

Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active. right and left heart catheterisations and more subtle imaging techniques<sup>12</sup> might be able to better differentiate pulmonary hypertension due to left heart disease from PAH.

## Christian Gerges, \*Irene M Lang

Department of Internal Medicine II, Division of Cardiology, Medical University of Vienna, Waehringer Guertel 18–20, 1090 Vienna, Austria

irene.lang@meduniwien.ac.at

CG declares that he has no conflicts of interest. IML has served as a consultant and as a member of scientific advisory boards for Actelion, Bayer-Schering, and Novartis. She has also been an investigator in trials involving these companies and has received research grants from these companies.

- 1 Pepke-Zaba J, Delcroix M, Lang I, et al. Chronic thromboembolic pulmonary hypertension (CTEPH): results from an international prospective registry. *Circulation* 2011; **124**: 1973–81.
- 2 Ghofrani HA, D'Armini AM, Grimminger F, et al. Riociguat for the treatment of chronic thromboembolic pulmonary hypertension. *New Engl J Med* 2013; 369: 319–29.
- 3 Tanabe N, Sugiura T, Tatsumi K. Recent progress in the diagnosis and management of chronic thromboembolic pulmonary hypertension. *Respir Investig* 2013; **51:** 134–46.
- 4 Pulido T, Adzerikho I, Channick RN, et al. Macitentan and morbidity and mortality in pulmonary arterial hypertension. *New Engl J Med* 2013; 369: 809–18.

- 5 Eyries M, Montani D, Girerd B, et al. *EIF2AK4* mutations cause pulmonary veno-occlusive disease, a recessive form of pulmonary hypertension. *Nature Genet* 2013; published online Dec 1. http://dx.doi.org/10.1038/ ng.2844.
- 6 Bonderman D, Ghio S, Felix SB, et al. Riociguat for patients with pulmonary hypertension caused by systolic left ventricular dysfunction: a phase IIb double-blind, randomized, placebo-controlled, dose-ranging hemodynamic study. *Circulation* 2013; **128**: 502–11.
- 7 Redfield MM, Chen HH, Borlaug BA, et al. Effect of phosphodiesterase-5 inhibition on exercise capacity and clinical status in heart failure with preserved ejection fraction: a randomized clinical trial. JAMA 2013; 309: 1268–77.
- 8 Naeije R, Vachiery JL, Yerly P, Vanderpool R. The transpulmonary pressure gradient for the diagnosis of pulmonary vascular disease. Eur Respir J 2013; 41: 217–23.
- 9 Gerges C, Gerges M, Lang MB, et al. Diastolic pulmonary vascular pressure gradient: a predictor of prognosis in "out-of-proportion" pulmonary hypertension. Chest 2013; 143: 758–66.
- 10 Rich S, Dantzker DR, Ayres SM, et al. Primary pulmonary hypertension. A national prospective study. Ann Intern Med 1987; **107**: 216–23.
- 11 Hoeper MM, Huscher D, Ghofrani HA, et al. Elderly patients diagnosed with idiopathic pulmonary arterial hypertension: results from the COMPERA registry. Int J Cardiol 2013; 168: 871–80.
- 12 Kraigher-Krainer E, Shah AM, Gupta DK, et al. Impaired systolic function by strain imaging in heart failure with preserved ejection fraction. J Am Coll Cardiol 2013; published online Oct 18. http://dx.doi.org/10.1016/j. jacc.2013.09.052.

## Emerging respiratory viral infections: MERS-CoV and influenza 🕢

Each year, the world faces the rising burden of viral respiratory infections. These infections are of major importance to public health because of the lack of specific therapeutic and preventive measures, and, more specifically, the lack of vaccines for most of these viruses.<sup>1</sup>

In 2013, the emergence of three new respiratory viruses—varian influenza virus (H3N2v), H7N9, and Middle East Respiratory Syndrome coronavirus (MERS-CoV)—was of particular concern. These viruses all have the potential to cause widespread pandemics with substantial morbidity and mortality. The two influenza viruses (variant influenza virus and H7N9) were reported to cause, potentially severe, disease in human beings.

