



Epidemic and pandemic viral infections: impact on tuberculosis and the lung

A consensus by the World Association for Infectious Diseases and Immunological Disorders (WAIDID), Global Tuberculosis Network (GTN), and members of the European Society of Clinical Microbiology and Infectious Diseases Study Group for Mycobacterial Infections (ESGMYC)

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This consensus statement describes the effects of the viral infections resulting in epidemics and pandemics affecting the lung (MERS, SARS, HIV, influenza A (H1N1)pdm/09 and COVID-19) and their interactions with TB, the top infectious disease killer <https://bit.ly/2UUjhGu>

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ABSTRACT Major epidemics, including some that qualify as pandemics, such as severe acute respiratory syndrome (SARS), Middle East respiratory syndrome (MERS), HIV, influenza A (H1N1)pdm/09 and most recently COVID-19, affect the lung. Tuberculosis (TB) remains the top infectious disease killer, but apart from syndemic TB/HIV little is known regarding the interaction of viral epidemics and pandemics with TB. The aim of this consensus-based document is to describe the effects of viral infections resulting in epidemics and pandemics that affect the lung (MERS, SARS, HIV, influenza A (H1N1)pdm/09 and COVID-19) and their interactions with TB. A search of the scientific literature was performed. A writing committee of international experts including the European Centre for Disease Prevention and Control Public Health Emergency (ECDC PHE) team, the World Association for Infectious Diseases and Immunological Disorders (WAIDID), the Global Tuberculosis Network (GTN), and members of the European Society of Clinical Microbiology and Infectious Diseases (ESCMID) Study Group for Mycobacterial Infections (ESGMYC) was established. Consensus was achieved after multiple rounds of revisions between the writing committee and a larger expert group. A Delphi process involving the core group of authors (excluding the ECDC PHE team) identified the areas requiring review/consensus, followed by a second round to refine the definitive consensus elements. The epidemiology and immunology of these viral infections and their interactions with TB are discussed with implications for diagnosis, treatment and prevention of airborne infections (infection control, viral containment and workplace safety). This consensus document represents a rapid and comprehensive summary on what is known on the topic.

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Introduction

The 21st century has been marked by major epidemics, including some that qualify as pandemics, caused by old diseases such as cholera, plague and yellow fever, as well as emerging diseases such as severe acute respiratory syndrome (SARS), Ebola, Zika, Middle East respiratory syndrome (MERS), HIV (although technically endemic), influenza A (H1N1)pdm/09 and most recently COVID-19. Several of these viruses affect the lung. Tuberculosis (TB) remains the top infectious disease killer caused by a single organism and was responsible for 1.5 million deaths in 2018 [1]. Apart from syndemic TB/HIV, little is known regarding the interaction of other viral epidemics with TB. This consensus-based document describes the effects of the main viral epidemics which predominately affect the lungs or cause systemic immunosuppression (MERS, SARS, HIV, influenza A(H1N1)pdm/09 and COVID-19) and their interactions with TB at a diagnostic, treatment and public health level. The document is the fruit of a collaborative project involving the European Centre for Disease Prevention and Control Public Health Emergency (ECDC PHE) team, the World Association for Infectious Diseases and Immunological Disorders (WAidid), the Global Tuberculosis Network (GTN), and members of the European Society of Clinical Microbiology and Infectious Diseases (ESCMID) Study Group for Mycobacterial Infections (ESGMYC).

Methods

We performed a rapid and nonsystematic search of the literature using the key words “COVID-19”, “tuberculosis”, “viral infection”, “HIV infection”, “SARS”, “lung”, “immunology”, “diagnosis”, “prevention”, “infection control” and “workplace” to identify a minimum set of references from an electronic database

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(PubMed), existing guidelines on TB, viral diseases and airborne diseases, and grey literature from their inception until April 29, 2020. Guidelines were retrieved from the websites of the main international health-related centres, whereas grey literature was accessed using the Google search engine.

A writing committee composed of international experts was established, including the ECDC PHE team, WAidid, GTN and ESGMYC.

Consensus on the content was achieved after multiple rounds of revisions between the writing committee and the larger group of experts [2].

A Delphi process involving the core group of authors, excluding the ECDC PHE team, identified the areas requiring review/consensus, followed by a second round to refine the definitive consensus elements.

As the review was not aimed at duplicating World Health Organization (WHO), ECDC and other existing guidelines, the GRADE (Grading of Recommendations, Assessment, Development and Evaluations) methodology was not used and no formal recommendations are provided. The available information on prevention, diagnosis and treatment of TB and pulmonary viral diseases was selected by the experts and summarised, and country examples were provided to critically discuss the public health response.

Viral diseases of the lung

Epidemiology

Viral respiratory infections are a major public health concern due to the capacity of viruses to spread from person to person directly *via* aerosols/droplet nuclei, small droplets or virus-laden secretions from larger droplets, or indirectly by contact with contaminated surfaces [3]. Large respiratory droplets are generated primarily during coughing, sneezing and talking, and during procedures such as suctioning and bronchoscopy, which can also generate droplet nuclei. Transmission occurs when droplets containing microorganisms from an infected person are expelled a short distance through the air and deposited on another individual's conjunctivae, nasal mucosa or mouth. Large droplets fall quickly onto surfaces close to the infected person, increasing the risk of contact transmission. Moreover, viral infections can also be transmitted *via* aerosol particles of small size (<5–10 µm) which may be infectious at a distance of several metres [4–7]. Recent evidence suggests the SARS coronavirus 2 (SARS-CoV-2) virus may be present in exhaled air while talking and breathing [8], being detected for several hours on different surfaces [9].

Respiratory infections can be classified by the causative virus (*e.g.* influenza) or clinically according to the clinical syndrome. Symptoms may include fever, nonproductive cough, coryza, sneezing, dyspnoea, myalgia, fatigue and nonexudative pharyngitis [10].

The clinical spectrum can encompass asymptomatic infection, upper respiratory tract infection and lower respiratory tract infection that can result in pneumonia or acute respiratory distress [11], and systemic infection [12].

The severity of viral respiratory illness varies widely and severe disease is more likely in older patients with or without comorbidities. Infants may have more severe disease for some organisms. Morbidity may result directly from viral infection, or may be due to exacerbation of other chronic medical conditions or bacterial superinfection [13, 14].

The spread of respiratory virus infections varies between countries and regions, depending on differences in population, geography, climate, immunisation coverage and socioeconomic status [15–17].

Immunology

The first line of defence against respiratory viral infections includes intrinsic defences such as mucus and antiviral peptides. When these are circumvented, viruses enter the epithelial cells by recognising viral components *via* Toll-like receptors and intracellular receptors (figure 1) [18], and initiate the inflammatory response. Innate cells such as dendritic cells, alveolar macrophages, natural killer cells and neutrophils are recruited. All these cells promote an antiviral response and are important for the establishment of adaptive responses. Concurrently, these inflammatory cells may be important in driving innate immune-mediated tissue damage, a process which also occurs in TB [19].

T-cells contribute to the generation of the B-cell response and cell-mediated immunity leading to viral clearance. In particular, B-cells produce antibodies that may neutralise the respiratory viruses directly by binding to viral surface proteins or activating the complement cascade (figure 1) [20, 21]. T-follicular helper cells, a specialised subset of CD4⁺ T-cells, play a critical role in protective immunity helping B-cells produce antibodies against foreign pathogens [22]. Viral clearance is also mediated by CD8⁺-specific T-cells with cytolytic activity. The protective antiviral T-cell response is a T-helper cell type 1 (Th1) response mainly mediated through interferon (IFN)- γ production [22–25]. Moreover, to prevent lung

tissue damage, all these responses are finely modulated; regulatory mechanisms adopted by T-cells such as cytokine secretion, upregulation of inhibitory receptors [26] or expansion of the T-regulatory cell subset lead to a balance between tissue damage and clearance of the virus. The immune systems of neonates, infants, children and adults are different, both in their composition and functional responsiveness to infectious diseases [27, 28].

Regarding the response to *Mycobacterium tuberculosis*, after mycobacterial dissemination to the lymph nodes, dendritic cells present bacterial antigens to T-cells and prime them [29, 30]. Priming occurs around 10 days post-infection in the mediastinal lymph nodes, and is followed by generation of effector T-cells [31] and Th1 CD4⁺ T-cells that lead to the formation of granulomas. Granulomas are organised structures where T-cells and B-cells surround innate immune cells (macrophages and neutrophils) with a fibrotic capsule to generate a hypoxic environment to prevent *M. tuberculosis* growth [32], with hypoxia potentially worsening tissue destruction in TB [33]. CD4 Th1 host responses are crucial [34], especially at the beginning for TB control [35, 36]. Regarding the CD8 T-cell component, mouse studies have revealed a relatively smaller role of CD8⁺ T-cells in protection against *M. tuberculosis* infection [37], and an even smaller contribution of B-cells and humoral immunity [38]. Differently, in human studies, *M. tuberculosis*-specific CD8 T-cells have been associated with active TB [39–44], both in HIV-uninfected and -infected patients [35, 42, 45], and in response to recent infections [43, 44]. Increased CD8 T-cell response is associated with *M. tuberculosis* load and longitudinal studies have shown a decrease of this CD8 T-cell response during anti-TB treatment [39, 40]. It has been shown that the cells are differently modulated over the course of TB disease, suggesting a role in TB pathogenesis that is not yet fully elucidated [46–48].

