

Feedback from operational stakeholders who manage or respond to outbreaks is that they are often too busy to review literature or obtain relevant background information to assist them with acute response. Unlike a traditional analytical outbreak investigation report, **Watching Briefs** are intended as a rapid resource for public health or other first responders in the field on topical, serious or current outbreaks, and provide a digest of relevant information including key features of an outbreak, comparison with past outbreaks and a literature review. They can be completed by responders to an outbreak, or by anyone interested in or following an outbreak using public or open source data, including news reports.

Watching brief	
<b>Title</b>	<b>Wuhan novel coronavirus 2019nCoV</b>
<b>Authors</b>	C R MacIntyre, The Biosecurity Program, Kirby Institute, University of New South Wales, Sydney, NSW, 2052, Australia
<b>Date of first report of the outbreak</b>	29 December 2019
<b>Disease or outbreak</b>	<b>Novel coronavirus 2019nCoV</b>
<b>Origin (country, city, region)</b>	Wuhan, Hubei, China
<b>Suspected Source (specify food source, zoonotic or human origin or other)</b>	Unknown. Snakes have been implicated (1), but the evidence for this is weak, and genetic analysis suggests a mammalian source such as bats. Coronaviruses arising from bats can have intermediary animal hosts.
<b>Date of outbreak beginning</b>	December 29 2019
<b>Date outbreak declared over</b>	Ongoing on January 24 <sup>th</sup> 2020
<b>Affected countries &amp; regions</b>	China (540 cases) Thailand (4 cases) Vietnam (2 cases) South Korea (1 case) Japan (1 case) Taiwan (1 case) Singapore (1 case) Hong Kong (1 case) Philippines (1 case) USA (1 case)

<b>Number of cases (specify at what date if ongoing)</b>	570
<b>Clinical features</b>	Fever, cough, shortness of breath, myalgia. Mild cases may present with a common cold like syndrome, whilst severe cases may develop severe acute respiratory syndrome and pneumonia.
<b>Mode of transmission (dominant mode and other documented modes)</b>	Coronaviruses are respiratory viruses, so can be found in the respiratory tract. Transmission is unknown yet and detailed risk factor analysis data have not yet been published. Most cases appear to be sporadic and are assumed to be from an animal source, with most cases localised to Wuhan and the initial outbreak linked to a fish market with other live animals. There has been some person to person spread, including a family in Guandong, and a single case which infected 14 health workers. One of the two cases in Vietnam also appears to be spread person-to-person from the index case in Vietnam. SARS was transmitted person-to-person, especially in the hospital setting. MERS CoV has mostly been sporadic, with some person-to-person spread and nosocomial outbreaks.
<b>Demographics of cases</b>	All cases are adults, with males and females equally affected. Anecdotal reports that people with chronic conditions are more at risk. No cases have been reported in children at this stage. The majority of the cases are in China, mainly in Wuhan.
<b>Case fatality rate (CFR)</b>	Ranges from 2-3% depending on changes in daily case and death counts.
<b>Complications</b>	Severe pneumonia, respiratory failure, death.
<b>Available prevention</b>	A vaccine is being developed by the National Institutes for Health in the USA and by other companies. A MERS CoV vaccine has been developed(2) and is a high priority for the WHO and Coalition for Epidemic Preparedness Initiatives. Whether the MERS vaccine has cross protection against 2019nCoV is unknown.
<b>Available treatment</b>	Supportive treatment only. Intensive care, ventilation and ECMO may be used for severe pneumonia and respiratory failure. Broad spectrum antivirals may have effectiveness against coronaviruses but are untested against 2019nCoV.(3)

<p><b>Comparison with past outbreaks</b></p>	<p>This is a new infection, so it can only be compared with SARS and MERS CoV.</p> <p>It appears less infectious than SARS, which had a R0 of 2. It may be closer to MERS CoV, which has a R0 close to 1, in terms of transmission. In terms of case fatality rate (CFR), the CFR with SARS was 12%, MERS CoV 26% and 2019nCoV appears to be about 2-3% based on informal reports of cases and deaths.</p> <p>The transmission appears mixed (like MERS CoV), with mostly sporadic or point-source cases and some propagated transmission from person to person. With SARS, travel-related cases in other countries frequently caused satellite epidemics with clear person to person transmission in Hong Kong, Vietnam, Singapore and Canada. This has not been seen so far with the new coronavirus. With MERS, the only outbreak outside of the Kingdom of Saudi Arabia was in South Korea, and over 60% of cases are sporadic.(4)</p>
<p><b>Unusual features</b></p>	<p>The source of infection remains unknown, although it is geographically in the city of Wuhan, Hubei, China.</p> <p>Transmission appears mainly point-source in the city of Wuhan, particularly linked to a seafood market which also sells other live animals. There has been some person-to-person spread, but like MERS CoV, most travel related cases imported to other countries have not caused satellite epidemics.</p> <p>China took the extraordinary measure of locking down Wuhan and other cities on January 23<sup>rd</sup> 2020, thereby reducing travel out of the disease epicentre. Given the timing of this epidemic around the Chinese Lunar New Year (Spring festival), when travel is at a peak, this would reduce the risk of travel-related importations of cases to other parts of China and the world.</p> <p>The phylogenetic analysis suggests low diversity (ie that the virus is not mutating rapidly) and a relatively recent origin of the virus.</p>
<p><b>Critical analysis and key questions</b></p>	<p>The key questions around this epidemic are:</p> <ol style="list-style-type: none"> <li>1. What is the source? Identifying the source can help curtail the epidemic.</li> <li>2. What is the dominant mode of transmission and what other modes of transmission are possible? This will inform optimal disease control strategies.</li> <li>3. What are the risk factors for disease? A case-control analysis is required to determine sociodemographic, clinical, behavioural and other risk factors.</li> </ol> <p>A modelling study suggests that, based on the number of travel-related cases, there could be 1000 to 9000 undetected cases of 2019nCoV. (5). A similar modelling estimation of a large proportion of undetected, asymptomatic or mild cases was made for MERS CoV,(6) but not supported by active screening studies or serological surveys of humans in affected areas.(7,8). Serological surveys in Wuhan will help determine how much mild or asymptomatic infection there may be.</p>

