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Research article

Impact of immunity loss on the optimal vaccination strategy for an age-structured epidemiological model

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Abstract: The pursuit of effective vaccination strategies against COVID-19 remains a critical endeavour in global public health, particularly amidst challenges posed by immunity loss and evolving epidemiological dynamics. This study investigated optimal vaccination strategies by considering age structure, immunity dynamics, and varying maximal vaccination rates. To this end, we formulated an SEIR model stratified into n age classes, with the vaccination rate as an age-dependent control variable in an optimal control problem. We developed an objective function aimed at minimising critical infections while optimising vaccination efforts and then conducted rigorous mathematical analyses to ensure the existence and characterization of the optimal control. Using data from three countries with diverse age distributions, in expansive, constrictive, and stationary pyramids, we performed numerical simulations to evaluate the optimal age-dependent vaccination strategy, number of critical infections, and vaccination frequency. Our findings highlight the significant influence of maximal vaccination rates on shaping optimal vaccination strategies. Under constant maximal vaccination rates, prioritising age groups based on population demographics proves effective, with higher rates resulting in fewer critically infected individuals across all age distributions. Conversely, adopting age-dependent maximal vaccination rates, akin to the WHO strategy, may not always lead to the lowest critical infection peaks but offers a viable alternative in resource-constrained settings.

Keywords: optimal control; epidemiology; long-term vaccination; COVID-19; age-structured model

1. Introduction

The quest for effective vaccination strategies against COVID-19, coupled with the challenges posed by immunity loss and evolving epidemiological dynamics, constitutes a multifaceted endeavour at the forefront of global public health efforts. This endeavour encompasses various facets, ranging from the durability of vaccine-induced immunity to the emergence of new viral variants and the imperative of sustained vaccination efforts [1].

The optimisation of control strategies, particularly for vaccination allocation in the context of COVID-19, has been extensively studied and discussed. Research has varied, with some focusing on managing the pandemic through non-pharmaceutical interventions (NPIs) like social distancing and mask-wearing [2, 3], while others have concentrated on vaccination strategies. Various studies offer insights into optimal vaccination strategies under different conditions. For instance, Gonzales-Parra, Cogollo, and Arenas [4] examined strategies considering factors like case fatality rates and population age structure. Hogan et al. [5] suggested targeting the elderly when vaccine supplies are scarce but recommended focusing on key transmitters to protect the susceptible population when supplies are plentiful and other interventions are ongoing. The study in [6] addressed the cost-effectiveness of combining vaccination with social distancing. Further research includes [7], which investigated disease control through combined vaccination, isolation, and treatment strategies. The authors of [8] explored the efficacy of single-dose vaccines under conditions of low transmission and high single-dose efficacy (SDE), showing that such a strategy could prevent up to 22% more deaths compared to a two-dose strategy prioritising the elderly. The impact of various control strategies on SARS-CoV-2 and simultaneous influenza infections was analyzed in [9]. Additionally, the implications of limited vaccine supplies and different vaccination rates and efficacies on virus transmission were explored in several studies, including those that assume a limited availability of effective COVID-19 rapid tests [10]. Urban et al. [11] assessed five vaccination approaches considering restricted vaccine access, daily rates, and efficacy. Finally, Zhou et al. [12] investigated the spatial heterogeneity of COVID-19 transmission in urban areas, optimising vaccine distribution strategies based on spatial priorities.

Central to this discourse is the concept of classical herd immunity, rooted in mathematical models that delineate the relationship between vaccination coverage and epidemic control. Defined as $h_C = 1-1/R_0$, where R_0 denotes the basic reproduction number, classical herd immunity underscores the pivotal role of vaccination in curtailing transmission dynamics [13].

However, the influence of age structure on epidemic transmission dynamics cannot be overstated. Age-dependent variations in susceptibility, transmission, and disease severity, as evidenced by empirical studies, underscore the need for tailored vaccination strategies that account for demographic nuances [14]. While younger individuals may exhibit lower risks of severe illness, they can serve as significant vectors of viral spread within communities. Conversely, older populations are at height-ened risk of severe outcomes, necessitating prioritized vaccination efforts to safeguard vulnerable cohorts [15–17].

Age-related factors extend beyond susceptibility and severity to encompass vaccine response dynamics, necessitating tailored dosing regimens and booster strategies to optimise long-term protection, particularly in older adults [18, 19]. Consequently, a comprehensive vaccination approach must intricately navigate these age-related nuances to optimise population-level immunity and mitigate the overall impact of COVID-19 [20]. Existing studies [20–22] have elucidated the role of age heterogeneity in achieving herd immunity thresholds and optimising vaccination prioritization; the transition to an endemic state underscores the need for sustained, long-term vaccination strategies. This necessitates a nuanced understanding of immunity loss dynamics, vaccine distribution logistics, and healthcare system constraints across

Addressing these challenges requires an interdisciplinary approach that integrates mathematical modeling and optimisation [23, 24]. Optimal control theory offers a principled framework for devising vaccination strategies that balance epidemiological efficacy with practical constraints. Previous research [6, 21] has explored short-term vaccination optimisation under resource constraints taking into consideration the age structure of the population. In [6], the authors presented an age-structured model to optimise vaccination and distancing for Brazil. However, several limitations arise. The vaccination compartment, condensed into a single equation, neglects the nuanced interplay of infection and recovery among vaccinated individuals. Additionally, assuming vaccination solely prevents infection without affecting transmission dynamics oversimplifies the complex relationship between vaccination and transmission. In [21], we provided a model to optimise the short-term vaccination strategy for the three age distributions where the vaccinated compartment undergoes the same epidemiological process of infection with different parameters. This study extends this paradigm to investigate the long-term implications of age and immunity loss on vaccination strategies.

In this study, we leverage mathematical optimisation techniques to formulate vaccination strategies tailored to the age demographics and immunity dynamics of three distinct age distributions. By categorizing age distributions into expansive, stationary, and constrictive pyramids, represented respectively by Senegal, USA, and Tunisia, we illuminate the profound influence of population age structure and immunity loss on disease transmission dynamics and optimal age-dependent vaccination [25].

In fact, we explore the impact of varying maximal vaccination rates on optimal vaccination strategies, ranging from uniform rates across all age groups to age-dependent approaches reminiscent of World Health Organization (WHO) guidelines [26]. Through rigorous mathematical analysis and numerical simulations, we elucidate the implications of these strategies on the optimal age-dependent vaccination strategy and the number of critical infections and vaccination frequencies, offering actionable insights for policymakers and healthcare professionals alike.

The remainder of this paper is structured as follows: Section 2 presents the formulation of the optimal control problem and underlying assumptions. In Section 3, we conduct mathematical analysis to establish the well-posedness of the model and characterise the optimal control. Section 4 delves into the analysis and discussion of numerical simulation results, while Section 5 concludes with key insights and avenues for future research.

