

Severe airport sanitarian control could slow down the spreading of COVID-19 pandemics in Brazil

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ABSTRACT

Background: We investigated a likely scenario of COVID-19 spreading in Brazil through the complex airport network of the country, for the 90 days after the first national occurrence of the disease. After the confirmation of the first imported cases, the lack of a proper airport entrance control resulted in the infection spreading in a manner directly proportional to the amount of flights reaching each city, following the first occurrence of the virus coming from abroad.

Methodology: We developed a Susceptible-Infected-Recovered model divided in a metapopulation structure, where cities with airports were demes connected by the number of flights. Subsequently, we further explored the role of the Manaus airport for a rapid entrance of the pandemic into indigenous territories situated in remote places of the Amazon region.

Results: The expansion of the SARS-CoV-2 virus between cities was fast, directly proportional to the city closeness centrality within the Brazilian air transportation network. There was a clear pattern in the expansion of the pandemic, with a stiff

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exponential expansion of cases for all the cities. The more a city showed closeness centrality, the greater was its vulnerability to SARS-CoV-2.

Conclusions: We discussed the weak pandemic control performance of Brazil in comparison with other tropical, developing countries, namely India and Nigeria. Finally, we proposed measures for containing virus spreading taking into consideration the scenario of high poverty.

Subjects Computational Biology, Ecology, Mathematical Biology, Virology, Statistics

Keywords SIR model, Metapopulation dynamics, Amazonia, Indigenous people, One-Ecohealth, SARS-CoV-2 pandemic

INTRODUCTION

The new disease COVID-19 has been spreading rapidly around the world since early January. It started in China at the end of 2019, been declared a “Public Health Emergency of International Concern” in 30 January and a pandemic in 11 March (*World Health Organization (WHO), 2020a; Zhu et al., 2020*). Its form of transmission is mainly through respiratory droplets of infected patients and contact with surfaces infected by aerosols (*Fathlzadeh et al., 2020*). However, the transmission dynamics has changed quickly in few months, with R_0 varying from 0.3 to 2.0 in some countries and close to 3.0 in others (*Lai et al., 2020a; World Health Organization (WHO), 2020a*). Large continental countries are likely to be very vulnerable to the occurrence of pandemics (*Morse et al., 2012; Dáttilo et al., 2020*). While the dissemination dynamics have varied between regions, country sanitary policies play a key role. For instance, two very large developing countries, India and Brazil, have a very different epidemical pattern. On March 18th, India had 137 cases and Brazil 621, as recorded in the Brazilian Ministry of Health (<https://covid.saude.gov.br/>, 2020) and John Hopkins (<https://gisanddata.maps.arcgis.com/>, 2020) monitoring sites dedicated to SARS-CoV-2 and COVID-19. From 17th to 18th March, Brazil had an increase of 31% in 1 day, with only four capitals exhibiting community transmission, which was the same in India. Nonetheless, a very distinct pattern in the ascending starting point for the reported disease exponential curve was observed in each country.

By enlarging the comparison to another developing tropical country in the Southern Hemisphere (thus, in the same season), we selected Nigeria since it was the first country to detect a COVID-19 case in sub-Saharan Africa (*World Health Organization (WHO), 2020d*). Nigeria reported eight confirmed cases during the same period of time (*Nigeria Centre for Disease Control, 2020*). Furthermore, Nigeria has a population similar to that of Brazil (201 million and 211.1 million, respectively, according to *United Nations, 2019*).

Both India (*Airports Authority of India, 2020*) and Nigeria (*Federal Airports Authority of Nigeria, 2020*) ensured severe entrance control, and close follow up of each confirmed case, as well as their living and working area, and people in contact with them, differently from Brazil. In Brazil, the Ministry of Health has developed a good monitoring network and a comprehensive preparation of the health system for the worst-case scenario. Nonetheless, apparently, the decisions from the Ministry of Health did not cover airport control, and only on March 19th, eventually too late, the government

decided to limit the number of flights coming from Europe or Asia. Hence, the entrance of potentially exposed passengers to SARS-CoV-2 in Brazil has been occurring with no control, at least until the aforementioned date. Moreover, after confirming that a person is infected with SARS-CoV-2, his/her monitoring is initiated but there is no monitoring of potential contacts.

For pandemic situations, the classical algebraic ecological models of species population growth from Verhulst, and species interaction models from Lotka-Volterra, are theoretical frameworks capable to describe the phenomenon and to propose actions to stop it (*Pianka, 2000*). In many aspects, social distancing is a way to severely reduce carrying capacity, that is, the resources available for the virus dissemination. This is the best action for within-city pandemic spreading of this new coronavirus (*Hellewell et al., 2020*), since the main form of transmission is direct contact between people or by contact with fomite, mainly in closed environments, such as classrooms, offices, etc. (*Rothe et al., 2020; Bedford et al., 2020*). Regardless of virulence, for a highly contagious virus such as SARS-CoV-2, the occurrence of the first case in a nation will result in a strongly and nearly uncontrollable exponential growth curve, depending only on the number of encounters between infected and susceptible people, and fueled by a high R_0 , which is ranging from 0.8 to 3.6 for COVID-19, depending on region and period analyzed (*Lai et al., 2020a, 2020b*).

