Standing Committee on Plants, Animals, Food and Feed Section Animal Health and Welfare

MARCH - APRIL 2023



Francesca Baldinelli Scientific Officer **Animal Health Team**



MONITORING HPAI OUTBREAKS

- Update of the avian influenza situation in poultry, captive and wild birds in Europe and beyond
 - Spatio-ten
 - Evolution
 - Character

Joint

Currently



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doi: 10.2903/j.efsa.2023.8039

Avian influenza overview March – April 2023

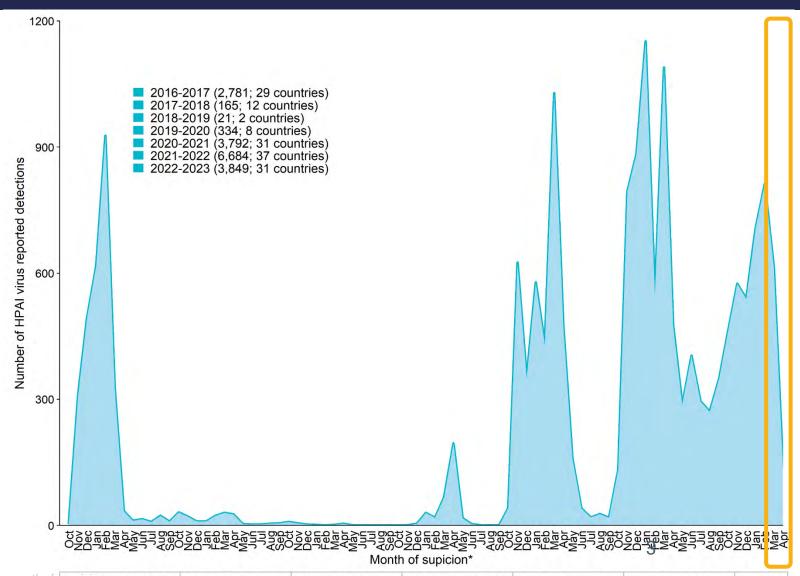
European Food Safety Authority, European Centre for Disease Prevention and Control, European Union Reference Laboratory for Avian Influenza, Cornelia Adlhoch, Alice Fusaro, José L Gonzales, Thijs Kuiken, Grazina Mirinaviciute, Éric Niqueux, Karl Stahl, Christoph Staubach, Calogero Terregino, Alessandro Broglia, Lisa Kohnle and Francesca Baldinelli



HPAI IN EUROPE IN BIRDS

Distribution of HPAI virus detections reported in EU/EEA and the UK by epidemic seasons and month of suspicion

