the vaccination
 217 is applied, which is a direct result of the long-term protection provided by the vaccine. The contrary
 218 is observed when only social distancing or other non-pharmaceutical interventions are applied, as new
 219 peaks maintain high number of deaths and hospitalizations.

220 As shown in Fig.4, there is a marked difference in the effectiveness of each intervention alone.
 221 Social distancing alone had a less pronounced effect than the quarantine of cases, except for the
 222 quarantine of only the severe cases (TS), which had a minimal delaying effect at the peak. A
 223 combination of mitigation policies makes significant impact in the peak of number of cases.



(a) Prevented deaths in different scenarios. Vaccination and no restrictions (yellow), vaccination and Lockdown (blue), vaccination and restriction (pink), and applying only the restrictions, without vaccination (green) .

(b) Cumulative deaths in different scenarios. No vaccination and no restrictions (yellow), vaccination and lockdown (blue), vaccination and restriction (pink), and applying only the restrictions, without vaccination (green).



(c) Prevented hospitalizations in different scenarios. No vaccination and no restrictions (yellow), vaccination and Lockdown (blue), vaccination and restriction (pink), and applying only the restrictions, without vaccination (green) .

(d) Cumulative hospitalizations in different scenarios. No vaccination and no restrictions (yellow), vaccination and lockdown (blue), vaccination and restriction (pink), and applying only the restrictions, without vaccination (green).

Figure 3: Different scenarios comparing prevented deaths and hospitalizations, and cumulative deaths and hospitalizations due to SARI. To calculate the prevented deaths and hospitalizations, we used our model to calculate a scenario where no restrictions and no vaccination were applied, the cumulative deaths and hospitalization curves of this scenario was our reference to calculate the absolute the number of prevention.

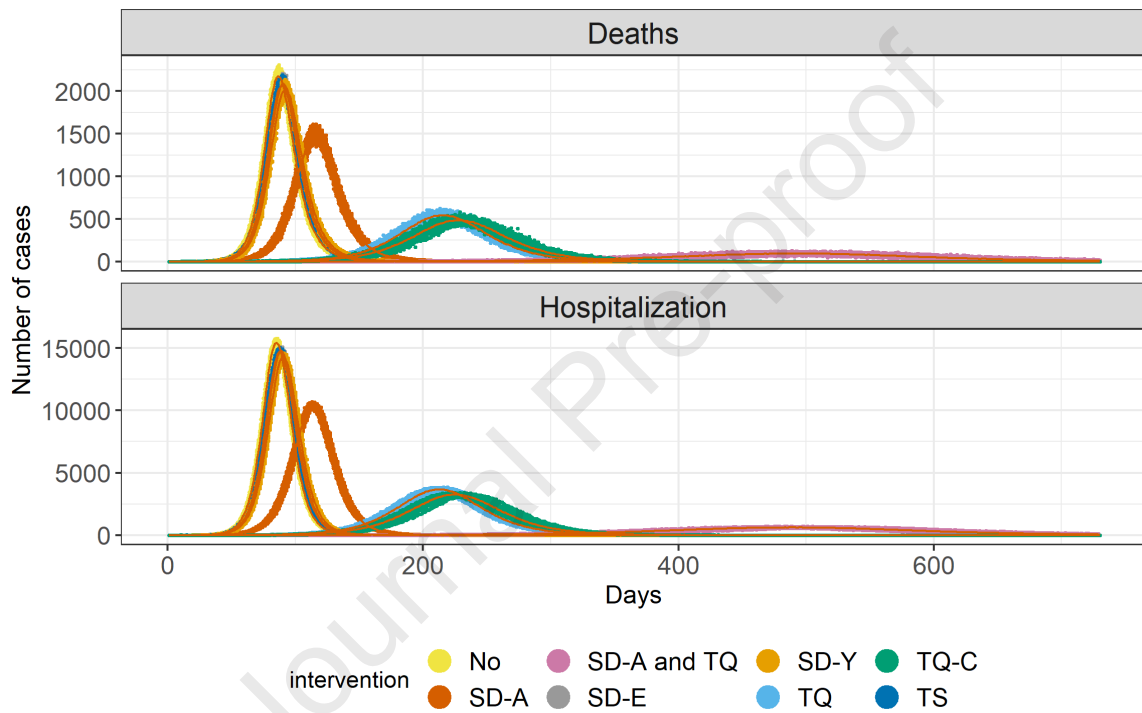


Figure 4: Different scenarios model comparison. A different color identifies each intervention. The points represent the stochastic calculation done with the model considering the given probabilities with 100 iterations per day. The red lines are means of each intervention. The used parameters are given in Table 1, with the exception of R_0 , which is 3.5.

224 As our model is stratified by age groups, we also observe how the different interventions change
 225 the number of deaths and hospitalizations by age, as shown by Fig.5. The quarantine of all cases, the
 226 social distancing of all individuals, and the combination of this intervention with the quarantine of
 227 symptomatic cases are the three most effective interventions, as also seen by Fig.4. In all cases, despite
 228 isolating or distancing different age groups, the pattern of hospitalizations and deaths regarding age
 229 groups is very similar. The major difference is observed in delaying the pandemic peak and the
 230 pandemic's length, broadening its profile through time but not through age groups. Hospitalizations
 231 are centered around older groups, mainly individuals around 60 years old and older, in all interventions.

232 Also, in Figure 5, despite profile similarity across age groups, some age groups are more affected
 233 since the beginning of the pandemic and at the end. There is a distortion of the profile's rectangular
 234 shape observed in almost all scenarios in favor of a more oval-oriented shape, which is more pronounced
 235 in the SD-A and TQ, only TQ, and only TQ-C scenarios.