On the other hand, the dynamics of disease spreading among cities are entirely distinct. This is a growing scientific area, an obliged interdisciplinary field, where neural network driven epidemiological phenomena is central (*Brockmann & Helbing, 2013*); however, it must be defined by metapopulation and closeness, as well as by an individual gene flow driven phenomenon (*Colizza & Vespignani, 2007*). In order to understand subpopulation flow between demes—which are cities—connected by roads or air flight networks, several complex details may be raised in the process of modelling such as reality (*Balcan et al., 2010*). For instance, probabilistic motivated effective distance, which is the actual proximity caused by business, culture, and investments between two cities, is much more important than real geography, especially for airlines network (*Brockmann & Helbing, 2013*). Nevertheless, for such theoretical approach to infection disease dissemination, an ecological concept is key: metapopulation (*Hanski, 1998*). However, ecologists are those who are least devoted to explore human–pathogen interaction at a global or continental scale. The ecological approach may focus on the most biological aspect of these dynamics, which is simply virus dissemination inside an infected person. As typically designed in Ecology, the first attempt will be based on the most neutral model (sensu *Hubbell, 2001*), that is, the directions taken through the most frequent and intensively used links.

In this work, we present an epidemiological model describing the free entrance of people coming from two highly infected countries with close links to Brazil: Italy and Spain. We show how SARS-CoV-2 had spread into the Brazilian cities by the international airports, and then to other, less internationally connected cities, through the Brazilian airport network. For exploring the dynamics of a continent size, nationwide spreading of

SARS-CoV-2, as it is the case of Brazil, we assumed cities connected by airports (simply cities hereafter) formed a metapopulation structure.

Each person in a city was taken as a component of a superorganism, that is, an interdependent entity where living individuals are not biologically independent between them in various subtle ways. By doing so, we dealt with cities as the sampling units, not the people, and, therefore, our model is slightly different from classic bolsonic models (Colizza & Vespignani, 2007; Nicolaidis et al., 2012). Flights coming from foreign countries with COVID-19 (namely Spain and Italy for this article) represent the probability of an external introduction of infection in each city. Additionally, we also further explored the vulnerability of the Amazon region, especially of those remote towns where indigenous and traditional communities predominate.

MATERIALS AND METHODS

In order to describe the pattern of air transportation and its role in the spreading of the disease, we built a Susceptible-Infected-Recovered (SIR) model (Hethcote, 1989; Anderson, 1991) split amongst the cities that are interconnected by flights. In this model, the population size inside each city is irrelevant. Moreover, the number of flights is highly correlated with city population ($R^2 = 0.76$; $p < 0.0001$) and, thus, a good proxy of city size. Similarly, to our purpose the time when the collective infection stage was reached inside each city was irrelevant. Thus, we assumed that the city was fully infected and became infectious to the whole system, and, therefore, became a source and not a sink of infection events, after a certain amount of arriving contaminated people sums up. Hence, the SIR model started having cities with only susceptible events and change of these to the infected stage was counted as proportion of the population. Infected events only appeared by migration, that is, travelers only from Italy and Spain, for sake of simplicity and proximity to the early facts.

After the first occurrence having been recorded in the country, infected people started to spread through the national airlines, and this spreading is proportional to the amount of infected people accumulating in a city (see the model explanation below). Because of that, the model causes a transient timing, when the amount of infected people arriving is greater than the community transmission. Afterwards, the city started to disseminate to other places by probability rules. As the transmission is quite likely to happen by the simple proximity person-to-person, due to the resistance of the virus in air droplets and surfaces, as hands or metals, we assumed that a simple encounter will cause infection. However, this became an inner trait of each city that after having local transmission, starts to transmit to other cities as explained above. In addition, the model assumes that there is no public control in people circulation after arriving of an infected person. Ethical approval was not necessary as human participants were not involved in the study.

We used a modified version of the SIR model, which took into account the topology of how the cities-demes were linked by domestic flights. In the SIR original model, the infection of susceptible cities occurs by probability β of a healthy being (S) encounters an

infected one (I). Conversely, the model has a probability of an infected one get recovered (R) given by a parameter γ . Analytically:

$$S_{t+1} = S_t - \frac{\beta}{N} S_t I_t$$

$$I_{t+1} = I_t + \frac{\beta}{N} S_t I_t - \gamma I_t$$

$$R_{t+1} = R_t + \gamma I_t$$

where the indexes t and $t + 1$ represent the present time and the next time, respectively, and $N = S + I + R$ is the total constant population. In this work, we proposed two modifications of the SIR model. The first one is related to the fact that we considered all the Brazilian cities that have an airport as subpopulations, meaning that each city has its own SIR variables set. Thus, we had S^i , I^i , and R^i where i was a given city. In our case study, $1 \leq i \leq 154$. The second important modification was related to the connections among the cities, which we used as a network of disease dissemination. This city network is given by the domestic flights among all the airports in Brazil, taking into account the number and the direction of flights. Therefore, we have a weighted network, where the weighted closeness centrality of each city was measured. Thus, the flights provided a dependent mechanism by which the between-cities connections caused a coupling of SIR equations based on the network flight structure. This coupling is provided by the last equation and is mediated by the new introduced parameter alpha. Using [Agência Nacional de Aviação Civil \(ANAC\) \(2020\)](#) data, it was possible to track all the domestic flights in Brazil ([Fig. 1](#)).