1 Oct 2016 – 28 Apr 2023 (17,626)

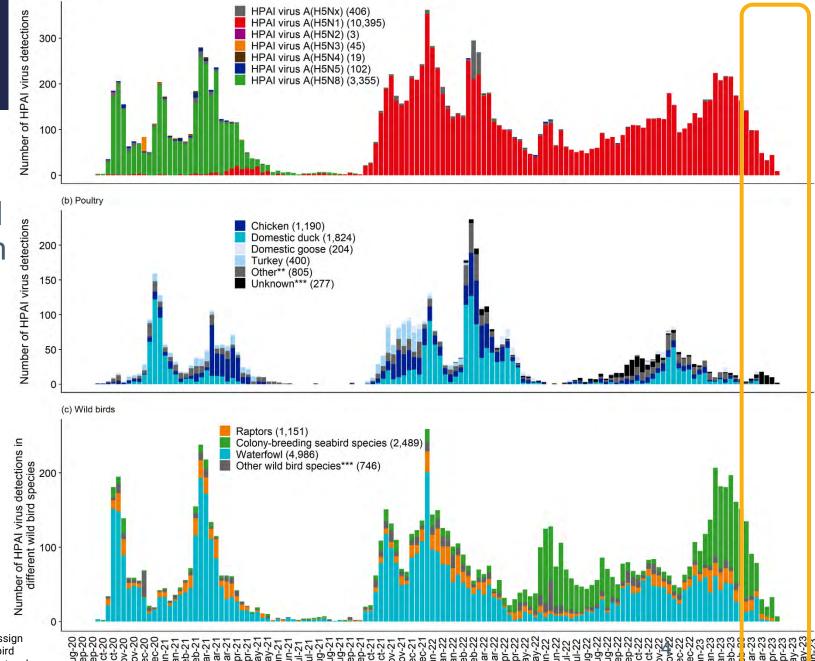


HPAI IN EUROPE IN BIRDS

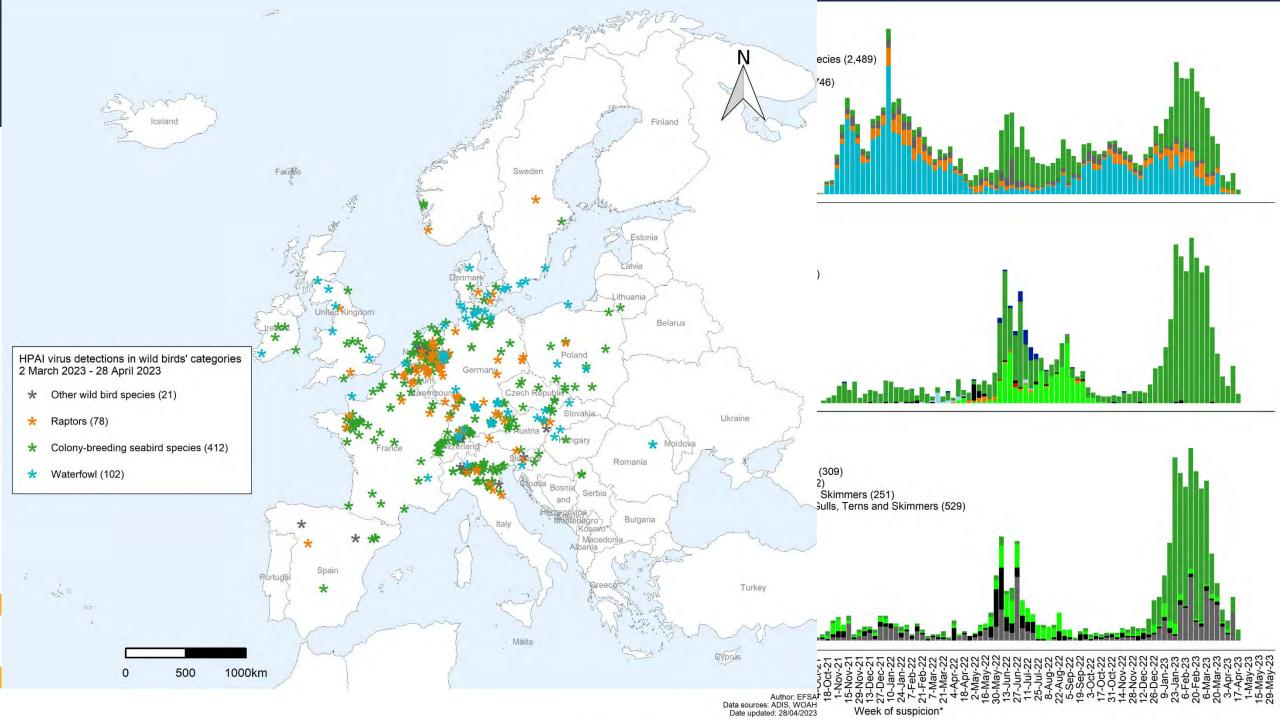
(a) All birds

Distribution of total number of HPAI virus detections reported in Europe by week of suspicion and

- a. virus subtype
- b. affected poultry categories
- c. affected wild bird categories
- 5 Oct 2020 28 Apr 2023