236 4 Discussion

237 The main objective of NPI interventions is to mitigate the effect of the pandemic for proper health
 238 care attention to mild and severe cases. As shown by Fig.4 independently from the nature of the
 239 intervention (social distancing or isolation of cases), as expected and seen in many studies (Matrajt
 240 and Leung, 2020; Ferguson et al., 2020; Flaxman et al., 2020; Prem et al., 2020), delaying the epidemic
 241 peak is a consequence of the reduction in transmission intensity.

242 As demonstrated in Fig.4, when comparing different interventions, there is a considerable difference
 243 between the isolation of both symptomatic and asymptomatic cases and isolating only the symptomatic
 244 cases, with the former intervention being a more successful application. Further, if we combine
 245 isolation with social distancing interventions, a better result is reached in reducing the number
 246 of hospitalizations and delaying the peak of new cases. This result highlights the importance of
 247 an enforced isolation measure, as the asymptomatic cases also impact the transmission dynamics.
 248 The correct identification and consequently isolation of these cases pose a problem which has been
 249 discussed in the actual pandemic (Gandhi et al., 2020; Nishiura et al., 2020), in some cases, following
 250 the correct procedure to identify and isolate these cases were responsible for ending the pandemic
 251 (Day, 2020). The isolation of only the severe cases did alter significantly the dynamics, demonstrating
 252 the importance of having a model in which mild and severe cases are studied separately, as they have
 253 marked differences in their epidemiology (Liu et al., 2020; Siordia Jr, 2020) besides having some
 254 studies indicating some similarities (Yilmaz et al., 2020; Wu and McGoogan, 2020). The isolation of
 255 only symptomatic cases was more effective when applied together with the social distancing of all age



Figure 5: Normalized death and hospitalization profiles for different intervention scenarios. Normalized values are calculated by the quotient of each daily new hospitalization or death by the highest hospitalization or death of the group with most hospitalizations or death through the pandemic.

256 groups. Therefore, it is imperative to recognize the importance of transmission by asymptomatic
257 individuals.

258 Comparing the applied social distancing measures, results here show a very marked difference
259 between the isolation of all age groups against the isolation of only young or elderly individuals and
260 the severity of SARS-CoV-2 among elderly individuals higher than younger individuals (Siordia Jr,
261 2020; Wu and McGoogan, 2020). However, there must be a very careful distinction between the
262 severity of cases and the epidemiological dynamic imposed by the different groups, the isolation of
263 only the elder individuals is not sufficient to significantly halt the pandemic. As shown in our
264 model, isolating the elderly group may give a false impression of protection to these individuals, as
265 this intervention is not sufficient to effectively stop the epidemic. Therefore, only the social distancing
266 of all age groups at an early stage acts to avoid severe cases.

267 The social distancing of all age groups had similar performance compared to the isolation of both
268 symptomatic and asymptomatic cases, as shown by Fig.4. This interesting result indicates that the
269 early recognition and application of broad interventions to the population are the most effective
270 measures to be studied. In regard to the social distancing, all age groups should be taken into account,
271 in agreement with other modeling studies (Ferguson et al., 2020; Flaxman et al., 2020). Regarding
272 the isolation intervention, all cases should be included in the measure, including asymptomatic cases,
273 which can only be reached through successful testing. This highlights the importance of mass testing
274 individuals exposed to the SARS-CoV-2 pandemic.

275 The value of 2.6 for the R_0 is within the range of the estimated value for other studies and even
276 other areas (Coelho et al., 2020; Yue et al., 2021; Abbott et al., 2020; Li et al., 2020; Wu et al., 2020).
277 Despite the significant number of interventions, either a social distancing or isolation intervention,
278 the best approach is clearly the combination of both measures. This is shown in 4 where the SD-A
279 intervention combined with the TQ isolation measure produced the best results.

280 Despite all of the interventions, combined or not, there is a growing concern about the social and
281 economic distress of a population during interventions (Ashraf, 2020; Fernandes, 2020). It is also
282 imperative to develop pharmaceutical interventions to reduce the posed threat by the virus infections.
283 Also, initiatives such as the vaccines being developed and the fundamental understanding of how the
284 virus acts biologically are essential to this end. Therefore, it is crucial to model beyond the dynamics
285 of only non-pharmaceutical interventions.

286 Non-pharmaceutical interventions also demonstrate through Fig.2 that they have the merit of
287 controlling the direction, evolution, and severity of the pandemic and should be studied and applied
288 whenever possible. However, pharmaceutical and non-pharmaceutical interventions need to be
289 considered altogether during the pandemic. Considering these results, it is clear that the vaccine has

290 a long-term effect on the population. Comparing the last peaks obtained by the results, although
291 vaccination did not control the direction of the pandemic, it was directly responsible to diminish the
292 number of cases and deaths effectively.

293 In all scenarios, the phased rollout of the vaccination program should be along with maintaining
294 social distancing and case isolation. Abandoning the quarantine shows to be a most critical scenario,
295 in which there is a considerable increase in the number of hospitalizations. The only condition where
296 the pandemic maintains its downward strategy during the vaccination program is combining social
297 distancing and isolation.

298 This is a crucial moment to study and show that we must yet consider the application of strict
299 interventions of social distancing, isolation, and vaccination as the risk of SARS-CoV-2 transmission is
300 present in multiple countries. The modelling in this work shows that effective control of the COVID-19
301 pandemic requires a combination of these efforts.

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Data Availability Statement

All SARI and ARI notification data are publicly available at OpenDataSUS database, maintained by the Ministry of Health, located at <https://opendatasus.saude.gov.br/>.

Competing interests

Authors declare no competing interests.

Ethics statement

No approval by an ethics committee was necessary, since the work involved only simulations and secondary anonymized data which are publicly available.

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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