The modified version of SIR model is then described as follows:

$$S_{t+1}^i = S_t^i - \frac{\beta}{N} S_t^i (I_t^i + \bar{I}_t^i)$$

$$I_{t+1}^i = I_t^i + \frac{\beta}{N} S_t^i (I_t^i + \bar{I}_t^i) - \gamma (I_t^i + \bar{I}_t^i)$$

$$R_{t+1}^i = R_t^i + \gamma (I_t^i + \bar{I}_t^i)$$

where the upper index i indicates the city, and t the time. The term \bar{I}_t^i represents the infection added to the i th city due to traveling diseases, and it is calculated as follow:

$$\bar{I}_t^i = \alpha \sum_{j=0}^{154} k_{j,i} I_j$$

where $k_{j,i}$ is the number of flights departing at city j and arriving at city i , and α is a newly introduced parameter, which represents the fraction of traveling infected population. For the time, considering the geographic of Brazil and the time-lag for the dissemination

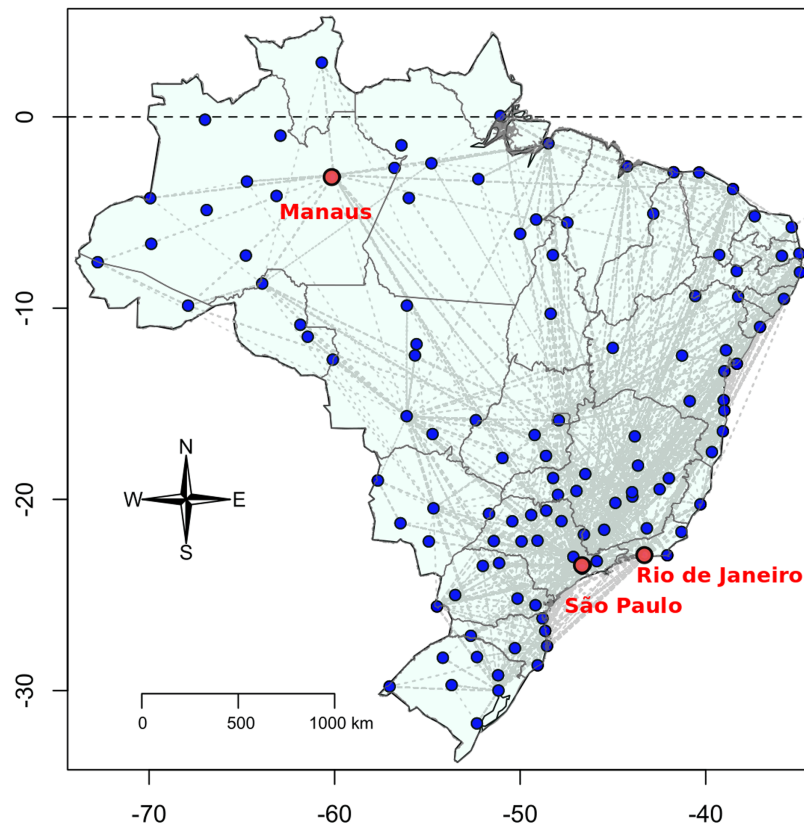


Figure 1 Brazilian flight network, taken from ANAC database.

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towards the most remote cities, we estimated 90 days of disease expansion and assumed γ as 0, in other words, no recovery. Despite the artificiality of this assumption, we considered that the amount of people still to be infected is larger than those recovered and, thus, becoming resistant. For instance, by 19th May the recovery rates in Brazil reached 85%, but this corresponded to only 100.5 thousand people around the country (www.worldometers.info/coronavirus/country/brazil/, 2020). Divided between each infected city, this makes mathematically small numbers, and thus resistance becomes demographically irrelevant to our output of early disease dissemination. Furthermore, by dealing with only the infected portion of the population (proportion of infected), we avoided the uncertainties related to the little known COVID-19 resistance development (*Li et al., 2020*).

The model was developed in C and is available as [Supplemental Material 1](#) (and the database as [Supplemental Material 2](#)). In addition, we also used a linear model to test whether those cities with higher city closeness centrality (i.e., important cities for connecting different cities within the Brazilian air transportation network) were more vulnerable to SARS-CoV-2 dissemination. The Beta parameter was defined by calibrating the model with real time series from Johns Hopkins Coronavirus Research Center (<https://coronavirus.jhu.edu/map.html>), which leads to $\beta = 0.3035$ ([Supplemental Material 1](#)). We used the value $\alpha = 0.0001$ based on the amount of flights

