
An ODE-PDE model for Covid-19 pandemic with vaccination and loss of immunity: well-posedness and simulation

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Résumé

The health crisis caused by the COVID-19 pandemic showed once again the need to model the propagation of epidemics. Compartmental models remain the dominant tool for meeting this need. The best-known compartmental model to date is the SIR model introduced by Kermack and McKendrick in 1927. Since then, a variety of SIR-like models have been studied.

In this talk, a SEHIRVD model is considered. In this model, at each time the population is divided into seven distinct compartments: the group S of Susceptible individuals, the group E of Exposed individuals that are infected but not infectious, the group H of Hospitalized individuals, the group I of Infectious individuals, the group R of Recovered individuals that are cured from the disease, are still immune and have not been vaccinated since they recovered, the group V of Vaccinated individuals that still hold an immunity against infection and the group D of Deceased individuals.

Our model is structured in age of infectivity and in age of immunity. The age of infectivity a is used in compartment I. Formally, $I(t,a)$ is the proportion of individuals at time t that have been infected for a time a . The age of immunity a' intervenes in compartments R and V. It counts the time spent in these compartments, where individuals are immune to infection. The model includes a progressive loss of immunity for both recovered and vaccinated individuals, parametrized by the age of immunity a' .

The system of equations driving the model is a system of coupled ordinary differential equations and partial differential equations with integral terms.

In this work, we state theoretical results on the well-posedness and bounds for the solution of the system.

Then we propose to solve numerically the equations with a discretized scheme, based on

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radial basis function-generated difference (RBF-FD) method to compute the differential operators in age a and a' . The integrals are approximated numerically with a Quasi-Monte Carlo (QMC) quadrature. The time evolution is calculated through a semi-implicit Euler scheme of order one.

Finally simulations allow to compare different scenarios.

Mots-Clés: Epidemiology, sir, age structured, vaccination, rbf, quasi monte carlo