Bacille Calmette–Guérin vaccination as a potential intervention against COVID-19

Bacille Calmette–Guérin (BCG), the current vaccine against TB, has important protective effects against other infections. In randomised trials, BCG reduced infant mortality by around 40% [49] and respiratory infections other than TB by 70% in adolescents [50]. These “nonspecific effects” of BCG vaccination are explained by epigenetic and metabolic reprogramming of innate immune cells, a process termed “trained immunity” [51]. Clinical evidence suggests that BCG may protect against viruses [52], and BCG-vaccinated healthy adults re-challenged with live yellow fever vaccine showed improved antiviral immunity and decreased viral loads [53]. In mice, BCG vaccination protects against influenza A, lowering viral replication and lung injury [54, 55].

A recent ecological analysis suggested that BCG vaccination may also protect against COVID-19 [56]. Countries without universal policies of BCG vaccination (e.g. Italy, the Netherlands and the USA) seem more severely affected by COVID-19 compared with countries with universal BCG policies. However, such ecological studies that relate country aggregate and individual data should be interpreted with caution. Also, COVID-19 has shown a recent increase since publication of the analysis [56] in low- and middle-income countries and may still be underreported, confounders such as age were not taken into account, and variable BCG policies over time affect individual BCG coverage [57–59]. Several large randomised controlled trials (RCTs) currently evaluating the effect of BCG vaccination against COVID-19 in thousands of healthcare workers and elderly, in the Netherlands, Australia and other countries, will provide evidence to support or refute BCG as a cheap and rapidly scalable preventive measure against COVID-19 and other viral respiratory infections.

Influenza H1N1 and lung disease

The two most serious impacts of influenza virus on the lung are the development of pneumonia and exacerbation of pre-existing pulmonary disease [60]. Such events seem rare and variable during most seasonal influenza periods, but may be more frequent and severe in pandemics. During the 2018–2019 season, there was an estimated 32 million cases of influenza resulting in 32000 deaths in the USA [61]. H1N1, the virus behind the 1918 and 2009 pandemics, appears to cause more rapid and severe pneumonia than other strains, with higher rates of bacterial superinfection [62]. H1N1 also affects the paediatric population [63, 64].

Primary viral pneumonia is characterised by rapid onset of nonproductive cough, headache, myalgias, dyspnoea, tachypnoea, hypoxia and ground-glass opacities on computed tomography (CT) scans. Secondary bacterial pneumonia, which may occur concurrently or following the development of viral pneumonia, is a frequent complication. Bacterial superinfection occurs through direct damage of the respiratory epithelium with modification of local and systemic immune defence. Bacterial superinfection, mainly due to *Staphylococcus aureus* and *Streptococcus pneumoniae* [65], is reported in 20–47% of influenza patients admitted to intensive care units (ICUs). Its global prevalence varies between 0.1% and 10% according to historical surveys, and 1.59% according to a recent Korean survey during the 2009

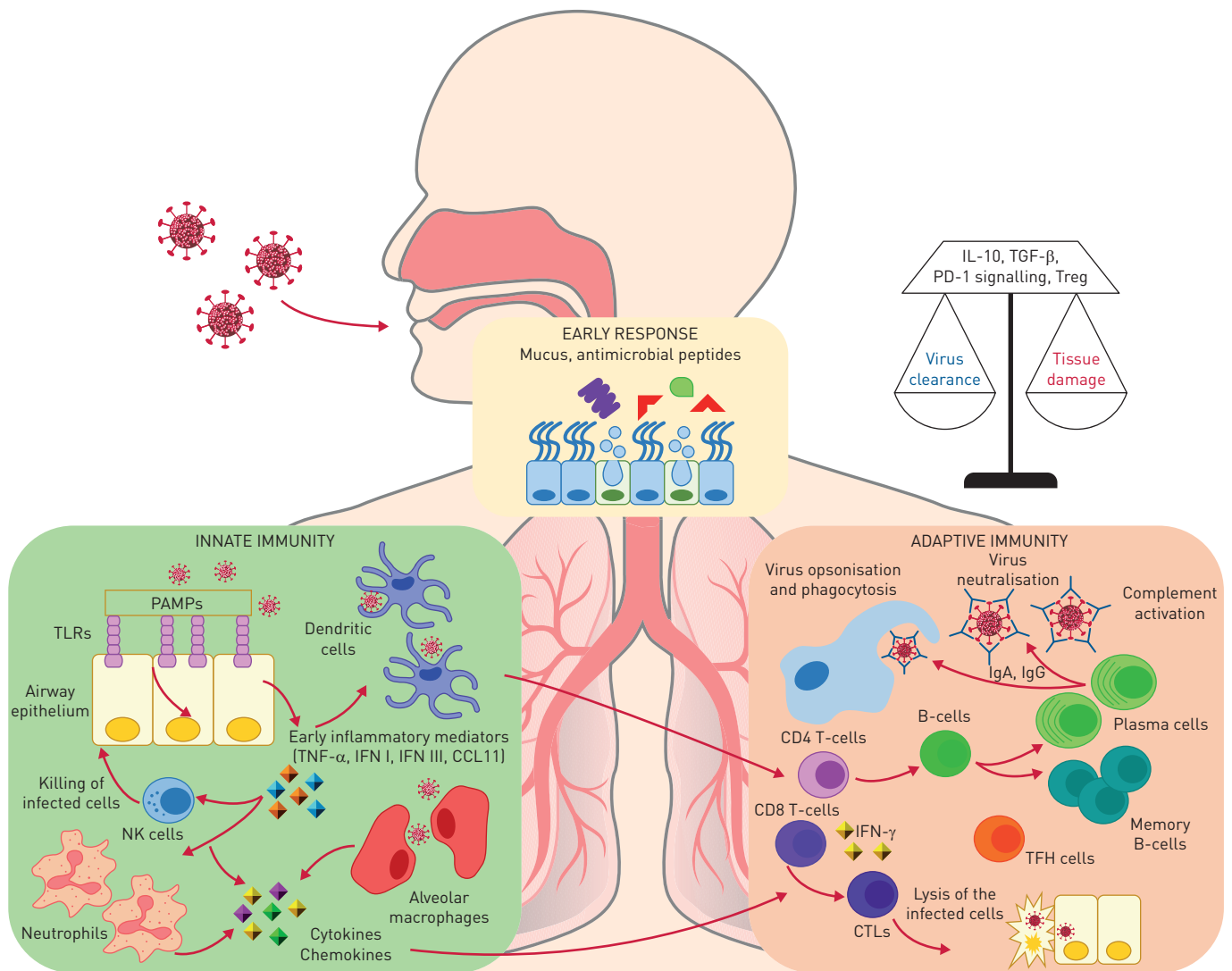


FIGURE 1 The lungs and gut are exposed to environmental substances and pathogens. The early protection response to respiratory viruses includes mucus, surfactants and antiviral peptides that can prevent initial attachment and viral entry. Respiratory viruses enter *via* the respiratory epithelium. Epithelial cells have a key role in initiating the immune response by recognising viral components [pathogen-associated molecular patterns (PAMPs)] *via* Toll-like receptors (TLRs) and intracellular receptors. These cellular sensors trigger a signalling cascade resulting in the upregulation of type I and III interferon (IFN) and the inflammatory response. This leads to differentiation of dendritic cells that mediate the induction of the adaptive immunity and promote the recruitment of innate immunity cells, in particular neutrophils and natural killer (NK) cells. NK cells have the ability to kill virus-infected cells *via* perforin–granzyme-dependent mechanisms or by the Fas–Fas ligand pathway. Moreover, alveolar macrophages, recruited monocytes and macrophages as well as dendritic cells pick pathogen components and contribute to the immune response. All of these cells produce cytokines and chemokines that are important for the establishment of the adaptive responses and of the antiviral state. The adaptive response to respiratory viruses is mediated by both T- and B-cell compartments. T-cells contribute to the generation of the B-cell response. B-cells produce antibodies that may neutralise the respiratory viruses directly by binding to viral surface proteins that are essential for entry of the virus into host cells or through the ligation of Fc receptors to trigger the complement cascade and antibody-dependent cell-mediated cytotoxicity. Antibodies are in the form of IgA, mainly in the upper respiratory tract, or IgG, in the lower respiratory tract. Viral clearance is also mediated by CD8⁺-specific T-cells with cytolytic activity. The protective antiviral T-cell response is mainly mediated by IFN- γ production and is therefore biased toward a T-helper cell (Th) 1 response, whereas other T-cell subsets such as Th2 cells and Th17 cells play a minor role and they may be responsible for lung tissue damage. Moreover, regulatory mechanisms adopted by T-cells such as interleukin (IL)-10 secretion, or upregulation of inhibitory receptors such as programmed cell death protein 1 (PD-1) or expansion of the T-regulatory (Treg) cell subsets, work to balance tissue damage and viral clearance. TNF: tumour necrosis factor; CTL: cytotoxic T-lymphocyte; TFH: T-follicular helper; TGF: transforming growth factor.

influenza A(H1N1)pdm/09 pandemic [66]. Acute respiratory distress syndrome (ARDS) and bacterial superinfection are two distinct clinical–pathological syndromes which have been described in the influenza pandemic of 1918 by MORENS and FAUCI [67]. ARDS was responsible for 10–15% of the fatal cases, while bacterial superinfection, with poorer prognosis with 85–90% fatal cases, manifests as acute bronchopneumonia, with pathogenic bacteria cultured on autopsy [60, 62, 68].

