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SARS-CoV-2 pandemic: An overview

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ABSTRACT

By the end of May 2020, SARS-CoV-2 pandemic caused more than 350,000 deaths worldwide. In the first months, there have been uncertainties on almost any area: infection transmission route, virus origin and persistence in the environment, diagnostic tests, therapeutic approach, high-risk subjects, lethality, and containment policies. We provide an updated summary of the current knowledge on the pandemic, discussing the available evidence on the effectiveness of the adopted mitigation strategies.

1. Introduction

On December 31, 2019, an outbreak caused by a new human pathogen, lately named Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) was notified in Wuhan, China (Gorbalenya et al., 2020). The virus caused a severe respiratory syndrome, generically defined as coronavirus disease (COVID-19) (Huang et al., 2020). Human-to-human transmission was first reported on January 22, 2020 and the World Health Organization (WHO) declared that SARS-CoV-2 infection had become a pandemic on March 11, when the 118,319 infections and 4292 deaths had been reported in 113 countries (World Health Organization, 2019, 2020a). By the end of May 2020, SARS-CoV-2 pandemic caused more than 350,000 deaths worldwide (Worldometers.info, 2020).

In the first months of the pandemic, there have been uncertainties on almost any area: infection transmission route, virus origin and persistence in the environment, diagnostic tests, therapeutic approach, high-risk subjects, lethality, and containment policies. An updated summary of the current knowledge on SARS-CoV-2 pandemic is therefore needed.

2. Virus origin, transmission routes, and persistence in the environment

As other coronaviruses (229E, NL63, OC43, HKU1, SARS-CoV, MERS-CoV), SARS-CoV-2 has enveloped virions that measure approximately 120 nm in diameter, and its genome consists of a single strand of RNA (Casella et al., 2020; Encyclopaedia Britannica., 2020). Due to wide genome affinities with coronaviruses affecting bats (Wu et al., 2020; Zhou et al., 2020a), the theories on the origin of the virus range from natural selection in an animal host before zoonotic transfer (possibly with an intermediate host),

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to selection during passage in laboratory cell cultures (this being the least accredited supposition). (Huang et al., 2020; Wu et al., 2020; Li et al., 2020a; Andersen et al., 2020; Xiao et al., 2020; Wang et al., 2020a).

SARS-CoV-2 genome has been detected in all of the following clinical specimens: nasopharyngeal swabs, sputum, bronchoalveolar lavage fluid, fibero bronchoscope brush biopsy, stool, ocular fluid, and blood (Wang et al., 2020a; Cheng et al., 2004; Peng et al., 2019; Colavita et al., 2020; Backer et al., 2020). However, according to current evidence, the virus is most commonly transmitted from infected persons through respiratory droplets and contact routes (Huang et al., 2020; Li et al., 2020a; Liu et al., 2020a; Chan et al., 2020). Droplet transmission occurs when a person is within 1–1.5 m from a contagious individual and is therefore at risk of having his/her mucosae (mouth and nose) or conjunctiva exposed to the droplets.

Airborne transmission is considered unlikely even when aerosol-generating procedures (e.g. tracheal intubation, non-invasive ventilation, etc.) are performed (Liu et al., 2020b; Tran et al., 2012), but contagion may occur by direct contact of the mucosae or conjunctiva with infected surfaces, like the skin or objects that have been exposed to infected droplets, sputum or sneeze (Liu et al., 2020b).

No evidence exists of fecal-oral transmission (Xu et al., 2020; Zhang et al., 2020; Tang et al., 2020a; Lamers et al., 2020), and although viral RNA was detected in the plasma or serum of 15% of the infected subjects (Huang et al., 2020), a review on diagnostic testing concluded that viral nuclear acid cannot be detected in serum and urine (Cheng et al., 2020), and transmission by blood transfusion remains unproven (Chang et al., 2020).

Finally, uncertainties remain on sexual and vertical transmissions. The limited available data suggest that vertical transmission is rare, with very few newborns infected to date, and no evidence of placental transmission (Yang and Liu, 2020). COVID-19, however, increased the risk of delivery complications and preterm birth (Yang and Liu, 2020; World Association of Perinatal Medicine Study Group on COVID-19, 2020; Hosier et al., 2020; Zamaniyan et al., 2020).

