

Elsevier has created a <u>Monkeypox Information Center</u> in response to the declared public health emergency of international concern, with free information in English on the monkeypox virus. The Monkeypox Information Center is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its monkeypox related research that is available on the Monkeypox Information Center - including this research content - immediately available in publicly funded repositories, 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 Monkeypox Information Center remains active.

Personal View

Mpox respiratory transmission: the state of the evidence

Amy Beeson, Ashley Styczynski, Christina L Hutson, Florence Whitehill, Kristina M Angelo, Faisal S Minhaj, Clint Morgan, Kaitlyn Ciampaglio, Mary G Reynolds, Andrea M McCollum, Sarah Anne J Guagliardo

The relative contribution of the respiratory route to transmission of mpox (formerly known as monkeypox) is unclear. We review the evidence for respiratory transmission of monkeypox virus (MPXV), examining key works from animal models, human outbreaks and case reports, and environmental studies. Laboratory experiments have initiated MPXV infection in animals via respiratory routes. Some animal-to-animal respiratory transmission has been shown in controlled studies, and environmental sampling studies have detected airborne MPXV. Reports from real-life outbreaks demonstrate that transmission is associated with close contact, and although it is difficult to infer the route of MPXV acquisition in individual case reports, so far respiratory transmission has not been specifically implicated. Based on the available evidence, the likelihood of human-to-human MPXV respiratory transmission appears to be low; however, studies should continue to assess this possibility.

Introduction

In 2022, human-to-human clade II monkeypox virus (MPXV) transmission occurred at an accelerated rate, resulting in tens of thousands of cases worldwide in nonendemic countries. Several factors probably contributed to the outbreak, including the virus's initial entry into a global network of sexually active men who have sex with men,¹ the waning of herd and individual immunity conferred by smallpox vaccination,² and increased recognition of and testing for the virus.

MPXV transmission can arise from many combinations of routes and sources of infection (figure). Based on its similarities to variola virus (smallpox), historical observations of MPXV, and a small amount of experimental evidence, routes of MPXV acquisition are widely thought to include percutaneous exposure, such as through direct exposure of skin, especially broken skin; direct exposure of mucous membranes, such as those found in the mouth, vagina, and rectum; and inhalation of infectious particles into the respiratory tract.3.4 Sources of infection include infected humans or animals, or alternatively, contaminated fomites (objects or materials that harbour infectious matter).5 In the ongoing outbreak that began in 2022, sexual contact among cases and the prevalence of anogenital lesions at diagnosis suggest that direct sexual contact is the primary route of acquisition;1 however, the role of inhalation of infectious viral particles in disease transmission remains uncertain and has substantial implications for public health recommendations.

In contrast to respiratory viruses such as SARS-CoV-2 and influenza that preferentially bind specific receptors found in human respiratory tract cells, orthopoxviruses such as MPXV do not bind to one specific receptor type for viral–cell surface attachment (although glycosaminoglycans have been implicated with vaccinia virus strains).⁶ Instead, for cell entry, orthopoxviruses are engulfed (at the plasma membrane or through macropinocytosis) and subsequently spread from cell to cell via actin tails.⁶

Respiratory transmission of infectious diseases has often been dichotomised into droplet and airborne transmission. Droplet transmission refers to the spread of pathogens through large particles (ie, droplets) produced during sneezing, coughing, or talking, and deposited onto the mucous membranes of a susceptible host. These large particles typically travel only short distances (ie, <2 m) before settling onto the ground or other surfaces. Airborne transmission refers to the spread of pathogens in very small particles (ie, aerosols) that can also be produced via sneezing, coughing, and talking, and remain suspended in the air for extended periods of time, thus not requiring close contact (<2 m) with an infected individual. Both types of particles can be



Lancet Microbe 2023; 4: e277–83

Published **Online** March 7, 2023 https://doi.org/10.1016/ S2666-5247(23)00034-4

Mpox Response Team (A Beeson MD, A Styczynski MD,

C L Hutson PhD, F Whitehill DVM, K M Angelo DO, F S Minhaj PharmD, C Morgan MS, K Ciampaglio MPH, M G Reynolds PhD, A M McCollum PhD, S A J Guagliardo PhD) and Epidemic Intelligence Service (A Beeson, F Whitehill, F S Minhaj), Centers for Disease Control and Prevention, Atlanta, GA, USA

Correspondence to: Dr Sarah Anne J Guagliardo, Mpox Response Team, Centers for Disease Control and Prevention, Atlanta, GA 30333, USA sguagliardo@cdc.gov



Figure: Main sources and potential acquisition routes of MPXV that have been documented historically Infection with MPXV is believed to arise from several combinations of routes of acquisition (percutaneous, mucosal, and respiratory) and sources (humans, animals, and fomites). MPXV=monkeypox virus.

infectious when they are inhaled or contact a person's mucous membranes. In reality, these types of transmission exist along a continuum, with particles of varying sizes being continually exuded, resulting in the potential for both long-range and short-range transmission for many infections. In the mpox (formerly known as monkeypox) context, lesion-derived particulate matter (eg, pulverised scab material) from non-oral lesions can also be inhaled when suspended in the air. Here, we collectively refer to transmission through the air (whether through droplet sprays or inhalation of aerosols or lesion-derived particulates) as respiratory transmission. Notably, we did not consider infection resulting from direct contact with upper respiratory tract mucosa (eg, during kissing or oral sex) to be respiratory transmission.

