

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. together with clonal lineage, and it was not possible to disentangle which genetic features influenced clinical outcome. Interestingly, Wang and colleagues¹ report that the O2v2 lipopolysaccharide O locus variant was associated with a lower 30-day mortality compared with other O loci in South America and the USA (adjusted odds ratio 0.34, 95% CI 0.15-0.78, after adjusting for age at culture, origin, Charlson comorbidity index, Pitt bacteremia score, region, and culture anatomical source).1 The O2v2 locus is highly conserved among ST258 CRKP7 and results in the production of O2afq-type lipopolysaccharide,⁸ which is considered less immunogenic than the other common lipopolysaccharide types, O2a and O1.9.10 Hence, there is a potential mechanism by which O2v2-harbouring CRKP infections (including most ST258, but few ST11) could carry a lower mortality risk. However, further work is needed to explore this hypothesis; the association should be replicated across distinct clones to confirm the specific effect of the O2v2-encoded antigen.

Wang and colleagues¹ conclude that the substantial regional differences between CRKP populations and their associated patient cohorts might preclude extrapolation of the findings to other settings. We agree that caution is required and advocate for further prospective, multi-site, multi-regional studies to characterise and understand the drivers of geographical variations. In particular, it will be important to include sites in so-far underrepresented regions in Africa, the Middle East, the Pacific islands, and south and southeast Asia, as the sparse data available suggest that the common CRKP clones and carbapenemases described here and elsewhere (namely ST258 and ST11 producing K pneumoniae carbapenemases) might be very rare or entirely absent

from many of these settings.^{2,4,5} The availability of data from diverse populations would facilitate analyses that are better powered to detect associations between clinical outcomes and the genomic features of CRKP, and would also support efforts to design novel, globally relevant control strategies targeting these important pathogens.

We declare no competing interests.

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- Wang M, Earley M, Chen L, et al. Clinical outcomes and bacterial characteristics of carbapenem-resistant Klebsiella pneumoniae complex among patients from different global regions (CRACKLE-2): a prospective, multicentre, cohort study. Lancet Infect Dis 2021; published online Nov 9. https://doi.org/10.1016/S1473-3099(21)00399-6.
- 2 Sands K, Carvalho MJ, Portal E, et al. Characterization of antimicrobialresistant Gram-negative bacteria that cause neonatal sepsis in seven low- and middle-income countries. Nat Microbiol 2021; 6: 512-23.
- David S, Reuter S, Harris SR, et al. Epidemic of carbapenem-resistant 3 Klebsiella pneumoniae in Europe is driven by nosocomial spread. Nat Microbiol 2019; 4: 1919-29
- Wyres KL, Lam MMC, Holt KE. Population genomics of Klebsiella pneumoniae. Nat Rev Microbiol 2020; 18: 344-59.
- Argimón S, David S, Underwood A, et al. Rapid genomic characterization 5 and global surveillance of Klebsiella using Pathogenwatch. bioRxiv 2021; published online June 23. https://doi.org/10.1101/2021.06.22.448967 (preprint).
- 6 Zhang Y, Jin L, Ouyang P, et al. Evolution of hypervirulence in carbapenemresistant Klebsiella pneumoniae in China: a multicentre, molecular epidemiological analysis. J Antimicrob Chemother 2020; 75: 327-36.
- Szijártó V, Guachalla LM, Hartl K, et al. Both clades of the epidemic KPC-producing Klebsiella pneumoniae clone ST258 share a modified galactan O-antigen type. Int J Med Microbiol 2016; 306: 89-98.
- 8 Clarke BR, Ovchinnikova OG, Kelly SD, et al. Molecular basis for the structural diversity in serogroup O2-antigen polysaccharides in Klebsiella pneumoniae. J Biol Chem 2018; 293: 4666-79.
- Bulati M, Busà R, Carcione C, et al. Klebsiella pneumoniae lipopolysaccharides 9 serotype O2afg induce poor inflammatory immune responses ex vivo. Microorganisms 2021: 9: 1-15
- 10 Pennini ME, De Marco A, Pelletier M, et al. Immune stealth-driven O2 serotype prevalence and potential for therapeutic antibodies against multidrug resistant Klebsiella pneumoniae. Nat Commun 2017; 8: 1991.

