



Simulating the effects of mobility restrictions in the spread of SARS-CoV-2 in metropolitan areas



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Transforming our Collective Urban Future: Learning from Covid-19 ISUH International Society for Urban Health 16:30 - 17:30 EDT [O10] Pandemic lockdowns in the city: capturing the effects of mobility restrictions

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In Portugal, strict lockdown periods have been established since March 2020:

- Imitations on inter-municipal flows
- suspension of all non-essential travel
- compulsory remote work



Simulations of daily new infections in metropolitan areas for different mobility scenarios, to test the potential effect of travel restrictions on virus dissemination.

Metropolitan areas:

- > the most densely populated areas in the country
- intense commuting flows
- virus dissemination across municipalities

Study area



Lisbon (LMA)

- 18 municipalities
- 3015 km²
- 2 840 005 inhabitants (28%)
- 81% of commuters
- 16% public transport



Povoa Varzim Via Conde Santo Tirso Trofa Maia Valongo Paredes Porto VNGaia Gondomar Espinho SMFeira Arouca Vale Cambra Oliv Azemeis

Porto (PMA)

- 17 municipalities
- 2041 km²

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50 km

- 1 721 038 inhabitants (17%)
- 79% of commuters
- 11% public transport







Based on the regulations implemented by the Portuguese government, essential sectors are:

- ✓ Agriculture, fisheries, and forestry; extraction and transformation industry;
- ✓ electricity and gas supply; water supply, sanitation, and residues;
- ✓ construction; trade and auto repair;
- ✓ transport and storage;
- ✓ human health.



All population is initially susceptible	<pre># set up SEIR matrix SEIR <- matrix(nrow = locs_len, ncol = 4) # initiate an empty SEIR matrix colnames(SEIR) <- c("S", "E", "I", "R") # rename the vectors SEIR[, "S"] <- N_k # assign the number of successible people in each cell SEIR[, "E"] <- 0 # assign the number of exposed people in each cell SEIR[, "I"] <- 0 # assign the number of infected people in each cell SEIR[, "R"] <- 0 # assign the number of recovered people in each cell</pre>
First infections based on nr. cases PT 23/03	<pre># first infection # assume no all are infectious and no one in incubation period at the beginning first_infections <- (pop_cases\$origin == zone0_id) * zone0_infected str(first_infections) str(zone0_infected)</pre>
First exposed calculated with model	<pre>#Calculate first exposed based on sigma (rate exposed to infected=0.2) and first infections first_exposed <- first_infections / sigma_vec first_exposed</pre>
Update of SEIR matrix based on initial cases	<pre>#Update the SEIR matrix with first infections and first exposed #remove first infections and exposed from susceptible population SEIR[, "S"] <- SEIR[, "S"] - (first_infections + first_exposed) #add first infections to infected population SEIR[, "I"] <- SEIR[, "I"] + first_infections #add first exposed to exposed population SEIR[, "E"] <- SEIR[, "E"] + first_exposed - first_infections</pre>
<pre>#Define OD to use OD < aml_all # New E infected_mat <- replicate(locs_len, SEIR_nsim[, "I"]) OD_all_infected_ flow_all_infected_ flow_all_infected_ total_low_all_infected) print(paste0("Total infected full flow: ", total_flow_all_infected)) new_exposed < beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * flow_all_infected / (N_k + colsums(aml_all)) + #exposed by contacting with imported infectious cases beta_vec * SEIR_sim[, "S"] * SEIR_sim[, "S"], new_exposed) #make sure the N exposed is not bigger than total susceptible</pre>	
<pre># New I new_infected <- sigma_vec * SEIR_sim[, "E"] total_new_infected <- round(sum(new_infected, na.rm = T)) print(paste0("New infected: ", total_new_infected))</pre> Calculation of new infected based on new exposed	
<pre># New R new_recovered <- gamma_vec * SEIR_sim[, "I"] total_new_recovered <- round(sum(new_recovered, na.rm = T)) print(paste0("New recovered: ", total_new_recovered))</pre> Calculation of new	w recovered based on new infected
SEIR_sim[, "S"] <- SEIR_sim[, "S"] - new_exposed SEIR_sim[, "E"] <- SEIR_sim[, "E"] + new_exposed - new_infected SEIR_sim[, "I"] <- SEIR_sim[, "I"] + new_infected - new_recovered SEIR_sim[, "R"] <- SEIR_sim[, "R"] + new_recovered SEIR_sim <- ifelse(SEIR_sim < 0, 0, SEIR_sim)	ау





New infections in LMA

New infections in PMA

Stricter lockdown periods (vertical dashed lines) cover the periods between 23rd March and 4th May 2020, 28th November to 8th December 2020, and from 27th December 2020 until 31st March 2021

RESULTS - SEIR modeling by mobility scenario – official records



- ✓ Daily new cases Estimations vs oficial records (23 March 2020 to 26 October 2020)
- ✓ 2 scenarios: Unrestricted mobility & Restricted to Essential activities

New infections in LMA



New infections in PMA

Estimated values considering the upper and lower thresholds of the confidence interval (95%) of R_t (dashed lines). Official recorded cases of new infections are represented as points, daily values until the 6th July, and cumulative weekly values afterwards.



Lisbon (LMA)

Porto (PMA)





Lisbon (LMA)

Porto (PMA)







- ✓ The spread of the virus has responded to the mobility restrictions implemented. The number of new infections decreases after lockdown periods, with a certain delay
- ✓ Timeline of virus spread differs among metropolitan areas. PMA, closer to the scenario of mobility restricted to essential activities, in LMA closer to the unrestricted mobility scenario (lower threshold). Linked to secondary outbreaks in July 2020, especially in LMA, likely due to the ease of restrictions
- ✓ After Christmas, there was a substantial increase in the number of infected cases and fatalities, and the capacity of the health system was challenged. New infections started to decrease in mid-January 2021, following the implementation of mobility and social restrictions. Trends are captured by the simulations.
- ✓ In LMA, Lisbon absorbs most of the flows regardless of the scenario considered, higher number of cases. In PMA, Porto and VNGaia share the highest number of cases, depending on the scenario.

 Further work: How these restrictions can be adjusted to define suitable reopening strategies; how other social distancing and hygiene measures can be of complement; how the mobility conditions in metropolitan areas can be improved beyond the pandemic.













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