2. Model formulation

diverse countries.

2.1. Epidemiological model description

This paper extends previous work [21] to explore the influence of age distribution and immunity loss on the optimal vaccination strategies over the long term. In our previous work [21], the model was developed to optimise an age-dependent vaccination in the short term, accounting for vaccine shortages. Building upon this foundation, we now introduce modifications to the model to incorporate immunity loss and vaccination across all compartments except for severely infected and deceased individuals.

In fact, similarly to the previous work, the population is divided into n age classes, with each age class $i, 1 \le i \le n$ divided into vaccinated and unvaccinated groups. Considering that vaccination reduces infection and death rates rather than stopping it, it is assumed that both the vaccinated and unvaccinated groups go through the same epidemiological process: individuals leave the susceptible compartments S_i (respectively vaccinated S_i^{ν}), toward the exposed E_i (respectively vaccinated E_i^{ν}) through contact with either exposed or infected individuals, whether they are vaccinated or not and of any age class j. The exposed can either develop or not a severe form of infection moving to one of the infected compartments I_{1i} (respectively vaccinated I_{1i}^{v}) for the non-severe form and I_{2i} (respectively vaccinated I_{2i}^{ν}) for the severe form. Subsequently, they either move toward the recovered compartments R_i (respectively vaccinated R_i^{v}) or the dead one D_i . However, we assume that for each age group *i*, all epidemiological compartments except severely infected and dead are subjected to vaccination at a timedependent rate τ_i . Additionally, we incorporate the concept that a segment of recovered individuals, both from vaccinated and unvaccinated groups, lose their immunity with respective rates ψ_1 and ψ_2 , and revert to the susceptible compartment. As both vaccine-induced immunity and infection-induced one are lost with similar time scales, we choose not to differentiate between the types and merely use the term immunity regardless of the gaining way. This aspect is depicted in Figure 1.

The model proposed is given by:

$$\dot{S}_{i} = -\sum_{j=1}^{n} p_{j} \left(\beta_{i} \frac{E_{j}}{N_{j}} + \beta_{i}^{v} \frac{E_{j}^{v}}{N_{j}} + \alpha_{1i} \frac{I_{1j}}{N_{j}} + \alpha_{2i} \frac{I_{2j}}{N_{j}} + \alpha_{1i}^{v} \frac{I_{2j}}{N_{j}} + \alpha_{2i}^{v} \frac{I_{2j}^{v}}{N_{j}} \right) d_{ij} S_{i} - \tau_{i} S_{i} + \psi_{1} R_{i} + \psi_{2} R_{i}^{v}$$
(2.1)

$$\dot{E}_{i} = \sum_{j=1}^{n} p_{j} \left(\beta_{i} \frac{E_{j}}{N_{j}} + \beta_{i}^{v} \frac{E_{j}^{v}}{N_{j}} + \alpha_{1i} \frac{I_{1j}}{N_{j}} + \alpha_{2i} \frac{I_{2j}}{N_{j}} + \alpha_{1i}^{v} \frac{I_{2j}}{N_{j}} + \alpha_{2i}^{v} \frac{I_{2j}^{v}}{N_{j}} + \alpha_{2i}^{v} \frac{I_{2j}^{v}}{N_{j}} \right) d_{ij} S_{i} - \tau_{i} E_{i} - (q_{1i} + q_{2i}) E_{i}$$
(2.2)

$$\dot{I}_{1i} = q_{1i}E_i - \tau_i I_{1i} - r_{1i}I_{1i}$$
(2.3)

$$I_{2i} = q_{2i}E_i - r_{2i}I_{2i} - \mu_i I_{2i}$$
(2.4)

$$\dot{R}_i = r_{1i}I_{1i} + r_{2i}I_{2i} - \tau_i R_i - \psi_1 R_i$$
(2.5)

$$\dot{S}_{i}^{\nu} = \tau_{i}S_{i} - \delta \sum_{j=1}^{n} p_{j} \left(\beta_{i} \frac{E_{j}}{N_{j}} + \beta_{i}^{\nu} \frac{E_{j}^{\nu}}{N_{j}} + \alpha_{1i} \frac{I_{1j}}{N_{j}} + \alpha_{2i} \frac{I_{2j}}{N_{j}} + \alpha_{1i}^{\nu} \frac{I_{1j}}{N_{j}} + \alpha_{2i}^{\nu} \frac{I_{2j}^{\nu}}{N_{j}} \right) d_{ij}S_{i}^{\nu}$$
(2.6)

$$\dot{E}_{i}^{\nu} = \delta \sum_{j=1}^{n} p_{j} \left(\beta_{i} \frac{E_{j}}{N_{j}} + \beta_{i}^{\nu} \frac{E_{j}^{\nu}}{N_{j}} + \alpha_{1i} \frac{I_{1j}}{N_{j}} + \alpha_{2i} \frac{I_{2j}}{N_{j}} + \alpha_{1i}^{\nu} \frac{I_{1j}^{\nu}}{N_{j}} + \alpha_{2i}^{\nu} \frac{I_{2j}^{\nu}}{N_{j}} \right) d_{ij} S_{i}^{\nu} + \tau_{i} E_{i} - (q_{1i}^{\nu} + q_{2i}^{\nu}) E_{i}^{\nu}$$
(2.7)

$$\dot{I}_{1i}^{v} = q_{1i}^{v} E_{i}^{v} + \tau_{i} I_{1i} - r_{1i} I_{1i}^{v}$$
(2.8)
$$\dot{I}_{i}^{v} = c_{i}^{v} E_{i}^{v} - r_{i} I_{i}^{v} - c_{i}^{v} I_{i}^{v}$$
(2.9)

$$\dot{P}_{2i} - q_{2i}L_i - q_{2i}L_i - \mu_i q_{2i}$$

$$\dot{P}_{2i} - \mu_i q_{2i}L_i - \mu_i q_{2i}$$
(2.9)

$$\dot{\mathbf{K}}_{i} = \mathbf{I}_{1i}\mathbf{I}_{1i} + \mathbf{I}_{2i}\mathbf{I}_{2i} + \mathbf{I}_{i}\mathbf{K}_{i} - \boldsymbol{\psi}_{2}\mathbf{K}_{i}$$
(2.10)

$$D_i = \mu_i I_{2i} + \mu_i^v I_{2i}^v \tag{2.11}$$

subject to positive initial conditions:

 $S_i(0) > 0, E_i(0) > 0, I_{1i}(0) > 0, I_{2i}(0) > 0, R_i(0) > 0, S_i^v(0) \ge 0, E_i^v(0) \ge 0, I_{1i}^v(0) \ge 0, I_{2i}^v(0) \ge 0$ and $R_i^v(0) \ge 0$, for all $1 \le i \le n$.