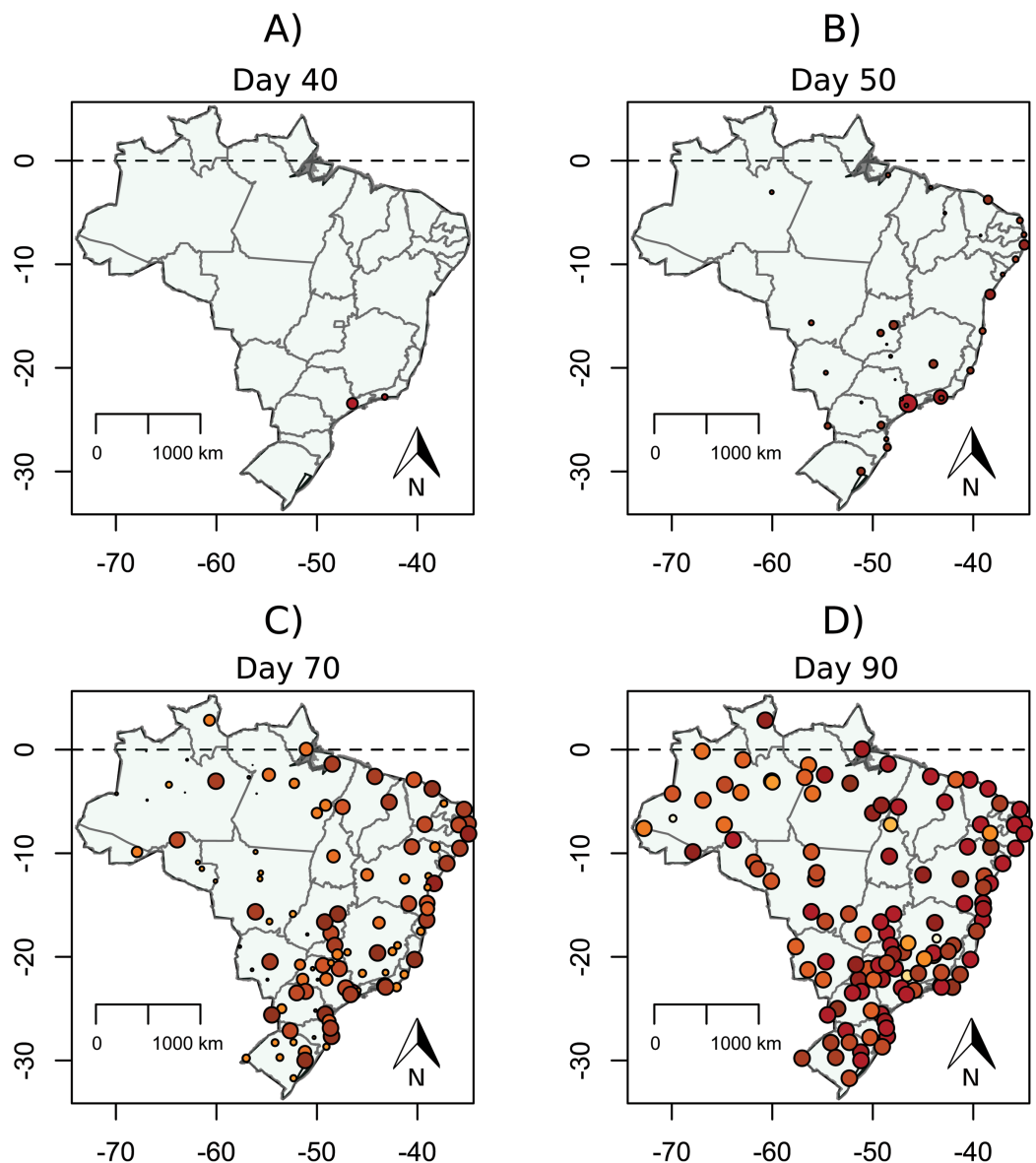


Figure 2 Proportion of infected population of each Brazilian city in 40 (A), 50 (B), 70 (C), and 90 (D) days. Circle colour temperature represents a gradient in percentage of the infected population. Circle size also reflects the size of the pandemics locally in the logarithm scale.

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needed from the peak of the disease in Italy until an infected person was recorded from that country in Brazil.

RESULTS

The expansion of the SARS-CoV-2 virus between cities was fast, directly proportional to the city closeness centrality within the Brazilian air transportation network. The disease spread from São Paulo and Rio de Janeiro to the next node-city by the flight network, and, in 90 days, virtually all the cities with airport(s) were reached; however, it occurred with a distinct intensity (Fig. 2; Supplemental Material 3). There was a clear pattern in the

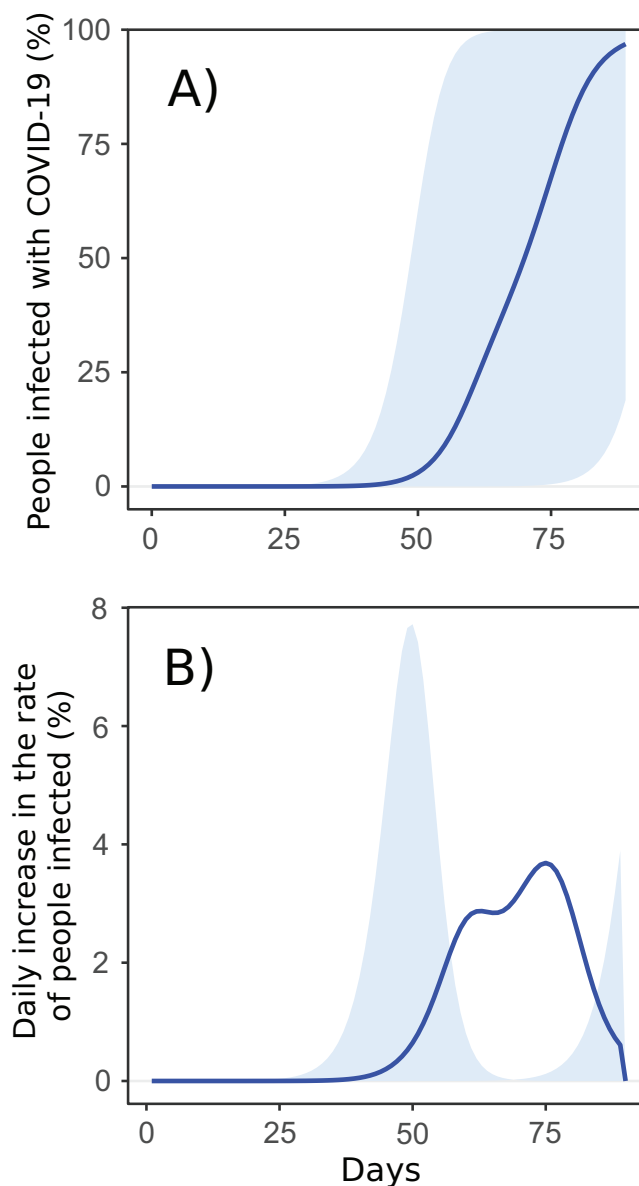


Figure 3 Proportion of infected people per cities until 90 days. (A) Cumulative increment rate. The blue line is the national average, and the shadow area is the summing up of minimum and maximum values of all the cities per time interval; (B) daily increment rate. The blue line is the average, showing the overall high rate of infection occurring from 50 to 80 days. Shadow shows the first and the highest peak in the hub cities, around 50 days, and, subsequently, a peripheric peak after 75 days.