	<p>Until the questions above are answered, the main disease control strategies should focus on</p> <ol style="list-style-type: none"> <li>1. Enhanced disease surveillance to detect new cases early</li> <li>2. Travel – the main route of global spread. Strategies include airport screening, health communication to passengers at risk, reduction or prevention of travel (such as the lock down of Wuhan which occurred on January 23<sup>rd</sup> 2020).</li> <li>3. Hospitals and the health system – both SARS and MERS CoV caused nosocomial outbreaks. Patients with 2019nCoV will present to the health system, and if they are not suspected as cases and isolated rapidly, they may infect others. Triage, isolation and infection control are key, as well as personal protective equipment (PPE) for health care workers. Health workers paid a heavy price with SARS, with many preventable deaths due to delayed diagnosis or inadequate PPE.</li> </ol>
<p><b>References</b></p>	<ol style="list-style-type: none"> <li>1. Ji, W., Wang, W., Zhao, X., Zai, J. and Li, X. (2020), Homologous recombination within the spike glycoprotein of the newly identified coronavirus may boost cross-species transmission from snake to human. <i>J Med Virol</i>. Accepted Author Manuscript. DOI: <a href="https://doi.org/10.1002/jmv.25682">https://doi.org/10.1002/jmv.25682</a></li> <li>2. Modjarrad K, Roberts CC, Mills KT, Castellano AR et al. Safety and immunogenicity of an anti-Middle East respiratory syndrome coronavirus DNA vaccine: a phase 1, open-label, single-arm, dose-escalation trial. <i>Lancet Infectious Diseases</i>. 2019;19: P1013-1022. DOI: <a href="https://doi.org/10.1016/S1473-3099(19)30266-X">https://doi.org/10.1016/S1473-3099(19)30266-X</a></li> <li>3. Sheahan TP, Sims AC, Leist SR, et al. Comparative therapeutic efficacy of remdesivir and combination lopinavir, ritonavir, and interferon beta against MERS-CoV. <i>Nat Commun</i>. 2020;11(1):222. doi:10.1038/s41467-019-13940-6. DOI: <a href="https://doi.org/10.1038/s41467-019-13940-6">https://doi.org/10.1038/s41467-019-13940-6</a></li> <li>4. Chen X, Chughtai AA, Dyda A &amp; MacIntyre CR (2017) Comparative epidemiology of Middle East respiratory syndrome coronavirus (MERS-CoV) in Saudi Arabia and South Korea, <i>Emerging Microbes &amp; Infections</i>, 6:1, 1-6, DOI: <a href="https://doi.org/10.1038/emi.2017.40">https://doi.org/10.1038/emi.2017.40</a></li> <li>5. Imai N, Dorigatti I, Cori A, Donnelly C, Riley C, Ferguson NM. Report 2: Estimating the potential total number of novel Coronavirus cases in Wuhan City, China. 22 January 2020 - Imperial College London. WHO Collaborating Centre for Infectious Disease Modelling. MRC Centre for Global Infectious Disease Analysis, J-IDEA, Imperial College London, UK. <a href="https://www.imperial.ac.uk/mrc-global-infectious-disease-analysis/news--wuhan-coronavirus/">https://www.imperial.ac.uk/mrc-global-infectious-disease-analysis/news--wuhan-coronavirus/</a></li> <li>6. Cauchemez S, Van Kerkhove MD, Riley S, Donnelly CA, Fraser C, Ferguson NM (2013) Transmission scenarios for Middle East respiratory syndrome coronavirus (MERS-CoV) and how to tell them apart. <i>Euro Surveill</i> 18(24):pii=20503</li> <li>7. Joseph C et al (2013) Highlights and conclusions from the technical consultative meeting on novel coronavirus infection, Cairo, Egypt, 14–16 January 2013. <i>East Mediterr Health J</i> 19(Suppl 1):S68–S74. DOI: <a href="https://doi.org/10.26719/2013.19.suppl1.S68">https://doi.org/10.26719/2013.19.suppl1.S68</a></li> <li>8. Aburizaiza AS, Mattes FM, Azhar EI, Hassan AM, Memish ZA, Muth D et al (2014) Investigation of anti-MERS-Coronavirus antibodies in blood</li> </ol>

	donors and abattoir workers in Jeddah and Makkah, Kingdom of Saudi Arabia, Fall 2012. <i>J Infect Dis</i> 209:243–246. DOI: <a href="https://doi.org/10.1093/infdis/jit589">https://doi.org/10.1093/infdis/jit589</a>
--	---

**How to cite this article:** MacIntyre CR. Wuhan novel coronavirus 2019nCoV. *Global Biosecurity*, 2019; 1(3).

**Published:** January 2020

**Copyright:** Copyright © 2019 The Author(s). This is an open-access article distributed under the terms of the Creative Commons Attribution 4.0 International License (CC-BY 4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited. See <http://creativecommons.org/licenses/by/4.0/> .

*Global Biosecurity* is a peer-reviewed open access journal published by University of New South Wales.