

# SC/69A/E/16

**Sub-committees/working group name: E**

**Emerging Pathogens in the Context of Marine Mammal Health SARS-COV- 2 and HPAI H5N1. Progress report IWC 2023 IWC SC Intersessional Correspondence Group on CDOC (Cetacean Diseases of Concern)**

**R. Stimmelmayer, T. Rowles, F. Gulland, R. Deaville, H.S. Ip, N. Davison, S. Mazzariol, A.J. Fernández Rodríguez, E.M. Sierra Pulpillo, G. Hernandez, M. Grigg, K. Colegrove, K. Groch, M. Uhart, M.A. Delaney, P. Holm, Y. Tajima**



INTERNATIONAL  
WHALING COMMISSION

Papers submitted to the IWC are produced to advance discussions within that meeting; they may be preliminary or exploratory.

It is important that if you wish to cite this paper outside the context of an IWC meeting, you notify the author at least six weeks before it is cited to ensure that it has not been superseded or found to contain errors.

**WORKING PAPER**  
**Emerging Pathogens in the Context of Marine Mammal Health**  
**SARS-COV- 2 and HPAI H5N1**

**Progress report IWC 2023 IWC SC**  
**Intersessional Correspondence Group on CDOC (Cetacean Diseases of Concern)**

Members: Raphaela Stimmelmayer, Teri Rowles, Frances Gulland, Rob Deaville, Hon S. Ip, Nick Davison, Sandro Mazzariol, Antonio Jesús Fernández Rodríguez, Eva María Sierra Pulpillo, Gaby Hernandez, Michael Grigg, Katie Colegrove, Kátia Groch, Marcela Uhart Martha A. Delaney, Patricia Holm, Yuko Tajima.

Paper prepared : Raphaela Stimmelmayer and Teri Rowles

## ABSTRACT

Cetacean diseases of concern (CDOC) are a standing topic for E. Special disease primer sessions during 2020/2021 focused on known marine mammal pathogens such as *morbillivirus (CeMV)*, *Herpes virus*, *Brucella ceti*, and *Toxoplasma gondii* that have been associated with morbidity and mortality in individual animals, but also have been implicated in UMEs. Recent emerging pathogens of concern include SARS-Cov2 and avian influenza. A primer session on SARS-Cov2 and implications from spillover into marine mammals was held during 2022. In response to the increasing evidence for Highly Pathogenic Avian Influenza (HPAI) spillover (natural infection) events in marine mammals in recent months the World Organization for Animal Health (WOAH) convened in March 2023 an ad hoc expert group to develop a Guidance Framework for HPAI outbreak management in marine mammals. Several members of the IWC CDOC intersessional working group are included in the WOAH working group as well as other experts at large. A final draft of the guidance document is expected by the end of May 2023.

**KEYWORDS:** CETACEAN DISEASES OF CONCERN; SARS-COV-2; HPAI H5N1 clade 2.3.4.4b; WOAH;

## INTRODUCTION

### SARS-COV-2 Update

Spill over (reverse zoonosis) of severe acute Respiratory Syndrome Coronavirus-2 (SARS-CoV-2) to various animal species have occurred during the height of the pandemic as well as now. As of 2023, 699 natural infections (spillovers) have been confirmed in the Americas, Africa, Asia, and Europe into twenty-six mammals. The majority of spillovers have occurred in terrestrial mammals, with a few semi aquatic and aquatic mammals (mink, eurasian otter, fishing cat, hippopotamus, West indian manatee). The majority of cases occurred in animals under human care including domestic pets, but natural infections in commercial fur industry (e.g. mink) and free-ranging wildlife species with onward animal to animal transmission (e.g. white-tailed deer), have also occurred. On 7 March 2022, the Tripartite Organization FAO/WHO/WOAH issued a joint statement on the prioritization of monitoring SARS-CoV-2 infection in wildlife and preventing the formation of animal reservoirs. Given the limited case material for aquatic and semi-aquatic species, health implications of Sars Cov2 with respect to morbidity and mortality in marine mammals remain difficult to judge, but both are part of the potential disease spectrum. That said, given the cumulative evidence for reverse zoonosis and our incomplete understanding of virus host range (see Audino et al. 2021; Li et al. 2022; Lean et al. 2022; 2023) surveillance of marine mammals under human care remains critical while virus circulation in humans remains ongoing. Furthermore, surveillance of stranded marine mammals and live animals during health assessments, especially coastal species, will provide a starting point to investigate epidemiological aspects of this novel viral agent within a marine environment. A recent stranding event of dolphins in New Zealand highlights marine mammal exposure pathways through waste water (Stocken pers. commun). Testing of wastewater has been used in evaluation of community cases in people in areas with lower resources to provide individual case evaluations. Briefly, buccal swabs of 3 dead animals tested positive for SARS-Cov2. Subsequent tests of blowhole swabs were not positive. Environmental cross contamination from waste water that had spilled into the coastal waters is considered the most parsimonious explanation. Buccal cavities of dolphins were likely exposed to the contaminated water while feeding. Though no spillover into dolphins occurred (animals were not the source of the SARS-Cov2 virus), it should be noted that false positives may be identified in stranded cetaceans due to environmental contamination, therefore additional pathology or internal testing should be done. In conclusion, though still considered a human disease, its potential for establishment of new terrestrial and aquatic animal host reservoirs, for viral adaptation etc. and for future spill backs clearly identifies it as a ONE-Health issue. (Banerjee et al. 2021; Clayton et al. 2022).