Other viral infections and lung disease (SARS and MERS)

In recent decades, previously unknown zoonotic respiratory tract infections with epidemic potential such as SARS and MERS have emerged. Human coronaviruses are usually classified into low and highly pathogenic [69]. The low pathogenic coronaviruses infect the upper respiratory tract and cause “flu-like” mild respiratory illness, while highly pathogenic coronaviruses (SARS and MERS) predominantly infect the lower airways, often causing fatal pneumonia [69].

Severe coronavirus pneumonia is often associated with rapid virus replication, massive inflammatory cell infiltration and elevated pro-inflammatory cytokine/chemokine responses producing acute lung injury and ARDS. Recent studies in experimentally infected animals strongly suggest a crucial role for virus-induced immunopathological events in causing fatal pneumonia following coronavirus infection [69].

High initial viral titres in the airways, age and comorbidities (*e.g.* hypertension, diabetes, obesity, heart failure, renal failure, *etc.*) are associated with worse outcomes [70–74].

SARS-CoV, which enters the human cell *via* the angiotensin-converting enzyme 2 receptor [75], usually presents with three phases [76]: 1) rapid viral replication with fever, cough and other nonspecific symptoms, disappearing in a few days; 2) high fever, hypoxaemia and progression to pneumonia-like symptoms, despite a progressive decline of viral replication; and 3) development of ARDS in around 20% of patients with mortality [77–79].

MERS, which enters the human cell *via* the dipeptidyl peptidase-4 receptor [80], usually starts with flu-like symptoms: fever, sore throat, nonproductive cough, myalgia, shortness of breath and dyspnoea, often progressing to pneumonia (ICU admission necessary) [73, 81]. It can also cause gastrointestinal symptoms (abdominal pain, vomiting and diarrhoea).

COVID-19 and lung disease

According to a recent report from China, COVID-19, the disease caused by SARS-CoV-2, is characterised by three clinical patterns: absence or paucity of symptoms, mild-to-moderate disease and severe pneumonia requiring admission to the ICU [82]. Dyspnoea develops after a median time of 8 days from illness onset, with a median time to ICU admission of 5 days. Up to 20% of patients require transfer to the ICU [83, 84], with consequent overwhelming of healthcare capacity. The evidence suggests that while 25% of COVID-19 patients have comorbidities including chronic obstructive pulmonary disease (COPD), diabetes mellitus, hypertension, coronary heart disease, cerebrovascular disease and malignancies, the proportion is more than 90% among those who die [83–87]. In children, COVID-19 symptoms are usually milder with better outcome than adults [88–92]. Frequency of clinical presentations and outcome appear different in Europe, with higher lethality compared with China, although this figure may change overtime due to better estimation of the total number of infections [93]. Besides virological diagnosis, imaging by chest radiography, ultrasound and CT are important for diagnosis and management. Main CT abnormalities include ground-glass opacity and consolidation [94]. The combination of CT scan findings, respiratory parameters (peripheral capillary oxygen saturation and arterial oxygen tension/inspiratory oxygen fraction ratio) and blood tests (C-reactive proteins, lymphocyte count, lactate dehydrogenase, triglycerides, ferritin, fibrinogen, D-dimer and interleukin (IL)-6) [84, 95] are important features to identify those at highest risk for ICU transfer. Lungs of dead COVID-19 patients showed oedema, proteinaceous exudate, focal reactive hyperplasia of pneumocytes with patchy inflammatory cellular infiltration and multinucleated giant cells with fibroblastic plugs in airspaces [96, 97]. A recent study reported autopsy cases contained diffuse alveolar damage with mononuclear response (CD4⁺ aggregates) surrounding thrombosed vessels, in the presence of associated haemorrhages [98–100].

HIV and lung disease

The spectrum of HIV-associated pulmonary diseases is broad and the lungs are one of the most frequently affected organ systems in HIV-infected persons regardless of age [101]. The absolute CD4 T-cell count, used as a surrogate marker of immunodeficiency, is important in guiding the aetiological evaluation of lung infections [102]. Pulmonary TB infection and reactivation are more likely with a CD4 count below 500 cells·mL⁻¹. Opportunistic infections, *Pneumocystis jirovecii*, bacterial infections, Kaposi sarcoma and extrapulmonary/disseminated forms of TB occur mainly in patients with CD4 T-cell counts below 200 cells·mL⁻¹. Cytomegalovirus infection, *Mycobacterium avium* complex and aspergillosis usually occur at CD4 counts below 50 cells·mL⁻¹. Risk factors to consider are the geographical origin that predisposes to specific disease (*e.g.* TB, coccidioidomycosis, paragonimiasis and histoplasmosis), adherence to antiretroviral therapy, prescription of *P. jirovecii* prophylaxis and presence of comorbidities. Community-acquired bacterial pneumonia occurs at all stages of HIV infection, but is more frequent in patients with profound CD4 T-cell depletion and decreases with antiretroviral therapy. Community-acquired pneumonia accounts for 35–50% of all hospital admission cases due to respiratory failure and is the main reason for ICU admission [103].

TB and respiratory viral diseases

TB and influenza

The association of TB and influenza could be bidirectional: TB may increase the susceptibility to influenza and the risk of complications, and influenza may increase the susceptibility to TB. The susceptibility to influenza appears to be greater in patients with pre-existing pulmonary disease (e.g. asthma and COPD). As a large proportion of post-TB treatment patients have long-term functional impairment, mainly COPD, which can be severe [104, 105], patients with such pulmonary sequelae may be predisposed and more susceptible to influenza infection and its complications, including mortality [106]. Furthermore, the temporary immunosuppression induced by TB may increase the susceptibility of patients to influenza infection. An excess mortality associated with influenza has been described among TB patients in South Africa [107]. TB patients have a similar prevalence of viral and bacterial co-infection as their household contacts, but TB patients often have more severe disease if they are co-infected [108].

As early as 1919, the occurrence of TB among patients surviving influenza or pneumonia, but without clear distinction between both diseases, was reported [109]. Influenza induces a temporary increase in the susceptibility to bacterial infections, exemplified by the frequent occurrence of bacterial pneumonia following viral pneumonitis [110]. Because influenza impairs the immune response, it may be expected that influenza could also promote the development of active TB among patients with latent TB infection (LTBI) [111]; however, the occurrence of TB may occur much later than the occurrence of influenza, thus making the temporal association difficult to demonstrate. There was an excess mortality from pulmonary TB during the influenza pandemics of 1889 and 1918 in Switzerland [112]. It has also been reported that summer influenza epidemics in Wuhan, China, may have contributed to the increase in reported TB cases [113]. Conversely, a report from Thailand did not demonstrate a worse outcome for patients with concurrent influenza and TB [114], and another report from Indonesia did not demonstrate a correlation between antibodies against influenza and the presence of TB, but there was an association between the level of antibody titres against influenza virus and the stage of TB [115]. Interestingly, influenza vaccination was reported to be a protective factor against TB in elderly persons in Taiwan [116]. The exact impact of concurrent influenza and TB remains uncertain.

TB and HIV

The interaction between TB and HIV is well known. Without antiretroviral treatment, the risk of LTBI progressing to active TB disease in people living with HIV and AIDS (PLWHA) is greater than in immunocompetent hosts. In PLWHA, the risk of developing TB is of the order of 10% per year [117, 118]. This elevated risk is behind the WHO recommendation of TB screening and/or preventive treatment for all PLWHA [119, 120]. New regimens as short as 1 month (daily rifapentine plus isoniazid) to 3 months (weekly rifapentine plus isoniazid) were recently recommended by the WHO, and are well tolerated and effective [119, 120]. Important programmatic implications for collaboration between TB and HIV/AIDS services exist: TB services should test for HIV (allowing treatment of TB patients with antiretrovirals and cotrimoxazole preventive therapy in patients with HIV/TB co-infection) and HIV/AIDS services should screen for LTBI, using the tuberculin skin test (TST) or IFN- γ release assay (IGRA), and initiate prompt treatment of TB or LTBI in PLWHA [119, 120]. Based on this rationale (two diseases, one patient), WHO promotes TB/HIV collaborative activities focused on three main pillars [121]: 1) establish TB/HIV collaborative mechanisms, 2) decrease the burden of TB among PLWHA and 3) decrease the burden of HIV among TB patients. Moreover, testing for other infections in addition to HIV in TB clinics may be indicated during epidemics and pandemics.

TB, SARS and MERS

TB co-infection with SARS is rare. A study of 83 patients with SARS found three patients with TB co-infection, where one patient with SARS subsequently developed TB, while the other two had TB and then developed SARS [122]. All three patients were on steroid therapy, which may have decreased viral- and/or TB-specific immunity and increased the risk of co-infection. In a different cohort of 236 SARS patients, two were diagnosed with pulmonary TB [123]. The development of TB in the presence of SARS may be due to CD4 lymphopenia during the viral infection [124], as CD4⁺ T-cells are crucial for TB-specific immunity [41, 125–129]. Lastly, one TB patient developed SARS co-infection because of a wrong admission to a cohort of SARS patients [130]. This highlights the importance of remaining vigilant to other communicable diseases, including TB, when epidemic or pandemic infections dominate media headlines [131]. TB with MERS-CoV co-infection is also rare. A report of 295 MERS-CoV patients found two TB patients [132], although it was unclear which the initial infection was.