The parameters of the model for all $1 \le i \le n$ are:

• $(d_{ij})_{1 \le i,j \le n}$ is the contact matrix, and each coefficient $d_{ij} > 0, \forall i, j \in \{1, ..., n\}$, denotes the proba-

bility that an individual of age *i* contacts an individual of age *j*.

- N_i is the total number of individuals of age class *i* given by: $N_i = S_i + E_i + I_{1i} + I_{2i} + R_i + S_i^v + E_i^v + I_{i1}^v + I_{i2}^v + R_i^v + D_i$.
- p_i is the probability of an individual to be of age class *i*. (i.e., $p_i = \frac{N_i}{N(0)}$), where $N(0) = \sum_{i=1}^n N_i$ is the total population.
- $\beta_i, \beta_i^v, \alpha_{1,i}, \alpha_{2,i}, \alpha_{1,i}^v$ and $\alpha_{2,i}^v$ are the infection rates of $E_i, E_i^v, I_{1i}, I_{2i}, I_{1i}^v$ and I_{2i}^v , respectively.
- δ denotes the infection reduction rate.
- $q_{1i}, q_{2i}, q_{1i}^{\nu}, q_{2i}^{\nu}$ are the rates at which an exposed develops severe or non-severe forms of infection, respectively, for vaccinated and unvaccinated groups. Noteworthy the rate of developing a severe infection is higher among the non-vaccinated than the vaccinated *i.e.*, $q_{2i} > q_{2i}^{\nu}$.
- r_{1i} , r_{1i}^{ν} , r_{2i} and r_{2i}^{ν} are the recovery rates for vaccinated and unvaccinated individuals, respectively.
- μ_i and μ_i^v are disease-induced mortality rates for the vaccinated and unvaccinated, respectively.

All parameters of the model are assumed to be positive constants. Additionally, the demographic process is not taken into account for this model. Therefore, the population size for each age class i, N_i , and the total population size, N, are supposed constant.





2.2. Objective function

Considering that vaccines do not stop infection, but rather reduce the rate of severe forms of it among the population, and that we are setting a long-term vaccination schedule, we aim to minimise the number of severe infections (vaccinated or not) and the cost of vaccination by controlling the agedependent vaccination rate. Therefore, the objective function is given by:

$$J(\tau) = \int_0^T A_1 \left(\|I_2(t)\|_2^2 + \|I_2^{\nu}(t)\|_2^2 \right) + A_2 \|\tau(t)\|_2^2 dt.$$
(2.12)

Mathematical Biosciences and Engineering

Volume 21, Issue 6, 6372–6392.

where:

- $I_2 = (I_{2i})_{1 \le i \le n}$ is the vector of unvaccinated critical infected and $I_2^{\nu} = (I_{2i}^{\nu})_{1 \le i \le n}$ is the vector of vaccinated critical infected individuals subject to ((2.1)–(2.11))
- $\tau = (\tau_i)_{1 \le i \le n}$ is the control variables vector.
- *T* denotes the time horizon of control.
- A_1 and A_2 are cost balancing factors and $|| ||_2$ stands for the Euclidean norm.

Minimising this function results in minimising the number of critical infections among both vaccinated and unvaccinated individuals, $||I_2(t)||_2^2 + ||I_2^v(t)||_2^2$, with a minimal vaccination effort, $||\tau(t)||_2^2$. Deaths are not explicitly present in the objective function, as minimising the number of critically infected individuals in hospitals has a direct impact on the minimisation of the number of disease-induced deaths. Due to the limitations that face various healthcare systems, we assume that there is a maximal threshold for the vaccination rate. And since we aim to control the vaccination rate according to age, we assume that the maximal vaccination rates are age-dependent, and then $0 \le \tau_i(t) \le \tau_i^{max}$ for all $i \in \{1, ..., n\}$ and for all $t \in [0, T]$.

This problem can be written as follows:

$$\min_{\tau \in U} J$$

subject to ((2.1)-(2.11)) where

$$U = \{\tau = (\tau_1, \dots, \tau_n) \in l^{\infty}([0, T], \mathbb{R}^n) : 0 \le \tau_i(t) \le \tau_i^{max} \,\forall i \in \{1, \dots, n\} \text{ and } \forall t \in [0, T]\}$$
(2.13)

3. Mathematical analysis

3.1. Positivity and boundedness of the solutions

To ensure that the model is biologically meaningful, it is important to prove that all solutions with non-negative initial conditions will remain non-negative at all positive times. The following proposition shows that the model is suitable to study living populations. Let $X(t) = (X_1(t), \ldots, X_n(t))$ be the vector of state variables, where

$$X_{i}(t) = (S_{i}(t), E_{i}(t), I_{1i}(t), I_{2i}(t), R_{i}(t), S_{i}^{v}(t), E_{i}^{v}(t), I_{1i}^{v}(t), I_{2i}^{v}(t), R_{i}^{v}(t), D_{i}(t)) = (X_{i}^{j})_{1 \le j \le 11}$$

for all $i \in \{1, \ldots, n\}$ and for all t > 0.

Proposition 1. For any non-negative initial condition $X_i(0)$, the solution $X_i(t)$ remains positive, for all t > 0 and $i \in \{1, ..., n\}$. Furthermore, the set Ω given by:

$$\Omega = \{X = (X_1, \dots, X_n) \in (\mathbb{R}^{11})^n : 0 < X_i^j \le N_i(0) \forall 1 \le j \le 11 \text{ and } 1 \le i \le n\}$$

is positively invariant under system ((2.1)–(2.11)).

Hint of the proof. [1)]

Existence and uniqueness of the solution: The right-hand side functions of the system are all of class C^1 . Thus, the mere application of the Cauchy-Lipchitz theorem yields the existence and uniqueness of the solution.

- Positivity: For all 1 ≤ j ≤ 11, if X_i^j = 0 and all other variables are positive i.e., X_i^k ≥ 0 for k ≠ j, X_i^j ≥ 0. Then, the vector field is always either pointing to the inside of the positive region or invariant on each of the previously mentioned hyperplanes. Consequently, the solution can never leave the positive region.
- **Boundedness:** For all $t \ge 0$, $X_i^j(t) \le N_i(t) = \sum_{j=1}^{11} X_i^j(t)$. And since $\dot{N}_i(t) = 0$, one has N_i constant

and consequently equals $N_i(0)$ as the age classes are supposed constant. Then, one has $X_i^j(t) \le N_i(0)$.

This proposition deals with both mathematical and biological existence problems. The first part makes sure that the model is well-posed in terms of having a solution whereas the second part of it deals with its suitability for application to living populations.

3.2. Optimal control problem analysis

Considering the loss of immunity, we formulate an optimal control problem to find the optimal vaccination rate under given constraints.