## LITERATURE CITED AND FURTHER READING

Audino T, Grattarola C, Centelleghes C, Peletto S, Giorda F, Florio CL, Caramelli M, Bozzetta E, Mazzariol S, Di Guardo G, Lauriano G, Casalone C. SARS-CoV-2, a Threat to Marine Mammals? A Study from Italian Seawaters. *Animals (Basel)*. 2021 Jun 3;11(6):1663.

Bae, J., Ro, C., Kang, Y., Ga, E., Na, W., & Song, D. (2023). Human-to-Animal Transmission of SARS-CoV-2, South Korea, 2021. *Emerging Infectious Diseases*, 29(5), 1066-1067. <https://doi.org/10.3201/eid2905.221359>.

Banerjee A, Mossman K, Baker ML. Zoonothroponotic potential of SARS-CoV-2 and implications of reintroduction into human populations. *Cell Host Microbe*. 2021 Feb 10;29(2):160-164.

Brnić D, Lojkić I, Škoko I, Krešić N, Šimić I, Keros T, Ganjto M, Štefanac D, Viduka B, Karšaj D, Štiller D, Habrun B, Jemeršić L. SARS-CoV-2 circulation in Croatian wastewaters and the absence of SARS-CoV-2 in bivalve molluscan shellfish. *Environ Res*. 2022 May 1;207:112638. Carmona, G., Burgos, T., Barrientos, R. et al. Lack of SARS-CoV-2 RNA evidence in the lungs from wild European polecats (*Mustela putorius*) from Spain. *Eur J Wildl Res* 69, 33 (2023).

Bui, V., Dao, T., Tran, L., Vu, T., Nguyen, T., Nguyen, G...Lee, H. (2023). SARS-CoV-2 Infection in a Hippopotamus, Hanoi, Vietnam. *Emerging Infectious Diseases*, 29(3), 658-661. <https://doi.org/10.3201/eid2903.220915>.

Chandler JC, Bevins SN, Ellis JW, Linder TJ, Tell RM, Jenkins-Moore M, Shriner SA (2021) SARS-CoV-2 exposure in wild white-tailed deer (*Odocoileus virginianus*). *Proc Natl Acad Sci* 118. <https://doi.org/10.1073/pnas.2114828118>

Clayton E, Ackerley J, Aelmans M, Ali N, Ashcroft Z, Ashton C, Barker R, Budryte V, Burrows C, Cai S, Callaghan A, Carberry J, Chatwin R, Davies I, Farlow C, Gamblin S, Iacobut A, Lambe A, Lynch F, Mihalache D, Mokbel A, Potamsetty S, Qadir Z, Soden J, Sun X, Vasile A, Wheeler O, Rohaim MA, Munir M. Structural Bases of Zoonotic and Zoonothroponotic Transmission of SARS-CoV-2. *Viruses*. 2022 Feb 17;14(2):418.

Gortázar C, Barroso-Arévalo S, Ferreras-Colino E, Isla J, Fuente G, Rivera B, Domínguez Rodríguez L, de la Fuente J, Sánchez-Vizcaíno J (2021) Natural SARS-CoV-2 infection in kept ferrets. *Spain Emerg Infect Dis* 27:1994–1996.

Giner J, Villanueva-Saz S, Tobajas AP, Pérez MD, González A, Verde M, Yzuel A, García-García A, Taleb V, Lira-Navarrete E, Hurtado-Guerrero R, Pardo J, Santiago L, Paño JR, Ruíz H, Lacasta D, Fernández A (2021) SARS-CoV-2 seroprevalence in household domestic ferrets (*Mustela putorius furo*). *Animals* 11:667.

Hofer, U. Dose-dependent COVID-19 symptoms. *Nat Rev Microbiol* 19, 682 (2021).  
<https://doi.org/10.1038/s41579-021-00634-4>

Jay C, Ratcliff J, Turtle L, Goulder P, Klenerman P. Exposed seronegative: Cellular immune responses to SARS-CoV-2 in the absence of seroconversion. *Front Immunol.* 2023 Jan 26;14:1092910. doi: 10.3389/fimmu.2023.1092910.

Le Guernic A, Palos Ladeiro M, Boudaud N, Do Nascimento J, Gantzer C, Inglard JC, Mouchel JM, Pochet C, Moulin L, Rocher V, Waldman P, Wurtzer S, Geffard A. First evidence of SARS-CoV-2 genome detection in zebra mussel (*Dreissena polymorpha*). *J Environ Manage.* 2022 Jan 1;301:113866.

Lean FZX, Cox R, Madslie K, Spiro S, Nymo IH, Brøjer C, Neimanis A, Lawson B, Holmes P, Man C, Folkow LP, Gough J, Ackroyd S, Evans L, Wrigglesworth E, Grimholt U, McElhinney L, Brookes SM, Delahay RJ, Núñez A. Tissue distribution of angiotensin-converting enzyme 2 (ACE2) receptor in wild animals with a focus on artiodactyls, mustelids and phocids. *One Health.* 2023 Jun;16:100492.

Lean FZX, Núñez A, Spiro S, Priestnall SL, Vreman S, Bailey D, James J, Wrigglesworth E, Suarez-Bonnet A, Conceicao C, Thakur N, Byrne AMP, Ackroyd S, Delahay RJ, van der Poel WHM, Brown IH, Fooks AR, Brookes SM. Differential susceptibility of SARS-CoV-2 in animals: Evidence of ACE2 host receptor distribution in companion animals, livestock and wildlife by immunohistochemical characterisation. *Transbound Emerg Dis.* 2022 Jul;69(4):2275-2286.