In this Personal View, we describe evidence for and against MPXV respiratory transmission available from key works published from 1961 to 2022, including controlled experiments with animal models, human case reports and outbreaks, and environmental studies for both clade I and the currently circulating clade II MPXV.⁷⁸

Animal models and MPXV respiratory transmission

Studies of clade I and clade II MPXV in prairie dogs and non-human primates have helped shape the current understanding of MPXV transmission. Prairie dogs are a useful animal model in laboratory experiments because they are susceptible to MPXV infection, exhibit a long incubation period, can transmit MPXV, and are the only small animal model that develops the characteristic skin rash seen in human mpox.⁹ Non-human primates are another valuable animal model because of their genetic proximity to humans, similar disease presentation (ie, characteristic skin rash), and susceptibility to infection with MPXV in the wild.¹⁰

Studies in prairie dogs have shown that inoculation via the upper respiratory tract results in experimentally infected animals with the ability to produce infectious secretions (and therefore potentially transmit virus without direct contact) with the recovery of viable virus from oropharyngeal secretions and oral and nasal swabs.³ Additionally, both the upper and lower respiratory tract can be involved in naturally occurring MPXV infection in animals: in a sooty mangabey (*Cercocebus atys*) infected in the wild in Côte d'Ivoire in 2012, viable clade II MPXV virus was isolated from a throat swab and lung tissue.¹⁰

Respiratory transmission was demonstrated with clade I MPXV (but not clade II MPXV) in a prairie dog experimental model in a 2013 study.¹¹ Eight prairie dogs (four challenged with intranasal MPXV and four naive) were housed separately in metal cages that were maintained 4 inches (roughly 10 cm) apart with multiple ventilation holes (1 inch [roughly 2.5 cm] in diameter) cut into walls that faced each other, with directional airflow

from the challenged to the naive animals. The animals were unable to touch. When challenged with clade II MPXV, three of four animals developed disease; however, no transmission occurred to the naive animals. When the authors repeated the experiment with clade I MPXV, all four challenged animals developed disease, and transmission occurred to one of four naive prairie dogs.ⁿ

Experiments with non-human primates in the 1970s also suggested possible respiratory transmission, leading scientists at the time to believe that this could be an important MPXV transmission pathway. In a 1971 laboratory study, several vellow baboons (Papio cynocephalus) were housed in separate cages in the same room with shared airflow.12 Two animals were inoculated intramuscularly with a high dose of MPXV of West African origin (clade II) and developed clinical illness consistent with mpox. Of six sentinel animals housed (in separate cages) in the same room, two (33%) became ill. The authors postulated that the two infected animals became sick during the third week of exposure. An important limitation of this study is that the distance between cages and the type of barrier between cages are not specified, leaving open the possibility of mucosal or percutaneous transmission (ie, if the animals were able to touch).

Although these studies used small numbers of animal subjects, experimental models with prairie dogs and non-human primates suggest that MPXV respiratory transmission is possible, at least for clade I MPXV. Importantly, however, the artificial approach to inoculation and simulated exposures might not replicate the conditions of human-to-human contact.

Human case reports and outbreaks

Respiratory manifestations of mpox in humans

The involvement of the respiratory tract in human mpox infections is important to consider when examining the possibilities for respiratory transmission. Although the presence of respiratory symptoms does not predict the route of acquisition, respiratory symptoms are likely to enhance the production of infectious respiratory secretions, which could, in turn, promote respiratory transmission.

Historically, case reports of human mpox have included upper or lower respiratory tract symptoms, but these can also be absent. Severe respiratory distress or bronchopneumonia have been noted late in the course of illness with clade I MPXV.¹³ Oral lesions, in addition to other oropharyngeal signs and symptoms (eg, tonsillar lesions, pharyngitis, odynophagia, and epiglottitis) have been reported both historically¹³ and during the 2022 clade II outbreak.^{14,15} Sore throat was reported in 17–37% of patients in 2022.^{14–17} Prior to 2022, cough was documented in approximately half of patients with both clade I and clade II MPXV,¹⁸ but in reports from the 2022 outbreak, this symptom, along with dyspnoea and nasal congestion, has been far less common, $^{14\!-\!17}$ with cough occurring in just 7% of patients in one cross-sectional study. 14

Both historical and recent reports also document isolation of MPXV from oral or respiratory anatomical locations. Pharyngeal swabs (n=37) and saliva (n=2) have yielded detectable MPXV via PCR in patients who have concurrent skin lesions.14 Adler and colleagues identified upper respiratory tract MPXV clade II DNA in seven patients described in a case series from the UK (2018-21) in the absence of respiratory symptoms and even after skin lesion resolution. In three patients, upper respiratory tract shedding occurred for longer than 3 weeks.¹⁹ Viral DNA was also detected during one patient's relapse of disease that occurred 6 weeks after hospital discharge and 10 weeks after initial symptom onset.19 The association between viral DNA presence and shedding of viable infectious virus from the respiratory tract leading to infection is unknown, since many of these studies did not include cell culture. However, in one study,²⁰ infectious virus was cultivated from some oropharyngeal swabs and measured by a plaque assay.

