### RESEARCH LETTERS

consistent with observations reported in similarly designed studies in Europe (7,8). Occult infection with oropharyngeal and rectal viral shedding might have contributed to the scale of the 2022 mpox outbreak, which spread through sexual networks. All but 1 patient with known MPXV infection in our cohort had detectable viral DNA in oropharyngeal or rectal swab samples, suggesting that the new infections we detected are likely true positives. Furthermore, all newly identified mpox patients in our study had >1 sample for which both qPCR targets were detected.

Current MPXV tests cleared for emergency use are indicated only for use on lesion samples (9). Further studies are needed to characterize viral shedding dynamics, particularly related to symptom onset and duration of infectivity. As data demonstrating mucosal viral shedding in mpox emerge, expanding testing to allow broader sample collection and expedite diagnostic validation of samples from various anatomic sites will be crucial.

In conclusion, during the ongoing mpox outbreak, clinicians should consider oropharyngeal and rectal MPXV qPCR testing for at-risk patients with pharyngitis or proctitis. In addition, asymptomatic screening in high-risk populations might be warranted if community prevalence is high or rising.

# **About the Author**

Dr. Contag is an infectious diseases fellow at Stanford University, Stanford, California, USA. Her primary research interests are emerging and reemerging infectious diseases and the management of critical illness in low-resource settings.

# References

- Thornhill JP, Barkati S, Walmsley S, Rockstroh J, Antinori A, Harrison LB, et al.; SHARE-net Clinical Group. Monkeypox virus infection in humans across 16 countries— April-June 2022. N Engl J Med. 2022;387:679-91. https://doi.org/10.1056/NEJMoa2207323
- Abbasi J. Reports of asymptomatic monkeypox suggest that, at the very least, some infections go unnoticed. JAMA. 2022;328:1023–5. https://doi.org/10.1001/jama.2022.15426
- Jezek Z, Marennikova SS, Mutumbo M, Nakano JH, Paluku KM, Szczeniowski M. Human monkeypox: a study of 2,510 contacts of 214 patients. J Infect Dis. 1986;154:551–5. https://doi.org/10.1093/infdis/154.4.551
- Li Y, Olson VA, Laue T, Laker MT, Damon IK. Detection of monkeypox virus with real-time PCR assays. J Clin Virol. 2006;36:194–203. https://doi.org/10.1016/j.jcv.2006.03.012
- Li Y, Zhao H, Wilkins K, Hughes C, Damon IK. Real-time PCR assays for the specific detection of monkeypox virus West African and Congo Basin strain DNA. J Virol Methods. 2010;169:223–7. https://doi.org/10.1016/ j.jviromet.2010.07.012

- Karan A, Styczynski AR, Huang C, Sahoo MK, Srinivasan K, Pinsky BA, et al. Human Monkeypox without viral prodrome or sexual exposure, California, USA, 2022. Emerg Infect Dis. 2022;28:2121–3. https://doi.org/10.3201/ eid2810.221191
- 7. Ferré VM, Bachelard A, Zaidi M, Armand-Lefevre L, Descamps D, Charpentier C, et al. Detection of monkeypox virus in anorectal swabs from asymptomatic men who have sex with men in a sexually transmitted infection screening program in Paris, France. Ann Intern Med. 2022;175:1491–2. https://doi.org/10.7326/M22-2183
- 8. De Baetselier I, Van Dijck C, Kenyon C, Coppens J, Michiels J, de Block T, et al.; ITM Monkeypox study group. Retrospective detection of asymptomatic monkeypox virus infections among male sexual health clinic attendees in Belgium. Nat Med. 2022;28:2288–92. https://doi.org/10.1038/s41591-022-02004-w
- US Food and Drug Administration. FDA monkeypox response: FDA's role in mpox preparedness and response, and information about mpox (formerly referred to as monkeypox) [cited 2023 Jan 7]. https://www.fda.gov/ emergency-preparedness-and-response/mcm-issues/ fda-monkeypox-response

Address for correspondence: Caitlin Contag, Stanford University, Department of Medicine, Division of Infectious Diseases and Geographic Medicine, 300 Pasteur Dr, L-134, Stanford, CA 94305, USA; email: cacontag@stanford.edu

# Human Metapneumovirus Infections during COVID-19 Pandemic, Spain

Maria L. García-García, Elena Pérez-Arenas, Pedro Pérez-Hernandez, Iker Falces-Romero, Sara Ruiz, Francisco Pozo, Inmaculada Casas, Cristina Calvo

Author affiliations: Severo Ochoa University Hospital Leganés, CIBERINFEC ISCIII, Madrid, Spain (M.L. García-García, S. Ruiz); La Paz University Hospital, Madrid, Spain (E. Pérez-Arenas, P. Pérez-Hernandez); La Paz University Hospital, IdiPaz Foundation, CIBERINFEC ISCIII, Madrid (I. Falces-Romero, C. Calvo); Severo Ochoa University Hospital Leganés, Madrid (S. Ruiz); National Center Microbiology, CIBEREST, Madrid (F. Pozo, I. Casas)

DOI: https://doi.org/10.3201/eid2904.230046

We describe an unusual outbreak of respiratory infections caused by human metapneumovirus in children during the sixth wave of COVID-19 in Spain, associated with the Omicron variant. Patients in this outbreak were older than usual and showed more hypoxia and pneumonia, longer length of stay, and greater need for intensive care.