Li S, Yang R, Zhang D, Han P, Xu Z, Chen Q, Zhao R, Zhao X, Qu X, Zheng A, Wang L, Li L, Hu Y, Zhang R, Su C, Niu S, Zhang Y, Qi J, Liu K, Wang Q, Gao GF. Cross-species recognition and molecular basis of SARS-CoV-2 and SARS-CoV binding to ACE2s of marine animals. *Natl Sci Rev.* 2022 Jun 23;9(9):nwac122.

Liew, A. Y., Carpenter, A., Moore, T. A., Wallace, R. M., Hamer, S. A., Hamer, G. L., Fischer, R. S. B., Zecca, I. B., Davila, E., Auckland, L. D., Rooney, J. A., Killian, M. L., Tell, R. M., Rekant, S. I., Burrell, S. D., Ghai, R. R., Behravesh, C. B., & the Companion Animals Working Group. (2023). Clinical and epidemiologic features of SARS-CoV-2 in dogs and cats compiled through

national surveillance in the United States, *Journal of the American Veterinary Medical Association*, 261(4), 480-489. Retrieved Apr 24, 2023, from <https://doi.org/10.2460/javma.22.08.0375>

Mastutik G, Rohman A, I'tishom R, Ruiz-Arrondo I, de Blas I (2022) Experimental and natural infections of severe acute respiratory syndrome-related coronavirus 2 in pets and wild and farm animals. *Vet World* 15:565.

Newman A, Smith D, Ghai RR, Wallace RM, Torchetti MK, Loiacono C, Behravesh CB (2020) First reported cases of SARS-CoV-2 infection in companion animals—New York, March–April 2020. *Morb Mortal Wkly Rep* 69:710. <https://doi.org/10.15585/mmwr.mm6923e3>

Padilla-Blanco M, Aguiló-Gisbert J, Rubio V, Lizana V, Chillida-Martínez E, Cardells J, Maiques E, Rubio-Guerri C (2022) The finding of the severe acute respiratory syndrome coronavirus (SARS-CoV-2) in a wild Eurasian river otter (*Lutra lutra*) highlights the need for viral surveillance in wild mustelids. *Front Vet Sci* 9. <https://doi.org/10.3389/fvets.2022.826991>

Sit TH, Brackman CJ, Ip SM, Tam KW, Law PY, To EM, Peiris M (2020) Infection of dogs with SARS-CoV-2. *Nature* 586:776–778.

Wang L, Mitchell PK, Calle PP, Bartlett SL, McAloose D, Killian ML, Torchetti MK (2020) Complete genome sequence of SARS-CoV-2 in a tiger from a US zoological collection. *Microbiol Resour Announc* 9:e00468–20. <https://doi.org/10.1128/MRA.00468-20>

## Highly Pathogenic Avian Influenza (HPAI)

The current HPAI outbreak (2020 ongoing) is unprecedented in global extent and diversity of avian species being affected. H5N1 clade 2.3.4.4b is currently the dominant virus strain subtype circulating globally. In contrast to previous HPAI variants the latter strain has caused significant mortality in wild birds especially among waterfowl, raptors, corvids, gulls, seabirds and in terrestrial mammals (predators; scavengers). Infection of the brain is common among both avian and mammalian species with live animals exhibiting a range of neurological signs. Two UME's involving pinnipeds, (harbor seal (*Phoca vitulina*); gray seal (*Halichoerus grypus*); Southern sea lion (*Otaria flavescens*) in the Americas (USA; PERU) have been linked to infection with H5N1 clade 2.3.4.4b. Both UME's were concurrent with significant regional seabird mortality events (Puryear et al. 2023; Leguia et al. 2023). Additional single cases of infection with the same HPAI H5N1 clade confirmed were reported as of March 17, 2023 from Europe (UK;SWE) and the Americas (USA; Canada; Chile, Peru), and include several small cetacean species (Thorsson et al. 2023) (common dolphin (*Tursiops truncatus*); white-sided dolphin (*Lagenorhynchus acutus*);

harbor porpoise (*Phocena phocena*); spiny porpoise (Burmeister porpoise, *Phocoena spinipinnis*); Chilean dolphin (*Cephalorhynchus eutropia*), a number of pinnipeds (gray seal; harbor seal; Southern sea lion; South American fur seal (*Arctocephalus australis*) and semi-aquatic mammals (marine (*Lontra felina*) and river otters (*Lontra canadensis*; *Lutra lutra*). Likely primary oral or inhalation exposure routes for affected aquatic mammals include environmental contamination, close contact between wild birds and marine mammals (pinnipedia) in haul-out areas, and scavenging on infected carrion. Survival of H5N1 virus pending tissue type and ambient temperature is prolonged (Yamamoto et al. 2017). Currently there is no evidence that mammal to mammal transmission has occurred, however the virus is rapidly evolving. In the past in the northeast US, low pathogenic avian influenza has been found in phocids and cetaceans associated with clusters or UMEs, however there is now strong evidence of pinniped to pinniped transmission and some strains have become endemic in the population. Therefore, a strong focus on tracking viral genome strains and animal disease is needed to inform whether animal to animal transmission is likely to occur. Direct contact transmission among pinnipeds is less likely thus far but cannot be ruled out as the viruses evolve.