Although respiratory symptoms seem to be an uncommon manifestation of MPXV in the 2022 outbreak, the presence of viral DNA in the respiratory tract has been documented even in individuals without respiratory symptoms.¹⁴ Whether respiratory manifestations of MPXV infection or isolation of MPXV from oral or respiratory secretions in humans have any correlation with the potential for respiratory transmission remains unclear.

Outbreaks in humans

Most of the mpox transmission studies in humans before 2022 were done in African households.^{5,21–25} Close contact primarily within households has been implicated in extended transmission chains ranging from four to eight generations.^{19,21–25} The nature of close contact in household settings might entail various combinations of acquisition routes, with confounding exposures such as caregiving, children's play, bed-sharing, shared eating utensils, and sexual contact.

The 2003 US human outbreak of the clade II strain arising from infected prairie dogs provides another illustration of confounding exposures but also offers examples of activities that did not result in infection. All 47 individuals who developed symptomatic MPXV infections had contact with prairie dogs or fomites contaminated by infected animals (eg, cleaning animal cages or touching animal bedding).^{26,27} Although excluding the possibility of animal-to-human respiratory transmission during this outbreak is difficult, simply being near an infected animal (defined by the investigators as within 6 feet [roughly 1·8 m] for >3 h) was not associated with infection.²⁸ Furthermore, no health-care personnel in proximity to the affected patients (<6 feet [roughly 1·8 m]) were infected, even though 81% (46/57) reported not consistently wearing an N95 respirator and 75% (43/57) reported not consistently wearing a surgical mask.²⁹

Data on the re-emergence of clade II MPXV in Nigeria in 2017, using viral genomic analyses, demonstrated instances of human-to-human transmission occurring among four people incarcerated in a prison and in a health-care worker.^{30,31} Three of the four incarcerated people who were infected had known contact with at least one other infected person (contact included sharing a prison cell and other unspecified contact). For the remaining individual, no link was disclosed. Without the details of the type of contact between people in the prison and the types of personal protective equipment worn or types of contact by health-care personnel, drawing conclusions about possible routes of exposure during this outbreak remains challenging.

Investigations of MPXV transmission associated with air travel also can provide insights about the risk of respiratory transmission. So far, there have been no cases of transmission on international or domestic flights. Between 2018 and 2021, eight individuals were diagnosed with clade II MPXV after travelling from Nigeria to the UK,^{19,32,33} Singapore,³⁴ Israel,³⁵ and the USA.^{36,37} Among these eight cases, four were symptomatic during international flights, and contact investigations were performed for all four (table 1). No additional cases were identified in any investigated contacts for ten different trips (flights, and car or train journeys), four of which

	Destination	Transportation type	Estimated trip duration	Estimated contacts investigated	Reference	
Case 1 (Sept 2, 2018)						
Abuja, Nigeria	London, UK	Flight	8 h 55 m	~41*	Vaughan et al (2018) ³²	
London, UK	Cornwall, UK	Train	5 h 20 m	NR		
Case 2 (Sept 4, 2018)						
Lagos, Nigeria	Paris, France	Flight	8 h 20 m	~41*	Vaughan et al (2018) ³²	
Paris, France	Manchester, UK (assumed)†	Flight	1 hr 15m	~41*		
Case 3 (July 8-9, 2021)						
Lagos, Nigeria	Atlanta, GA, USA	Flight‡	13 h 2 m	144§	Rao et al (2022) ³⁷	
Atlanta, GA, USA	Dallas, TX, USA	Flight‡	2 h 14 m	5		
Dallas, TX, USA	Dallas, TX, USA	Car	NR	1		
Dallas, TX, USA	Dallas, TX, USA	Car	NR	1		
Case 4 (November, 2021)						
Lagos, Nigeria	Maryland, USA	Flight‡	13 h 55 m	9	(Kreuze M, Centers for Disease Control and Prevention, personal communication)	
Maryland, USA	Maryland, USA	Car	NR	8		

No transmission was identified in any of the contact investigations. NR=not reported. *Assumptions: each row has two sets of three seats; people sitting on both sides of the aisle were contacted. †Based on the location of Blackpool Teaching Hospital. ‡Masks worn on flights. §Those investigated included flight crew, close exposures, and far exposures who could have used the toilets.