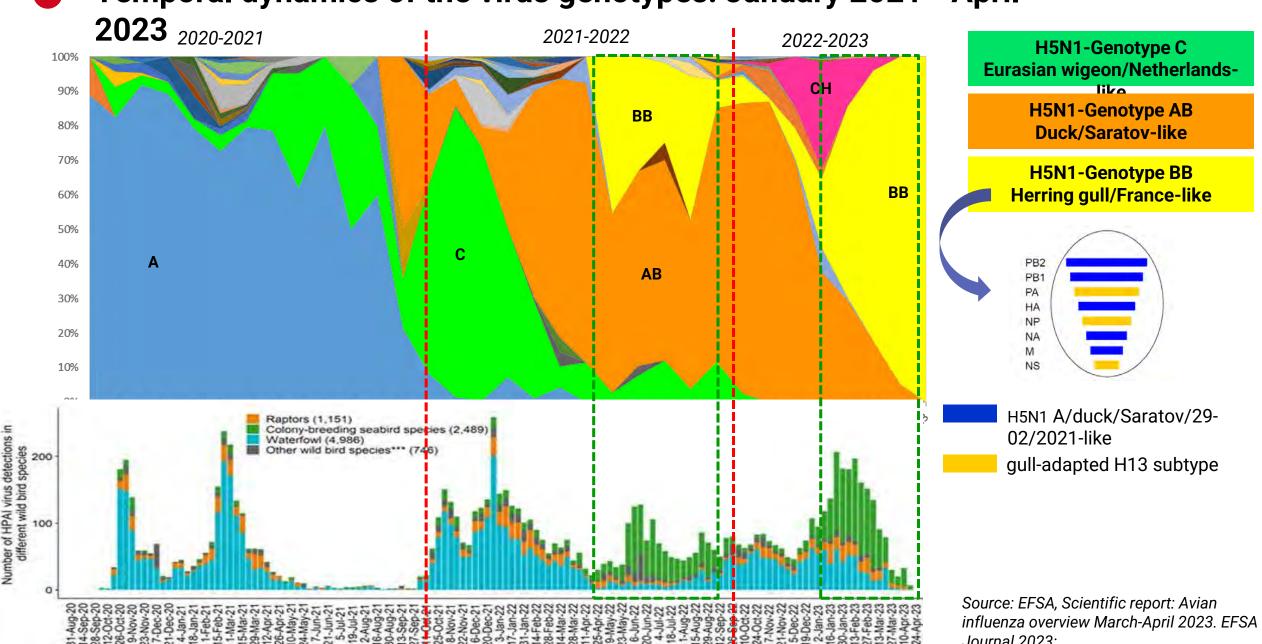
^{*} When the date of suspicion is not available then the date of confirmation is used to assign the week of suspicion. ** 'Other domestic species' category contains mixed, unknown bird species, or categories different from those displayed (i.e guinea fowl, peacock, pheasant and quail). *** 'Other wild species' category contains mixed, unknown bird species, or categories different from those displayed





Temporal dynamics of the virus genotypes: January 2021 - April

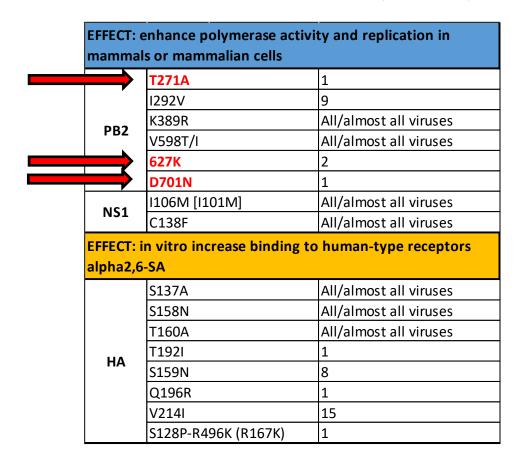
Week of suspicion



Journal 2023: https://doi.org/10.2002/j.ofca.2022.8020



Mutations identified in A(H5N1) from avian species in Europe (2022-2023)