## World Organization for Animal Health

In response to the increasing evidence for HPAI spillover (natural infection) in marine mammals the World Organization for Animal Health (WOAH) convened during March 2023 an ad hoc expert group to develop a *Guidance framework for HPAI outbreak management in marine mammals*. Several members of the IWC CDOC intersessional working group are included in the WOAH working group as well as other experts at large. A final draft of the guidance document is expected by the end of May 2023. In the interim given the emerging nature of HPAI in non-avian species, and the many unknowns (transmission, pathogenesis, pathology, host susceptibility) thorough investigation of morbidity and mortality events (single, clustered, UME) in marine mammal species including those under human care is becoming more important. From a field response perspective, both clinical presentation and gross abnormalities on post-mortem examination are among the important gateway keepers to alert and raise suspicion regarding future potential HPAI cases in novel and known species (Table. 1). Standardized assessment of presenting symptoms in alive/moribund animals plus complete life history information, and succinct and complete description of gross abnormalities are required to inform our evolving knowledge base. Furthermore a detailed stranding history with associated spatial and temporal history of concurrent and prior events of HPAI (suspect/confirmed) in avian and non-avian species will be equally helpful. Finally neurologic signs in marine mammals are also a hallmark of infection with several other cetacean diseases of concern (Herpes virus, morbilli virus, toxoplasma gondii, brucella spp.) and marine biotoxins (domoic acid). Clinical samples collected from animals with neurological disease should be concurrently tested for HPAI and these CDOCs including SARS-COV2.

## Reported Cetacean Cases (as of April 24, 2023)

Avian influenza Americas (Canada)	mortality	White sided dolphin (!)	2022	H5N1
Avian influenza Americas (USA)	mortality	common bottlenose dolphin (1)	2022	H5N1
Avian influenza Europe (Sweden)	mortality	harbor porpoise (1)	2022	H5N1
Avian influenza Americas (canada)	mortality	harbor porpoise (1)	2022	H5
Avian influenza Americas (PERU)	mortality	dolphin (1)	2023	H5N1
Avian influenza Americas (Chile)*	mortality	chilean dolphin (2); spiny porpoise (2)	2023	H5N1
Avian influenza Europe (UK)	mortality	dolphin (2); harbor porposie (1)	2022/2023	H5N1

- Both cetacean species are endemic to South america. The Chilean dolphin (*Cephalorhynchus eutropia*) “ tonino” is a small protected cetacean species along the coasts of Chile.



Table. 1. Summary of HPAI associated clinical and gross pathologic observations in Mammals (terrestrial/aquatic). source: see references below note: cat and dog cases are from media; black bear media; **purple** Sea Lion; **green** Harbor Porpoise; **red** Harbor seal/Gray seal.

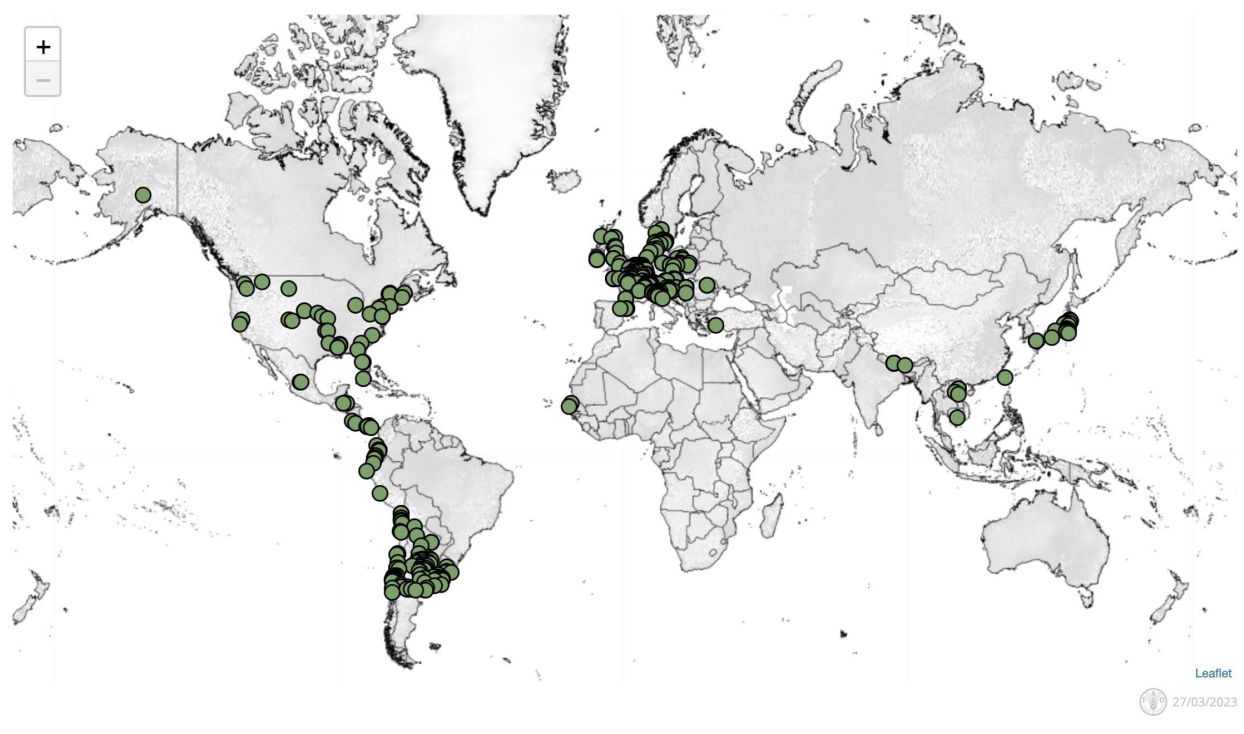
<b>DATA</b>	<b>Composite of HPAI associated OBSERVATIONS in Mammals (terrestrial/aquatic) source: see references below note: cat and dog cases are from media; black bear media; purple Sealion; green Harbor Porpoise; red Harbor seal/Gray seal</b>	NOTE: Harbor seal/Greyseal /sealion pathology not published; Note: Different observers -clinics/postmortem =Absence of evidence is not evidence of absence.;
<b>Out of habitat</b>	stranded, hauled out,	
<b>Status</b>	<i>alive, dead, moribund, euthanized</i>	
<b>Carcass condition</b>	fresh to autolized	
<b>Predator scavenger clade</b>	mesocarnivores; piscivorous; opportunistic feeders	
<b>Species</b>	red fox, bobcats, coyotes, grey foxes, fishers, raccoons, striped skunks, Virginia opossums, black bear, brown bear; Grizzly bear; Kodiak brown bear; mountain lion, stone marten, <b>eurasian otter, marine otter</b> , dogs, cats, mink, bush dogs; lynx, <b>harbor porpoise, sealion, furseal, common dolphin, atlantic white sided dolphin, harbor seal, gray seal</b>	
<b>Age class</b>	mixed; possible demographic bias - juvenile, male	
<b>Body condition index</b>	poor to fair to good	
<b>HPAI Confirmation</b>	PCR/IHC	
<b>HPAI Variant</b>	A/H5N1 lineage 2.3.4.4b (some mammalian adaptive mutations) e.g. mutation PB2-627K increases the replication of the virus in mammalian cell lines and at the mammalian temperature	
<b>Samples</b>	brain other tissues; oropharyngeal swabs; nasal swabs, rectal swabs	
<b>ORGAN SYSTEM/BEHAVIOR</b>	<b>CLINICAL PRESENTATION</b>	
Neurological symptoms	<b>seizures</b> , tremors, nystagmus, grimace; <30 minutes lip retraction, rapid opening and closing of mouth, skin twitching, head shaking; <b>neurological</b>	

