

Mathematical Modeling of COVID-19 Transmission by a k Phases SEIR Model

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Background

- Mathematical models can provide insights on the spread of infectious diseases, such as the novel SARS-CoV-2 (COVID-19).
 - This work applied a SEIR epidemiological compartmental model (susceptible-exposed-infected-recovered) with k phases to predict the actual spread of the COVID-19 virus.

Methods

- During the mathematical modeling of the COVID-19 spread, four parameters of the SEIR model (Fig. 1) were obtained by international experiences: the incubation period = 3.7 in days, the proportion of critical cases = 0.05, the overall case-fatality rate = 0.023, and the asymptomatic proportion of COVID-19 = 0.18.
- These values can be modified for a specific region, but, the critical step in the prediction of COVID-19 by the model is the value of R0 (the basic reproduction number) and T_infectious (the infectious period, in days).
- To solve this difficult, R0 and T_infectious were calculated by mathematical constrained optimization, a numerical method. Differently from a statistical modelling, a numerical method is a type of mathematical modelling that is not dependent on a probability distribution.
- In this numerical modelling process the objective function that measures the model error is minimized with respect to R0 and T_infectious in the presence of constraints on those variables. For R0, constraints are valid range of values ($0.5 \leq R0 \leq 20$). For T_infectious (days), constraints also are related to its range of values ($2 \leq T_{\text{infectious}} \leq 14$).
- A Solver from Excel or NEOS Server, for example can be used for finding numerically minimum of a function Z, that represents the sum of squared errors between each COVID-19 new case observed in one day, and the COVID-19 cases predicted by the SEIR model (Fig. 2).

Fig 1 - The SEIR model for COVID-19.

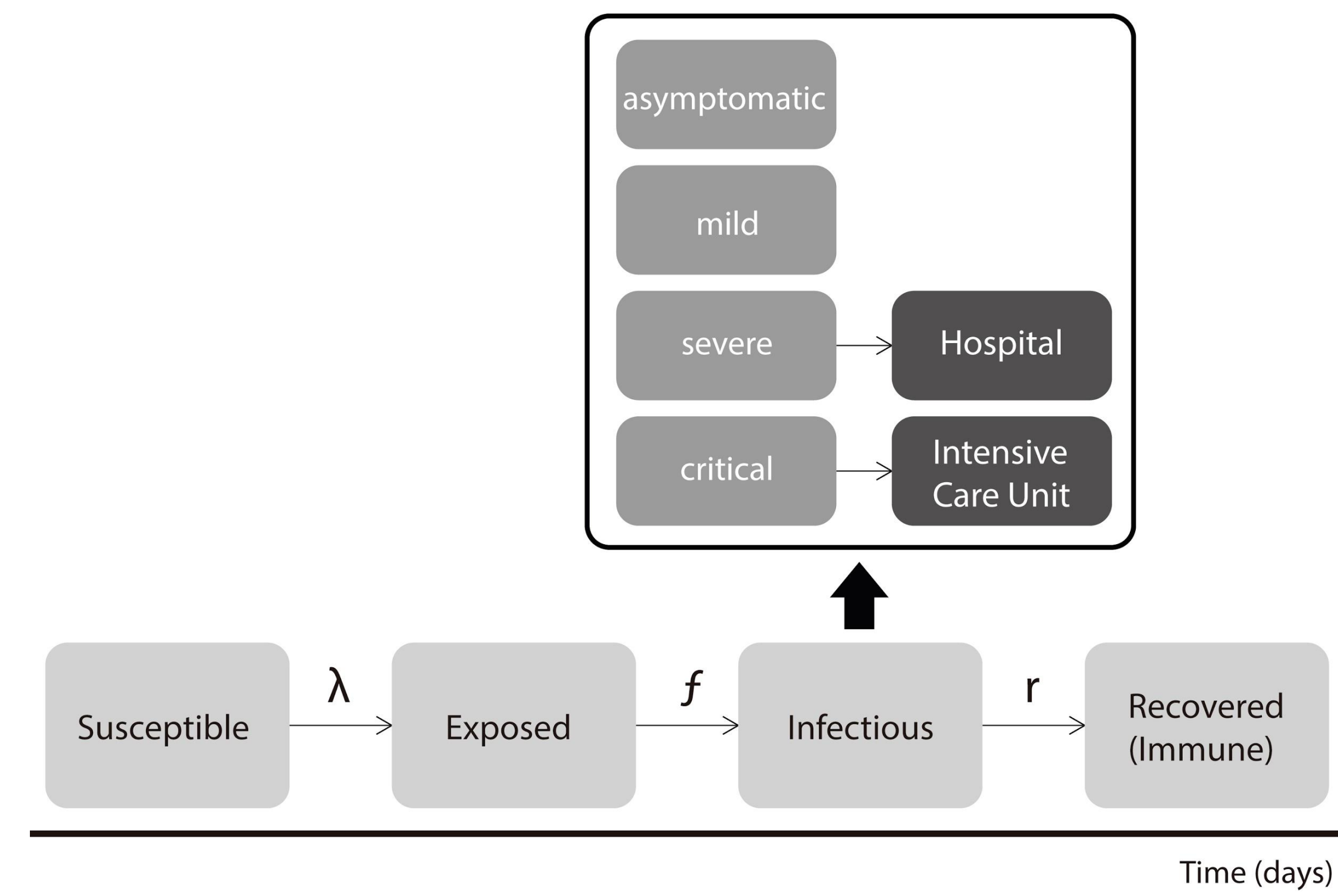


Fig 2 - Algorithm for the SEIR model applied to COVID-19: initialization and calculation of new COVID-19 cases day-by-day.