Table 1: Transportation routes and duration of travel for symptomatic mpox patients, 2018–21, ordered by origin and date of arrival at destination

were flights longer than 8 h in duration. Importantly, masks were worn during the 2021 flights because of requirements related to the COVID-19 pandemic and might have protected against respiratory transmission.³⁷

Finally, reports from health-care and congregate settings for the 2022 outbreak are becoming available and provide useful case studies for examining the possibility of respiratory transmission in the absence of sexual or close, intimate contact. In endemic areas, health-care personnel are known to be at higher risk for MPXV infection relative to the general population;³⁸ however, in a 2022 investigation in Colorado, USA, none of 313 health-care personnel exposed to patients with mpox became infected, including seven reported to be exposed during aerosol-generating procedures (four of whom did not wear an N95 respirator).39 Additionally, transmission without close, intimate contact has not yet been documented in congregate settings such as shelters, correctional facilities, and schools. After a symptomatic resident with mpox spent 7 days in congregate housing in a prison in Chicago, IL, USA, sharing a dormitory with 57 individuals, no additional cases were identified among a subset of those exposed who were actively monitored for symptoms or consented to serological testing.40 Of ten distinct instances in which a child or adolescent with mpox attended a childcare facility or school while symptomatic, none resulted in secondary cases.41

Clinical and outbreak data show that transmission has primarily occurred during close, prolonged contact. So far, no outbreaks that clearly implicate human-to-human respiratory transmission have been reported in other congregate or residential group settings.

MPXV in the environment

Orthopoxviruses, including MPXV, are stable in the environment, and several studies have assessed the ability of the virus to persist in the air, as well as the role of fomites as a vehicle for transmission. For example, MPXV viral particles were recently found in three of four air samples taken during a bedding change in a healthcare setting, including replication-competent virus (Ct value range: 32.7-36.2).42 Air samples from which MPXV was identified were located more than 15 m away and at a height of up to 2 m and were collected over the course of 10 min. In the same study, a high degree of surface contamination of viral DNA was found in a patient room and bathroom, including surfaces unlikely to have been directly touched (eg, the air vent above the door), which could indicate non-contact contamination through droplets or aerosolised viral particles, although viral culture was not attempted for these samples (Ct value range: 25.9-33.6).42 Another investigation conducted continuous air sampling for viral DNA for 4-h sessions in a clinic suite housing patients suspected of having mpox.43 MPXV DNA was detected in six of six sampling sessions (Ct value range: $32 \cdot 0 - 38 \cdot 0$) that

occurred with confirmed patients with mpox in the room; culture was not attempted. Notably, attending health-care personnel were protected with N95 respirators and patients wore surgical masks, and transmission within the facility did not occur. A laboratory study of suspension of viral particles in the air also demonstrated the persistence of replication-competent viral particles for at least 90 h in a small (10.7 L) rotating chamber.⁴⁴

Evidence also suggests that MPXV can remain viable on surfaces for long periods, although respiratory transmission under these circumstances has not been definitively proven. A 2021 US environmental sampling study detected culturable virus on household surfaces 15 days after the infected person had left,45 which could pose a potential risk for respiratory transmission through inhalation of viral material resuspended in the air (eg, during handling of contaminated linens). A case report from the UK in 2018 described a health-care worker who was infected while handling the used bedding and clothing of a patient with mpox, despite wearing a disposable apron and gloves.46 No face mask or respirator was worn. Although respiratory exposure (eg, inhalation or direct mucosal inoculation of lesion-derived particulate matter) from a fomite source has been hypothesised, the exact nature of exposure to the bedding was unknown and any of the three acquisition routes (percutaneous, mucosal, or respiratory) was possible.

Key findings and public health implications

In this Personal View, we summarise different types of key evidence to evaluate the contribution of respiratory transmission to the spread of MPXV. The available types of evidence regarding respiratory transmission of MPXV are summarised in table 2. Laboratory experiments in prairie dogs and non-human primates attempt to isolate the respiratory route of acquisition (by eliminating other possibilities) but are challenging to extrapolate to real-life settings. Respiratory symptoms and the presence of virus in the upper respiratory tract suggest transmission potential by the respiratory route, but these data cannot definitively demonstrate respiratory transmission, nor do they allow assessment of the relative importance of the respiratory route compared with other routes. Environmental sampling studies can show the presence of replication-competent virus in the air or on surfaces, but the presence of virus does not equate to infectivity or person-to-person transmission. Outbreak reports offer insight into the likelihood of transmission under realistic conditions, but often do not have detailed exposure histories.

Equally important is what the outbreak data do not show. If respiratory transmission of MPXV between humans were commonplace, we would expect to see many more infections of uncertain origin, in which there is no physical contact with cases. Case investigations involving air travel and congregate settings would be expected to yield secondary cases with greater frequency. Furthermore, we would expect a higher secondary attack rate in households—for MPXV clade I, the secondary attack rate is estimated to be between 0% and 11%.⁵⁶ These figures contrast with those of other viruses (eg, SARS-CoV-2 and respiratory syncytial virus) for which respiratory transmission is the dominant route of spread: for these viruses, household secondary attack rates range from 19% to 70%.⁵⁷ From these observations, we conclude that although respiratory transmission of MPXV is possible, it is not the primary mode of spread and is unlikely to be a substantial contributor to person-to-person transmission of the virus in the current mpox outbreak.

Moving forward, outbreak investigations that include detailed documentation of exposure histories could help to elucidate whether respiratory, percutaneous, or mucosal exposures occurred, and whether or not fomites were involved in transmission. Additional data from contact investigations from air travel exposures and congregate settings would also be informative, including documentation of exposures that did not result in secondary cases.