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expansion of the pandemic, with a stiff exponential expansion of cases (measured as the cumulative percentage of infected people per city) for all the cities. On average, the model showed an ascendant curve starting at day 50 (around 15 April), with the most connected cities starting their ascendant curve just after 25 days, and the most isolated ones from day 75 (10th May; Fig. 3A). Looking at the daily increment rates, it is clear a fast and high peak of infections in the hub cities, happening around 50 days and, starting from 75 days, a new peripheric peak (Fig. 3B).

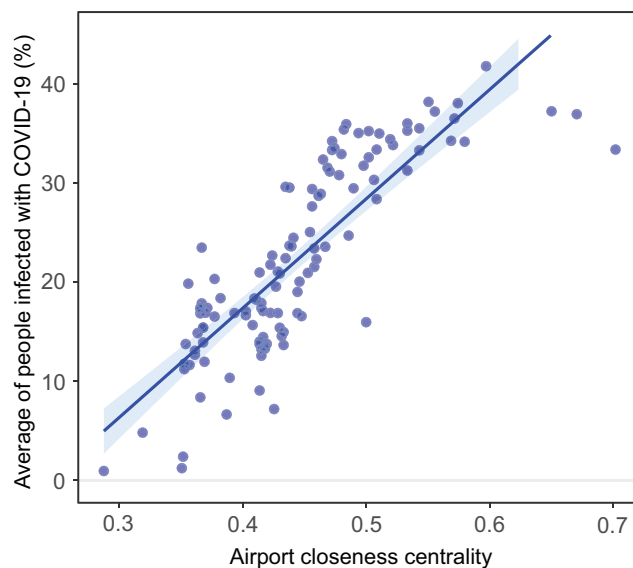


Figure 4 Airport closeness centrality within the Brazilian air transportation network, and its effect on the vulnerability of each city. Correlation between airport closeness centrality within the Brazilian air transportation network, and its effect on the vulnerability of each city (represented by the average of the percentage of cases per city for the whole 90 days running: $r^2 = 0.71$ $p < 0.00001$).

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The first ten cities to ascend infection rates (São Paulo, Rio de Janeiro, Salvador, Recife, Brasília, Fortaleza, Belo Horizonte, Porto Alegre, Curitiba, and Florianópolis) will actually reach this point about the same time, which is a concerning pattern for the saturation of the public health services. Moreover, this peak in those cities will saturate all the best hospitals in the country simultaneously.

Therefore, we defined the average proportion of infected people for the 90 days as a measure of the vulnerability to COVID-19 dissemination. Henceforth, we found the more a city shows closeness centrality within the air transportation network, the greater was its vulnerability to disease transmission (Fig. 4). This scenario confirmed the importance of a city connecting different cities within the Brazilian air transportation network and, thus, acting as the main driver for the pandemic spreading across the country.

Consequences for the Amazonian cities and indigenous people

Herein, we showed that an uncontrolled complex airport system made a whole country vulnerable in few weeks, allowing the virus to reach the most distant and remote places, in the most pessimistic scenario. According to our model, any connected city will be infected after 3 months. As the number of flights arriving in a city is the driver for the proportion of infected people, Manaus, which is a relevant regional clustering, was infected sooner. Indeed, on the 17th of March, Manaus was the first Amazonian city with confirmed cases (without community transmission yet, according to the Ministry of Health: <https://covid.saude.gov.br/>, 2020), and it is a node that is one or two steps to all the Amazonian cities. Thus, according to our model, Manaus may reach 1% of the infected

population by the 44th day, while, for instance, the far west Amazonian Tabatinga will take 61 days to reach the same 1% of the population infected. By day 60, Manaus may have an average of 50% of its population infected if nothing is done to prevent it. Tabatinga may also reach the aforementioned value by day 78, if nothing is done to avoid it. To sum up, within 46 days, all the Amazonian cities will have 1% of their population infected and a mean of 50% by day 70.

DISCUSSION

Our model suggests that Brazil must be prepared for an exponential rise in COVID-19 cases within 3 months from end of February, starting synchronously by the wealthiest cities. Such increase is expected based only on the dissemination among cities by the commercial airports and may get worse without the measures of social distancing proposed by the WHO (*Coelho et al., 2020*). The Country has failed to contain COVID-19 in airports and to closely monitor those infected people coming from abroad, as well as their living network. According to the Brazilian Airport Authority (*Agência Nacional de Aviação Civil (ANAC), 2020*), Brazil has the second-largest flight network in the world (just after the USA), with a total of 154 airports registered to commercial flights of which 31 are considered international. In comparison, airport control may be much easier to set up in Nigeria (31 airports of which only five are international: Federal Airport Authority of Nigeria, 2020). Nonetheless, with a population 6.4 times higher than Brazil (United Nations 1919), India, in turn, has a similar sized airport network to Brazil, harboring a total of 123 airports of which 34 are considered international (*Airports Authority of India, 2020*).