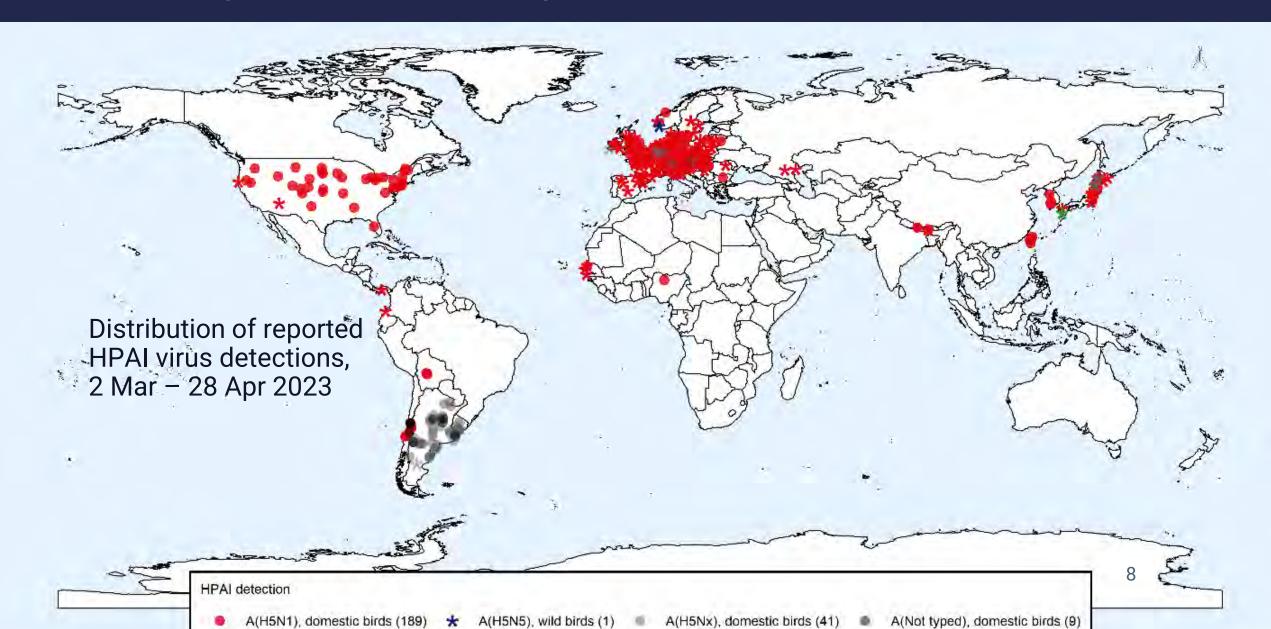


EFFECT: increase virulence					
	P42S	All/almost all viruses			
	V149A	All			
NS1	L103F, I106M	All/almost all viruses			
	K55E, K66E, C138F	All/almost all viruses			
	227ESEV230 (PDZ domain	All/almost all viruses			
EFFECT: resistance toward antiviral drugs					
M2	271	3			
PA	A36V	3			
EFFECT: decrease antiviral response in ferrets					
NS1	N205S (with NS2: T47A) All/almost all viruses				
EFFECT: disruption of the second sialic acid binding site (2SBS)					
NA	S369I	266	BE		
	S369C/N/R	17			
	K432E/N	33			
EFFECT: evade human BTN3A3 (a potent inhibitor of avian					
but not human influenza A viruses replication)					
NP	Y52N	354	BE		

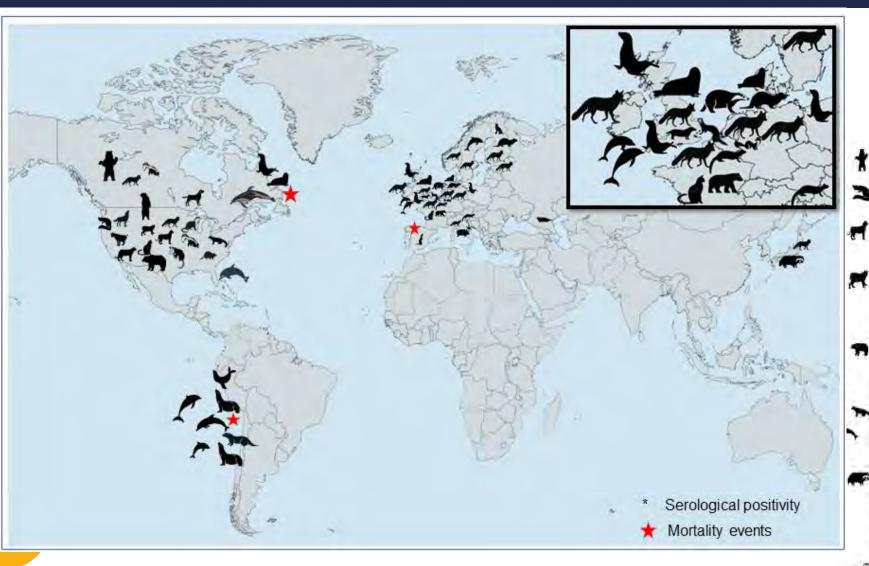
Molecular analyses of the studied A(H5N1) viruses circulating in birds in Europe during the 2022–2023 epidemiological year indicate that these viruses continue to be well-adapted to avian species, as they retain a preferential binding for avian-like receptors