Experimental approaches with animal models could further address questions about the frequency of different acquisition routes, and even help elucidate the probability of airborne spread versus large respiratory droplet spread by repeating experiments at various distances. Studies in prairie dogs examining respiratory transmission were done only at very short distances (eg, 4 inches [roughly 10 cm]),¹¹ and could be repeated at further distances using impermeable barriers (to exclude close contact while also allowing for airflow). The efficiency of fomites as vehicles for various inoculation routes, including the respiratory pathway, could also be addressed. MPXV clade II should be evaluated for both aerosol and surface stability.⁵⁸

Historically, MPXV has been presumed to be less transmissible than its most famous congener, variola virus (smallpox), which was thought to transmit efficiently between hosts via the respiratory route.4 In recent years, although both clade I and clade II MPXV have proven to be more transmissible than was previously believed,59,60 accumulating data continue to support a minimal role of respiratory transmission. However, this possibility should continue to be thoroughly and repeatedly examined as phenotypic changes due to viral evolution are a constant threat. While DNA viruses such as MPXV mutate with less frequency than RNA viruses (such as coronaviruses and influenza), a high number and frequency of mutations have been noted in the two dominant strains causing the 2022 outbreak. The phenotypic implications of such mutations are so far unknown.^{55,61} Another notable discovery from the current outbreak is the effect of a human enzyme (APOBEC3) that appears to cause mutations in the MPXV genome.⁶¹ Poxviruses were not previously thought to be subject to APOBEC3 editing, but recent findings suggest APOBEC3

	Examples in the scientific literature	Study limitations	
Animal models			
Experiments showing transmission between separately housed animals with shared airflow	Huston et al (2013; clade I only), ¹¹ Heberling et al (1971) ¹²	Animals might be infected following laboratory challenge methods that differ from exposures in natural settings; small numbers of animals	
Respiratory inoculation with monkeypox virus	Hutson et al (2009), ³ Hutson et al (2011), ⁹ Hutson et al (2013), ¹¹ Saijo et al (2009), ⁴⁷ Stittelaar et al (2005), ⁴⁸ Stittelaar et al (2006), ⁴⁹ Estep et al (2011), ⁵⁰ Goff et al (2011) ⁵¹	Animals might be infected with relatively large viral doses (eg, $\sim 1 \times 10^4$ -3·53 $\times 10^7$ plaque- forming units) not likely to be observed in natural settings; respiratory droplet size produced by nebuliser systems in some studies might not mimic natural systems; small numbers of animals	
Identification of virus from saliva or upper respiratory tract, evidence of oral lesions, inflammation of the lungs	Hutson et al (2011), ⁹ Radonic et al (2014), ¹⁰ Dyall et al (2011), ⁵² Hanon and McGavran (1961), ⁵³ Nalca et al (2010), ⁵⁴ Hutson et al (2015) ⁵⁵	Identification of virus on mucosal surfaces does not necessarily indicate that infective respiratory secretions could be expelled; small numbers of animals	
Human case reports			
Identification of replication- competent virus from saliva, upper respiratory tract, or evidence of oral lesions	Angelo et al (2022), ¹⁴ Thornhill et al (2022), ¹⁵ Adler et al (2022) ¹⁹	Identification of virus on mucosal surfaces does not necessarily indicate that infective respiratory secretions could be expelled	
Human outbreaks			
Observed transmission between people with shared airspace, no direct contact, and no mucosal or percutaneous transmission via fomites (eg, adjacent hospital beds or prison cells)	No evidence to date	Settings where this is possible are often confounded by the possibility of transmission via fomites, which itself could result in mucosal, percutaneous, or inhalational inoculation	
Environmental studies			
Aerosolisation of contaminated droplets, detection of monkeypox virus on surfaces unlikely to have been touched (eg, air vents on ceilings)	Gould et al (2022), ⁴² Mellon et al (2022), ⁴³ Verreault et al (2013) ⁴⁴	Viability and infectivity of respiratory particles in uncontrolled settings is probably more variable	

activity has been important in clade II MPXV evolution.⁶¹ The potential emergence of more highly transmissible MPXV variants through this or other mechanisms deserves careful vigilance and will require additional study.

In the absence of more definitive data, public health authorities have recommended broad respiratory transmission reduction strategies, including both source control (containing infectious particles that a person breathes, coughs, or sneezes out) and respiratory protection (filtering out infectious particles in inhaled air), as an adjunct to other protective measures focused on reducing transmission through close contact. Specifically, the UK Health Security Agency and the US Centers for Disease Control and Prevention recommend that people with mpox wear a well-fitting medical mask if close contact with others cannot be avoided, and that contacts of an individual with mpox wear a respirator or well-fitting medical mask when in proximity with infected people for longer than a brief encounter.^{62,63} Furthermore, it is also recommended that health-care personnel wear a respirator with N95 or higher-level filters, in addition to a gown, gloves, and eye protection, when entering the rooms or care areas of patients with suspected or confirmed mpox.^{63,64}

Despite an increase in human-to-human transmission of mpox during the past 40 years, Dr Zdenek Jezek's remarks about a 1988 Democratic Republic of the Congo outbreak could still hold true today: "The absence of illness among neighbours who had no direct face-to-face contact with a [mpox] patient suggests that there is no (or only minimal) risk of airborne transmission."⁴ Only time (and meticulous epidemiological investigation) will tell whether or not this observation prevails.

