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Avian influenza overview December 2023–March 2024

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HPAI IN DOMESTIC AND WILD BIRDS IN EUROPE





HPAI IN BIRDS | DECEMBER 2023 - MARCH 2024



HPAI IN DOMESTIC BIRDS | DECEMBER 2023 - MARCH 2024





Author: EFSA Data sources: ADIS, WOAH Date updated: 15/03/2024



Temporal distribution in poultry and wild birds

- a) HPAI virus subtypes
- b) Poultry categories
- c) Wild bird categories





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Spatial distribution in wild birds

- Waterfowl all over Europe and particularly in southeastern parts
- Colony-breeding seabirds only along coastlines
- Still a few common cranes ('Others') in southern parts



EFSA'S BIRD FLU RADAR

https://app.bto.org/hpai







Most frequently affected wild bird species

- Waterfowl: swans and geese
- Colony-breeding seabirds: gulls
- Raptors: peregrine falcons
- <u>'Others'</u>: common cranes

No new wild bird species affected In more than 20% of HPAI virus detections, the species was not identified

HPAI IN BIRDS WORLDWIDE | DECEMBER 2023 – MARCH 2024





HPAI IN MAMMALS | DECEMBER 2023 – MARCH 2024

FARMED	PET	WILD
 Fur farms (Finland): Arctic fox, red fox, common raccoon dog A(H5) antibodies were found in 44 out of all 346 (12.7%) surveyed establishments Goat (USA) Cattle (USA) 	 Cat (Canada, USA) 	 Red fox (Germany, Sweden*) Red fox (Norway) A(H5N5) Mountain lion (USA) Raccoon (USA) Striped skunk (USA) Abert's squirrel (USA)* Polar bear (USA)* South American fur seal (Antarctic region) Southern elephant seal (Antarctic region)



*previous reporting period

HPAI IN HUMANS | DECEMBER 2023 – APRIL 2024



Subtype	New cases reported (deaths)	Total cases (deaths)	Countries reporting human cases
	1 Dec 2023–8 Apr 2024		
A(H3N8)	-	3 (1), since 2022	China
A(H5N1)	Cambodia: 5 (1)	889* (463)	23 countries, including one EU/EEA country: Spain*
	Vietnam: 1 (1)	Since 2004	
	United States: 1		
A(H5N6)	China: 2 (1)	90 (35)	No EU/EEA country; China (89), Laos (1)
		Since 2014	
A(H9N2)	China: 7	136 (2)	No EU/EEA country; China (122), Egypt (4), Bangladesh (3), Cambodia (2), Oman (1), Pakistan (1), India (1), Senegal (1), Vietnam (1)
	Vietnam: 1	Since 1998	
A(H10N3)	China: 1	3, since 2021	China (3)
A(H10N5)	China: 1 (1)	1 (1), first reported in 2024	China (1)
*includes detections due to suspected environmental contamination from Spain (2) and the USA (1) in 2022, and from the UK (3) in 2023		Source: ECDC line list; WHO; Cumulative number of confirmed human cases for avian influenza A(H5N1) reported to WHO, 2003-2024 (2024_feb_tableh5n1.pdf (who.int)); Highly Pathogenic Avian Influenza A (H5N1) Virus Infection Reported in a Person in the U.S. CDC Online Newsroom CDC; HCMC: recorded the	

country's first human case of avian influenza A(H9N2) (hcdc.vn)



HPAI IN HUMANS | RISK ASSESSMENT

- Transmission to humans is rare but disease can be severe
- No sustained human-to-human transmission
- Sporadic human infections likely to continue occurring in settings where people are exposed to infected animals or their environment

Risk assessment:

- General public in the EU/EEA: the risk of human infection with avian influenza A(H5) clade 2.3.4.4b viruses currently circulating in Europe is assessed as <u>low</u>
- The risk is considered <u>low-to-moderate</u> for people who are occupationally or otherwise exposed to birds or mammals infected with avian influenza



GENETIC CHARACTERISTICS OF HPAI VIRUSES IN EUROPE



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ZOONOTIC POTENTIAL OF HPAI VIRUSES IN EUROPE

- A(H5N1) viruses characterised in the current epidemiological year continue to be well-adapted to avian species, as they retain a preferential binding for avian-like receptors
- Mutations in the PB2 protein associated with virus adaptation in mammals (E627K or D701N), and therefore increased zoonotic potential, have recently been detected in viruses collected from birds:
- <u>A(H5N5) viruses (genotype EA-2021-I)</u> collected in Norway and Germany from wild birds
- <u>A(H5N1) viruses</u> collected from separate outbreaks in domestic birds in Poland (<u>genotypes EA-2022-CH and EA-2022-AB</u>) and Denmark (<u>genotype EA-2023-DB</u>)
- About 43% of European viruses collected from mammalian species contain molecular markers of mammalian adaptation in the PB2 protein → these mutations can be rapidly acquired by the virus during infection in mammalian species



European Reference Laboratory for Avian Influenza



OPTIONS FOR RESPONSE

- Maintaining high levels of biosecurity in poultry establishments
- Surveillance:
 - Active surveillance in wild birds to estimate viral prevalence and seroprevalence (to assess whether some level of flock immunity has been acquired)
 - Inclusion of fur farms in national surveillance programmes
 - Increased surveillance in mammals, especially in those in or around affected poultry establishments
- Accurate and comprehensive recording, investigation and reporting of HPAIassociated mortality events in wild birds and mammals
 - Timely sharing of viral genomic sequence data
- Improving wild bird species identification
- Continuously monitoring the dynamics of HPAI A(H5N5) virus