TB and COVID-19

There may be interaction between COVID-19 and TB [133], but long-term observations are lacking [134]. Only two studies have investigated the interactions so far. In the first ever cohort of 49 patients from eight

countries, COVID-19 was diagnosed before, simultaneously or after TB [135]. In the second study including 69 patients, mortality was investigated [136]. In a separate anecdotal report, Liu *et al.* [133] reported an increased prevalence of LTBI in patients with severe COVID-19 infection and concluded that infection with *M. tuberculosis* may influence the progression and outcome of COVID-19. Evidence on the interactions between TB and COVID-19 is needed.

Respiratory viral diseases and TB in the elderly, prisoners and other vulnerable groups

The elderly (aged 65 years old or older), prisoners and other vulnerable groups such as forced migrants may reside in high-density communal settings which can perpetuate rapid infectious disease transmission during an epidemic or pandemic. Immunosenescence is an additional risk factor in the elderly [137].

Clinical presentation of these infections in the elderly can be subtle, with atypical manifestation such as delirium, and may present with complications. SARS and COVID-19 respiratory failure are well documented in the elderly [138, 139]. Conversely to influenza A (H1N1)pdm/09, a study of 4962 patients found elderly patients had less risk of respiratory failure, ICU admission or mechanical ventilation [140]. For TB, old age is a risk factor for active TB with poorer treatment outcomes [1, 141], while TB symptoms are indistinguishable from symptoms of malignancies. Moreover, the elderly may also suffer from abnormal drug absorption and/or drug toxicities due to polypharmacy for comorbidities. Similarly, immunocompromised and pregnant women may also present with complications including respiratory failure when infected with pandemic H1N1 influenza or TB [142–144].

Diagnostic tests for each group depend on available resources and on accessibility, which may be limited to none for the homeless and the incarcerated, although there should be equity in the availability of testing and treatment. The elderly may further have technical difficulties in providing quality respiratory samples for testing, such as for TB, when they have an impaired cough response. A poor quality respiratory sample inevitably delays diagnosis and contact tracing efforts.

Treatment and management of viral infections and TB includes prompt isolation of presumptive cases and of microbiologically confirmed cases depending on available resources, or even controlled release of prisoners [145]. This is part of the comprehensive strategy to mitigate transmission in nursing homes for the elderly, in refugee camps and in correctional facilities [146]. Prognosis of TB, SARS and COVID-19 tends to be worse, with higher mortality in the elderly [77, 138, 147].

Diagnostic challenges in viral diseases and TB

For prompt diagnosis of viruses causing severe acute respiratory infections [148], such as SARS-CoV, MERS-CoV and SARS-CoV-2 [149–152], and differentiation from other common bacterial infections, a strategic laboratory approach is needed. This approach requires integrating conventional virology assays, molecular platforms combining nucleic acid extraction and PCR or real-time PCR, and rapid molecular tests (RDTs) used in point-of-care minilabs (table 1). Positive results using single or multiplex RDTs may lead to adequate cohorting and management of infected patients [153]. Negative results are often less conclusive because of a lack of sensitivity and nonstandardised collection of specimens. Using metagenomic next-generation sequencing, pathogens not included in the tests can be detected including both known and novel viruses. Genomic data provides information on virulence genes [154], resistance mutations and clusters using phylogenetic approaches [155, 156]. Specific antibody detection remains useful for seroprevalence studies in selected populations and in vaccine studies. The recurrence of old pathogens and emergence of new pathogens like SARS-CoV-2 underlines the importance of worldwide virus surveillance systems [157]. For this purpose, developing protein microarrays to respiratory virus serology is useful [153].

Diagnosis of active TB relies on direct detection of *M. tuberculosis*, most often in respiratory specimens. Although culture remains the “gold standard” in terms of sensitivity and specificity, effective molecular assays to detect *M. tuberculosis* DNA are also used on platforms and in point-of-care tests [158]. Moreover, these *M. tuberculosis* molecular assays can detect mutations associated with resistance, rapidly detecting multidrug-resistant TB strains resistant to rifampin and isoniazid, allowing appropriate therapies and curbing transmission of these strains [159]. Testing immune memory to a previous TB exposure (*i.e.* LTBI), performed with TSTs and IGRAs, cannot be used as a surrogate for protection, but identifies persons who have been previously infected and can be useful to guide the TB diagnostic algorithm [160].

The massive use of molecular assays to diagnose COVID-19 introduced similar molecular platforms that could be used for detecting pathogens in respiratory specimens. The challenge lies in sample processing and RNA/DNA extraction protocols rather than cross-reactivity resulting in false-positive results. This is an opportunity to strengthen the diagnostic potential of microbiological laboratories, producing an

TABLE 1 Performance characteristics of diagnostic approaches to respiratory infection

	Respiratory viruses					<i>Mycobacterium tuberculosis</i> complex				
	Sensitivity	Specificity	Time to result [#]	User friendly	Unknown or uncommon viruses	Sensitivity	Specificity	Time saving	User friendly	Drug susceptibility testing
Molecular diagnosis										
Manual NAAT	+++	+++	-	-	-	+++	++	+	-	+
Automated NAAT	+++	+++	+	+	-	+++	+++	++	++	++
POCT-NAAT	+	++	+++	+++	-	+++	+++	+++	+	+
NGS [¶]	-	-	-	-	++	+	+++	-	-	+++
Microscopy[*]										
Culture [§]	-	+	-	-	+	+++	+++	-	-	+++
Antigen detection										
POCT	-	+	+++	++	-	^f	++	+++	+++	NA
ELISA	++	++	-	-	-	NA	NA	NA	NA	NA
Immunodiagnosis										
Serology	NRU	NRU	-	-	-	Not recommended	Not recommended	NRU	NRU	NA
IGRA	ES	ES	-	-	-	++	+	++	+	NA
TST	NA	NA	NA	NA	NA	++	+	++	++	NA

Quantitation: -: very poor; +: poor; ++: good; +++: excellent. NAAT: nucleic acid amplification test; POCT: point-of-care test; NGS: next-generation sequencing; NA: not applicable; NRU: not routinely used; ES: experimental settings only; IGRA: interferon-γ release assay; TST: tuberculin skin test (Mantoux test); TB: tuberculosis. [#]: considering only the time of the procedure <2 h; [¶]: metagenomics and whole-genome sequencing; ^{*}: immunofluorescence microscopy on respiratory samples to detect the most common viruses, or Ziehl-Neelsen or auramine/rhodamine staining to detect acid-fast bacilli; [§]: viral culture established in several eukaryotic cell lines and mycobacterial culture in liquid or solid media; ^f: the only approved antigen POCT for TB detects lipoarabinomannan in urine samples and has been licensed to diagnose TB in HIV-infected patients and to monitor therapy.

invaluable asset to improve diagnosis against other infections including TB. At the same time, the expertise, tools and networks developed for TB diagnosis could aid the rapid implementation of molecular diagnosis of COVID-19 and other viral infections. Tests (lateral flow assay and ELISA based) to rapidly detect antigens in swabs or respiratory secretions and to determine serological evidence of recent and past infection and evidence of neutralising antibodies are currently being evaluated.

Impact of new, potential and existing drugs for viral diseases and COVID-19 on TB therapy

80% of COVID-19 cases are generally mild and self-limiting, and may require no treatment. Lacking licensed drugs for SARS-CoV-2, therapeutic approaches in severely ill patients are limited to supportive care and empirical use of antibiotics to prevent or treat secondary infections [84, 86, 161]. To provide active treatment for SARS-CoV-2, drugs potentially inhibiting viral replication are of interest (figure 2) [162].

TB patients with COVID-19 may require an adapted therapeutic approach compared with patients without COVID-19. Switching to intravenous anti-TB drugs is recommended for patients in intensive care in order to optimise drug exposure in critically ill patients [163]. Therapeutic solutions for COVID-19 in TB patients need to be considered in the perspective of anti-TB treatment. However, more evidence ideally from clinical trials is necessary.

Lopinavir/ritonavir (Kaletra)

Lopinavir/ritonavir is a widely studied treatment for COVID-19, although evidence of efficacy is still limited [164]. In non-TB patients, the combination, studied in an open-label RCT, did not show any virological or clinical response compared with standard of care [165]. For the treatment of TB/HIV co-infected patients, lopinavir/ritonavir is not recommended in combination with rifampicin due to cytochrome P450 (CYP) induction. Superboosting of lopinavir/ritonavir by additional ritonavir in children on rifampicin-based TB treatment could be attempted as comparable drug exposure was achieved in situations without rifampicin [166]. Alternatively, rifabutin at a dose of 150 mg once daily has been used in combination with lopinavir/ritonavir [167].