The real effect of these mutations on the biological characteristics of the viruses is still unknown and further studies are needed to improve existing knowledge

HPAI WORLDWIDE IN BIRDS



HPAI IN MAMMAL SPECIES OTHER THAN HUMANS



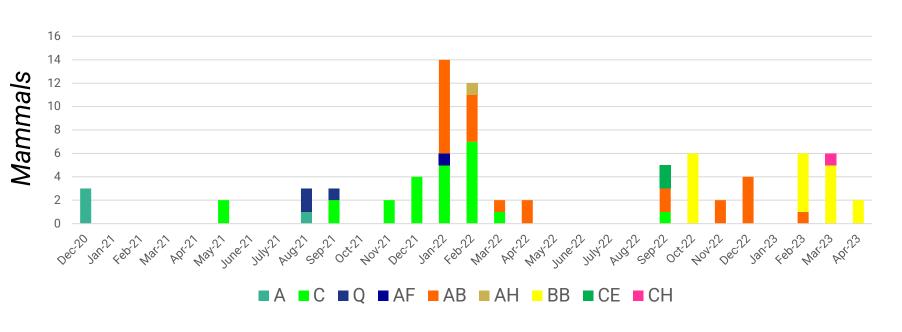
American black bear (Ursus americanus) Burmeister's porpoise (Phocoena spinip American mink (Neogale vison) Caspian seal (Pusa caspica) American pine marten (Martesamericana) Cat (Felis catus) Amur leopard (Panthera pardus orientalis) Chilean dolphin (Cephalorhynchus eutro Amur tiger (Pantheratigris) Common dolphin (Delphinus delphi) Asiatic black bear (Ursus thibetanus) Coyote (Canis latrans) Bobcat (Lynxrufus) Dog (Canisfamiliaris) Beech marten (Martesfoina) Eurasian badger (Melesmeles) Bottlenose dolphin (Tursiops truncatus) A Eurasian lynx (Lynx lynx) Rown bear (Ursus arctos) Eurasian otter (Lutra lutra) European polecat (Mustela putorius) Porpoise (Phocoena phocoena) Ferret (Mustela furo) Raccoon (Procyon lotor) Fisher cat (Pekania perinanti) Red fox (Vulpes vulpes) Grey seal (Halichoerus grypus) Skunk (Mephitismephitis) South America fur seal (Arctophoca Harbour seal (Phoca vitulina) South American bush dogs (Speothos Japaneseraccoon dog (Nyctereutes viverrinus) South American sea lion (Otaria Kodiak grizzly bear (Ursus arctos horribilis) Marine otter (Lontra felina) Virginia opossum (Didelphis virginiana) White-sided dolphin (Lagenorhynchus Mountain lion (Puma concolor) acutus)

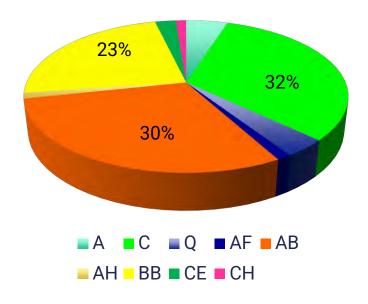
Pig (Sus scrofa)

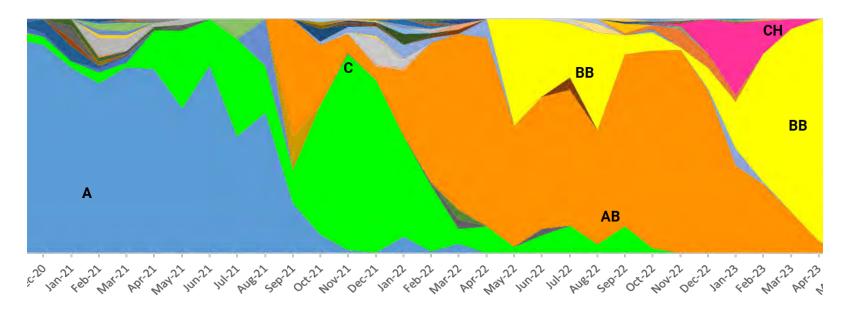
North American river otter (Lontra canadensis)