Nevertheless, the situation of COVID-19 in India is currently much milder than in Brazil, and it is hard to blame the complexity of the airport networks for the contrasting exponential curve of these two countries. In 20 days, from the first infection in Brazil (February 26th) against 47 days after the first Indian case (January 30th), Brazil has already had 5.4 more confirmed cases than India (<https://covid.saude.gov.br/>, 2020; <https://gisanddata.maps.arcgis.com/>, 2020). Clearly, one country is doing much better in preventing the entrance of cases and the spreading of the disease by controlling infected citizens. Indeed, according to the WHO, the Ministry of Health and Family Welfare (MoHFW) of India has taken an early action, and “aggressively stepped up the response measures—find, isolate, test, treat and trace” (*World Health Organization (WHO), 2020b*).

Nigeria, a country poorer (27th world GDP position with \$446,543 billions nominal) than Brazil (9th world GDP position with \$1.85 trillion nominal) but with a similar population (*United Nations, 2019*), had imposed a very successful control so far, with guidelines and laws for within all the cities embracing social distancing and obliging wearing of face masks, starting from 31st March, 2 weeks before than Brazil (*Nigeria Centre for Disease Control, 2020*). As in most of the African countries, screening at the points of entries have been conducted in Nigeria’s airports (*World Health Organization (WHO), 2020c*). The *Federal Airports Authority of Nigeria (2020)* had established precautionary measures to be observed by inbound passengers, the opposite that Brazilian authorities did (*Agência Nacional de Aviação Civil (ANAC), 2020*).

In order to find and isolate is, from an ecological perspective, the most efficient way to reduce the carrying capacity of a new disease, and thus, restrict its wide spreading, and this must start at the airports. As Brazil is just struggling to impose social distancing, a State-to-State decision with little support from the Federal Government, the scenario is evolving more severely. Regardless of flaws in the comparisons of confirmed cases between the countries, by 16th April, India still had 12,380 cases and 414 deaths while Brazil had 25,262 cases and 1,532 deaths (*World Health Organization (WHO), 2020c*). Nigeria has 373 cases and 11 deaths (*World Health Organization (WHO), 2020c*). Despite the clearly more severe airport sanitarian control in Nigeria than in Brazil, one needs to be aware of the unexpectedly low number of cases in the continent, which may be related to a very young population or to other situations not well understood yet (*Vaughan, 2020*). For Brazil, on the contrary, enhanced case detection would make the scenario even worse. Considering the high probability of a synchronizing SARS-CoV-2 spreading in various capitals, the country may face a quick health service collapse.

Besides the within-city pattern of virus spreading, one must take into account the pattern of dispersion between cities after the virus has invaded. Additionally, for the Brazilian case, one cannot ignore that, eventually, the occurrence of the first case may have occurred nearly 1 month before official records, during the carnival period. This is the largest popular street party on the planet, with 6.4 million people in Rio de Janeiro, and 16.3 million in Salvador, and the *Brazilian Ministry of Tourism (2020)* revealed that 86,000 foreigners from France, Germany, Spain, Italy, UK, and the US had visited Brazil in this period. As airport control might have been even more lax in small airports, it might unavoidably result in strengthening of the capability of an infected city to infect the next new one, if no public policy is adopted.

Without a social distancing policy, virus propagation may result in chaotic dynamics, sensu *May (1976)*. The lack of control for these situations may result in a dramatic rate of host infection, and an eventual collapse of the host-parasite interaction in a given population, depending on the amount of susceptible, infected, and recovered events. Nonetheless, if the population is split into deme-cities, in a metapopulation structure, the collapse takes longer, and a much greater amount of people in different locations may eventually be infected, as found in our model. It is worthwhile to mention that this model, already pessimistic, did not consider the Brazilian road network, one of the largest on the planet. Most importantly, the best road-connected cities are exactly those mostly connected by airport, and that will be vulnerable earlier, and thus, probably spreading the disease faster than our model can predict, unless roads are soon blocked for people. Another weakness of the model is that it cannot account for a great number of small airports not registered for commercial flights, very common in the Amazonian and Western regions. Taking this into a global scale, for a highly interconnected human population, the consequences may be catastrophic, as it was for the influenza pandemic (Spanish flu) in 1918 (*Ferguson, Alison & Bush, 2003*). Furthermore, one aspect that must not be neglected is the way an increasing number of infected people in a city drives the pandemic towards the next city or country. In this context, the complex and large flight network of Brazil, which is also key for the whole Latin America, if not properly

monitored and controlled, may cause a window of opportunity for the virus to spread over the entire continent.

The consequences of this uncontrolled SARS-CoV-2 spreading is particularly serious if one takes into consideration the chances of a mutant virulent strain appearing and spreading into poorer and little monitored places of the world. Specifically, for the Amazon region, the lack of any control will make the city of Manaus a very sensitive cluster for public health, due to predominantly poor and indigenous-dominated cities in the region, which are connected to Manaus and will be rapidly infected. Reaching isolated regions means reaching indigenous or traditional communities, whose individuals are classically more susceptible to new pathogens than western-influenced or mixed urban populations. Therefore, a way to prevent such spreading, if still there is time, would be to deal with airports as entrances that need severe infection barriers.

CONCLUSIONS

The Brazilian media and the Ministry of Health have announced that 50 days from the introduction of SARS-CoV-2 in the country, it has spread across the regions and cities quite like we had predicted, even slightly faster. For instance, the time taken for Manaus to be compromised by the disease was as short as we predicted. A combination of being an important regional clustering in the airport network, and relatively limited hospital capability, resulted in a fast saturation of intensive care units, in cities as Manaus, Recife, and São Luis. As Manaus will disseminate the disease across the region, quickly reaching far remote indigenous communities, it should be on lockdown right now, to cause an invasion threshold, sensu *Colizza & Vespignani (2007)*. The first indigenous person, a teenager girl, has been already killed by COVID-19.