There have also been controversies over who is contagious, when, and for how long. The viral load, and in turn contagiousness, is highest in the pharynx of symptomatic individuals in the first week after symptoms onset (Wang et al., 2020a; Tang et al., 2020a; Wölfel et al., 2020; Lauer et al., 2020; To et al., 2020; Byrne et al., 2020). It is thought, however, that the infection could be spread also by those who do not develop symptoms (He et al., 2020a; Aguilar et al., 2020), although the proportion of infective asymptomatic individuals is likely to be low (Gao et al., 2020). Once infected, persons usually remain contagious for approximately 14 days, although a low proportion of subjects remain infectious for more than 30 days (63 days being the longest documented duration of the infection). (Hosier et al., 2020; Byrne et al., 2020; Pan et al., 2020).

Because of the above characteristics, SARS-CoV-2 is considered highly contagious, with an estimated basic reproduction number (the now famous “ R_0 ”) ranging from 2 to 6 without containment (which means that, without protection, a single individual infects two to six persons on average), and a household secondary attack rate close to 12% (which means that household or close contacts of an infected persons have a 12% probability of being infected). (He et al., 2020a; Jing et al., 2020; Bi et al., 2020; Streeck et al., 2020).

Finally, when considering virus transmission, it cannot be ignored whether and how long the active virus can persist in the environment. It has been documented, under experimental conditions with high viral load, that SARS-CoV-2 may remain infectious in the aerosol for 3 h, and on the surfaces (especially plastic and stainless steel) for up to three days (van Doremalen et al., 2020).

3. Symptoms of disease

The clinical manifestations and lethality of SARS-CoV-2 infection widely vary across individuals, depending on a number of factors (already recognized or just hypothesized), that have been discussed in a dedicated paragraph. Approximately half of the infected subjects remain asymptomatic, and most of the remaining only experience influenza-like symptoms, including fever, cough, sore throat, runny nose, weakness, myalgia, headache and, less frequently, conjunctivitis, hemoptysis, olfactory and taste dysfunction, and diarrhea (Casella et al., 2020; He et al., 2020a; Aguilar et al., 2020; ECDC, 2020; Whitcroft and Hummel, 2020). Without early treatment, approximately 10%–15% of the infected persons develop a severe disease (COVID-19), which may include any of the followings: pneumonia, acute respiratory distress syndrome, pulmonary embolism, cardiomyopathy, disseminated intravascular coagulation, ultimately leading to death in most critical cases (Huang et al., 2020; Tang et al., 2020b; Klok et al., 2020). Although the likelihood of severe disease is lower in children, the clinical scenario is similar (Su et al., 2020a; Liguoro et al., 2020).

The average time from exposure to symptom onset (incubation period) is 5 days, and the time from symptoms to death usually ranges between 10 and 18 days, with a median of 14 (He et al., 2020a; Wang et al., 2020b).

4. Diagnosis

The knowledge on the diagnostic criteria and tests to detect SARS-CoV-2 infection, and COVID-19, is continuously evolving (Sethuraman et al., 2020; Zhai et al., 2020). Although several novel or complementary diagnostic methods are being developed, including serologic and point-of-care molecular diagnostic tests, real-time reverse transcriptase polymerase chain reaction (RT-PCR)–based assays performed in the laboratory on respiratory specimens are currently the cornerstone of the diagnosis of SARS-CoV-2 infection (Cheng et al., 2020; Sethuraman et al., 2020; World Health Organization, 2020b). Various RT-PCR assays are used around the world: the CDC has developed an assay containing PCR primer–probe sets for 2 regions of the viral nucleocapsid gene (N1 and N2), and for the human RNase P gene to ensure the RNA extraction was successful. This assay differs from WHO primer–probe sets (originally proposed by the Charité-Universitätsmedizin Berlin Institute of Virology) (Corman et al., 2020a), which target the SARS-CoV-2 RNA dependent RNA polymerase (*RdRP*) and envelope (*E*) genes (World Health Organization, 2020b; Corman et al., 2020b). Both assays have high analytic sensitivity and specificity, with minimal cross reactivity with other circulating strains of

coronaviruses. However, a number of laboratories also obtained an emergency use authorization from regulatory agencies for their own laboratory developed assays (Cheng et al., 2020). As of May 8, the European Union has granted the CE mark to 40 PCR and nine serology tests; the United States Food and Drug Administration has approved 74 PCR, twelve serology, and two isothermal amplification tests, while additional PCR tests have been approved by other competent institutions around the World (Garg et al., 2020). More recently, droplet digital PCR (ddPCR) has shown significantly better performances than RT-PCR, especially for the diagnosis of the subjects with low viral load (Liu et al., 2020c; Suo et al., 2020).