The optimal control problem consists of finding a piece-wise continuous control

$$\tau^*(t) = (\tau_1^*(t), \tau_2^*(t), \dots, \tau_n^*(t))$$

and the associated state variables $X^* = (S_i^*, E_i^*, I_{1i}^*, I_{2i}^*, R_i^*, S_i^{\nu*}, E_i^{\nu*}, I_{1i}^{\nu*}, I_{2i}^{\nu*}, R_i^{\nu*}, D_i^*)$, for $i \in \{1, \dots, n\}$ to minimise the given objective functional (2.12), i.e.,

$$J(\tau^*) = \min_{\tau \in \mathcal{U}} J(u) = \min_{\tau \in \mathcal{U}} \int_0^T \|I_2(t)\|_2^2 + \|I_2^v(t)\|_2^2 + \|\tau(t)\|_2^2 dt.$$

subject to ((2.1)-(2.11)) and their initial conditions.

The following proposition concerns the existence of an optimal control for the vaccination.

Proposition 2. There exists an optimal control variable $\tau^* \in \mathcal{U}$ such that $J(\tau^*) = \min_{\tau \in \mathcal{U}} J(\tau)$, subject to the controlled system ((2.1)–(2.11)).

Hint of the proof. To prove the existence of the optimal control, we use the results by Fleming and Rishel (1975) and by Lukes (1982) [27]. One can easily verify that:

- 1) The set of controls and corresponding state variables is nonempty.
- 2) The admissible set U is convex and closed.
- 3) The right-hand side of the state variables system ((2.1)–(2.11)) is bounded by a linear function in the state and control variables.
- 4) The integrand of the objective functional *J* is convex on \mathcal{U} and there exist constants $\omega_1 > 0, \omega_2 > 0$ and $\rho > 1$ such that

$$J(\tau) \le \omega_2 + \omega_1 \left(\sum_{i=1}^n |\tau_i|^2\right)^{\frac{\rho}{2}}$$

where $\rho = 2, \omega_1 = \frac{\epsilon}{2\sqrt{n}\tau_{max}} + 1, \omega_2 = \frac{\epsilon}{2} \text{ and } \epsilon = \sum_{i=1}^n (I_{2i}^2(0) + I_{2i}^{v2}(0)) e^{-2r_{2i}T}.$

Mathematical Biosciences and Engineering

Volume 21, Issue 6, 6372-6392.

The previous proposition answers the mathematical requirements of not characterising the control before ensuring its existence. Once the existence of optimal control has been proven, we use the minimum principle of Pontryagin [28], which transforms the optimality problem into a problem of minimisation of the Hamiltonian H with respect to the controls τ , to characterise this optimal control.

Therefore, we define the Hamiltonian given by:

$$H(t,\tau,X,\lambda) = <\lambda(t), \dot{X}(t) > +A_1(||I_2(t)||_2^2 + ||I_2^{\nu}(t)||_2^2) + A_2||\tau(t)||_2^2.$$

Proposition 3. Given an optimal control $\tau^* \in \mathcal{U}$ and solution $X^* = (S_i^*, E_i^*, I_{1i}^*, I_{2i}^*, R_i^*, S_i^{\nu*}, E_i^{\nu*}, I_{1i}^{\nu*}, I_{2i}^{\nu*}, R_i^{\nu*}, D_i^{\nu*}), i = (1, ..., n)$ of the corresponding state system ((2.1)–(2.11)), there exists a vector of adjoint variables $\lambda = (\lambda_i^1, ..., \lambda_i^{11})$, for all $i \in \{1, ..., n\}$, satisfying:

$$\begin{cases}
\frac{d\lambda_{i}^{1}}{dt} = \lambda_{i}^{1} \left(\sum_{j=1}^{n} p_{j} F_{ij} d_{ij} + \tau_{i}\right) - \lambda_{i}^{2} \sum_{j=1}^{n} p_{j} F_{ij} d_{ij} - \lambda_{i}^{6} \tau_{i} \\
\frac{d\lambda_{i}^{2}}{dt} = \lambda_{i}^{2} \left(\tau_{i} + q_{1,i} + q_{2,i}\right) - \lambda_{i}^{3} q_{1,i} - \lambda_{i}^{4} q_{2,i} - \lambda_{i}^{7} \tau_{i} + \frac{\beta_{i}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{3}}{dt} = \lambda_{i}^{3} \left(\tau_{i} + r_{i}^{1}\right) - \lambda_{i}^{5} r_{i}^{1} - \lambda_{i}^{8} \tau_{i} + \frac{\alpha_{1,i}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{4}}{dt} = -2A_{1} I_{2,i} + \lambda_{i}^{4} \left(r_{i}^{2} + \mu_{i}\right) - \lambda_{i}^{5} r_{i}^{2} - \lambda_{i}^{11} \mu_{i} + \frac{\alpha_{2,i}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{5}}{dt} = -\lambda_{i}^{1} \psi_{1} + \lambda_{i}^{5} \left(\tau_{i} + \psi_{1}\right) - \lambda_{i}^{10} \tau_{i} \\
\frac{d\lambda_{i}^{6}}{dt} = \delta \sum_{j=1}^{n} p_{j} F_{i,j} d_{i,j} \left(\lambda_{i}^{6} - \lambda_{i}^{7}\right) \\
\frac{d\lambda_{i}^{7}}{dt} = \lambda_{i}^{7} \left(q_{1,i}^{v} + q_{2,i}^{v}\right) - \lambda_{i}^{8} q_{1,i}^{v} - \lambda_{i}^{9} q_{2,i}^{v} + \frac{\beta_{i}^{v}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{8}}{dt} = r_{i}^{1} \left(\lambda_{i}^{8} - \lambda_{i}^{10}\right) + \frac{\alpha_{1,i}^{v}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{4}}{dt} = -2A_{1} I_{2,i}^{v} + \lambda_{i}^{9} \left(r_{i}^{2} + \mu_{i}^{v}\right) - \lambda_{i}^{10} r_{i}^{2} - \lambda_{i}^{11} \mu_{i}^{v} + \frac{\alpha_{2,i}^{v}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{4}}{dt} = r_{i}^{1} \left(\lambda_{i}^{8} - \lambda_{i}^{10}\right) + \frac{\alpha_{1,i}^{v}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{4}}{dt} = -2A_{1} I_{2,i}^{v} + \lambda_{i}^{9} \left(r_{i}^{2} + \mu_{i}^{v}\right) - \lambda_{i}^{10} r_{i}^{2} - \lambda_{i}^{11} \mu_{i}^{v} + \frac{\alpha_{2,i}^{v}}{N} d_{i,i} \left(\lambda_{i}^{1} S_{i} - \lambda_{i}^{2} S_{i} + \delta\lambda_{i}^{6} S_{i}^{v} - \delta\lambda_{i}^{7} S_{i}^{v}\right) \\
\frac{d\lambda_{i}^{4}}{dt} = -2A_{1} I_{2,i}^{v} + \lambda_$$

with transversality conditions

$$\lambda_i^j(T) = 0, i \in \{1, \dots, n\}, j \in \{1, \dots, 11\}.$$

Furthermore, the optimal control $\tau^* = (\tau_1, \dots, \tau_n)$ is given by:

$$\tau_{i}^{*} = \min\left\{\max\left\{0, \frac{1}{2A_{2}}\left[S_{i}(\lambda_{i}^{1} - \lambda_{i}^{6}) + E_{i}(\lambda_{i}^{2} - \lambda_{i}^{7}) + I_{1,i}(\lambda_{i}^{3} - \lambda_{i}^{8}) + R_{i}(\lambda_{i}^{5} - \lambda_{i}^{10})\right]\right\}; \tau_{i,max}\right\}, \forall i \in \{1, \dots, n\}.$$
(3.2)

Mathematical Biosciences and Engineering

Volume 21, Issue 6, 6372–6392.