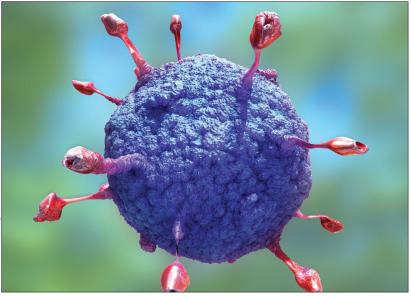
Infection with MERS-CoV was initially described in a patient from Saudi Arabia and then retrospectively identified in patients from Zarqa, Jordan.<sup>2,3</sup> As of Dec 2, 2013, 163 cases of infection with MERS-CoV have been reported, with 71 fatalities.<sup>4</sup> MERS-CoV can cause sporadic infection, infection among families, and, of particular concern, infection among healthcare workers.<sup>5</sup> The largest outbreak of MERS-CoV was described in Al-Hasa, the eastern province of Saudi Arabia.<sup>5</sup> Fever and cough was present in most cases, with shortness of breath in almost half of all cases, and gastrointestinal symptoms in about a third. A study of the largest reported outbreak of MERS-CoV<sup>6</sup> estimated the median incubation period to be 5.2 days (95% CI 1.9-14.7), and reported a high rate of person-to-person transmission in 21 of 23 cases in health-care settings.6 Case-fatality rate was high (65%) in this outbreak.7 Of more than 417 household and health-care contacts, symptoms of MERS-CoV developed in only seven people.<sup>6</sup> A subsequent largescale phylogenetic analysis of 21 genome sequences and inclusion of the previous nine published MERS-CoV genomes showed that multiple introductions of MERS-CoV and lower R<sub>o</sub> values were possible.<sup>7</sup> Thus MERS-CoV has not yet reached pandemic potential. Transmission within Saudi Arabia was consistent with movement of an animal reservoir, animal products, or infected people. The source of the infection has yet to be identified, although bats and camels have been implicated.

A clinical and epidemiological analysis of 47 cases showed that infection with MERS-CoV occurred



\$2213-2600(13)70255-8

predominantly in men,<sup>5</sup> although this finding did not hold after an interim analysis of 133 cases.<sup>8</sup> Most patients who were initially infected with MERS-CoV had underlying comorbid medical conditions and laboratory testing showed that most of these cases had raised concentrations of lactate dehydrogenase and aspartate aminotransferase associated with thrombocytopenia and lymphopenia.<sup>5</sup> In preparation for the Hajj this year, the largest recurring religious mass gathering worldwide, the Saudi Ministry of Health recommended that certain individuals postpone their participation. Groups particularly at risk from infection include pregnant women, children younger than 12 years, adults older than 65 years, and those with chronic or acute diseases. The concern over the pandemic potential of MERS-CoV was estimated in two recent publications.9 In one optimistic view,9 the estimated MERS-CoV R<sub>o</sub> was 0.69 compared with the R<sub>o</sub> for prepandemic severe acute respiratory syndromecoronavirus of 0.80. This optimistic estimate downplayed the possibility of a MERS-CoV pandemic, which supported the recommendations of the Third Meeting of the International Health Regulations Emergency Committee,<sup>10</sup> that MERS-CoV does not warrant international measures to curtail Hajj-related travel. Sporadic cases MERS-CoV continue to be reported, and thus continued vigilance and further studies are needed to close the knowledge gap in MERS-CoV epidemiology and clinical presentations.<sup>5</sup>