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Algorithm COVID
Read N (population's size)
Read T_phase_I (duration of the first phase of the epidemic (days))
Read R0_phase_I (basic reproduction number for COVID-19 from the first phase of the epidemic)
Read T_infectious_phase_I (the infectious period from the first phase of the epidemic (days))
Read T_phase_II (duration of the second phase of the epidemic (days))
Read R0_phase_II (basic reproduction number for COVID-19 from the second phase of the epidemic)
Read T_infectious_phase_II (the infectious period from the second phase of the epidemic (days))
Read R0_phase_III (basic reproduction number for COVID-19 from the third phase of the epidemic)
Read T_infectious_phase_III (the infectious period from the third phase of the epidemic (days))
T_incubacao = 3.7 (COVID-19 the incubation period, in days)
p_CTI = 0.05 (spectrum of disease: proportion of critical COVID-19 cases)
letalidade = 0.023 (the overall case-fatality rate)
p_asintomaticos = 0.18 (asymptomatic proportion of COVID-19)
f = 1/T_incubacao (the rate at which individuals move from the latent class to the infected class)
t = 0 (first day = "zero day")
t_Max = 180 (last day of simulation: 180 days after the first COVID-19 case)
Susceptivel[t] = (N - COVID) (susceptible individuals)
Freinfec[t] = 0 (exposed or latent patients)
COVID[t] = 1 (infected patients or COVID-19 cases)
Imunes[t] = 0 (recovered or immune patients)
CTI[t] = 0 (critical cases)
Obitos[t] = 0 (case-fatality)

Repeat
  t = t + 1
  If (t < T_phase_I) then
    ecr = R0_phase_I/T_infectious_phase_I (effective contact rate for phase 1)
    recupera = 1/T_infectious_pahse_I (recovery rate for phase 1)
  else se (t < T_phase_II) then
    ecr = R0_phase_II/T_infectious_phase_II (effective contact rate for phase 2)
    recupera = 1/T_infectious_pahse_II (recovery rate for phase 2)
  else
    ecr = R0_phase_III/T_infectious_phase_III (contact rate for phase 3)
    recupera = 1/T_infectious_pahse_III (recovery rate for phase 3)

  Beta = ecr/N (transmission rate)
  Susceptivel[t] = Susceptivel[t-1] - Beta*COVID[t-1]*Susceptivel[t-1]
  Freinfec[t] = Freinfec[t-1] + Beta*COVID[t-1]*Susceptivel[t-1] - f*Freinfec[t-1]
  COVID[t] = COVID[t-1] + f*Freinfec[t-1] - recupera*COVID[t-1]
  Imunes[t] = Imunes[t-1] + recupera*COVID[t-1]
  CTI[t] = p_CTI*COVID[t]
  Obitos[t] = letalidade*Imunes[t]*p_asintomaticos
  Write t, Susceptivel[t],Freinfec[t],COVID[t],Imunes[t],CTI[t],Obitos[t]
  If (t = t_Max)
    Then break
  End If
End Repeat
End Algorithm
  
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Results

- Based on real COVID-19 data until April 11, R0 and the infectious periods were calculated for the third phase of the epidemic by the mathematical constrained optimization used for finding the numerically minimum of a function Z (Fig. 3) for 15 countries (Fig. 4).

Conclusion

- The k phases SEIR model proved to be a useful to measure the COVID-19 transmission in a City, State or Country. A four-phase model can be applied to fit a scenario with a new second COVID wave.

Fig 3 - Mathematical constrained optimization model.

$$\text{Minimize } Z = \sum_{i=1}^D |I_i - \hat{I}_i| \quad (1)$$

Subject to

$$\begin{aligned} &T_{\text{phase_I}} \leq D \\ &0.5 \leq R0_{\text{phase_I}} \leq 20 \\ &2 \leq T_{\text{infectious_phase_I}} \leq 14 \\ &T_{\text{phase_II}} \leq T_{\text{phase_I}} \\ &0.5 \leq R0_{\text{phase_II}} \leq 20 \\ &2 \leq T_{\text{infectious_phase_II}} \leq 14 \\ &0.5 \leq R0_{\text{phase_III}} \leq 20 \\ &2 \leq T_{\text{infectious_phase_III}} \leq 14 \end{aligned}$$

Where: I_i = number of COVID-19 new cases observed in a city, state or country during the day i;
 $\hat{I}_i$ = number of COVID-19 new cases predicted by the model in the day i;
 D = total days of the epidemic in a city, state or country;
 $T_{\text{phase_I}}$ = duration of the first phase of the epidemic (days);
 $T_{\text{infectious_phase_I}}$ = the infectious period from the first phase of the epidemic (days);
 $T_{\text{phase_II}}$ = duration of the second phase of the epidemic (days);
 $R0_{\text{phase_II}}$ = basic reproduction number for COVID-19 from the second phase of the epidemic;
 $T_{\text{infectious_phase_II}}$ = the infectious period from the second phase of the epidemic (days);
 $R0_{\text{phase_III}}$ = basic reproduction number for COVID-19 from the third phase of the epidemic;
 $T_{\text{infectious_phase_III}}$ = the infectious period from the third phase of the epidemic (days);

Fig 4 - Mathematical Modeling of COVID-19 Transmission by a 3 Phases SEIR Model in 15 countries.