The absence of an established reference standard, varying sample collection (including nasal, nasopharyngeal, oropharyngeal, mid-turbinate specimens, and endotracheal aspirates), differing preparation methods, and infection phase at the time of testing hamper rigorous assessment of the diagnostic accuracy of the many newly introduced SARS-CoV-2 assays (Tang et al., 2020a; Cheng et al., 2020; Mallapaty, 2020; Lippi et al., 2020a). In preliminary studies, the sensitivity and specificity widely varied, with four reports showing concerning rates of false negative results, higher than 20%, for some RT-PCR assays (Hu et al., 2020; He et al., 2020b; Fang et al., 2020; Meng et al., 2020; Wen et al., 2020; Zhifeng et al., 2020). The accuracy may also differ by time of infection: viral RNA becomes detectable as early as day 1 of symptoms and peaks within the first week of symptom onset, declining thereafter. Viral RNA has been detected even beyond week 6 following the first positive test, thus, at least two negative samples collected at ≥ 24 -h intervals are required to document SARS-CoV-2 clearance. A few cases have been found positive after 2 consecutive negative PCR tests within 24 h, but it is unclear if these were testing errors, re-infections, or reactivations (Sethuraman et al., 2020; Zheng et al., 2020). Finally, as conventional RT-PCR tests could not be performed in low infrastructure settings (such as on cruise ships or in remote communities), rapid point-of-care molecular diagnostic tests have also been developed, although the sensitivity is likely to be lower than laboratory-based tests (Cheng et al., 2020; Sethuraman et al., 2020; Corman et al., 2020b).

With these limitations, RT-PCR tests are crucial to identify the subjects with circulating viruses, thus being essential to integrate the clinical diagnosis of severe COVID-19. However, they are unable to assess whether an individual was previously infected and has recovered, that could rather be detected using serologic tests, which identify antibodies to SARS-CoV-2 (IgA, IgM and IgG) from blood or saliva (Cheng et al., 2020; Perera et al., 2020). Serologic tests can be of four types: RDTs, chemiluminescent immunoassays, enzyme linked immunosorbent assays (ELISA), and neutralization assays. The Johns Hopkins University compiled a list of the tests approved for diagnostics or research, finding sensitivities ranging from 82% to 100%, and specificities ranging from 96% to 100%, with false negatives mainly due to cross-reactivity with other coronavirus proteins (Johns Hopkins Bloomberg School of Public Health - Center for Health Security, 2020).

Clearly, antibody responses to infection take weeks to be detectable (Johns Hopkins Bloomberg School of Public Health - Center for Health Security, 2020), thus serologic tests cannot be used for rapid diagnoses of symptomatic subjects (Cheng et al., 2020; Sethuraman et al., 2020). However, they will be of paramount importance for epidemiologic studies assessing SARS-CoV-2 lethality, vaccine studies, ongoing surveillance, and risk assessment and stratification of the population and health care workers (Cheng et al., 2020; Sethuraman et al., 2020; Stadlbauer et al., 2020; Hains et al., 2020; Goudsmit, 2020).

Besides RT-PCR and serologic tests, some biomarkers have been associated with COVID-19, including decreased albumin, elevated C-reactive protein, elevated lactate dehydrogenase, and lymphopenia (Rodriguez-Morales et al., 2020). To date, however, no biomarker or combination of biomarkers can be used to diagnose the infection or predict the clinical course (Cheng et al., 2020).

In light of the above mentioned limitations of the available tests, and of their accessibility, the diagnosis of COVID-19 is often made presumptively based on a compatible clinical presentation and the exclusion of other potential etiologies, in the setting of an exposure risk (residence in or travel to an area with widespread community transmission or known contact). (McIntosh, 2020).

For patients suffering from fever, sore throat, fatigue, coughing or dyspnea that is coupled with recent exposure, the diagnosis should be made using chest computerized tomography (CT) characteristics (typically, bilateral pulmonary parenchymal ground-glass and consolidative pulmonary opacities) despite negative RT-PCR results, which should be repeated 24–48 h after the initial test (Cheng et al., 2020; Zhai et al., 2020; Xie et al., 2020). Given that chest imaging is highly sensitive (78%–100%), but lacks specificity, the clinicians should always keep in mind the other pulmonary, cardiac, or systemic differentials because if missed, they could even lead to more severe consequences (Wu et al., 2007).