A novel avian-origin influenza A virus, H7N9, was initially described in human beings in China on March 30, 2013. At that time, three patients developed fatal pneumonia and were subsequently diagnosed as infected with H7N9.11,12 Since this initial description, H7N9 has resulted in 139 infections and 45 fatalities.13 Review of the initial 111 patients with H7N9 infection<sup>14</sup> showed a high rate of admission to intensive-care units (76.6%) with a mortality rate of 27%. Infected patients were older adults with a median age of 61 years, with twice as many men infected than women. It is also interesting to note that most patients infected with H7N9 had a pre-existing medical condition. Subsequent analysis of 136 laboratory confirmed cases showed a mortality bias toward men older than 50 years.<sup>15</sup> Of the total cases, 7% were reported in individuals younger than 20 years, with no fatal cases in this age group.<sup>15</sup>

The emergence of these viral respiratory infections (H7N9 and MERS-CoV) showed a similar initial pattern: the predominant involvement of older men and the presence of comorbid conditions in most cases. This pattern seems to be due to recognition bias, because younger cases are now being identified who have no underlying medical conditions. In facing future challenges of emerging respiratory viruses such as influenza there is a clear need for the development of effective influenza vaccines that target the conserved antigenic structures of influenza virus. Continued contact of humans with animals creates an added risk of development of zoonotic diseases, adaptation of the new virus rendering it infectious to humans, and possible efficient transmission of these viruses among the human population.

The emergence of respiratory viruses that cause significant disease in human beings is a major risk to the global economy and the health of the human population. The potential effect of newly discovered viruses calls for a better understanding of the human-animal interface, the development of rapid diagnostic tests, and effective antiviral and immunomodulatory therapies.<sup>13</sup> The eradication of respiratory viruses is not possible and thus the development of effective vaccines directed against the conserved antigens of these viruses would be extremely welcome. Finding predictors of severe disease and the initiation of antiviral drugs early in the course of many respiratory viral infections might prove to be beneficial.

## Jaffar A Al-Tawfiq, \*Ziad A Memish

Saudi Aramco Medical Services Organization, Saudi ARAMCO, Dhahran, Saudi Arabia (JAA-T); Department of Medicine, Indiana University School of Medicine, Indianapolis, IN, USA (JAA-T); Global Centre for Mass Gathering Medicine, Public Health Directorate, Ministry of Health, Riyadh 11176, Saudi Arabia (ZAM); and Department of Medicine, Al-Faisal University, Riyadh 11176, Saudi Arabia (ZAM)

zmemish@yahoo.com

We declare that we have no conflicts of interest

- 1 WHO. Research needs for the Battle against Respiratory Viruses (BRaVe). http://www.who.int/influenza/patient\_care/clinical/BRaVe\_Research\_ Agenda\_2013.pdf (accessed Nov 18, 2013).
- 2 Zaki AM, van Boheemen S, Bestebroer TM, Osterhaus AD, Fouchier RA. Isolation of a novel coronavirus from a man with pneumonia in Saudi Arabia. N Engl J Med 2012; 367: 1814–20.
- 3 Hijawi B, Abdallat M, Sayaydeh A, et al. Novel coronavirus infections in Jordan, April 2012: epidemiological findings from a retrospective investigation. East Mediterr Health J 2013; 19 (suppl 1): S12–18.
- 4 WHO. Middle East respiratory syndrome coronavirus (MERS-CoV) update. http://www.who.int/csr/don/2013\_12\_02/en/index.html (accessed Dec 7, 2013).
- 5 Assiri A, Al-Tawfiq JA, Al-Rabeeah AA, et al. Epidemiological, demographic, and clinical characteristics of 47 cases of Middle East respiratory syndrome coronavirus disease from Saudi Arabia: a descriptive study. *Lancet Infect Dis* 2013; **13**: 752–61.