Hint of the proof. The adjoint equations and transversality conditions are determined using Pontryagin's minimum principle such that:

$$\dot{X}_{i}^{j}(t) = \frac{dH}{d\lambda_{i}^{j}}(t, X(t), \lambda(t), \tau(t))$$
(3.3)

$$\dot{\lambda}_i^j(t) = -\frac{dH}{dX_i^j}(t, X(t), \lambda(t), \tau(t))$$
(3.4)

where $X_i = (X_i^j)_{1 \le j \le 11}^T = (S_i, E_i, I_{1i}, I_{2i}, R_i, S_i^v, E_i^v, I_{1i}^v, I_{2i}^v, R_i^v, D_i)^T$, $\lambda_i = (\lambda_i^j)^T$ for all $i \in \{1, ..., n\}$. By minimising the Hamiltonian H, for all $i \in \{1, ..., n\}$, the optimal control τ_i^* can be obtained from

By minimising the Hamiltonian H, for all $i \in \{1, ..., n\}$, the optimal control τ_i^* can be obtained from the optimality condition:

$$\frac{dH}{d\tau_i} = 0$$

In fact, since the optimal control $\tau^* \in \mathcal{U}$, it is easy to obtain τ^* in the form given in (3.2).

4. Numerical simulations and discussion

The objective of this section is to investigate the impact of different age distributions and immunity loss on the optimal vaccination strategy of model ((2.1)–(2.11)). Age distributions are classified into three major categories *. The first category is expansive, where most of the population is constituted of young individuals and the older classes get, the fewer individuals there are. The second type is the stationary age distribution. Such distribution tends to remain constant, exhibiting a more or less equal distribution for all age categories. And the third type is the constrictive one, where middle-aged individuals form the majority of the population and fewer individuals are to be found in the oldest and youngest classes. For that, for the numerical simulations, we have chosen three countries presenting the previous three different age distributions: Senegal, the USA, and Tunisia (See Figure 2 for age class distribution per country).



Figure 2. The age pyramid for Senegal, Tunisia and the United States of America.

For each one of the studied countries, the initial conditions were estimated based on data from its own population[†]. All populations were divided into six age groups : $G_1 = [0, 14], G_2 = [15, 29],$

^{*}https://populationeducation.org/what-are-different-types-population-pyramids/

 $^{^{\}dagger}https://raw.githubusercontent.com/CSSEGIS and Data/.../time_series_covid19_confirmed_global.csv$

 $G_3 = [30, 44], G_4 = [45, 59], G_5 = [60, 74], and G_6 \ge 75, according to data from WHO database[‡].$

Moreover, considering that these countries have distinct economical classification, their healthcare systems face disparate limitations. Hence, we would add to the age distribution the impact of maximal age-dependant vaccination rate $\tau_{i,max}$, for all $1 \le i \le n$. We study four scenarios according to the value of $\tau_{i,max}$. For the first three scenarios $\tau_{i,max} = \tau_{max}$, $\forall 1 \le i \le 6$, is constant for all age classes and corresponds respectively to 0.005, 0.01, and 0.05, whereas the fourth is age-dependant as it is given by a vector ($\tau_{i,max}$)_{1 \le i \le 6} = (0,005; 0.005; 0.01; 0.01; 0.05; 0.05).

The problem is solved numerically using the fourth-order Runge–Kutta method. The used parameters are listed in the Appendix A.

In what follows, we present the numerical simulations' results in two parts. The first one corresponds to the effect of constant maximal vaccination rate (τ_{max}) and age distribution on the optimal vaccination strategy for the three studied age distributions, while the second part displays the impact of setting an age-dependent maximal vaccination rate.

4.1. Impact of constant maximal vaccination rate

In what follows, we evaluate the optimal vaccination strategy and its impact on the number of critical infections for the three countries and with three maximal vaccination rates, $\tau_{max} = 0.005$, 0.01, and 0.05. To do so, we plot the optimal vaccination (3.2) resulting from solving the optimal control problem ((2.1)–(2.11)) (Figure 3), and the total number of critical infections without vaccination, $\tau = 0$, and with an optimal vaccination (Figure 4).

For $\tau_{max} = 0.005$, the optimal vaccination strategies for the three countries exhibit consistent features: they remain constant over time and are set equal to the maximal vaccination rate across all age classes. Indeed, Figure 3(a)–(c) shows that the age distribution does not affect the determination of an age-tailored optimal vaccination strategy.

This induces a first peak of critical infections among the unvaccinated individuals that mimics the no-vaccination peak (see Figure 4(a)-(c)). This peak is lowest in Senegal, gets higher for Tunisia, and is highest for the USA. However, in the case of Senegal, no second peak is observed, unlike Tunisia where its beginnings start to appear by the end of the year, or the USA where it is more clearly noticed. In fact, in the absence of vaccination, we notice that all countries register a high peak of infection followed by a lower one.

For $\tau_{max} = 0.01$, (see Figure 3(d)–(f)), a slight difference in the vaccination strategies of the various age classes becomes noticeable. For the three countries, vaccination starts at the same time at its maximum simultaneously for all age classes. However, differences start to be noted as the various age classes start to show disparate decreasing times of vaccination curve corresponding to various age classes.

We observed that the optimal strategy is to vaccinate the most representative age groups for both Senegal and Tunisia. Indeed, in Senegal (see Figure 3(d)), older age groups are vaccinated at a maximum rate for a shorter duration compared to younger age groups. The duration of vaccination increases with age. Similarly, in Tunisia (refer to Figure 3(e)), a similar pattern is observed, with the duration of vaccination overlapping across age groups. However, priority is given to middle-aged groups (classes 2–4), which are the most represented in the country's constrictive population pyramid.