	<b>signs (TBD)</b>	
Respiratory symptoms	<b>dyspnoe; nasal discharge, respiratory signs (TBD)</b>	
Digestive symptoms	diarrhea; vomiting; <b>sialorrhea (drooling) salivation;</b> foaming	
Ocular symptoms	ocular discharge; no response to sight stimuli (blindness), nystagmus	
Hearing & Smell	not responding to sound (Black bear), responded only slightly to smells of people (black bear)	
Touch Reception	loss of proprioception	
Animal grooming/preening behavior	<b>data gap</b> (Two foxes had a shiny coat, and one fox (Oosterbeek) was covered with mud)	
Appetite	inappetence; showed no interest in water	
Posture/Motor Coordination	<b>swam in circles, was unable to right itself (drowned) disorientation; "stargazing" posture; ataxia,</b> torticollis, circling, stumbling; crouching down with convex back and legs tucked under	
Reflexes	diminished swallowing reflex	
Responsiveness	inappropriate or lack of fear of humans; lethargy; fatigue/sleep; aggression; somnolent	
Withdrawal & Hiding behavior	approaching and trying to get under cars (black bear)	
Animal communication	vocalization (moaning),	
Body temperature	fever	
Body mass	weight loss	
<b>mortality</b>		
<b>PATHOLOGY_GROSS DESCRIPTION</b>		
CNS	hemorrhage, congestion; congestion with hydrocephalus and malacia	
Sensory system	no lesions but behavior consistent with blindness reported	
Respiratory system	lung congestion, <b>edema</b> , failure to collapse, hemorrhage, and pleural effusion	
Cardiovascular system	Pericardial effusion, petechiae , and myocardial pallor	
Immune System	<b>data gap (on histopathology lesions)</b>	
Hepatobiliary system	pallor, congestion , enlargement , and hemorrhage , liver necrosis (Mountain lion)	

Urinary system	kidney congestion and cortical hemorrhage	
Body cavities	hemoperitoneum & hemothorax	
Reproductive system	<b>data gap</b>	
Digestive system	congestion, hemorrhage, and loose feces, pancreatitis	
Skeletal muscle system	petechial haemorrhages in the muscles	
Integumentary system	petechial hemorrhages subcutaneous tissue	
Trauma	road traffic accident;	
Co-Infection	rare (viral co-infection) including adenovirus, parvovirus, rabies, parasites common	

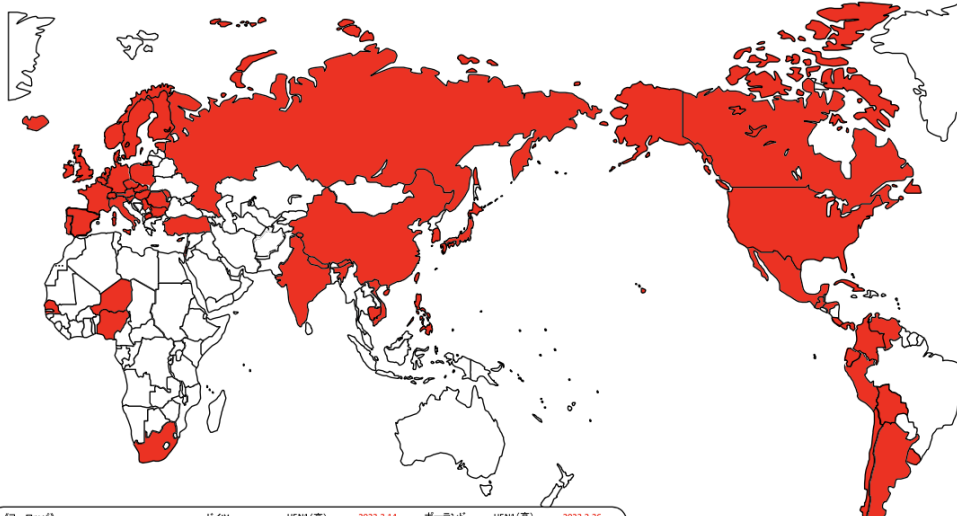
Fig.1-3. Maps by the FAO ( Jan 01 22 - Mar 27, 2023 wild birds), the Japanese Ministry of Agriculture, Fisheries and Forestry (MAFF) (wild birds & mammals 2023), and USGS (USA) (cumulative HPAI detection in the Americas 2021-2023).