(IV) COVID-19 vaccine inequity, dependency, and production capability in low-income and middle-income countries: the case of Bangladesh

Published Online January 28, 2022 https://doi.org/10.1016/ \$1473-3099(22)00028-7 Bangladesh is listed among the 57 human resource for health (HRH) crisis countries as identified by the Global Health Workforce Alliance.^{1,2} Despite shortages and maldistribution of health-care personnel, Bangladesh was known for successful immunisation

campaigns even before the COVID-19 pandemic.³ The COVID-19 vaccination drive initially was not smooth for Bangladesh, but the country has, since July 2021, regained its reputation for strong vaccination efforts. COVID-19 vaccination started on Feb 7, 2021, with

Comment

the ChAdOx1-nCoV-19 vaccine known as Covishield, (Serum Institute India [SII]). Vaccination was initially postponed while the SII suspended the export of vaccines as the number of COVID-19 infections in India increased. The Government of Bangladesh, however, managed to procure vaccines from alternative sources, which helped to alleviate its dependency on a single source. The Government intends to administer four different vaccines (Covishield, BBIBP-CorV [Sinopharm], Comirnaty [Pfizer-BioNTech], and Spikevax [Moderna]) to 170 million people in Bangladesh. As of December 13, 2021, Bangladesh has administered 86.5 million first doses and 43.3 million second doses (figure),⁴ equating to roughly 50% of the total Bangladeshi population having received at least one dose and roughly 25% having been administered two doses of COVID-19 vaccines. Despite vaccine hesitancy and commonly held misconceptions about vaccines globally, vaccine acceptance among this population is encouraging. Like other low-income and middle-income countries (LMIC), Bangladesh has been supported by COVAX, which is co-led by the Coalition for Epidemic Preparedness Innovations, Gavi, the Vaccine Alliance, WHO, and UNICEF to ensure equitable access to COVID-19 vaccines. After Indonesia, COVAX vaccine roll-out in Bangladesh ranks second highest among southeast Asian countries. This achievement is a monumental feat for Bangladesh given that highincome countries have already administered 69 times more doses than Bangladesh, showing a stratified and inequitable vaccine procurement and roll-out.⁵

Tedros Adhanom Ghebreyesus, Director-General of WHO, notes that inequitable global distribution of the COVID-19 vaccine will burden LMICs most, resulting in a catastrophic moral failure.⁶ The possibility of this moral failure accentuates the importance of sharing technologies among manufacturers to enhance the capacity of vaccine manufacturing facilities in LMICs, which could then allow vaccines to be manufactured locally and complement global efforts to ensure equitable COVID-19 vaccine supply. Both Bangladeshi Prime Minister Sheikh Hasina and Muhammad Yunus, the only Nobel Laureate in Peace (2006) from Bangladesh, have been instrumental in a campaign to declare COVID-19 vaccines a global common good. This endeavour is a joint appeal from 3000 people including international personalities, Nobel Laureates, global



Figure: COVID-19 vaccination profile in Bangladesh as of December 13, 2021

leaders, international organisations, pharmaceutical companies, and governments.⁷ In addition to procuring vaccines from a diverse portfolio of manufacturers, Bangladesh has adopted a model to become selfreliant in vaccine manufacturing. A Bangladeshi biotech company, Globe Biotech, has completed non-human primate trials of its own mRNA-based vaccine and received approval for human trials.8 Additionally, other Bangladeshi vaccine manufacturing companies are capable of addressing the void in vaccine antigen production. Experts predict that the global COVID-19 vaccine supply could be substantially increased if vaccine manufacturers would share their efforts and technological knowledge with external manufacturers capable of vaccine production.⁵ For example, Incepta Vaccine Limited (IVL), the largest human vaccine manufacturing facility in Bangladesh, has signed a memorandum of understanding with Sinopharm, China, for production of the Sinopharm BBIBP-COVID-19 vaccine in Bangladesh. This collaboration will undoubtedly expedite the efforts of Bangladesh to produce viable COVID-19 vaccines in local settings. Along with contract manufacturing, IVL is also in the race to develop its own research and development based COVID-19 vaccine in collaboration with international institutes.9 Furthermore, Bangladesh has agreed to participate in a clinical trial for a novel nasal-route COVID-19 vaccine, developed by Karolinska Institute, Sweden.