Contributors

AB, AS, and SAJG contributed to conceptualisation, project administration, writing the original draft, and review and editing of the manuscript. CLH, FW, KMA, FSM, and CM contributed to writing the original draft, and review and editing of the manuscript. KC contributed to data curation, and review and editing of the manuscript. MGR and AMM contributed to conceptualisation, and review and editing of the manuscript.

Declaration of interests

We declare no competing interests.

Acknowledgments

We thank the current and past members of the Centers for Disease Control and Prevention (CDC) Poxvirus and Rabies Branch and the CDC Multi-national Mpox Response. The findings and conclusions in this report are those of the authors and do not necessarily represent the official position of the CDC.

References

- 1 Vivancos R, Anderson C, Blomquist P, et al. Community transmission of monkeypox in the United Kingdom, April to May 2022. *Euro Surveill* 2022; **27**: 2200422.
- 2 Rimoin AW, Mulembakani PM, Johnston SC, et al. Major increase in human monkeypox incidence 30 years after smallpox vaccination campaigns cease in the Democratic Republic of Congo. *Proc Natl Acad Sci USA* 2010; 107: 16262–67.
- 3 Hutson CL, Olson VA, Carroll DS, et al. A prairie dog animal model of systemic orthopoxvirus disease using West African and Congo Basin strains of monkeypox virus. J Gen Virol 2009; 90: 323–33.
- 4 Jezek Z, Grab B, Szczeniowski MV, Paluku KM, Mutombo M. Human monkeypox: secondary attack rates. Bull World Health Organ 1988; 66: 465–70.
- 5 Learned LA, Reynolds MG, Wassa DW, et al. Extended interhuman transmission of monkeypox in a hospital community in the Republic of the Congo, 2003. *Am J Trop Med Hyg* 2005; 73: 428–34.
- 6 Cudmore S, Cossart P, Griffiths G, et al. Actin-based motility of vaccinia virus. *Nature* 1995; 378: 636–38.
- 7 Likos AM, Sammons SA, Olson VA, et al. A tale of two clades: monkeypox viruses. J Gen Virol 2005; 86: 2661–72.
- 8 Chen N, Li G, Liszewski MK, et al. Virulence differences between monkeypox virus isolates from West Africa and the Congo basin. *Virology* 2005; 340: 46–63.
- 9 Hutson CL, Carroll DS, Gallardo-Romero N, et al. Monkeypox disease transmission in an experimental setting: prairie dog animal model. PLoS One 2011; 6: e28295.
- 10 Radonic A, Metzger S, Dabrowski PW, et al. Fatal monkeypox in wild-living sooty mangabey, Côte d'Ivoire, 2012. *Emerg Infect Dis* 2014; 20: 1009–11.
- 11 Hutson CL, Gallardo-Romero N, Carroll DS, et al. Transmissibility of the monkeypox virus clades via respiratory transmission: investigation using the prairie dog-monkeypox virus challenge system. *PLoS One* 2013; 8: e55488.