The same criteria are adopted to select the individuals to test. In fact, the characteristics of who should be tested have been controversially changing over time (European Centre for Disease Prevention and Control, 2020a; European Centre for Disease Prevention and Control, 2020b; Tsang et al., 2020). Although, theoretically, all symptomatic patients with suspected infection should undergo testing (Cheng et al., 2020; World Health Organization, 2020b), the CDC currently recommends to test the following, symptomatic subjects: hospital patients; residents in long-term care facilities or other congregate living settings, including prisons and shelters; healthcare workers; and workers in congregate living settings (CDC CfDCaP, 2020a). Testing is not recommended in asymptomatic individuals, unless they are prioritized by health departments or clinicians. Priority criteria may include, but are not limited to: the close contact with a confirmed or suspected COVID-19 case in the prior 14 days, residence in or travel to an area with widespread community transmission, and high risk of poor outcomes (CDC CfDCaP, 2020a).

5. Infection rates and lethality

By the end of May 2020, SARS-CoV-2 infection spread all over the world, with more than 5 million detected cases (0.7×1000 inhabitants worldwide) (Worldometers.info, 2020). The reported incidence rates ranged from 0.01 to approximately 16 cases per 1000 persons across nations with at least 100,000 residents (Worldometers.info, 2020). Besides a generally low incidence in African countries, no clear trends in the infection rates emerged by climate zone or per-capita domestic gross product (Worldometers.info, 2020; Berumen et al., 2020; Su et al., 2020b). Clearly, the incidence rates are influenced by the capacity to detect the infected persons

through RT-PCR tests, the availability of which widely varied across nations. In fact, high income countries tested more than 4% of their population, while most African countries tested less than 5/1000 residents (Worldometers.info, 2020; Lachmann et al., 2020).

The variability in diagnostic testing does not only complicate the assessment of incidence rates of infection, it introduces an even higher likelihood of bias in the evaluation of the lethality. In many countries, during the weeks of the epidemic peak, because of the limited availability of PCR tests, these were administered only to the persons with severe symptoms (Wong et al., 2013). Thus, a large proportion of the infected subjects - most of those asymptomatic or with mild disease - remained undetected (Henegan et al., 2020; Ioannidis, 2020a). This creates a very large margin of uncertainty in the estimates of the case-fatality rate, as the numerator may be solid (n. of deaths), but the denominator (n. of infected persons) could be underestimated by a factor of 2, up to 50 (Ioannidis, 2020a; Oke and Henegan, 2020). In other terms, the real lethality of the disease, which is currently estimated at 6.2% worldwide, may range from 0.2% to 3.1%, and the summary estimates from two meta-analyses were 0.8% (Meyerowitz-Katz and Merone, 2020) and 3.3% (He et al., 2020a). Indeed, SARS-CoV-2 case-fatality rate widely varied across nations, even within the same continent and geographic area (Belgium, 16.2% and France, 15.5%, versus Germany, 4.7%, and Austria, 3.9%). To further complicate the correct estimation of the lethality, it has been shown that the risk of death largely depends upon age and comorbidities, thus the age distribution of a country, in addition to the detection rate, might have played a major role in determining current case-fatality rates. Also, the procedures of case recording (Iosa et al., 2020), the preparedness of the healthcare system, which in turn is affected by the intensity of the spread (Iosa et al., 2020; Boccia et al., 2020), and, especially, mitigation policies, certainly contributed to determine current case-fatality rates (Meyerowitz-Katz and Merone, 2020). Finally, some preliminary analyses are suggesting that, with current, more tailored therapies, COVID-19 lethality decreased over time (Flacco et al., 2020), and attenuated virus variants have been reported (Lau et al., 2020).

Some interesting mathematical models were developed to help estimating infection and fatality rates, and have been briefly summarized in a dedicated paragraph (Meyerowitz-Katz and Merone, 2020; Ferguson et al., 2020; Imperial College London MRC Centre for Global Infectious Disease Analysis, 2020; Remuzzi and Remuzzi, 2020; University of Washington Institute of Health Metrics and Evaluation (IHME), 2020). Although technically refined, they were made in the earlier phase of the pandemic, with an incomplete, varying set of rapidly evolving assumptions, the less tenable of which was the absence of mitigation strategies.

All considered, given the large variability among countries, age class, risk profile and, possibly, time, providing an overall, single estimate of SARS-CoV-2 case-fatality rate is currently impossible. It is however possible to compute lethality estimates stratified by the main risk factors, that are essential to support the clinicians in defining the prognosis and therapeutic strategies. We briefly discuss the main predictors of severe/lethal COVID-19 in the next paragraph.