Respiratory infections caused by human metapneumovirus (hMPV) are typically epidemic during February–April. hMPV infections cause bronchiolitis and recurrent wheezing episodes and are responsible for hospitalizations mainly in children <2 years of age (1). The COVID-19 pandemic has changed the epidemiology of respiratory viral infections, modifying the classic seasonality of respiratory syncytial virus (RSV), influenza, and other viruses (2–4). In Spain, an outbreak of pediatric hMPV infections was identified in November–December 2021, coinciding with the sixth wave of COVID-19 in the country. This study aimed to describe this outbreak and to compare it with previous hMPV outbreaks, before the COVID-19 pandemic.

We retrospectively collected data from respiratory hMPV infections requiring hospitalization in children <18 years of age during October-December 2021 in La Paz University Hospital (Madrid, Spain) and analyzed clinical characteristics. We performed multiple PCR respiratory panels as standard clinical practice in children in whom infection

by RSV, influenza, and SARS-CoV-2 had previously been ruled out by rapid test or PCR.

We obtained data from a systematic prospective study conducted over 15 years in all hospitalized children with respiratory infections in Severo Ochoa University Hospital (Madrid) using multiple PCR panels performed at the National Center for Microbiology. This study was approved by the hospital ethics committee. We compared clinical data from hospitalized patients in 2021 with our historical series of hMPV infections collected during 2005–2020, before the CO-VID-19 pandemic.

We analyzed data from 48 patients with hMPV infection during October-December 2021; of those, 56.3% were male and 54% were >2 years of age, and median age was 22.7 (interquartile range [IQR] 7.1-34.9) months. Twenty-nine (60%) of the cases were detected in December and 1 in October. In 19 cases (39%), we detected co-infection with other viruses, mainly adenoviruses and human coronaviruses. A total of 34 (70.8%) case-patients had fever >38°C; in addition, 41 (85.4%) had hypoxia, and 34 (70.8%) had an infiltrate visible in their chest radiograph; radiography was not performed in 11 cases. The most frequent diagnoses were pneumonia (35%), bronchiolitis (27%), and episodes of wheezing (16%). Antimicrobial drugs were prescribed in 26 cases (54%). Seven patients (14%) required pediatric intensive care unit (PICU) admission; 2 needed mechanical ventilation. The median duration of fever was 3 (IQR 2.7-5)

hospitalization during the COVID-19 pandemic (20 Clinical data	2021, n = 48	2005–2020, n = 498	OR (95% CI)	p value
Age, mo	22 +15.9	13 +17.2	NA	0.002
Sex, no. (%)				
M	27 (56.2)	289 (58.0)	NA	0.875
F	21 (43.8)	209 (42.0)	NA	
Temperature >37.9°C	34 (70.8)	342 (68.7)	NA	0.734
Highest temperature	39.0 <u>+</u> 0.5	38.7 <u>+</u> 0.6		0.008
Hypoxia, oxygen saturation <93%, no. (%)	41 (85.4)	331 ( <del>6</del> 6.5)	3.9 (1.4-9.0)	0.002
Chest infiltrate, no. (%)	34 (70.8)	184 (36.9)	10.2 (3.1-32.9)	<0.001
Antimicrobial treatment, no. (%)	26 (54.2)	123 (24.7)	3.2 (1.8-5.6)	<0.001
Viral co-infection, no. (%)	19 (39.6)	208 (41.8)	NA	0.632
Time of illness, Nov-Dec, no. (%)	47 (97.9)	9 (1.8)	NA	<0.001
Diagnosis, no. (%)				< 0.001
Wheezing episode	8 (16.7)	288 (57.8)	NA	NA
Bronchiolitis	13 (27.1)	144 (28.9)	NA	NA
Pneumonia	17 (35.4)	44 (8.8)	NA	NA
Blood tests				
Leucocytes, cells/mm <sup>3</sup>	11,140 <u>+</u> 5,530	12,112 <u>+</u> 4,620	NA	0.589
C-reactive protein	40 <u>+</u> 44	37 <u>+</u> 54	NA	0.698
Outcome				
Duration of stay, d	6.8 <u>+</u> 4.8	4.6 <u>+</u> 7.1	NA	0.004
Duration of fever, d	3.9 <del>+</del> 2.4	2.7 <u>+</u> 1.8	NA	0.001
Duration of hypoxia, d	6.2 <del>+</del> 4.2	2.9 <u>+</u> 2.1	NA	<0.001
PICU admission, no. (%)	7 (14.6)	13 (2.6)	5.1 (1.8-8.0)	0.004

<sup>\*</sup>Values are mean (SD) except as indicated. Bold text indicates statistically significant differences. NA, not applicable; OR, odds ratio; PICU, pediatric intensive care unit.

days, hypoxia 5 (IQR 3.5–7.5) days, and admission, 6 (IQR 4–8) days.