- 12 Heberling RL, Kalter SS. Induction, course, and transmissibility of monkeypox in the baboon (*Papio cynocephalus*). J Infect Dis 1971; 124: 33–38.
- 13 McCollum AM, Damon IK. Human monkeypox. Clin Infect Dis 2014; 58: 260–67.
- 14 Angelo KM, Smith T, Camprubi-Ferrer D, et al. Epidemiological and clinical characteristics of patients with monkeypox in the GeoSentinel Network: a cross-sectional study. *Lancet Infect Dis* 2022; 23: 196–206.
- 15 Thornhill JP, Barkati S, Walmsley S, et al. Monkeypox virus infection in humans across 16 countries - April-June 2022. N Engl J Med 2022; 387: 679–91.
- 16 Patel A, Bilinska J, Tam JCH, et al. Clinical features and novel presentations of human monkeypox in a central London centre during the 2022 outbreak: descriptive case series. *BMJ* 2022; 378: e072410.
- 17 Tarin-Vicente EJ, Alemany A, Agud-Dios M, et al. Clinical presentation and virological assessment of confirmed human monkeypox virus cases in Spain: a prospective observational cohort study. *Lancet* 2022; 400: 661–69.
- 18 Benites-Zapata VA, Ulloque-Badaracco JR, Alarcon-Braga EA, et al. Clinical features, hospitalisation and deaths associated with monkeypox: a systematic review and meta-analysis. Ann Clin Microbiol Antimicrob 2022; 21: 36.
- 19 Adler H, Gould S, Hine P, et al. Clinical features and management of human monkeypox: a retrospective observational study in the UK. *Lancet Infect Dis* 2022; 22: 1153–62.
- 20 Paran N, Yahalom-Ronen Y, Shifman O, et al. Monkeypox DNA levels correlate with virus infectivity in clinical samples, Israel, 2022. Euro Surveill 2022; 27: 2200636.
- 21 Breman JG, Kalisa R, Steniowski MV, Zanotto E, Gromyko AI, Arita I. Human monkeypox, 1970-79. Bull World Health Organ 1980; 58: 165–82.
- 22 Doshi RH, Guagliardo SAJ, Doty JB, et al. Epidemiologic and ecologic investigations of monkeypox, Likouala Department, Republic of the Congo, 2017. Emerg Infect Dis 2019; 25: 281–89.
- 23 Hutin YJ, Williams RJ, Malfait P, et al. Outbreak of human monkeypox, Democratic Republic of Congo, 1996 to 1997. *Emerg Infect Dis* 2001; 7: 434–38.
- 24 Jezek Z, Arita I, Mutombo M, Dunn C, Nakano JH, Szczeniowski M. Four generations of probable person-to-person transmission of human monkeypox. *Am J Epidemiol* 1986; 123: 1004–12.
- 25 Mukinda VB, Mwema G, Kilundu M, Heymann DL, Khan AS, Esposito JJ. Re-emergence of human monkeypox in Zaire in 1996. Monkeypox Epidemiologic Working Group. *Lancet* 1997; 349: 1449–50.
- 26 Centers for Disease Control and Prevention. Update: multistate outbreak of monkeypox–Illinois, Indiana, Kansas, Missouri, Ohio, and Wisconsin, 2003. MMWR Morb Mortal Wklγ Rep 2003; 52: 642–46.
- 27 Sejvar JJ, Chowdary Y, Schomogyi M, et al. Human monkeypox infection: a family cluster in the midwestern United States. J Infect Dis 2004; 190: 1833–40.
- 28 Reynolds MG, Davidson WB, Curns AT, et al. Spectrum of infection and risk factors for human monkeypox, United States, 2003. *Emerg Infect Dis* 2007; 13: 1332–39.
- 29 Fleischauer AT, Kile JC, Davidson M, et al. Evaluation of human-tohuman transmission of monkeypox from infected patients to health care workers. *Clin Infect Dis* 2005; 40: 689–94.
- 30 Ogoina D, Izibewule JH, Ogunleye A, et al. The 2017 human monkeypox outbreak in Nigeria–report of outbreak experience and response in the Niger Delta University Teaching Hospital, Bayelsa State, Nigeria. *PLoS One* 2019; 14: e0214229.
- 31 Yinka-Ogunleye A, Aruna O, Dalhat M, et al. Outbreak of human monkeypox in Nigeria in 2017–18: a clinical and epidemiological report. *Lancet Infect Dis* 2019; **19**: 872–79.
- 32 Vaughan A, Aarons E, Astbury J, et al. Two cases of monkeypox imported to the United Kingdom, September 2018. *Euro Surveill* 2018; 23: 1800509.
- 33 Hobson G, Adamson J, Adler H, et al. Family cluster of three cases of monkeypox imported from Nigeria to the United Kingdom, May 2021. Euro Surveill 2021; 26: 2100745.
- 34 Ng OT, Lee V, Marimuthu K, et al. A case of imported monkeypox in Singapore. *Lancet Infect Dis* 2019; 19: 1166.

- 35 Erez N, Achdout H, Milrot E, et al. Diagnosis of imported monkeypox, Israel, 2018. *Emerg Infect Dis* 2019; **25**: 980–83.
- 36 Minhaj FS, Rao AK, McCollum AM. Imported monkeypox from international traveler, Maryland, USA, 2021. *Emerg Infect Dis* 2022; 28: 1738.
- 37 Rao AK, Schulte J, Chen TH, et al. Monkeypox in a traveler returning from Nigeria – Dallas, Texas, July 2021. MMWR Morb Mortal Wkly Rep 2022; 71: 509–16.
- 38 Petersen BW, Kabamba J, McCollum AM, et al. Vaccinating against monkeypox in the Democratic Republic of the Congo. Antiviral Res 2019; 162: 171–77.
- 39 Marshall KE, Barton M, Nichols J, et al. Health care personnel exposures to subsequently laboratory-confirmed monkeypox patients – Colorado, 2022. MMWR Morb Mortal Wkly Rep 2022; 71: 1216–19.
- 40 Hagan LM, Beeson A, Hughes S, et al. Monkeypox case investigation – Cook County Jail, Chicago, Illinois, July–August 2022. MMWR Morb Mortal Wkly Rep 2022; 71: 1271–77.
- 41 Hennessee I, Shelus V, McArdle CE, et al. Epidemiologic and clinical features of children and adolescents aged <18 years with monkeypox — United States, May 17–September 24, 2022. MMWR Morb Mortal Wkly Rep 2022; 71: 1407–11.
- 42 Gould S, Atkinson B, Onianwa O, et al. Air and surface sampling for monkeypox virus in a UK hospital: an observational study. *Lancet Microbe* 2022; **3:** e904–11.
- 43 Mellon G, Rubenstein E, Antoine M, et al. Air detection of monkeypox virus in a dedicated outpatient clinic room for monkeypox infection diagnosis. J Infect 2022; published online Dec 28. https://doi.org/10.1016/j.jinf.2022.12.025.
- 44 Verreault D, Killeen SZ, Redmann RK, Roy CJ. Susceptibility of monkeypox virus aerosol suspensions in a rotating chamber. *J Virol Methods* 2013; **187**: 333–37.
- 45 Morgan CN, Whitehill F, Doty JB, et al. Environmental persistence of monkeypox virus on surfaces in household of person with travelassociated infection, Dallas, Texas, USA, 2021. *Emerg Infect Dis* 2022; 28: 1982–89.
- 46 Vaughan A, Aarons E, Astbury J, et al. Human-to-human transmission of monkeypox virus, United Kingdom, October 2018. *Emerg Infect Dis* 2020; 26: 782–85.
- 47 Saijo M, Ami Y, Suzaki Y, et al. Virulence and pathophysiology of the Congo basin and West African strains of monkeypox virus in non-human primates. J Gen Virol 2009; 90: 2266–71.
- 48 Stittelaar KJ, Neyts J, Naesens L, et al. Antiviral treatment is more effective than smallpox vaccination upon lethal monkeypox virus infection. *Nature* 2006; **439**: 745–48.
- 49 Stittelaar KJ, van Amerongen G, Kondova I, et al. Modified vaccinia virus Ankara protects macaques against respiratory challenge with monkeypox virus. J Virol 2005; 79: 7845–51.
- 50 Estep RD, Messaoudi I, O'Connor MA, et al. Deletion of the monkeypox virus inhibitor of complement enzymes locus impacts the adaptive immune response to monkeypox virus in a nonhuman primate model of infection. J Virol 2011; 85: 9527–42.