6. Predictors of severe/lethal disease

In a number of descriptive studies, the infected persons who died were most frequently older than 70 years, males, hypertensive, diabetics, and affected by serious comorbidities such as congestive heart failure, chronic obstructive pulmonary diseases or renal diseases (Garg et al., 2020; Yang et al., 2020; Patel and Verma, 2020; Jordan et al., 2020; Castagnoli et al., 2020; Wu and McGoogan, 2020; Zhou et al., 2020b; Zuin et al., 2020; Bravi et al., 2020). Many of the unadjusted reports were confirmed when more reliable, adjusted estimates became available from cohort analyses, although the magnitudes of the associations were reduced (as typically happens) (Matsushita et al., 2020). As compared with the subjects younger than 50 years, those older than 70 years showed a 5- to 10-fold increase in the risk of severe/lethal COVID-19 (Bravi et al., 2020; Matsushita et al., 2020; Williamson et al., 2020). Males, hypertensive, diabetics, and those with COPD or major cardiovascular diseases all showed an approximately doubled risk of death or severe disease (Matsushita et al., 2020; Williamson et al., 2020; Mehra et al., 2020). Obesity is also emerging as a significant, independent predictor of severe/lethal disease (Garg et al., 2020; Williamson et al., 2020; Simonnet et al., 2020; Lighter et al., 2020), while fewer data are available on other potential risk factors such as tobacco smoking (Hu et al., 2020; Williamson et al., 2020; Alqahtani et al., 2020), socioeconomic status (Rose et al., 2020; Wadhera et al., 2020), ethnicity, (Williamson et al., 2020; Teo et al., 2020), immune system dysfunctions (ECDC, 2020; Tay et al., 2020), and exposure to pollution (Lippi et al., 2020b). Notably, no association was found between COVID-19 progression and use of Angiotensin converting enzyme inhibitors or Angiotensin II receptor blockers (Bravi et al., 2020; Mancina et al., 2020; Mehta et al., 2020; Reynolds et al., 2020).

With regard to the children/adolescents, few deaths ($n < 20$) have been reported worldwide among children aged 0–9 years, and a low case-fatality rate (0.2%) has been observed for those aged 10–18 years (Worldometers.info, 2020; Castagnoli et al., 2020; CDC CfDcCaP, 2020b). As for the adults, a higher risk of severe/lethal COVID-19 was observed among children with underlying comorbidities (Worldometers.info, 2020; Shekerdemian et al., 2020).

7. Accuracy of model predictions

As anticipated, a number of interesting statistical models were developed to predict (or just estimate the real) infection and lethality rates (Meyerowitz-Katz and Merone, 2020; Ferguson et al., 2020; Imperial College London MRC Centre for Global Infectious Disease Analysis, 2020; Remuzzi and Remuzzi, 2020; University of Washington Institute of Health Metrics and Evaluation (IHME), 2020). Given that the assumptions were inevitably affected by underreporting of cases (with test shortages, asymptomatic transmission, emergency situations that impaired rigorous contact tracing), and could not take into account unprecedented containment strategies (Li et al., 2020b), it was inevitable that most model estimates turned out to be incorrect.

As an example, a model predicted that all 5200 Italian ICU beds would have been occupied by March 14, when 1518 beds were actually employed for COVID-19 patients, and the peak in ICU beds (4068) was reached on April 3 (Remuzzi and Remuzzi, 2020).

However, the model assumed no mitigation strategies, while lockdown measures were gradually implemented in Italy since February 21. Without the containment, the prediction might have been consistent. Another model, probably the most renowned, was developed by experts of the Imperial College of London (Ferguson et al., 2020). It predicted that, if the pandemic was left unmitigated, 81% of the population would be infected, with 510,000 deaths in the UK and 2.2 millions in the USA. Again, it is impossible to verify these estimates. However, 2009 H1N1 influenza pandemic infected no more than 27% of Americans, thus an overall infection rate of 81% may be overestimated (Yamey and Gonsalves, 2020; McKee, 2020; Van Kerkhove et al., 2013). Another model from China predicted that without implementation of any measures, China (without Hubei) would have had more than 800 million COVID-19 cases and an epidemic duration of 477 days (Chen et al., 2020).

In any case, as frequently occurs in preventive medicine (Sackett, 2002), a precise evaluation of these models cannot be made, which further complicates the choice of the most adequate containment strategies to adopt, in the next few months as well as for the next pandemics. Beyond accuracy, to increase their utility for public health decisions makers, the future models will have to incorporate separate estimates according to the most likely containment policies (including a strongly needed estimation of the health consequences of lockdown measures) and, possibly, produce estimates with smaller uncertainty margins (Ferguson et al., 2020; Karanikolos et al., 2013; Parmar et al., 2016; Fong et al., 2020). Notably, the last model from the Imperial College provided different estimates according to various combinations of mitigation strategies (Ferguson et al., 2020).