- 6 Assiri A, McGeer A, Perl TM, et al. Hospital outbreak of Middle East respiratory syndrome coronavirus. N Engl J Med 2013; **369:** 407–16.
- 7 Cotten M, Watson SJ, Kellam P, et al. Transmission and evolution of the Middle East respiratory syndrome coronavirus in Saudi Arabia: a descriptive genomic study. *Lancet* 2013; published online Sept 20. http://dx.doi. org/10.1016/S0140-6736(13)61887-5.
- 8 Penttinen PM, Kaasik-Aaslav K, Friaux A, et al. Taking stock of the first 133 MERS coronavirus cases globally—is the epidemic changing? Euro Surveill 2013; 18: 20596.
- 9 Breban R, Riou J, Fontanet A. Interhuman transmissibility of Middle East respiratory syndrome coronavirus: estimation of pandemic risk. *Lancet* 2013; **382**: 694–99.
- 10 WHO. WHO statement on the third meeting of the IHR Emergency Committee concerning MERS-CoV. http://www.who.int/mediacentre/ news/statements/2013/mers\_cov\_20130925/en/index.html (accessed Nov 29, 2013).
- 11 Gao R, Cao B, Hu Y, et al. Human infection with a novel avian-origin influenza A (H7N9) virus. N Engl J Med 2013; **368**: 1888–97.
- 12 Chen Y, Liang W, Yang S, et al. Human infections with the emerging avian influenza A H7N9 virus from wet market poultry: clinical analysis and characterization of viral genome. *Lancet* 2013; **381**: 1916–25.
- 13 WHO. Number of confirmed human cases of avian influenza A(H7N9) reported to WHO. Report 9—data in WHO/HQ as of 12 August 2013, 14:45 GMT+1. http://www.who.int/influenza/human\_animal\_interface/ influenza\_h7n9/09\_ReportWebH7N9Number.pdf (accessed Nov 18, 2013).
- 14 Gao HN, Lu HZ, Cao B, et al. Clinical findings in 111 cases of influenza A (H7N9) virus infection. N Engl J Med 2013; 368: 2277-85.
- 15 Dudley JP, Mackay IM. Age-specific and sex-specific morbidity and mortality from avian influenza A(H7N9). J Clin Virol 2013; 58: 568–70.

## Tuberculosis: progress and challenges in product development and delivery

Although global tuberculosis control is improving, progress remains slow; about 1.3 million people died of tuberculosis in 2012, more deaths than from any other single infectious agent (other than HIV).<sup>1</sup> WHO emphasised two major challenges in 2013: identifying and treating the "missing 3 million" people who develop active tuberculosis every year and whose condition never becomes known to national tuberculosis control programmes; and the crisis of multidrug-resistant (MDR) tuberculosis, in which three of four people with MDR tuberculosis are never diagnosed, and less than half of those diagnosed are successfully treated.<sup>1</sup>

New diagnostics, drugs, and vaccines are crucial for transformational progress. 2013 heralded both progress and disappointment in these areas. In diagnostics, progress was made in the global scale-up of Xpert MTB/RIF (Cepheid Inc, Sunnyvale, CA, USA), a molecular test for tuberculosis that also enables rapid detection of resistance to rifampicin. More than 4 million MTB/RIF cartridges had been procured under concessional pricing as of September, 2013, and a US\$25.9 million UNITAID project was announced in March, 2013, for rollout in 21 countries.<sup>2</sup> Based on new evidence, WHO recommended in October, 2013, that Xpert MTB/RIF be used as the initial diagnostic test in adults and children with presumed MDR tuberculosis or HIV-associated tuberculosis (and, conditionally, for all forms of tuberculosis), as well as for diagnosis of certain types of extrapulmonary tuberculosis.<sup>3</sup>

The first randomised trial of Xpert MTB/RIF was published in October, 2013.<sup>4</sup> This pragmatic, multicentre trial of nurse-led Xpert MTB/RIF versus sputum-smear microscopy in people with symptoms of tuberculosis presenting to primary care clinics in four African countries showed Xpert MTB/RIF to be more accurate than smears and more effective in leading to same-day treatment initiation. However, the primary outcome of tuberculosis-related morbidity was equivalent between the two arms, as was treatment initiation by 56 days.<sup>4</sup> The authors postulated that this lack of difference in morbidity was the result of high empirical treatment rates among smear-negative individuals with ongoing symptoms. Results from additional trials in Brazil and

(W

Published **Online** December 23, 2013 http://dx.doi.org/10.1016/ S2213-2600(13)70256-X