Chloroquine phosphate and hydroxychloroquine

Chloroquine and hydroxychloroquine are being evaluated in several clinical trials for therapy and prophylaxis against SARS-CoV-2 following promising *in vitro* results [162, 168]. In the absence of results from well-designed clinical studies, clinical benefit is currently unknown [169, 170]. Both drugs have immunomodulatory properties [171, 172]. Chloroquine has been shown to reduce tumour necrosis factor- α production and receptor-mediated signalling in monocytes, which could prevent SARS-CoV-2-induced severe inflammatory response [173]. TB physicians should be careful when combining these drugs with TB drugs such as moxifloxacin, bedaquiline, delamanid and clofazimine due to the risk of increased QTc prolongation. An ongoing trial was halted due to irregular heart rates and

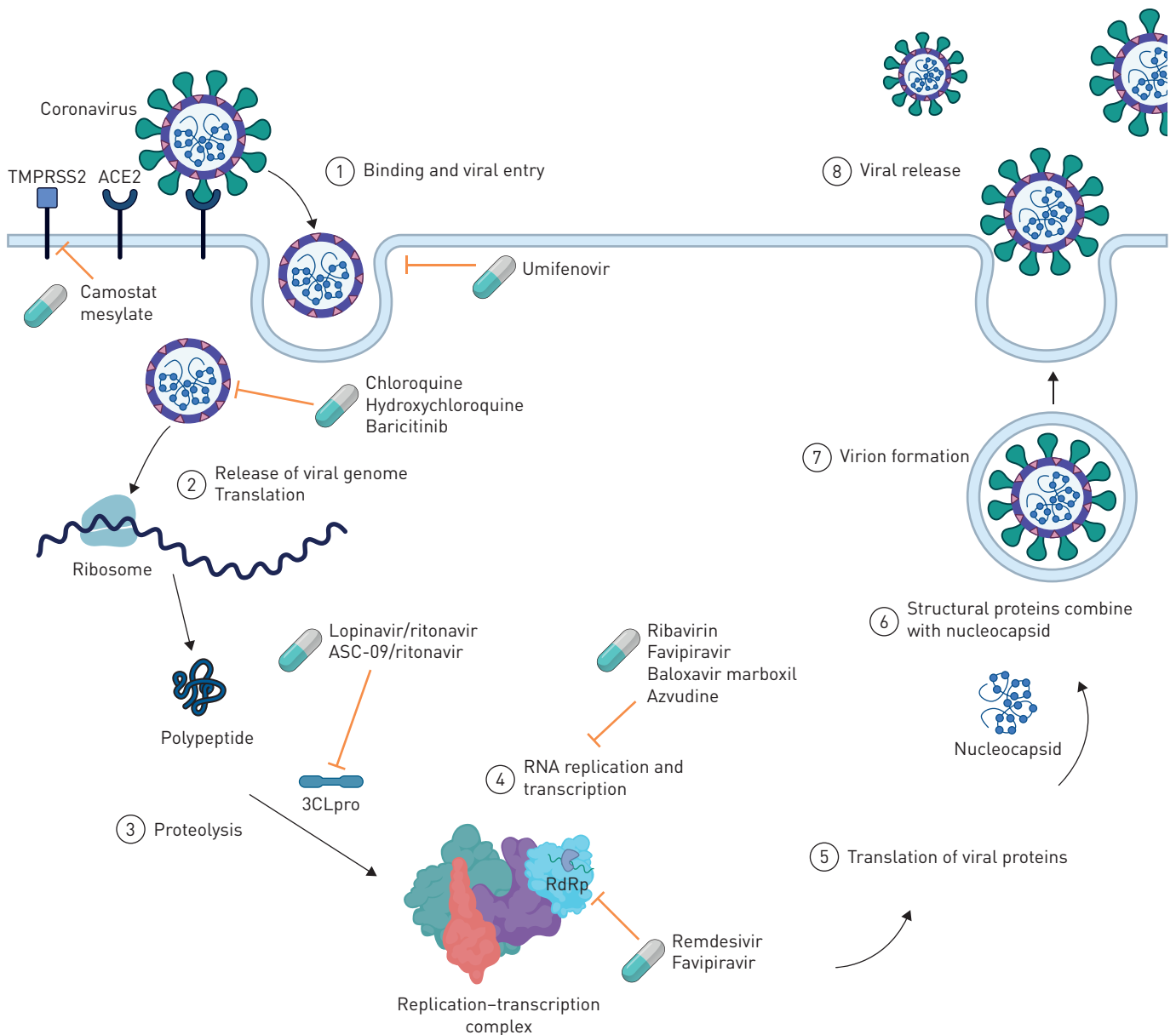


FIGURE 2 Proposed mechanism of action of drugs used for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). SARS-CoV-2 can enter the cell through angiotensin-converting enzyme 2 (ACE2) and type II transmembrane serine protease (TMPRSS2). Camostat mesylate acts as an inhibitor of TMPRSS2 and umifenovir can inhibit the viral entry to the cell [180, 228, 229]. Chloroquine, hydroxychloroquine and baricitinib mechanisms of action are not fully understood; however, it is proposed that these drugs affect viral entry. Baricitinib also inhibits the AP-2-associated protein kinase [173, 180, 230]. Lopinavir/ritonavir and ASC-09/ritonavir as protease inhibitors inhibit the proteolysis. Lopinavir/ritonavir inhibits specifically the proteinase 3CLpro [231]. Ribavirin and favipiravir both have wide antiviral activity and have the potential to inhibit SARS-CoV-2 RNA replication [232–234]. Azvudine, a nucleoside reverse transcriptase inhibitor, also inhibits RNA replication [235]. A probable mechanism of action for baloxavir marboxil is the inhibition of transcription through inhibiting cap-dependent endonuclease [236]. Favipiravir and remdesivir inhibit the RNA-dependent RNA polymerase (RdRp), which results in reduced RNA synthesis [180, 233, 234, 237]. Adapted from “Coronavirus Replication Cycle” (2020; <https://app.biorender.com/biorender-templates>).

increased risk of fatal heart arrhythmia [174]. The US Food and Drug Administration (FDA) has also issued a warning on these two drugs [175]. In addition, rifampicin increases chloroquine CYP3A4- and CYP2D6-mediated metabolism to desethylchloroquine and bisdesethylchloroquine [176]. It is unclear whether these metabolites are active against SARS-CoV-2.

Steroids

Intravenously administered steroids have been recommended for selected non-TB patients with ARDS, preferably in a trial setting [86, 161]. However, the role of steroids to reduce ARDS in TB patients is limited as data of good quality to support the use of steroids outside the treatment of TB meningitis is scarce [163]. Evidence on the use of steroids in COVID-19 is awaited.

Drug interactions

Potential drug–drug interactions (DDI) are presented in table 2. The summary of interactions is largely based on evaluations made from pharmacokinetics and toxicity profiles of drugs given alone and comparatively, when co-administered with other drugs in a separate study, in the absence of real dedicated DDI studies. The summary includes effects on drug exposure, monitoring/action and potential mechanisms.

Immunomodulatory drugs

To reduce the inflammatory response, inhibitors of IL-1 and IL-6 and the Janus kinase JAK1/JAK2 inhibitor baricitinib are being studied [177, 178]. Azithromycin may be of potential interest for its immunomodulatory effect [179], although data on its efficacy are lacking and its effect on antimicrobial resistance should be considered. IFNs are being tested because of their stimulatory activities for innate antiviral responses [162].

Novel drugs

Antiviral candidates such as azvudine, baloxavir marboxil, favipiravir, remdesivir, ribavirin and umifenovir are being tested for COVID-19 (figure 2) [162, 180–182]. Remdesivir is a nucleotide analogue showing *in vitro* activity against SARS-CoV-2 [182]. Remdesivir is being studied in two large phase 3 RCTs (ClinicalTrials.gov: NCT04252664 and NCT04257656), of which one multicentre trial conducted in Hubei, China, in severe COVID-19 showed no difference in time to clinical improvement or mortality benefit [183]. Umifenovir has shown *in vitro* activity against SARS-CoV-1 [184] and improved radiological findings when added to lopinavir/ritonavir in a small RCT [185], and seems suitable for further development.

Controlling viral diseases and TB: strengths and opportunities

Principles of viral containment

Globalisation, increased urbanisation resulting in large vastly populated and overcrowded cities, and the development of fast mass transit networks and consequent ease of travelling means that a virus can spread across a country or a continent in just a few hours. In the absence of a vaccine or an effective treatment, the tools to control a new viral infection have remained the same as during the 1918 influenza pandemic, *i.e.* early public health interventions designed to reduce the risk of transmission and spread of infection such as increased respiratory hygiene, cough etiquette and hand washing, voluntary isolation of infected individuals or households as well as quarantine of their contacts, followed by voluntary or mandatory physical distancing measures, restrictions on travel and transportation, and dissemination of basic infection prevention and control messages and advice to the general population [93]. National lockdown is an extreme measure that, while potentially reducing transmission, may also result in the collapse of the economy of a country. These nonpharmaceutical countermeasures aim at reducing the impact of COVID-19 by minimising the number of contacts that result in disease transmission and, thus, reducing the effective reproduction number R_0 to below 1. The reduction of the number of cases during the epidemic peak is crucial to reduce the burden on the healthcare services and other related sectors, and aims to flatten the curve by spreading cases over a longer period of time. This approach, while not necessarily reducing the total number of cases, gains time necessary for the development, production and distribution of effective and safe pharmaceuticals (*i.e.* vaccines and antiviral drugs), implementation of adequate hospital response and obtaining necessary ICU equipment as well as more sensitive diagnostic tests.