8. Therapy

There still is no evidence from randomized trials to recommend any specific drug or other therapeutic to prevent or treat COVID-19 (Zhai et al., 2020; CDC CfDCaP, 2020c). However, the U.S. National Institutes of Health (NIH) have published interim guidelines for the medical management of COVID-19, that are regularly updated and contain information about investigational therapeutics (National Institutes of Health, 2020). Besides supplemental oxygen and mechanical ventilatory support when indicated (CDC CfDCaP, 2020c), and although no drug has been proven to be safe and effective (National Institutes of Health, 2020), commonly tested treatments include antiviral agents (remdesivir, lopinavir, ritonavir - most of which, however, are discouraged by NIH if not administered in the context of a clinical trial), anticoagulants (mostly low molecular weight heparin - which is however not recommended by NIH for non-hospitalized subjects), and immune-based therapies (COVID-19 convalescent plasma, SARS-CoV-2 immune globulins, and interleukin 1 or 6 inhibitors such as sarilumab or tocilizumab) (National Institutes of Health, 2020; AIFA, 2020; Bhimraj et al., 2020; Sanders et al., 2020; American Society of Hematology, 2020). As for the other treatments, despite some preliminary, promising results (Capra et al., 2020; Di Giambenedetto et al., 2020; Mazzitelli et al., 2020), there are insufficient data to recommend either for or against the use of immune-based therapies (National Institutes of Health, 2020).

More than 800 randomized trials on treatments for COVID-19 have been registered worldwide (ECDC, 2020; Zhai et al., 2020; Covid-19 - living NMA initiative, 2020; World Health Organization, 2020c). Of them, more than 400 were actively recruiting patients by the end of May 2020. With such an unprecedented commitment, it is plausible to suppose that safe and effective therapeutic approaches will be available in a next future.

9. Containment policies

In the absence of vaccines and specific treatments, all the uncertainties on transmission routes, virus persistence in the environment, diagnostic tools accuracy, infection and lethality rates, therapies, and even the duration of the immunity (He et al., 2020b) contribute with a multiplier effect in delineating an extraordinary critical scenario at the end of the process, when adequate containment strategies have to be, or should have been chosen. Virtually all countries adopted preventive and containment measures directed to (a) reduce as much as possible human-to-human transmission, and (b) identify and isolate infected subjects through epidemiological investigations and extensive testing, and 14-day quarantine of COVID-19 cases and contacts (AIFA, 2020; Sanders et al., 2020; American Society of Hematology, 2020; Capra et al., 2020; Mazzitelli et al., 2020; European Centre for Disease Prevention and Control, 2020c).

To decrease the likelihood of transmission and slow the pandemic, a wide range of measures were gradually adopted: closure of the borders and block of passenger transportation across cities, obligation of social distancing (usually 1–1.5 m) and use of masks, respirators and/or gloves, shut down of schools, public services, social events, and inessential job activities, continuous indoor disinfection, up to complete lock-down with prohibition to leave the house, if not for essential needs (such as in Wuhan province and in Italy). (Sanders et al., 2020; Nussbaumer-Streit et al., 2020; Signorelli et al., 2020; Italian Prime Minister Italian Government, 2020; Italian Prime Minister Government I, 2020).

Besides some oddities, such as suspending 90% of the lawsuits and leaving tobacconists wide open (Italian Prime Minister Italian Government, 2020; Italian Prime Minister Government I, 2020), sending flawed diagnostic kits (Grady, 2020), or leaving healthcare workers untested and with no personal protective equipments in the first month of the pandemic, (Editor. Coronavirus, 2020; Errante, 2020), there are few doubts that the adopted containment measures were effective in reducing the spread of the infection and, in turn, drastically decreasing the death toll (Nussbaumer-Streit et al., 2020; Koo et al., 2020; Lai et al., 2020).

What is in fact entirely unknown, and urgently requires to be discerned, is which of the adopted strategies were more effective, and which were less, or even ineffective, and could thus be avoided in the next phases or next pandemics, in order to reduce as much as possible the tremendous socio-economic (and health) burden of an extensive lock-down (Nussbaumer-Streit et al., 2020; Ioannidis, 2020b; Parmet and Sinha, 2020; Morgan, 2020).