I



## 高病原性鳥インフルエンザの発生状況(2022年7月以降)

■ : 2022年7月以降に継続発生  
または新規発生した国



国名	発生状況	発生日	国名	発生状況	発生日
ドイツ	HSN1(高)	2023.3.14	ポーランド	HSN1(高)	2023.2.26
アイスランド	HSN1(高)	2022.10.17			2023.3.10
アイルランド	HSN1(高)	2022.11.18	ノルウェー	HSN1(高)	2022.11.9
		2023.2.14			2022.11.9
イタリア	HSN1(高)	2022.3.6	ハンガリー	HSN1(高)	2023.3.6
		2023.3.10			2023.3.4
英国	HSN1(高)	2023.3.2	フィンランド	HSN1(高)	2022.8.16
オランダ	HSN1(高)	2023.1.26			2022.9.17
		2022.12.23	フェロー諸島	HSN1(高)	2022.10.2
北マケドニア	HSN1(高)	2022.11.3			2022.9.22
スイス	HSN1(高)	2023.3.9	フランス	HSN1(高)	2023.3.1
スウェーデン	HSN1(高)	2023.2.19			2023.3.6
スペイン	HSN1(高)	2023.2.4	ブルガリア	不明(高)	2022.10.20
		2023.2.24			2023.1.24
スロベニア	HSN1(高)	2023.2.24			2023.1.24
		2023.2.25	ベルギー	HS(高)	2023.3.1
セルビア	HSN1(高)	2023.1.23			2023.3.12
デンマーク	HSN1(高)	2023.3.16			不明(高)
		2022.2.10			2023.3.6

国名	発生状況	発生日
日本	HSN1(高)	2023.3.14
		2023.3.10
中国	HSN1(高)	2022.7.9
韓国	HSN1(高)	2023.3.9
		2022.10.20
台湾	HSN1(高)	2023.3.16
		2023.2.8
香港	HSN5(高)	2023.1.18
イスラエル	HSN1(高)	2022.12.5
		2023.1.11
フィリピン	HSN1(高)	2023.1.11
ベトナム	HSN1(高)	2022.12.16
インド	HSN1(高)	2022.10.3
ネパール	HSN1(高)	2023.2.23
カンボジア	HSN1(高)	2023.2.24

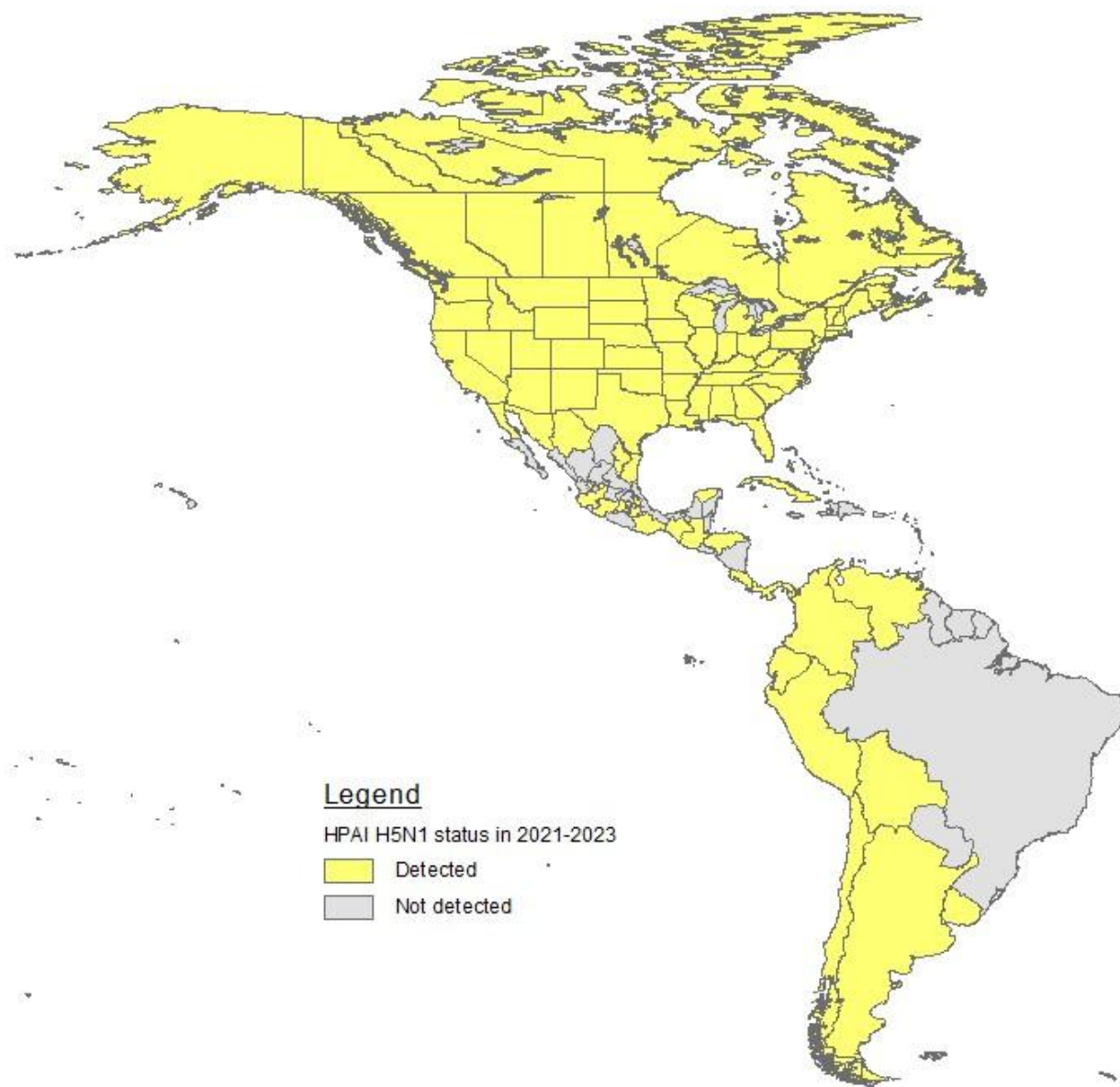
国名	発生状況	発生日
米国	HSN1(高)	2023.3.8
		2023.2.28
		2022.9.10
カナダ	HSN4(高)	2023.3.10
		2023.3.9
		2023.1.11
メキシコ	HS(高)	2023.1.10
		2022.12.6
パナマ	HSN1(高)	2023.3.10
エクアドル	HSN1(高)	2023.2.28
コロンビア	HSN1(高)	2023.1.12
		2023.2.20
		2023.3.3
ペネズエラ	HSN1(高)	2022.11.17
ペルー	HS(高)	2023.1.6
		2022.12.10*
		2023.2.15
ホンジュラス	不明(高)	2023.2.22
チリ	HSN1(高)	2023.3.10
		2023.3.9
コスタリカ	HS(高)	2023.2.16
ウルグアイ	HS(高)	2023.1.11
		2023.3.14
グアテマラ	HSN1(高)	2023.2.16
アルゼンチン	HS(高)	2023.3.26
ボリビア	HSN1(高)	2023.2.3
		2023.2.1
		2023.2.4*