[‡]https://covid19.who.int



(a) Optimal vaccination strategy for (b) Optimal vaccination strategy for (c) Optimal vaccination strategy for Senegal with $\tau_{max} = 0.005$ Tunisia with $\tau_{max} = 0.005$ USA with $\tau_{max} = 0.005$



(d) Optimal vaccination strategy for (e) Optimal vaccination strategy for (f) Optimal vaccination strategy for Senegal with $\tau_{max} = 0.01$ Tunisia with $\tau_{max} = 0.01$ USA with $\tau_{max} = 0.01$



(g) Optimal vaccination strategy for (h) Optimal vaccination strategy for (i) Optimal vaccination strategy for Senegal with $\tau_{max} = 0.05$ Tunisia with $\tau_{max} = 0.05$ USA with $\tau_{max} = 0.05$

Figure 3. Optimal administration of the vaccine to the different age classes from the youngest class 1 to the oldest class 6 of each country and with different constant values of maximal vaccination rate $\tau_{max} = 0.005, 0.01, 0.05$. We observe that for $\tau_{max} = 0.005$, optimal vaccination strategies across three countries show uniformity over time, aligning with maximal vaccination rates for all age classes, suggesting minimal age-based impact on strategy. As τ_{max} increases to 0.01, slight variations emerge, with priority given to younger age classes in Senegal and middle age classes in Tunisia. $\tau_{max} = 0.05$ exacerbates disparities, emphasizing consistent prioritization of the most represented age class within each population, particularly evident with higher thresholds.

However, in the case of USA (see Figure 3(f)), despite doubling the maximal vaccination rate, we continue to observe a simultaneous and constant vaccination equal to its maximal rate over the whole vaccination time horizon.

Moreover, the first peak of unvaccinated critical infected individuals decreases considerably for the case of Senegal and Tunisia and a little less for the case of the USA, while the second peaks vanish for the three countries (see Figure 4(d)-(f)).

For a maximal vaccination rate of $\tau_{max} = 0.05$ (Figure 3(g)–(i)), the previously forming disparities



(a) Total number of critical infec- (b) Total number of critical infec- (c) Total number of critical infections for Senegal with and without tions for Tunisia with and without tions for USA with and without concontrol for $\tau_{max} = 0.005$ control for $\tau_{max} = 0.005$ trol for $\tau_{max} = 0.005$



(d) Total number of critical infec- (e) Total number of critical infec- (f) Total number of critical infections tions for Senegal with and without tions for Tunisia with and without for USA with and without control for control for $\tau_{max} = 0.01$ control for $\tau_{max} = 0.01$ $\tau_{max} = 0.01$



(g) Total number of critical infec- (h) Total number of critical infec- (i) Total number of critical infections tions for Senegal with and without tions for Tunisia with and without for USA with and without control for control for $\tau_{max} = 0.05$ control for $\tau_{max} = 0.05$ $\tau_{max} = 0.05$

Figure 4. The impact of optimal vaccination at different constant maximal vaccination rates on critical infections across three countries, compared to no vaccination. Orange curves represent infections without vaccination, while green and purple curves show unvaccinated and vaccinated critical infections for $\tau_{max} = 0.005, 0.01, 0.05$. In the absence of vaccination (orange curves), we notice that all countries register a high peak of infection followed by a lower one. With $\tau_{max} = 0.005$, an initial peak of severe infections among the unvaccinated resembles the no-vaccination peak, followed by a decrease in the second peak. Increasing τ_{max} to 0.01 notably reduces the initial peak, particularly for expansive and constrictive distributions, and eliminates the second peak across all distributions. At $\tau_{max} = 0.05$, the expansive distribution exhibits no peaks, while the constrictive and stationary distributions display minimal peaks.

become even more visible.

Like in previous cases, Senegal and Tunisia exhibit similar behaviors in their vaccination strategies,

with a shorter common vaccination period observed for both countries (see Figure 3(h),(g)). However, in Senegal, the optimal vaccination of the fifth and sixth age classes (the oldest) start decreasing since t = 0.

Additionally, in the case of the USA, optimal vaccination curves begin to show a similar pattern to those of Senegal and Tunisia. However, priority is given to the fourth age class (aged between 45 and 59), with the next level of priority granted to age classes 2 and 3. This results in no peaks at all in Senegal, while a little bump being observed for Tunisia, and a considerably small peak noticed in the USA (see Figure 4(g)–(i)).

These findings underscore the common feature of emphasizing the vaccination of the most represented class among the population. This, also, implies that vaccinating the young age classes can protect old classes for old and middle-aged populations. However, the impact of the chosen maximal vaccination rate differs according to the age distribution considered. The older the population is, the higher the vaccination rate needed to control critical infections is.



(g) Optimal vaccination proportion (h) Optimal vaccination proportion (i) Optimal vaccination proportion for Senegal with $\tau_{max} = 0.05$ for Tunisia with $\tau_{max} = 0.05$ for USA with $\tau_{max} = 0.05$

Figure 5. Comparison between optimal vaccination proportions (in green) and corresponding age classes (in blue) across the three countries for varying values of τ_{max} . At lower τ_{max} values, vaccination proportions closely align with age distributions, prioritising overrepresented age groups. As τ_{max} increases, vaccination proportions rise proportionally, maintaining this trend. Notably, in expansive age distributions, younger age groups receive height-ened priority compared to older ones.

Figure 5 displays a comparison between the age distribution of vaccinated individuals for the opti-

mal strategy (in green) and the corresponding age distribution class (in blue) for each country and for the three different values of the maximal vaccination rate τ_{max} . For $\tau_{max} = 0.005$, the optimal vaccination proportions show a similar appearance to that of the age distribution, with frequencies that are nearly equal to twice the corresponding age classes for the three countries (see Figure 5(a)–(c)) with an emphasis on the over-represented class of the distribution. Setting higher $\tau_{max} = 0.01$ and $\tau_{max} = 0.05$ does not change this aspect but rather increases the proportions to three or four times the age class (see Figure 5(d)–(f) and (i)).

However, we notice that for the expansive age distribution and for $\tau_{max} = 0.05$, the frequency of vaccinating the youngest age class is now twice that of the oldest class. The three constant maximal vaccination rates τ_{max} generated an age-distribution-like schedule, prioritising not only vulnerable age classes but also giving significant attention to classes with the highest numbers of individuals.

4.2. Impact of age-dependent maximal vaccination rate

The inclusion of an age-dependant maximal vaccination rate generates different responses, and the optimal strategy shows a drastic change (see Figure 6). We notice a similar vaccination strategy for Senegal and Tunisia (see Figure 6(a),(b)) despite the difference in their age distributions. In fact, as in the constant maximal vaccination scenario, we start by vaccinating all classes with a maximal rate; however, the common vaccination period is shorter for the two oldest classes. The vaccination of these two oldest classes stops nearly at the middle of the time horizon. Conversely, the vaccination curves for the USA show a constant vaccination rate equal to its maximum over the whole time horizon for all age classes, which can be explained by the stationary aspect of the age distribution (see Figure 6(c)).