Unfortunately, no evidence is available except for modeling studies, but the analyses of the impact of various mitigation measures

on other respiratory transmitted viral diseases, such as SARS, MERS or influenza, might certainly be helpful (Fong et al., 2020; Nussbaumer-Streit et al., 2020; Mahtani et al., 2020). In brief, there is evidence in favor of isolating symptomatic individuals and exposed contacts (as early as possible), avoiding crowds and restricting movements, social distancing and remote working (Fong et al., 2020; Nussbaumer-Streit et al., 2020; Lai et al., 2020; Mahtani et al., 2020; Rashid et al., 2015). However, controversies remain on the cost-benefit of school closures, methods for active case-finding (screening and contact tracing), tools for indoor disinfection, which masks to wear (and where), and types of quarantine (van Doremalen et al., 2020; Fong et al., 2020; Nussbaumer-Streit et al., 2020; Rashid et al., 2015; Radonovich et al., 2019; Smith et al., 2016; Setti et al., 2020; Patel et al., 2020).

Concerning screening tools, in Italy more than a million airport passengers were screened with thermal scanners in the months preceding the pandemic, with minimal or no effect over the course of the disease (Signorelli et al., 2020; Health Protection Agency IG, 2020). In fact, it is hard to support a screening test that misses almost half of the infected subjects (all those that are asymptomatic or in the incubation period), and inappropriately classifies as “dangerous” the much larger amount of persons with body temperature higher than 37.5 C° because of many other reasons (Quilty et al., 2020; O’Grady et al., 2008; Sund-Levander et al., 2002). Notwithstanding, the Italian National Institute for Insurance Against Accidents at Work recommended the use of thermal scanners in all workplaces (Italian National Institute for Insurance Against Accidents at Work (INAIL), 2020).

With regard to contact tracing, there are no doubts on its effectiveness, but limited evidence exists on cost-effectiveness, as considerable resources are required (Mahtani et al., 2020). One of the proposed solutions, a smartphone application to decrease health officials burden, raises concerns on its accuracy in detecting contacts, lack of standardization, potential danger for privacy and risk of use for social control in the future (Ioannidis, 2020b; McGee et al., 2020).

Even more important is the debate over the measures of social isolation for the general population (non-sick, non-exposed). A model has estimated - and may be logical to suppose - that the combination of case isolation, household quarantine, social distancing of the entire population, and school and university closures would achieve the greatest effect (Ferguson et al., 2020). However, besides some critiques on the plausibility of these estimates (Nussbaumer-Streit et al., 2020), the same model predicted that a less extreme combination of case isolation and voluntary quarantine for three months, and social distancing of people 70 years or older for four months, could prevent 49% of deaths. Other researchers suggested an alternative, lower-impact approach, that they called “inverse quarantine”, where isolation would be requested only to high-risk subjects (Standl et al., 2020). Also, an analysis of the official data indicate that the infection rates similarly decreased after the end of the lockdown in the vast majority of countries, thus suggesting that more stringent lockdown might have not substantially altered pandemic dynamics (Kolanovic and Kaplan, 2020).

This is not a trivial difference, as the most aggressive approaches are having huge economic consequences, are likely to cause psychological and health harms (potentially larger than the pandemic) (Kolanovic and Kaplan, 2020), and pose serious ethical issues (Ioannidis, 2020b; Parmet and Sinha, 2020; Gostin and Hodge, 2020). Italy represents a case study, as it was the first Western country to be seriously hit by the pandemic, and the most draconian restrictions were imposed (Italian Prime Minister Italian Government, 2020; Italian Prime Minister Government I, 2020). For almost two months, people throughout the country were not allowed to exit from their houses, unless for working (although only essential activities were maintained) or well-motivated and verifiable health needs or food supplying (Italian Prime Minister Italian Government, 2020; Italian Prime Minister Government I, 2020). Thousands of fines were imposed, with the support of drones and helicopters, and the prohibition was valid for any individual in any context, even alone, wearing masks, in open spaces (Ziniti, 2020). However, based on current evidence, the latter, extreme measures do not seem justified, as airborne transmission has not been proven (Liu et al., 2020b; Tran et al., 2012), the real infectivity of asymptomatic infected individuals appears low (Gao et al., 2020), public health authorities define a significant exposure to SARS-CoV-2 as face-to-face contact within 2 m with a symptomatic infected subject that is sustained for at least a few minutes, and the chances of catching SARS-CoV-2 in open spaces, wearing masks, are therefore minimal (or even negligible in many provinces, considering disease prevalence at the time of the lockdown). (Qian et al., 2020; Protezione Civile. COVID-19 Italia - Monitoraggio della situazione, 2020; Klompas et al., 2020).