国名	発生状況	発生日
ロシア	HSN1(高)	2023.1.26
		2023.2.2

国名	発生状況	発生日
南アフリカ共和国	HSN1(高)	2023.1.6
		2022.11.29
		2022.12.1
ニジェール	HSN1(高)	2022.12.18
ナイジェリア	HSN1(高)	2022.12.19
セネガル	HSN1(高)	2023.3.8

**2023年3月17日現在**  
出典: OIE等

※日付は発生日又は検体採取日に基づく  
※[ ]は野鳥及び養殖鳥等における発生を示す  
※本圖は発生の有無を示したもので、  
その後の清浄性確認については記載していない  
※個別に最新の発生事例を記載  
※( )は病態性  
※韓国、野鳥については最新の情報更新が10月20日  
※白色の国、地域であっても継続発生で報告されていない可能性もある。



## LITERATURE CITED AND FURTHER READING

Alkie TN, Cox S, Embury-Hyatt C, Stevens B, Pople N, Pybus MJ, Xu W, Hisanaga T, Suderman M, Koziuk J, Kruczkiewicz P, Nguyen HH, Fisher M, Lung O, Erdelyan CNG, Hochman O, Ojkic D, Yason C, Bravo-Araya M, Bourque L, Bollinger TK, Soos C, Giacinti J, Provencher J, Ogilvie S, Clark A, MacPhee R, Parsons GJ, Eaglesome H, Gilbert S, Saboraki K, Davis R, Jerao A, Ginn M, Jones MEB, Berhane Y. Characterization of neurotropic HPAI H5N1 viruses with novel genome constellations and mammalian adaptive mutations in free-living mesocarnivores in Canada. *Emerg Microbes Infect.* 2023 Dec;12(1):2186608. doi: 10.1080/22221751.2023.2186608.

Bordes L, Vreman S, Heutink R, Roose M, Venema S, Pritz-Verschuren SBE, Rijks JM, Gonzales JL, Germeraad EA, Engelsma M, Beerens N. Highly Pathogenic Avian Influenza H5N1 Virus Infections in Wild Red Foxes (*Vulpes vulpes*) Show Neurotropism and Adaptive Virus Mutations. *Microbiol Spectr.* 2023 Feb 14;11(1):e0286722. doi: 10.1128/spectrum.02867-22. Epub 2023 Jan 23.

Elsmo EJ, Wünschmann A2, Beckmen, KB3, Broughton-Neiswanger LB4, BucklesEL5, Ellis J6, Fitzgerald SD6, Gerlach R7, Hawkins S8, Ip HS9, Lankton JS9, Lemley EM8,10,Lenoch JB11, Killian ML12, Lantz K12, Long L13, Maes R6, Mainenti M14, Melotti J15, MoriartyME15, Nakagun S5, Ruden RM14,16, Shearn-Bochsler V9, Thompson D6, Torchetti MK12, VanWettere AJ17, Wise AG6, Lim AL1 Pathology of natural infection with highly pathogenic avian influenza virus (H5N1) clade 2.3.4.4b in wild terrestrial mammals in the United States in 2022  
<https://www.biorxiv.org/content/10.1101/2023.03.10.532068v1.full.pdf>