(a) Optimal vaccination strategy for (b) Optimal vaccination strategy for (c) Optimal vaccination strategy for Senegal with age-dependent τ_{max} Tunisia with age-dependent τ_{max} USA with age-dependent τ_{max} **Figure 6.** Optimal administration of the vaccine to the different age classes for each country with age-dependent maximal vaccination rate $\tau_{max} = (0.005, 0.005, 0.01, 0.01, 0.05, 0.05)$. In the case of Senegal and Tunisia, all age classes are initially vaccinated at the maximum rate but over different periods. In the case of the USA, a constant vaccination rate equal to its maximum is maintained throughout the entire time horizon across all age groups.

The effect of optimal vaccination on the number of critical infections (see Figure 7(a)–(c)) shows, for the three studied countries, a peak higher than that of the scenario with $\tau_{max} = 0.05$ (see Figure 4(g)–(i)), the scenario with the lowest peak among the four considered scenarios. Despite this fact, the age-dependent vaccination strategy manages to contain infections similarly to the maximal vaccination rate of 0.01. Therefore, it represents a viable alternative to the latter strategy, which helps avoid overwhelming medical staff and expanding the vaccination cost. These results suggest that, in the long term, age-dependent vaccination rates can be an effective alternative to maximal vaccination strategies





(a) Total number of critical infec- (b) Total number of critical infec- (c) Total number of critical infections for Senegal with and without tions for Tunisia with and without tions for USA with and without concontrol for τ_{max} control for τ_{max} trol for τ_{max}

Figure 7. The impact of the optimal vaccination at an age-dependent maximal vaccination rate $\tau_{max} = (0.005, 0.005, 0.01, 0.01, 0.05, 0.05)$ across three countries, compared to no vaccination. Orange curves represent infections without vaccination, while green and purple curves show unvaccinated and vaccinated critical infections. For the three countries, one peak of severe infections among the unvaccinated that resembles the no-vaccination peak is observed. The second peak observed for the no-vaccination case is eliminated for the three countries.

In the last figure (see Figure 8), we plot the proportion of the vaccinated individuals for each age class for the three countries using an age-dependent maximal vaccination rate. We observe a huge alteration in the proportions compared to the constant maximal vaccination rate. Vaccination frequencies across all countries exhibit a trend of decreasing frequency as individuals become younger, with the most pronounced effect observed in populations with stationary age distributions, where vaccination frequency for older classes is twice that of younger ones. These results indicate that the age-dependent maximal vaccination strategy aligns with a WHO-like approach.



(a) Optimal vaccination proportion (b) Optimal vaccination proportion (c) Optimal vaccination proportion for Senegal with age-dependent τ_{max} for Tunisia with age-dependent τ_{max} for USA with age-dependent τ_{max}

Figure 8. Comparison between optimal vaccination proportions (in green) and corresponding age classes (in blue) across three countries for an age-dependent $\tau_{max} = (0.005, 0.005, 0.01, 0.01, 0.05, 0.05)$. Notably, vaccination frequencies decrease as age decreases, with the most prominent effect observed in countries with stationary age distributions, where vaccination frequencies for older age groups are double those for younger ones.

5. Conclusions

In this endeavour, our objective is to investigate the impact of age demographics and immunity loss on long-term vaccination strategies. To achieve this goal, we established a controlled age-structured SEIRD mathematical model, ensuring its biological validity. This model provided the framework needed to scrutinise the repercussions of varying maximal vaccination rates on disease propagation across distinct age distributions.

Having established the mathematical validity of our model, we formulated an optimal control problem and utilised the Pontryagin maximum principle to delineate the optimal course of the control. Subsequently, through numerical simulations applied to three disparate age distributions—expansive, constrictive, and stationary—we contrasted the outcomes of three distinct constant maximal vaccination rates (0.005, 0.01, and 0.05) with an age-dependent vaccination rate for each country.

We showed that the vaccination is most effective when applied to old and young age groups. However, our study shows also that in the long-term vaccination approach, two optimal strategies emerge depending on the chosen maximal vaccination rate: constant or age-dependent. In our previous work [21], the short-term vaccination strategy prioritised also the oldest age classes as well as the most significant age class, irrespective of any constraints. This was also confirmed by the age-structured model presented in [6], a study conducted for Brazil, which has a constrictive pyramid.

Furthermore, in this work, the results revealed that the optimal vaccination strategy is highly influenced by the choice of the maximal vaccination rate. When examining constant maximal vaccination rates, the significance of prioritising age groups based on population demographics became evident, with more attention given to the most represented age classes within each population. Moreover, augmented maximal vaccination rates led to a reduction in the number of critically infected individuals across all countries.

Conversely, an age-dependent approach yielded a WHO-like strategy, with infection peaks of greater magnitude than the maximal vaccination rate of 0.05 for all the nations under consideration. This suggests the need for careful consideration of both age distribution and vaccination strategy when designing public health interventions.

In summation, this research study contributes valuable insights into the field of optimal vaccination planning, emphasising the importance of considering the nuances of age groups and the impact of maximal vaccination efforts. The conclusions drawn from this research can inform policymakers and decision-makers in their ongoing efforts to reduce the spread of infectious diseases. As perspectives to this study, further exploration into the impact of memory on the dynamics of the model and the optimal control would provide valuable avenues for future research.

Use of AI tools declaration

The authors declare they have not used Artificial Intelligence (AI) tools in the creation of this article.

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Conflict of interest

On behalf of all authors, the corresponding author states that there is no conflict of interest.

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A. Appendix

The parameters of the model are either based on literature or estimated using the data of the countries [32, 33]. All these parameters are defined in Table A1.

The infection rates by unvaccinated type-one infected α_{1i} , $1 \le i \le n$, were estimated, with respect to the value of R_0 ($R_0 = 2.19$), in a way to satisfy the following equations:

$$\alpha_{1i} + \alpha_{2i} + \beta_i + \alpha_{1i}^v + \alpha_{2i}^v + \beta_i^v = 1, \ 1 \le i \le n.$$
(A.1)