Infection control refers to the different methods and strategies deployed to reduce or prevent the incidence and/or transmission of infections (see the following subsection on “Airborne infection control and workplace safety”). Containment, through early detection, investigation and reporting of cases, together with contact tracing with self-isolation, aims at containing, preventing or delaying the spread of the disease in the community. Geographical containment in a defined area relies on measures to restrict the virus spreading beyond the “containment zone” or “cordon sanitaire”, including pharmacological and public health interventions such as intensified surveillance and laboratory testing, movement restrictions in and

TABLE 2 Drug interactions between tuberculosis (TB) and potential COVID-19 medications

	WHO first-line TB drugs				WHO second-line TB drugs													
	INH	RIF	EMB	PZA	Group A			Group B			Group C							
					LFX	MFX	BDQ	LZD	CFZ	Cs	DLM	IMI/CIS	MEM	AMI	STR	ETO	PTO	PAS
Antivirals																		
Atazanavir [238]	-	↓X	-	-	-♥	↑♥	↑♥	-	-♥	-	↑♥	-	-	-	-	-	-	-
Baloxavir marboxil [239]	-	↓X#	-	-	-	-	△+	-	△ ^s	-	△+	-	-	-	-	-	-	-
Favipiravir [240, 241]	-	-	-	-U	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Galidesivir	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
Lopinavir/ritonavir [242, 243]	-	↓X	-	-	-♥	↓♥	↑♥	-	-♥	-	↑♥	-	-	-	-	-	-	-
Oseltamivir	-	-	-	-	-	-	-	-	-	-	-	-	-	△ ^f	△ ^f	-	-	-
Remdesivir [242]	-	↓X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Ribavirin [242]	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Umifenovir [244, 245]	-	↓X#	-	-	-	△ ^{fl}	△+	-	△ ^s	-	△+	-	-	-	-	-	-	-
Antibacterials																		
Azithromycin [246]	-	-	-	-	-♥	-♥	-♥	-	-♥	-	-♥	-	-	-	-	-	-	-
Antiprotozoals																		
Chloroquine [247-249]	-	↓X	-	-	-♥	-♥	-♥	-	-♥	-	-♥	-	-⚡	-⚡	-🌀	-🌀	-	-
Nitazoxanide	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

Continued

TABLE 2 Continued

	WHO first-line TB drugs				WHO second-line TB drugs														
	INH	RIF	EMB	PZA	Group A				Group B				Group C						
					LFX	MFX	BDQ	LZD	CFZ	Cs	DLM	IMI/CIS	MEM	AMI	STR	ETO	PTO	PAS	
Immunomodulators																			
Anakinra	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Baricitinib	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Hydroxychloroquine [247, 250]	-	↓X	-	-	-♥	-♥	-♥	-	-♥	-	-♥	-⚡	-⚡	-	-	-	-	-	-
Interferons [251–253]	-	-	-	-	-♥	-♥	-♥	-	-♥	-	-♥	-⚡	-⚡	-⌘	-⌘	-	-	-	-
Tocilizumab [242, 254]	-	-	-	-	-	-	-	-♯	-	-	-	-	-	-	-	-	-	-	-

Interaction symbols:

- ↓ Decreased exposure to the TB drug and action required (action: dose adjustment or monitoring)
- ↑ Increased exposure to the TB drug and action required
- ↓ Decreased exposure to the COVID-19 drug and action required
- ↑ Increased exposure to the COVID-19 drug and action required
- No significant interaction predicted based on metabolic pathway (does not mean absence of interaction)
- △ Possible interaction based on metabolism and clearance, but no specific data available
- NA No available pharmacokinetic data

Monitoring/action symbols:

- ♥ Requires ECG monitoring due to the risk of QT and/or PR prolongation, or other cardiac abnormalities
- ♯ Requires full blood count monitoring
- U Uric acid monitoring
- ⚡ Monitor for potential seizures (rare)
- ⌘ Monitor for ototoxicity
- X Should not be administered together

Mechanism symbols:

- ♥ CYP-mediated mechanism
- ⌘ UGT enzyme-mediated glucuronidation

WHO: World Health Organization; INH: isoniazid; RIF: rifampicin; EMB: ethambutol; PZA: pyrazinamide; LFX: levofloxacin; MFX: moxifloxacin; BDQ: bedaquiline; LZD: linezolid; CFZ: clofazimine; Cs: cycloserine; DLM: delamanid; IMI/CIS: imipenem/cilastin; MEM: meropenem; AMI: amikacin; STR: streptomycin; ETO: ethionamide; PTO: prothionamide; PAS: p-aminosalicylic acid; CYP: cytochrome P450; UGT: UDP glucuronosyltransferase. #: recommended based on predicted interaction; †: UGT 1A1 is involved in moxifloxacin metabolism and could be involved in umifenovir metabolism (mainly UGT 1A9); *: both drugs are metabolised by CYP3A4; ‡: CYP3A4 is involved in the metabolism of baloxavir (minor extent) and umifenovir, and clofazimine is a CYP3A4 inhibitor; †: both drugs primarily undergo renal excretion.

out of the containment zone, and monitoring the area immediately surrounding the containment zone (buffer zone) for secondary infections [186].

Delaying the spread of infection can be achieved by early identification and treatment of cases, monitoring and follow-up of contacts, physical distancing measures such as proscribing public or religious gatherings and closing schools (impacting working parents), sports events or businesses, reducing contact between people. It aims at lowering the epidemic peak, reducing the burden of cases at a given time and maintaining it below healthcare capacity.

A systematic analysis of the responses to the 1918 influenza pandemic showed that in the USA, cities that introduced early social isolation measures experienced a significant reduction of viral spread, approximately 50% lower peak death rates and nearly 20% lower cumulative excess mortality than cities that did not, with a consequent reduction on healthcare pressures [187].

Airborne infection control and workplace safety

Airborne infection control in healthcare settings uses a hierarchy of control measures based on elimination of sources of infection, engineering controls, administrative controls and personal protective equipment (*i.e.* surgical masks for infectious patients and respirators for healthcare workers and visitors) [188–190]. This approach is described in detail in TB guidelines [188, 189], but can be extended also for viral infections including COVID-19 [191, 192]. As presently under discussion in countries under a post-lockdown phase perspective, the concept is valid also to ensure workplace safety in nonhealthcare settings.

While N95/N99 and FFP2/FFP3 respirators or higher-level respirators including disposable filtering face piece respirators, powered air purifying respirators, elastomeric respirators (defined as per USA and European standards) and eye protection are recommended to protect healthcare staff and other exposed individuals at the workplace level (after adequate training), the use of surgical masks is debated [193, 194]. Although there is agreement on the use of surgical masks to limit the spread of droplet nuclei for isolated symptomatic patients, the potential mass use of surgical masks to limit the community spread of COVID-19 during the early stages of infection and from asymptomatic individuals is strongly discussed [191, 193, 194]. Arguments against their wide use are based on the false sense of protection (*e.g.* the individual feels the surgical mask protects him/her from acquiring infection) as well as the potential risks of moisture retention, long mask re-use and limited filtration capacity [195]. While the WHO is revising its recommendations, the use of masks among community members has been re-evaluated [194, 196, 197]. Recent ECDC guidance states that face masks used by the general population may reduce the spread of the infection in the community by minimising the excretion of respiratory droplets from infected individuals who have not yet developed symptoms or who remain asymptomatic [198]. In general, all infection control measures are important to prevent infections and render workplaces safe.

The stability of SARS-CoV-2 is similar to SARS-CoV-1, and studies indicate that aerosol and fomite transmission of SARS-CoV-2 is plausible [9], and can be associated with nosocomial spread and superspreading events since it can remain viable and infectious in aerosols for hours and on surfaces up to days [199].

Current evidence provides support for direct contact and respiratory droplets as predominant routes of SARS-CoV-2 transmission [200], and highlights the importance of environmental surface cleaning with a hospital-grade disinfectant and meticulous hand hygiene.

SARS-CoV-2 is inactivated by common disinfection measures such as 5 min contact with household bleach [201]. The following disinfectants kill the virus: ice-cold acetone (90 s), ice-cold acetone/methanol mixture (40/60, 10 min), 70% ethanol (10 min), 100% ethanol (5 min), paraformaldehyde (2 min) and glutaraldehyde (2 min). Commonly used brands of hand disinfectants also inactivate SARS-CoV (30 s) [202]. The ECDC guidance on disinfection of environments in healthcare and nonhealthcare settings potentially contaminated with SARS-CoV-2 recommends products with virucidal activity licensed in the national markets or 0.05% sodium hypochlorite (dilution 1/100, if household bleach is used, which is usually at an initial concentration of 5%). For surfaces that can be damaged by sodium hypochlorite, products based on ethanol of at least 70% can be used [203]. The virus is sensitive to heat (60°C for 30 min) [202] and UV radiation (60 min) [204].

Outside the host, the virus can survive for 4 days in diarrheal stool samples with an alkaline pH [202], more than 7 days in respiratory secretions at room temperature, for at least 4 days in undiluted urine, faeces and human serum at room temperature [201], up to 9 days in suspension, 60 h in soil/water, more than 1 day on hard surfaces such as glass and metal [202], up to 48 h on plastic surfaces [205], and 6 days in the dried state [202].