Floyd, T. Ashley C. Banyard, Fabian Z. X. Lean, Alexander M. P. Byrne, Edward Fullick, Elliot Whittard, Benjamin C. Mollett, Steve Bexton, Vanessa Swinson, Michele Macrelli, Nicola S. Lewis, Scott M. Reid, Alejandro Núñez, J. Paul Duff, Rowena Hansen, Ian H. Brown Systemic infection with highly pathogenic H5N8 of avian origin produces encephalitis and mortality in wild mammals at a UK rehabilitation centre  
bioRxiv 2021.05.26.445666; doi: <https://doi.org/10.1101/2021.05.26.445666>

Frymus, T., Sándor Belák, Herman Egberink, Regina Hofmann-Lehmann, Fulvio Marsilio, Diane D. Addie, Corine Boucraut-Baralon, Katrin Hartmann, Albert Lloret, Hans Lutz, Maria Grazia Pennisi, Etienne Thiry, Uwe Truyen, Séverine Tasker, Karin Möstl, and Margaret J. Hosie. 2021. "Influenza Virus Infections in Cats" *Viruses* 13, no. 8: 1435.  
<https://doi.org/10.3390/v13081435>

Gamarra-Toledo, V., Pablo I. Plaza, Roberto Gutiérrez, Giancarlo Inga-Diaz, Patricia Saravia-Guevara, Oliver Pereyra-Meza, Elver Coronado-Flores, Antonio Calderón-Cerrón, Gonzalo Quiroz-Jiménez, Paola Martinez, Deyvis Huamán-Mendoza, José C. Nieto-Navarrete, Sandra Ventura, Sergio A. Lambertucci Mass Mortality of Marine Mammals Associated to Highly Pathogenic Influenza Virus (H5N1) in South America  
 bioRxiv 2023.02.08.527769; doi: <https://doi.org/10.1101/2023.02.08.527769>

Jimenez-Bluhm, P. Jurre Y. Siegers, Shaoyuan Tan, Bridgett Sharp, Pamela Freiden, Katherinne Orozco, Soledad Ruiz, Cecilia Baumberger, Pablo Galdames, Maria Antonieta Gonzalez, Camila Rojas, Erik A. Karlsson, Christopher Hamilton-West, Stacey Schultz-Cherry Detection and Phylogenetic Analysis of Highly Pathogenic A/H5N1 Avian Influenza Clade 2.3.4.4b Virus in Chile, 2022  
 bioRxiv 2023.02.01.526205; doi: <https://doi.org/10.1101/2023.02.01.526205>

Lean, F.ZX., Marco Falchieri, Natalia Furman, Glen Tyler, Caroline Robinson, Paul Holmes, Scott M Reid, Ashley C Banyard, Ian H Brown, Catherine Man, Alejandro Núñez. Pathology of naturally acquired high pathogenicity avian influenza virus H5N1 infection in seabirds  
 bioRxiv 2023.02.17.528990; doi: <https://doi.org/10.1101/2023.02.17.528990>

Leguia, M Alejandra Garcia-Glaessner, Breno Muñoz-Saavedra, Diana Juarez, Patricia Barrera, Carlos Calvo-Mac, Javier Jara, Walter Silva, Karl Ploog, Lady Amaro, Paulo Colchao-Claux, Marcela M. Uhart, Martha I. Nelson, Jesus Lescano Highly pathogenic avian influenza A (H5N1) in marine mammals and seabirds in Peru  
<https://www.biorxiv.org/content/10.1101/2023.03.03.531008v1>

Puryear W, Sawatzki K, Hill N, Foss A, Stone JJ, Doughty L, et al. Highly pathogenic avian influenza A(H5N1) virus outbreak in New England seals, United States. *Emerg Infect Dis.* 2023 Apr [date cited]. <https://doi.org/10.3201/eid2904.221538>

Reperant LA, van Amerongen G, van de Bildt MW, Rimmelzwaan GF, Dobson AP, Osterhaus AD, Kuiken T. Highly pathogenic avian influenza virus (H5N1) infection in red foxes fed infected bird carcasses. *Emerg Infect Dis.* 2008 Dec;14(12):1835-41. doi: 10.3201/eid1412.080470.



Tammiranta N, Isomursu M, Fusaro A, Nylund M, Nokireki T, Giussani E, Zecchin B, Terregino C, Gadd T. Highly pathogenic avian influenza A (H5N1) virus infections in wild carnivores connected to mass mortalities of pheasants in Finland. *Infect Genet Evol.* 2023 Mar 6;111:105423. doi: 10.1016/j.meegid.2023.105423. Epub ahead of print.

Thorsson, E., Zohari, S., Roos, A., Banihashem, F., Bröjer, C., & Neimanis, A. (2023). Highly Pathogenic Avian Influenza A(H5N1) Virus in a Harbor Porpoise, Sweden. *Emerging Infectious Diseases*, 29(4), 852-855. <https://doi.org/10.3201/eid2904.221426>..

Vreman S, Kik M, Germeraad E, Heutink R, Harders F, Spierenburg M, Engelsma M, Rijks J, van den Brand J, Beerens N. Zoonotic Mutation of Highly Pathogenic Avian Influenza H5N1 Virus Identified in the Brain of Multiple Wild Carnivore Species. *Pathogens.* 2023 Jan 20;12(2):168. doi: 10.3390/pathogens12020168.

Yamamoto Y, Nakamura K, Mase M. Survival of Highly Pathogenic Avian Influenza H5N1 Virus in Tissues Derived from Experimentally Infected Chickens. *Appl Environ Microbiol.* 2017 Aug 1;83(16):e00604-17. doi: 10.1128/AEM.00604-17.