An eventual lesson to take for the whole country is that inflexible, severe, and easy to repeat protocols must be applied to all the cities with airports. Likewise, the follow-up monitoring of suspicious individuals and their living network should be reinforced as a national strategy to prevent a large territory to be taken over by a pandemic in a short period of time. In other words, internationally accepted procedures must be taken and even be reviewed to adjust to complex national flight networks of any country. Such procedures must be considered as a priority for national remote airports too, in order to keep poorer and worse equipped cities away from a rapid spread of a pandemic disease.

It is clear, at this point, that a fast spread of the SARS-CoV-2 is a reality in Brazil, and across most of the country. We proposed this model in order to emphasize the fragility of Brazilian surveillance in the airport network, in an attempt to cause some policy change in time to preserve at least the most remote regions, which are also the most vulnerable, with a weaker health service. Moreover, most of the Eastern part of the country must stay in social distancing in order to prevent a health public collapse by mid-May, as the Brazilian Ministry of Health predicted. So far, this has been a State-to-State decision, similarly to the United States, and those States that have been stronger in isolation measures, are indeed delaying the peak. Moreover, even if it is too late for the first

wave of infection in many cities, one must be prepared for a second likely wave, mainly considering the lack of a central government policy for social distancing.

In addition, we also could consider the generalized poverty of Brazil as a further problem that our model did not deal with. The chances to produce home-to-home isolation, even legally imposed, is impossible for these poor communities. Nonetheless, considering the few main entrances of most of the Brazilian shanty towns and communities, a similar to airport entrance severe control must be considered to protect a larger but closely connected set of people, eventually following the protocols used for the control of Ebola during the last epidemic in Africa (*Lau et al., 2017*).

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Author Contributions

- Sérgio Pontes Ribeiro conceived and designed the experiments, performed the experiments, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the paper, and approved the final draft.
- Alcides Castro e Silva conceived and designed the experiments, performed the experiments, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the paper, and approved the final draft.
- Wesley Dáttilo conceived and designed the experiments, performed the experiments, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the paper, and approved the final draft.
- Alexandre Barbosa Reis conceived and designed the experiments, authored or reviewed drafts of the paper, and approved the final draft.
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Data Availability

The following information was supplied regarding data availability:

Raw data and code are available in the [Supplemental Files](#).

Supplemental Information

Supplemental information for this article can be found online at <http://dx.doi.org/10.7717/peerj.9446#supplemental-information>.

REFERENCES

- Airports Authority of India. 2020.** Airports Authority of India Flight Schedule. Available at <https://www.aai.aero/en/airports/flights-schedule/allAirports/hadapsar.jsp> (accessed 13 April 2020).
- Agência Nacional de Aviação Civil (ANAC). 2020.** Histórico De Voos. Available at <https://www.anac.gov.br/assuntos/dados-e-estatisticas/historico-de-voos> (accessed 13 April 2020).
- Anderson RM. 1991.** Discussion: the Kermack–McKendrick epidemic threshold theorem. *Bulletin of Mathematical Biology* 53(1–2):1–32 DOI 10.1007/BF02464422.
- Balcan D, Gonçalves B, Hu H, Ramasco JJ, Colizza V, Vespignani A. 2010.** Modelling the spatial spread of infectious diseases: the Global Epidemic and Mobility computational model. *Journal of Computing Science* 1(3):132–145 DOI 10.1016/j.jocs.2010.07.002.
- Bedford J, Enria D, Giesecke J, Heymann DL, Ihekweazu C, Kobinger G, Lane HC, Memish Z, Oh M, Sall AA, Schuchat A, Ungchusak K, Wielwer LH. 2020.** COVID-19: towards controlling of a pandemic. *Lancet* 395(10229):1011–1088.
- Brazilian Ministry of Tourism. 2020.** Ministério Do Turismo: Página Inicial. Available at <http://www.turismo.gov.br/> (accessed 13 April 2020).
- Brockmann D, Helbing D. 2013.** The hidden geometry of complex, network-driven contagion phenomena. *Science* 342(6164):1337–1342 DOI 10.1126/science.1245200.
- Coelho FC, Lana RM, Cruz OG, Villela D, Bastos LS, Piontti AP, Davis JT, Vespignani A, Codeço CT, Gomes MFC. 2020.** Assessing the potential impact of COVID-19 in Brazil: mobility, morbidity, and the burden on the health care system. *medRxiv preprint* DOI 10.1101/2020.03.19.20039131.
- Colizza V, Vespignani A. 2007.** Invasion threshold in heterogeneous metapopulation networks. *Physical Review Letters* 99(14):148701 DOI 10.1103/PhysRevLett.99.148701.