The virus does not survive well after drying on paper, but lasts longer on disposable gowns compared with cotton gowns [201]. Human coronavirus 229E can remain infectious on high-touch environmental surfaces (polyvinylchloride, laminate, wood and stainless steel) for at least 7 days at ambient temperature (24°C) and relative humidity conditions (around 50%) [206].

The specific features of COVID-19, which spreads very rapidly with a short incubation period and infects exponentially thousands of individuals in all age groups [207], calls for the implementation of specific containment measures as discussed in the previous subsection on “Principles of viral containment”.

Human resources, equipment and new approaches to clinical management

The COVID-19 pandemic is, first and foremost, a health crisis [208]. However, it is rapidly also becoming an economic crisis. In a vicious circle, the reduction in economic activities reduces money circulation, tax revenues and finances available for establishing the public health countermeasures needed to control the pandemic. At the same time, social protection measures to ensure a minimum salary to the many workers who cannot be supported by their employers increases the financial constraints at the government level. The poverty generated by the economic crisis is likely to have medium- and long-term consequences, particularly in resource-limited countries, with increases in malnutrition and poverty-related diseases, which include TB.

To mitigate the consequences of this or future pandemics it is important countries develop specific plans with adequate human and financial resources [209]. This will prevent or limit resources currently reserved for other purposes (e.g. for TB programmes) becoming diverted to the emergency [210], including the shortage of PCR reagents being used for COVID-19 diagnosis which may impact on molecular TB diagnosis. Moreover, the emergency plan should be able to ensure rapid procurement and distribution of diagnostics, drugs, ventilators, masks, personal protective equipment and respirators needed to ensure an adequate response and adequate human resources [209, 211]. Telemedicine would be an important approach to deliver care, especially as a means to reduce the risk of cross-contamination caused by close contact [212]. To be effective, as part of an emergency response, telehealth would need to be routinely used in the health system. This would imply a change in the management and the redesign of existing models of care. Moreover, a central system of controlling the pandemic is needed (e.g. in Italy, the Italian Civil Protection Dept [213], which is normally dedicated to seismic hazard or natural disaster assessment and intervention). A central system will ensure prompt coordination of the emergency response and implementation of the emergency plan [211].

Impact of “fear” of viral infections on health services and TB services

There are many factors affecting access to healthcare: affordability and physical/geographical accessibility are essential; sociodemographic factors (ethnicity, sex, age, marital and socioeconomic status) and psychological factors such as fear can significantly hinder or delay access to TB diagnosis and care.

Fear, defined as an instinctive emotional reaction to a specific, identifiable and immediate threat such as a dangerous animal, infection, deportation or imprisonment, has a protective function associated with defensive behaviours such as hiding or fight-or-flight responses. Fear of TB itself, fear of discrimination (either self-stigma or fear of being stigmatised by others, including healthcare providers) and fear of factors associated with healthcare such as the fear that receiving a diagnosis of TB or TB treatment could affect the way they are perceived by society and even lead to deportation or exclusion are well-recognised barriers to timely access to care [214].

In addition, physical distancing measures imposed to reduce the transmission of COVID-19 (SARS-CoV-2) such as self-quarantine, closure of all nonessential services including small clinics, movement restrictions and limited access to public transport, police patrols, and enforced isolation measures all have a potential deleterious impact on access to TB care. These factors affect all groups, but disproportionately impact those minorities more often afflicted by TB such as migrants, refugee and asylum seekers, ethnic minorities or the poor. The effect on management of nonpandemic conditions including TB, strokes and myocardial infections is not only because of the unwillingness of individuals to attend healthcare facilities for fear of catching infection, but also because anything associated with fever may wrongly be assumed to be caused by the pandemic organism.

Lessons on COVID-19 and TB: lessons learned and common solutions

Country response to COVID-19

The response of European Union (EU)/European Economic Area (EEA) countries and the UK to the COVID-19 epidemic is provided in table 3 [215]. By April 3, 2020, 25 out of 31 (81%) countries had closed all educational institutions, higher education and secondary schools, primary schools, and day care/nurseries; in some countries primary schools (two out of 31 (6%)) and/or day care/nurseries (six

out of 31 (19%)) remained open. Public spaces were closed in 30 out of 31 (97%) countries, including closure of museums, theatres, cafes and restaurants, and gyms and sport facilities. Mass gatherings had been cancelled in all 31 countries. In some countries this meant that indoor and outdoor gatherings of more than five persons were forbidden and in others only larger gatherings of more than 1000 persons were banned. Finally, more than half of EU/EEA countries and the UK (18 out of 31 (58%)) implemented a full national lockdown, *i.e.* stay-at-home orders for regions or entire countries. Of the 13 countries without national lockdown, one country issued stay-at-home recommendations to the general population and nine countries issued stay-at-home recommendations for risk groups. The assessment of the response shows that there has been a lack of uniformity in the implementation of different measures. An analysis of the efficacy of the different measures at the end of the emergency should guide future policies.

TB and COVID-19 interactions

There are multiple interactions at the different levels of prevention, diagnosis and treatment between the responses to TB and COVID-19 [135, 136, 216]. Similarities and differences related to infection control and workplace safety have been discussed.

For diagnosis, the possibility of using a platform like Xpert offers synergies, although the risk is that during the COVID-19 emergency the existing equipment will be diverted from TB diagnosis. Personal protective equipment is needed to protect laboratory personnel handling viral specimens [135, 136].

Cough is the pivotal symptom to diagnose TB. In the absence of adequate diagnostic tools, the presence of cough, fever and other nonspecific symptoms complicates differential diagnosis of TB, COVID-19 and other respiratory infections [135].

The necessary physical distancing policies are likely to negatively affect active case finding as well as community-based activities which are important to manage TB (and HIV) in high TB burden countries [216]. However, they may reduce transmission of TB, which has already been shown for influenza [217, 218]. Both clinical and programmatic approaches to TB treatment are also affected in high-burden countries and drug procurement (particularly for second-line drugs) is also likely to suffer.

Last but not least, the expected major economic crisis which will follow the COVID-19 pandemic may result in increased poverty, social disturbances and malnutrition with a profound impact increasing TB incidence and mortality [219–221].

What we can and cannot learn from TB programmes in response to the COVID-19 pandemic

Political commitment, strategic planning, community mobilisation, and research and development are key elements in the battle against either TB or COVID-19. While a sustainable and effective treatment programme is the key to successful control of TB, good pandemic preparation with adequate emergency response capacity is needed when confronted with viruses with pandemic potential such as SARS-CoV-2 or influenza. Early and determined action to contain the COVID-19 epidemic at its localised or importation stage will give more time for reinforcement of care facilities, manpower and supplies [222]. Triage of suspected cases and patients according to their isolation and treatment needs will help to maximise the healthcare throughput with limited resources. Simultaneous measures to contain the spread of the epidemic is necessary to limit the mounting patient load and avoid major breaches in nosocomial infection control and total collapse of the healthcare system.

Breaking the transmission link is the main method for containing an infectious disease in the absence of an effective vaccine. Controlling a respiratory infection at source is often more cost-effective than targeting multiple intervention points downstream for environmental control or personal protection [223]. For TB, wearing a surgical mask can reduce the transmission by 56% [224]. The evidence for the protective effect of cloth masks is limited and contradictory [195, 225]. For both diseases, early case detection is important for source control, but case finding may be delayed by inconspicuous or nonspecific symptoms [135]. TB can be stopped at source by effective treatment. Isolation for the whole infectious period is needed to contain the spread of COVID-19. Like TB, contact tracing may only pick up a small portion of COVID-19 cases when widespread local transmission occurs with unclear transmission links [226]. Physical distancing [227] and adjunctive use of face masks for unavoidable person-to-person contact are then the only effective measures to slow the otherwise exponential growth of the COVID-19 epidemic [193]. The current physical distancing and stay-at-home measures imposed due to the COVID-19 pandemic will pose challenges to TB programmes to provide the necessary diagnosis, treatment and care for the people and communities affected by TB. To avoid disruption of TB services, innovative approaches (*e.g.* virtual care, digital health, community monitoring solutions, *etc.*) will be necessary [196].