Daramatar Nacorintion	•	Countre	
		Country	
	Tunisia	Senegal	USA
Population size [29]	11818619	16743928	331002647
Age class distribution P_i [29]	24, 3%, 21, 3%,	42,5%,27,3%,	18%, 20%,
	23,4%,17,7%,	16.7%, 8.6%,	20%, 19%,
	10,2%, 3,1%	4%,0.7%	16%, 7%
Infection rate α_{1i}	0,326797	0,326797	0,326797
Infection rate α_{2i}	$0, 1. lpha_{1i}$	$0, 1.\alpha_{1i}$	$0, 1.\alpha_{1i}$
Infection rate β_i	$(3/5).\alpha_{1i}$	$(3/5).\alpha_{1i}$	$(3/5).lpha_{1i}$
Infection rate β_i^v	$0, 8.eta_i$	$0, 8, \beta_i$	$0, 8.eta_i$
Infection rate α_{1i}^{ν}	$0, 8.\alpha_{1i}$	$0, 8. \alpha_{1i}$	$0, 8.\alpha_{1i}$
Infection rate $\alpha_{2i}^{\overline{v}}$	$0, 8. \alpha_{2i}$	$0, 8.\alpha_{2i}$	$0, 8.\alpha_{2i}$
Infection probability among δ	0.3	0.3	0.3
Incubation rate λ [33]	0.2	0.2	0.2
Hospitalization rate η [30]	0.0027, 0.0137, 0.03,	0.0027, 0.0137, 0.0301,	0.0027, 0.0137, 0.03005,
	0.0792, 0.1585, 0.7158	0.0792, 0.1584, 0.7158	0.0792, 0.1585, 0.7158
Infection rate $q1$	0.19999, 0.19995, 0.19988	0.19999, 0.19995, 0.19988	0.19999, 0.19995, 0.19988
	0.19968, 0.19937, 0.19714	0.19968, 0.19937, 0.19714	0.19968, 0.19937, 0.19714
Infection rate q^2	$1.09 \ 10^{-5}, 5.46 \ 10^{-5}, 1.202 \ 10^{-4}$	$1.09 \ 10^{-5}$, $5.46 \ 10^{-5}$, $1.202 \ 10^{-4}$	$1.09 \ 10^{-5}, 5.46 \ 10^{-5}, 1.202 \ 10^{-4}$
	$3.16910^{-4}, 6.33910^{-4}, 2.863410^{-3}$	$3.169\ 10^{-4}, 6.339\ 10^{-4}, 2.8634\ 10^{-3}$	$3.169 \ 10^{-4}, 6.339 \ 10^{-4}, 2.8634 \ 10^{-3}$
Infection rate q_1^{v}	0.199998, 0.199995, 0.199988	0.199998, 0.199995, 0.199988	V
•	0.199968, 0.199937, 0.199714	0.199968, 0.199937, 0.199714	0.199968, 0.199937, 0.199714
Infection rate q_2^{v}	$1.09 \ 10^{-6}, 5.46 \ 10^{-6}, 1.202 \ 10^{-5}$	$1.09 \ 10^{-6}, 5.46 \ 10^{-6}, 1.202 \ 10^{-5}$	$1.09 \ 10^{-6}, 5.46 \ 10^{-6}, 1.202 \ 10^{-5}$
ı	$3.169 \ 10^{-5}, 6.339 \ 10^{-5}, 2.8634 \ 10^{-4}$	$3.169\ 10^{-5}, 6.339\ 10^{-5}, 2.8634\ 10^{-4}$	$3.169 \ 10^{-5}, 6.339 \ 10^{-5}, 2.8634 \ 10^{-4}$
Immunity loss rate ψ [33]	1/180	1/180	1/180
Recovery rate of I_{1i} , r_{1i} [33]	1/8	1/8	1/8
Recovery rate of I_2 , r_2 [17]	1/13.5, 1/13.5, 1/14,	1/13.5, 1/13.5, 1/14,	1/13.5, 1/13.5, 1/14,
	1/14.5, 1/14.5, 1/15	1/14.5, 1/14.5, 1/15	1/14.5, 1/14.5, 1/15
Natural mortality rate [34]	0, 0, 0.0001, 0.0005,	0, 0, 0.0001, 0.0005,	0, 0, 0.0001, 0.0005,
•	0.034, 0.072	0.034, 0.072	0.034, 0.072
Basic reproduction number	2.1	2.1	2.1
Balancing cost factors A_1 and A_2	1000, N/2	1000, N/2	1000, N/2
τ_{max} age-dependent	0.005, 0.005, 0.01, 0.01, 0.05, 0.05	0.005, 0.005, 0.01, 0.01, 0.05, 0.05	0.005, 0.005, 0.01, 0.01, 0.05, 0.05
Time horizon T	1 year	1 year	1 year

Table A1. Epidemiological and demographic data.

Mathematical Biosciences and Engineering

	0 0		1 0
Parameter description		Country	
	Tunisia	Senegal	USA
Date of first case	Mars 2, 2020	Mars 2, 2020	26 February 2020
Date of first vaccination t_0	Mars 13, 2021	Feb 22, 2021	10 December 2020
Number of cumulative cases at t_0	242,124	16,529	16,217,128
Number of active case at t_0	24,554	6451	9,669,611

Table A2. Data at the beginning of the vaccination campaign.

Initial condition per country	Tunisia	Senegal	USA
$S(t_0)$	$P_i - (272124)P_i / Pop$	$P_i - (23529)P_i / Pop$	$P_i - (25916216)P_i / Pop$
$E(t_0)$	$(30000P_i/Pop$	$(7000P_i/Pop$	$(1000000P_i/Pop$
$I_1(t_0)$	$(1 - 0.02\eta)(24554)P_i/Pop$	$(1 - 0.02\eta)(6451)P_i/Pop$	$(1 - 0.02\eta)(9669611)P_i/Pop$
$I_2(t_0)$	$0.02\eta(24554)P_i/Pop$	$0.02\eta(6451)P_i/Pop$	$0.02\eta(9669611)P_i/Pop$
$R(t_0)$	217570 <i>P_i</i> / <i>Pop</i>	$10078P_i/Pop$	6246605 <i>P_i</i> / <i>Pop</i>
$S^{\nu}(t_0)$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)
$E^{\nu}(t_0)$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)
$I_{1}^{v}(t_{0})$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)
$I_2^{\nu}(t_0)$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)
$\overline{R^{\nu}(t_0)}$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)
$D(t_0)$	(0,0,0,0,0,0)	(0,0,0,0,0,0)	(0,0,0,0,0,0)

 Table A3. The initial condition of the various countries.

$$\alpha_{2i} = 0.1 \,\alpha_{1i}, \, \beta_i = 0.6 \,\alpha_{1i}, \, \forall \, 1 \le i \le n \tag{A.2}$$

$$\alpha_{ki}^{\nu} = 0.8 \,\alpha_{ki} \,, \, k = 1, 2 \,, \, \beta_{i}^{\nu} = 0.8 \,\beta_{i} \,, \, \forall \, 1 \le i \le n \tag{A.3}$$

Noteworthy that R_0 is not directly included in this study and is only used for parameter estimation. More details are available in [21].

For the three different countries, Tunisia, Senegal, and the USA, the contact matrices, *D*, used are estimated for 6 age groups based on [31].

The initial conditions used are estimated based on real data of the populations of the different countries mentioned, at the beginning of the vaccination campaign see Table A2. And using the age distribution of each country, the initial conditions for each infectious state and age per country are then defined in Table A3.



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