- Dáttilo W, Silva AC, Guevara R, MacGregor-Fors I, Ribeiro SP. 2020. COVID-19 most vulnerable Mexican cities lack the public health infrastructure to face the pandemic: a new temporally-explicit model. *medRxiv* DOI 10.1101/2020.04.10.20061192.
- Fathlzadeh H, Maroufi P, Momen-Heravi M, Dao S, Köse S, Ganbarov K, Pagliano P, Espolito S, Kafli HS. 2020. Protection and disinfection policies against SARS-CoV-2 (COVID-19). *Le Infezioni in Medicina* 2:185–191.
- Federal Airports Authority of Nigeria. 2020. Federal Airports Authority Of Nigeria. Available at <https://www.faan.gov.ng/> (accessed 13 April 2020).
- Ferguson NM, Alizon P, Bush RM. 2003. Ecological and immunological determinants of influenza evolution. *Nature* 422(6930):428–433 DOI 10.1038/nature01509.
- Hanski I. 1998. Metapopulation dynamics. *Nature* 396(6706):41–49 DOI 10.1038/23876.
- Hellewell J, Abbott S, Gimma A, Bosse NI, Jarvis CI, Russell TW, Munday JD, Kucharski AJ, Edmunds WJ, Funk S, Eggo RM, Centre for the Mathematical Modelling of Infectious Diseases COVID-19 Working Group. 2020. Feasibility of controlling COVID-19 outbreaks by isolation of cases and contacts. *Lancet* 8(4):e488–e496 DOI 10.1016/S2214-109X(20)30074-7.
- Hethcote HW. 1989. Three basic epidemiological models. *Applied Mathematical Ecology* 18:119–144.
- Hubbell SP. 2001. *The unified neutral theory of biodiversity and biogeography*. Princeton: Princeton University Press.
- Lai A, Bergna A, Acciarri C, Galli M, Zehender G. 2020a. Early phylogenetic estimates of the effective reproduction number of SARS-CoV-2. *Journal of Medical Virology* 92(6):675–679 DOI 10.1002/jmv.25723.
- Lai C, Ship T, Ko W, Tang H, Hsueh P. 2020b. Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and coronavirus disease-2019 (COVID-19): the epidemic and the challenges. *International Journal of Antimicrobial Agents* 55(3):105924 DOI 10.1016/j.ijantimicag.2020.105924.
- Lau MSY, Dalziel BD, Funk S, McClelland A, Tiffany A, Riley S, Jessica C, Metcalf E, Grenfell BT. 2017. Spatial and temporal dynamics of superspreading events in the 2014–2015 West Africa Ebola epidemic. *Proceedings of the National Academy of Sciences of the United States of America* 114(9):2337–2342 DOI 10.1073/pnas.1614595114.
- Li G, Fan Y, Lai Y, Han T, Li Z, Zhou P, Pan P, Wang W, Hu D, Liu X, Zhang Q, Wu J. 2020. Coronavirus infections and immune responses. *Journal of Medical Virology* 92(4):424–432 DOI 10.1002/jmv.25685.
- May RM. 1976. Simple mathematical models with very complicated dynamics. *Nature* 261(5560):459–467 DOI 10.1038/261459a0.
- Morse SS, Mazet JAK, Woolhouse M, Parrish CR, Carroll D, Karesh WD, Zambrana-Torrel C, Lipkin WL, Daszak P. 2012. Prediction and prevention of the next pandemic zoonosis. *The Lancet* 380(9857):1956–1965 DOI 10.1016/S0140-6736(12)61684-5.
- Nicolaides C, Cueto-Felgueroso L, Gonzales MC, Juanes R. 2012. A metric of influential spreading during contagion dynamics through the air transportation network. *PLOS ONE* 7(7):e40961 DOI 10.1371/journal.pone.0040961.
- Nigeria Centre for Disease Control. 2020. NCDC coronavirus (COVID-19) information minisite. Available at <https://covid19.ncdc.gov.ng/> (accessed 14 April 2020).
- Pianka ER. 2000. *Evolutionary ecology*. Sixth Edition. San Francisco: Addison Wesley Longman.
- Rothe C, Schunk M, Sothmann P, Bretzel G, Froeschl G, Wallrauch C, Zimmer T, Thiel V, Janke C, Guggemos W, Seilmaier M, Drosten C, Vollmar P, Zwirgmaier K, Zange S,

Wölfel R, Hoelscher M. 2020. Transmission of 2019-nCoV infection from an asymptomatic contact in Germany. *New England Journal of Medicine* **382(10)**:970–971
DOI 10.1056/NEJMc2001468.

United Nations. 2019. World population: prospects 2019, Volume II: demographic profiles (ST/ESA/SER.A/427). Available at <https://population.un.org/wpp/DataQuery/> (accessed 17 April 2020).

Vaughan A. 2020. We don't know why so few Covid-19 cases have been reported in Africa: new scientist. Available at <https://www.newscientist.com/article/2236760-we-dont-know-why-so-few-covid-19-cases-have-been-reported-in-africa/> (accessed 17 April 2020).

World Health Organization (WHO). 2020a. Novel coronavirus (2019-nCoV). Available at https://www.who.int/docs/default-source/coronaviruse/situation-reports/20200121-sitrep-1-2019-ncov.pdf?sfvrsn=20a99c10_4 (accessed 14 April 2020).

World Health Organization (WHO). 2020b. Coronavirus disease (COVID-19). Available at <https://www.who.int/india/emergencies/novel-coronavirus-2019> (accessed 16 April 2020).

World Health Organization (WHO). 2020c. Coronavirus disease 2019 (COVID-19). Available at https://www.who.int/docs/default-source/coronaviruse/situation-reports/20200416-sitrep-87-covid-19.pdf?sfvrsn=9523115a_2 (accessed 16 April 2020).

World Health Organization (WHO). 2020d. COVID-19: Situation update for the WHO African Region. Available at https://apps.who.int/iris/bitstream/handle/10665/331330/SITREP_COVID-19_WHOAFRO_20200304-eng.pdf (accessed 14 April 2020).

Zhu N, Zhang D, Wang W, Li X, Yang B, Song J, Zhao X, Huang B, Shi W, Lu R, Niu P, Zhan F, Ma X, Wang D, Xu W, Wu G, Gao GF, Tan W. 2020. A novel coronavirus from patients with pneumonia in China, 2019. *New England Journal of Medicine* **382(8)**:727–733
DOI 10.1056/NEJMoa2001017.