TABLE 3 Response measures undertaken in European Union (EU)/European Economic Area (EEA) Member States and the UK at the national level as of April 3, 2020

	Closure of educational institutions			Closure of public spaces	Mass gathering cancellations	Stay-at-home order (enforced)	Stay-at-home recommendation (general population)	Stay-at-home recommendation (risk group)
	Higher education and secondary school	Primary school	Day care or nursery					
Austria	✓	✓	✓	✓	✓	✓	✓	
Belgium	✓	✓		✓	✓	✓		
Bulgaria	✓	✓	✓	✓	✓			✓
Croatia	✓	✓	✓	✓	✓			
Cyprus	✓	✓	✓	✓	✓	✓		
Czech Republic	✓	✓	✓	✓	✓	✓		
Denmark	✓	✓	✓	✓	✓			✓
Estonia	✓	✓		✓	✓	✓		
Finland	✓	✓		✓	✓			✓
France	✓	✓	✓	✓	✓	✓		
Germany	✓	✓	✓	✓	✓			✓
Greece	✓	✓	✓	✓	✓	✓	✓	
Hungary	✓	✓	✓	✓	✓			
Iceland	✓			✓	✓			✓
Ireland	✓	✓	✓	✓	✓	✓	✓	✓
Italy	✓	✓	✓	✓	✓	✓		✓
Latvia	✓	✓		✓	✓	✓		
Liechtenstein	✓	✓	✓	✓	✓		✓	✓
Lithuania	✓	✓	✓	✓	✓			
Luxembourg	✓	✓	✓	✓	✓	✓		✓
Malta	✓	✓	✓	✓	✓			✓
Netherlands	✓	✓	✓	✓	✓	✓		✓
Norway	✓	✓	✓	✓	✓			✓
Poland	✓	✓	✓	✓	✓	✓		
Portugal	✓	✓	✓	✓	✓	✓		✓
Romania	✓	✓	✓	✓	✓	✓		✓
Slovakia	✓	✓	✓	✓	✓			
Slovenia	✓	✓	✓	✓	✓	✓		
Spain	✓	✓	✓	✓	✓	✓	✓	✓
Sweden	✓				✓			✓
UK	✓	✓	✓	✓	✓	✓	✓	✓

The data on response measures are based on information available from official public sources as of Friday April 3, 2020 at 18:00 and may not capture measures being taken by countries that are not reported on publicly available websites. The situation is evolving rapidly and this represents a snapshot of the measures that countries in the EU/EEA and the UK reported to date. The response measures displayed are national measures, reported on official public websites. Response measures collected include: mass gathering cancellations (for specific events or a ban on gatherings of a particular size); closure of public spaces (including restaurants, entertainment venues, nonessential shops, etc.); closure of educational institutions (including day care or nursery, primary schools, and secondary schools and higher education); stay-at-home recommendations for risk groups or vulnerable populations (e.g. the elderly, people with underlying health conditions, physically disabled people, etc.); stay-at-home recommendations for the general population (which are voluntary or not enforced); and stay-at-home orders for the general population (these are enforced and also referred to as "lockdown"). The data on response measures has several limitations. First, there is substantial heterogeneity in physical distancing policies and their implementation between countries. For instance, the level of enforcement of measures may vary between countries and there may be specific rules and exceptions to the measures, making interpretation of the data challenging. The measures displayed in these figures are measures reported at a national level and it should be noted that due to the evolution of the outbreak in certain regions, regional or local measures often preceded national ones. The exact dates of introduction were often available from official sources but delays in their implementation may have occurred. Additionally, availability of public data from official government sources varies among countries. For some countries, data are no longer available on official websites concerning measures that are no longer in force, which may result in the data for more recent measures being more complete.

Consensus statements

The consensus statements derived from the Delphi process and the level of agreement achieved are summarised in table 4. The ECDC PHE did not participate in the Delphi process and development of table 4.

The scores, from the 37 questionnaires received, exceeded 3 (from intermediate to highest score) in all of the 18 statements, ranging from 3.1 to 4.8 out of a maximum of 5. The lowest scores were found for the statements on hydroxychloroquine/chloroquine and possible protective effect of BCG (Statements 5 and 12), where less

TABLE 4 Consensus statements derived from the Delphi process and the level of agreement achieved

Consensus statement	1 (lowest relevance)	2	3	4	5 (highest relevance)	Mean±sd
1. Large droplets increase the risk of respiratory viral infection through direct transmission.	0 (0.0)	6 (16.2)	6 (16.2)	10 (27.0)	15 (40.5)	3.9±1.1
2. Respiratory viral infections are more likely to occur in older patients (with or without comorbidities) and infants.	1 (2.7)	5 (13.5)	10 (27.0)	12 (32.4)	9 (24.3)	3.6±1.1
3. Elderly patients are more likely to develop acute respiratory distress syndrome and there is an age-related death risk.	0 (0.0)	0 (0.0)	4 (10.8)	10 (27.0)	23 (62.2)	4.5±0.7
4. Antibodies might neutralise respiratory viruses and, then, decrease the risk of recurrent infections.	0 (0.0)	4 (10.8)	7 (18.9)	17 (46.0)	9 (24.3)	3.8±0.9
5. BCG vaccination might offer protection against COVID-19. RCTs are needed.	4 (10.8)	7 (18.9)	10 (27.0)	10 (27.0)	6 (16.2)	3.2±1.2
6. Severe COVID-19 is associated with rapid virus replication, massive inflammatory cell infiltration in the lung and elevated pro-inflammatory cytokine/chemokine response.	0 (0.0)	0 (0.0)	1 (2.7)	7 (18.9)	29 (78.4)	4.8±0.5
7. High initial SARS-CoV-2 load in the airways, older age (≥65 years) and comorbidities of the infected individual are associated with worse COVID-19 outcome and thus patients with these risk factors need close attention.	0 (0.0)	0 (0.0)	3 (8.1)	7 (18.9)	27 (73.0)	4.7±0.6
8. The combination of CT scan findings (ground-glass opacity and consolidation), clinical presentation respiratory parameters (S_{pO_2} and P_{aO_2}/F_{IO_2}) and blood tests (C-reactive proteins, lymphocyte number, fibrinogen, D-dimers, IL-6) helps identifying COVID-19 patients at highest risk for ICU transfer.	0 (0.0)	1 (2.7)	0 (0.0)	10 (27.0)	26 (70.3)	4.6±0.6
9. CD4 T-cell count is key to guide the aetiological evaluation of lung infections in HIV-infected individuals.	0 (0.0)	3 (8.1)	7 (18.9)	18 (48.7)	9 (24.3)	3.9±0.9
10. Temporary immunosuppression induced by TB might increase the susceptibility to influenza viruses.	2 (5.4)	6 (16.2)	8 (21.6)	18 (32.5)	9 (24.3)	3.5±1.2
11. An excess mortality associated with influenza is found among TB patients.	1 (2.7)	5 (13.5)	8 (21.6)	15 (40.5)	8 (21.6)	3.7±1.1
12. Chloroquine and hydroxychloroquine have potential to improve the treatment success rate of COVID-19 patients. RCTs are needed.	4 (10.8)	8 (21.6)	10 (27.0)	9 (24.3)	6 (16.2)	3.1±1.3
13. Public and social distancing reduce the risk of SARS-CoV-2 transmission.	0 (0.0)	0 (0.0)	3 (8.1)	6 (16.2)	28 (75.7)	4.7±0.6
14. Appropriate use of facial masks (surgical masks in the general population; N95 for HCWs performing aerosol-producing activities) on symptomatic patients and their contacts can reduce the risk of SARS-CoV-2 infection by limiting the spread of droplet nuclei from isolated symptomatic patients.	0 (0.0)	0 (0.0)	1 (2.7)	9 (24.3)	27 (73.0)	4.7±0.5
15. SARS-CoV-2 virus remains infectious in the environment on different surfaces for days.	0 (0.0)	5 (13.5)	7 (18.9)	5 (13.5)	20 (54.1)	4.1±1.1
16. Social protection measures and specific national centralised emergency plans can reduce the healthcare and socioeconomic burden of respiratory viral infections resulting in epidemics/pandemics.	0 (0.0)	0 (0.0)	4 (10.8)	14 (37.8)	19 (51.4)	4.4±0.7
17. Stigma and social discrimination affect all virus-infected population groups but disproportionately the minorities.	0 (0.0)	5 (13.5)	7 (18.9)	12 (32.4)	13 (35.1)	3.9±1.1
18. Late implementation of national lockdown can itself alone be effective in reducing the burden of COVID-19 but it has serious impact on the society and the economy.	3 (8.1)	3 (8.1)	6 (16.2)	8 (21.6)	17 (46.0)	3.9±1.3

Data are presented as n (%), unless otherwise stated. BCG: bacille Calmette–Guérin; RCT: randomised controlled trial; SARS-CoV-2: severe acute respiratory syndrome coronavirus 2; CT: computed tomography; S_{pO_2} : arterial oxygen saturation measured by pulse oximetry; P_{aO_2} : arterial oxygen tension; F_{IO_2} : inspiratory oxygen fraction; IL: interleukin; ICU: intensive care unit; TB: tuberculosis; HCW: healthcare worker.

evidence is available. The highest scores were found for the statements on risk factors, outcomes and transmission (Statements 6 and 7), and for public health and prevention measures (Statements 13 and 14).

The majority of experts scored 4 or 5 (high scores) in all statements except Statements 5 and 12.

The Delphi process suggests the experts tended to agree on the relevance of the statements where more solid and high-quality evidence is available.

Conclusions

Altogether in this consensus-based document we describe the effects of epidemic and pandemic viral infections (SARS, MERS, influenza A (H1N1)pdm/09, HIV and COVID-19) and the interactions with TB and the lung; the majority of these diseases are droplet-borne. The diagnostic tools range from nucleic acid detection and molecular techniques to immunoassays and traditional cultures for *M. tuberculosis*. Core management issues were discussed, including drugs, DDIs, novel therapies, and principles of infection control and workplace safety. COVID-19 and TB interactions were discussed in-depth based on the scant information available so far, which requires broad and in-depth research into the subject. New approaches to clinical management and country responses specifically to COVID-19 were elaborated by the expert panel, as well as opportunities and lessons for future responses.

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