At the very end, the forced isolation of the entire population of a country for months is a quarantine on such an unprecedented scale to raise serious concerns on the balance between public health protection and civil liberties, requiring a particularly careful evaluation. The Constitutions of most Western countries require civil confinement to be temporary and applied only when individuals pose a significant risk of spreading dangerous, infectious diseases, with a use of coercion proportionate to the risk (Gostin and Hodge, 2020; Republic of Italy Italy Ro, 1947; United States of America America USo, 1788). In this view, probability remains the beacon that should guide governments, as the precautionary principle can be applied in both directions in the case of a lockdown (Ioannidis, 2020b). Establishing whether a disease has a lethality of 3% or 0.3%; whether the risk of transmission in some situations is 10% or 0.1%; and whether the risk of encountering a contagious person is 1% or 0.1%, is not just an exercise for epidemiologists, it is the fundamental basis for achieving the most correct, effective and less restrictive measures to mitigate a pandemic, possibly saving billions, balancing health protection and civil rights (Parmet and Sinha, 2020; Gostin and Hodge, 2020). A good example is provided in Fig. 1: even with the most pessimistic assumptions (altogether implausible), it is hard to conclude that the general Italian population forced in quarantine posed “a significant risk of spreading dangerous, infectious diseases, with a use of coercion proportionate to the risk”. It might have been so in some provinces (e.g. Bergamo or Brescia), certainly not in the entire country. With no blame for governments that faced an unprecedented challenge, let us learn from experience. For the next pandemic, and for the next phases of the current one.

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Average risk of causing severe/lethal COVID-19 of the general, asymptomatic Italian population, in open spaces, during SARS-CoV-2 epidemic months (March, April, 2020), under the most pessimistic assumptions

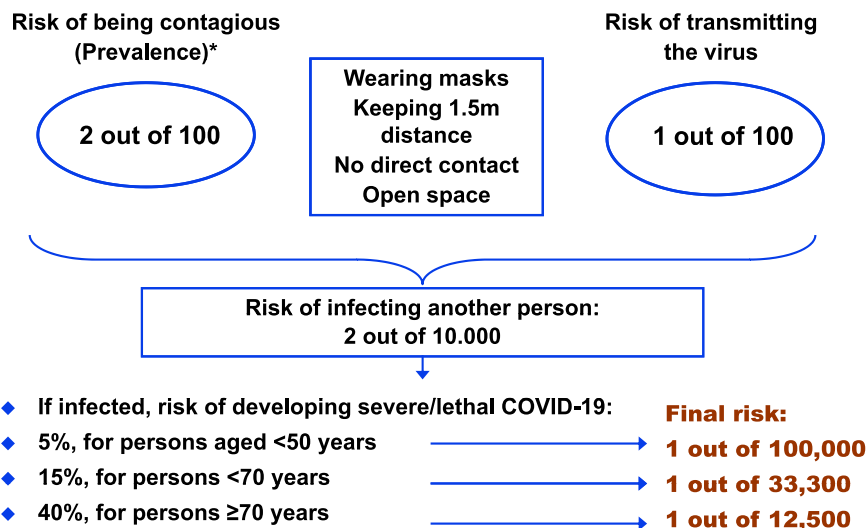


Fig. 1. Average risk of causing severe/lethal COVID-19 of the general, asymptomatic Italian population, in open spaces, during SARS-CoV-2 epidemic months (March, April 2020), under the most pessimistic assumptions.

* Official data indicate that, in the first three months of infection (up to May 25, 2020), the overall rate of infected subjects was 0.4 out of 100. Even increasing tenfold this value, asymptomatic persons are approximately half of all the infected subjects. Also, with an overestimated average disease duration of 30 days, only half of the overall number of infected subjects could be actively contagious on a single day. Importantly, these are intended as average prevalence values for the entire country, while some provinces had a completely different epidemiologic scenario, with official cumulative incidence rates as high as 10.8% (Bergamo), or 7.4% (Brescia); and as low as 0.05% (Taranto) or 0.08% (L'Aquila). As the lockdown was introduced in the entire country, these estimates are referred to the whole nation. In some provinces, the final risk of infecting other persons might have been 20 times higher, or 10 times lower.

CRedit authorship contribution statement

Cecilia Acuti Martellucci: Conceptualization, Investigation, Data curation, Writing - original draft. **Maria Elena Flacco:** Conceptualization, Investigation, Data curation, Validation, Writing - original draft. **Rosaria Cappadona:** Data curation, Writing - review & editing. **Francesca Bravi:** Data curation, Writing - review & editing. **Lorenzo Mantovani:** Data curation, Writing - review & editing. **Lamberto Manzoli:** Conceptualization, Methodology, Validation, Writing - review & editing, Visualization, Supervision, Project administration.

Declaration of competing interest

All authors declare they have no potential competing interests.

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