We compared the data from those recent patients with data from 498 cases of hMPV infection at the hospital during 2005–2020. During that period, 9 (1.8%) cases were detected in November–December and 453 (90.9%) during February–May. Patients in the 2021 season were older than those from the previous 15-year period and had significantly higher rates of hypoxia, pneumonia, antimicrobial drug treatment; they also had longer durations of fever, hypoxia, hospital admission, and PICU admission (Table 1).

In November–December 2021, during the sixth wave of COVID-19 in Spain, the country experienced an extemporaneous hMPV outbreak at the time when RSV epidemic was usually observed. This outbreak of hMPV infections affected children older than were usually affected in previous years; we also observed a more severe clinical course and higher rates of hypoxia, pneumonia, and admission to PICU than historically.

We considered that there may have been competition between respiratory viruses that could justify the delay in the RSV outbreak (5), which occurred in summer (June-July) 2021 in Spain; such competition was not the case for the hMPV outbreak, which coincided with spread of the Omicron variant of SARS-CoV-2. One possible explanation is relaxation of social distancing measures or the extreme contagiousness of Omicron. The increased severity of illness could be partly explained by the absence of hMPV infections in the previous 2 years, resulting in a susceptible population of older children who had not had previous hMPV infections and therefore had no immunity. In previous seasons, children >1 year of age were immunized by previous infections or even through residual maternal protection; this protection did not exist in 2021, and older children were infected. In conclusion, this outbreak illustrates that clinicians should be aware of potential differences in the epidemiology of other viral respiratory infections during and after the COVID-19 pandemic.

This study was partially funded by FIS (Fondo de Investigaciones Sanitarias – Spanish Health Research Fund), grant nos. PI06/0532, PI09/0246, PI12/0129, PI-18CIII/00009, PI21CIII/00019, and PI21/00377.

# **About the Author**

Dr. García-García is the head of the pediatrics department at Severo Ochoa Hospital, Leganés, Spain. She is an expert in respiratory viral infections and pediatric pneumology.

# References

- García-García ML, Calvo C, Rey C, Díaz B, Molinero MD, Pozo F, et al. Human metapnuemovirus infections in hospitalized children and comparison with other respiratory viruses. 2005–2014 prospective study. PLoS One. 2017;12:e0173504. https://doi.org/10.1371/ journal.pone.0173504
- Yeoh DK, Foley DA, Minney-Smith CA, Martin AC, Mace AO, Sikazwe CT, et al. Impact of coronavirus disease 2019 public health measures on detections of influenza and respiratory syncytial virus in children during the 2020 Australian winter. Clin Infect Dis. 2021;72:2199–202. https://doi.org/10.1093/cid/ciaa1475
- Flores-Pérez P, Gerig N, Cabrera-López MI, de Unzueta-Roch JL, Del Rosal T, Calvo C; COVID-19 Study Group in Children. Acute bronchiolitis during the COVID-19 pandemic. Enferm Infecc Microbiol Clin (Engl Ed). 2022;40:572–5. https://doi.org/10.1016/j.eimc.2021.06.012
- Torres-Fernandez D, Casellas A, Mellado MJ, Calvo C, Bassat Q. Acute bronchiolitis and respiratory syncytial virus seasonal transmission during the COVID-19 pandemic in Spain: a national perspective from the pediatric Spanish Society (AEP). J Clin Virol. 2021;145:105027. https://doi.org/ 10.1016/j.jcv.2021.105027
- 5 Dee K, Goldfarb DM, Haney J, Amat JAR, Herder V, Stewart M, et al. Human rhinovirus infection blocks severe acute respiratory syndrome coronavirus 2 replication within the respiratory epithelium: implications for COVID-19 epidemiology. J Infect Dis. 2021;224:31–8. https://doi.org/ 10.1093/infdis/jiab147

Address for correspondence: Cristina Calvo, Pediatrics and Infectious Diseases Dept, Hospital Universitario La Paz, P° Castellana, 261, 28046 Madrid, Spain; email: ccalvorey@gmail.com

# Highly Pathogenic Avian Influenza A(H5N1) Virus in a Harbor Porpoise, Sweden

Elina Thorsson, Siamak Zohari, Anna Roos, Fereshteh Banihashem, Caroline Bröjer, Aleksija Neimanis

Author affiliations: National Veterinary Institute (SVA), Uppsala, Sweden (E. Thorsson, S. Zohari, F. Banihashem, C. Bröjer, A. Neimanis); Swedish Museum of Natural History, Stockholm, Sweden (A. Roos)

DOI: https://doi.org/10.3201/eid2904.221426