- 51 Goff AJ, Chapman J, Foster C, et al. A novel respiratory model of infection with monkeypox virus in cynomolgus macaques. *J Virol* 2011; 85: 4898–909.
- 52 Dyall J, Johnson RF, Chen DY, et al. Evaluation of monkeypox disease progression by molecular imaging. J Infect Dis 2011; 204: 1902–11.
- 53 Hahon N, McGavran MH. Air-borne infectivity of the variolavaccinia group of poxviruses for the cynomolgus monkey, *Macaca irus. J Infect Dis* 1961; 109: 294–98.
- 54 Nalca A, Livingston VA, Garza NL, et al. Experimental infection of cynomolgus macaques (*Macaca fascicularis*) with aerosolized monkeypox virus. *PLoS One* 2010; 5: e12880.
- 55 Isidro J, Borges V, Pinto M, et al. Phylogenomic characterization and signs of microevolution in the 2022 multi-country outbreak of monkeypox virus. *Nat Med* 2022; 28: 1569–72.
- 56 Beer EM, Rao VB. A systematic review of the epidemiology of human monkeypox outbreaks and implications for outbreak strategy. *PLoS Negl Trop Dis* 2019; 13: e0007791.
- 57 Leung NHL. Transmissibility and transmission of respiratory viruses. *Nat Rev Microbiol* 2021; **19**: 528–45.
- 58 van Doremalen N, Bushmaker T, Morris DH, et al. Aerosol and surface stability of SARS-CoV-2 as compared with SARS-CoV-1. N Engl J Med 2020; 382: 1564–67.
- 59 Charniga K, McCollum AM, Hughes CM, et al. Updating reproduction number estimates for monkeypox in the Democratic Republic of Congo using surveillance data. American Society of Tropical Medicine and Hygiene; Oct 30–Nov 3, 2022 (abstr 1810).
- 60 Centers for Disease Control and Prevention. Technical report 3: multi-national mpox outbreak, United States, 2022. Jan 6, 2023. https://www.cdc.gov/poxvirus/monkeypox/cases-data/technicalreport/report-3.html (accessed Jan 18, 2023).
- 61 Gigante CM, Korber B, Seabolt MH, et al. Multiple lineages of monkeypox virus detected in the United States, 2021–2022. *Science* 2022; 378: 560–65.
- 62 Centers for Disease Control and Prevention. Hand hygiene, source control, and personal protective equipment. Aug 11, 2022. https:// www.cdc.gov/poxvirus/monkeypox/clinicians/infection-controlhome.html#hand-hygiene (accessed Oct 12, 2022).
- 63 UK Health Security Agency. Principles for monkeypox control in the UK: 4 nations consensus statement. Nov 3, 2022. https://www. gov.uk/government/publications/principles-for-monkeypox-controlin-the-uk-4-nations-consensus-statement/principles-for-monkeypoxcontrol-in-the-uk-4-nations-consensus-statement (accessed Dec 2, 2022).
- 64 Centers for Disease Control and Prevention. Infection prevention and control of monkeypox in healthcare settings. Oct 31, 2022. https://www.cdc.gov/poxvirus/monkeypox/clinicians/infectioncontrol-healthcare.html (accessed Nov 1, 2022).

Copyright © 2023 The Author(s). Published by Elsevier Ltd. This is an Open Access article under the CC BY-NC-ND 4.0 license.