With the increasing concern of waning vaccine immunity 10 and effective results from heterologous

vaccine priming and boosting,11,12 Bangladesh should determine its policy and act quickly. Bangladesh has local vaccine research and production capacity at its disposal, along with four distinct COVID-19 vaccines. Looking forward, Bangladesh should focus on hosting clinical trials, including heterologous prime boost trials, to determine vaccine efficacy and long-term protection for its population. Although there is a debate on whether to administer a third vaccine dose as booster, evidence to support recommended schedules and target groups is scarce.13 A third dose for individuals who have already received two doses should be carefully considered only after two doses of the vaccine have been administered to the remaining unvaccinated people. To ensure maximum vaccine coverage, countries such as Bangladesh should allocate resources to local manufacturers and support large-scale research efforts, clinical trials, cold-chain technology, and international collaboration. Support could be in the form of subsidies, duty-free import of research items, or tax incentives for vaccine manufacturers and research organisations.

To bring this pandemic under control, more than 75% of the global population must be vaccinated.¹⁴ This goal becomes increasingly challenging given the unequal distribution and hoarding of vaccines by high-income countries. This inequality could be obviated, however, by strict adherence to the procurement and supply operation under the COVAX initiative. This mandate should serve as a reference for ensuring equitable global access to the COVID-19 vaccine. Discrete high vaccination performance and travel bans from high-income countries will not only be futile in bringing the pandemic under control but will also put all nations at risk unless the global population is universally vaccinated.¹⁵ Initiatives should be undertaken to acquire and distribute vaccines among LMICs. Without such initiatives there remains a growing concern for the persistent presence of COVID-19 hot spots in low-income countries, which increases the risk of escape variants of SARS-COV-2.

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- WHO. The human resources for health crisis. 2011. https://www.who.int/ workforcealliance/countries/bgd/en/ (accessed Nov 10, 2021).
- 2 WHO. Bangladesh. https://www.who.int/workforcealliance/countries/bgd/ en/ (accessed Aug 30, 2021).
- 3 Sarma H, Budden A, Luies SK, et al. Implementation of the world's largest measles-rubella mass vaccination campaign in Bangladesh: a process evaluation. BMC Public Health 2019; 19: 925.
- 4 DGHS. COVID-19 dashboard for Bangladesh. http://103.247.238.92/ webportal/pages/covid19-vaccination-update.php (accessed Aug 23, 2021).
- Burki T. Global COVID-19 vaccine inequity. Lancet Infect Dis 2021; 21: 922–23.
- 5 WHO. WHO Director-Generals opening remarks at 148th session of the Executive Board. Jan 18, 2021. https://www.who.int/director-general/ speeches/detail/who-director-general-s-opening-remarks-at-148thsession-of-the-executive-board (accessed Oct 10, 2021).
- 7 Yunus M, Donaldson C, Perron J-L. COVID-19 vaccines a global common good. Lancet Healthy Longev 2020; 1: e6–8.
- 8 The Business Standard. Globe biotech gets BMRC nod for human trials. https://www.tbsnews.net/coronavirus-chronicle/covid-19-bangladesh/ globe-biotech-gets-bmrc-nod-human-trials-333622 (accessed Nov 24, 2021).
- 9 Baylor College of Medicine. Baylor College of Medicine and Texas Children's Hospital collaborate with Incepta Vaccine Ltd. to develop a COVID-19 vaccine for Bangladesh. https://www.bcm.edu/news/baylor-college-of-medicineand-texas-childrens-hospital-collaborate-with-incepta-vaccine-ltd-todevelop-a-covid-19-vaccine-for-bangladesh (accessed Dec 14, 2021).
- Pegu A, O'Connell S, Schmidt SD, et al. Durability of mRNA-1273 vaccine-induced antibodies against SARS-CoV-2 variants. *Science* 2021; 373: 1372–77.
- 11 Barros-Martins J, Hammerschmidt SI, Cossmann A, et al. Immune responses against SARS-CoV-2 variants after heterologous and homologous ChAdOx1 nCoV-19/BNT162b2 vaccination. Nat Med 2021; 27: 1525-29.
- 12 Shaw RH, Stuart A, Greenland M, Liu X, Nguyen Van-Tam JS, Snape MD. Heterologous prime-boost COVID-19 vaccination: initial reactogenicity data. Lancet 2021; **397**: 2043–46.
- 3 The Lancet Infectious Diseases. COVID-19 vaccine equity and booster doses. Lancet Infect Dis 2021; 21: 1193.
- 14 Kreier F. Unprecedented achievement: who received the first billion COVID vaccinations? https://www.nature.com/articles/d41586-021-01136-2 (accessed Oct 10, 2021).
- 15 Mendelson M, Venter F, Moshabela M, et al. The political theatre of the UK's travel ban on South Africa. *Lancet* 2021; **398:** 2211–13.