Genetic diversity of A(H5Nx) in mammals in Europe (2020-2023)



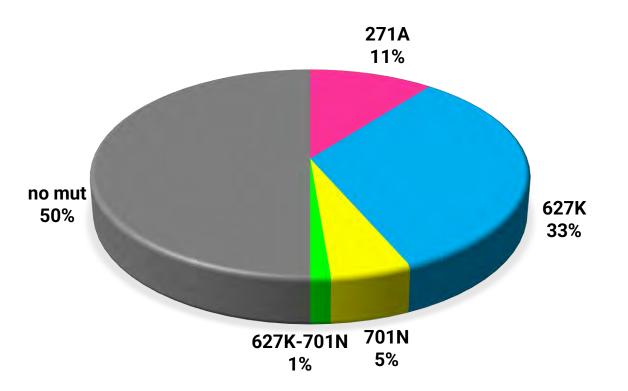




All the characterized viruses belong to 9 different A(H5N1) and A(H5N8) genotypes previously identified in birds, with most of the viruses (86%) belonging to the 3 most widespread genotypes in birds in Europe: C, AB and BB



Mutations identified in A(H5Nx) from mammals In Europe (2020-2023)



50% of the characterized viruses from mammals contain at least **one of the adaptive markers** associated with an **increased virulence and replication in mammals** in the **PB2** protein (E627K, D701N or T271A) (Suttie et al., 2019).

These mutations have rarely been identified in the HPAI A(H5) viruses of clade 2.3.4.4b collected in birds in Europe since October 2020 (<0.5% of viral sequences from birds).

HUMAN CASES DUE TO AVIAN INFLUENZA, 2023



Subtype	New cases in 2023	Total cases (deaths)	Countries reporting human cases
A(H3N8)	One case:	3 (0)	China
	China	Since 2022	
A(H5N1)	Seven cases:	876 (457)	23 countries reported cases,
	Cambodia, Chile, China and Ecuador 2 United Kingdom (asymptomatic, involved in culling during outbreak)	Since 2004	EU/EEA: Spain with virus fragment detections in two poultry workers considered contamination and no productive infections
A(H5N6)	No new case	84 (33)	China (83), Laos (1)
		Since 2014	
A(H9N2)	Five cases:	123 (2)	No EU/EEA country;
	China	Since 998	China (107), Egypt (4), Bangladesh (3), Cambodia (2), Oman (1), Pakistan (1), India (1), Senegal (1)

RISK ASSESSMENT REMAINS VALID



The risk of human infection due to avian influenza viruses of the currently circulating clade 2.3.4.4b in Europe is assessed as low for the general population and low to moderate for people occupationally or otherwise HPAI virus exposed

But...

Sporadic human infections with severe disease in the EU/EEA cannot be excluded

OPTIONS FOR RESPONSE

- Passive surveillance in poultry could be complemented by active surveillance (bucket sampling) and testing of birds before slaughter is highly recommended to increase early detection
- Close monitoring of seabird breeding colonies, including black-headed gulls, for unusual mortality will allow for the early detection of HPAI virus and, if appropriate, to remove carcasses for the reduction of environmental contamination and therefore reduced mortality
- Serological surveillance in seabirds, in particular in endangered species, is urgently needed to evaluate the real level of virus circulation in those species as well as their immune protection
- Species identification remains of utmost importance for the correct interpretation of passive surveillance efforts in wild birds



OPTIONS FOR RESPONSE

- Better, more accurate and timely reporting of HPAI virus detections in mammals in a way that reliable numbers of infected animals could be used as quantitative information for risk assessment
- Avoid exposure of carnivore pets to dead or diseased animals (mammals and birds) in areas where mortality in gulls or other potentially HPAI virus-infected animals are reported
- Extended and enhanced surveillance of both wild mammals (particularly carnivores and cetacean) and farmed mammals (particularly American mink and domestic pigs) in risk areas where HPAI is present in wild birds and poultry is recommended
- Thoroughly investigate the dynamic of the infection in case of mass mortality events associated with HPAI virus detected in mammalian species



EXPERT INVOLVED

Member State representative for avian influenza

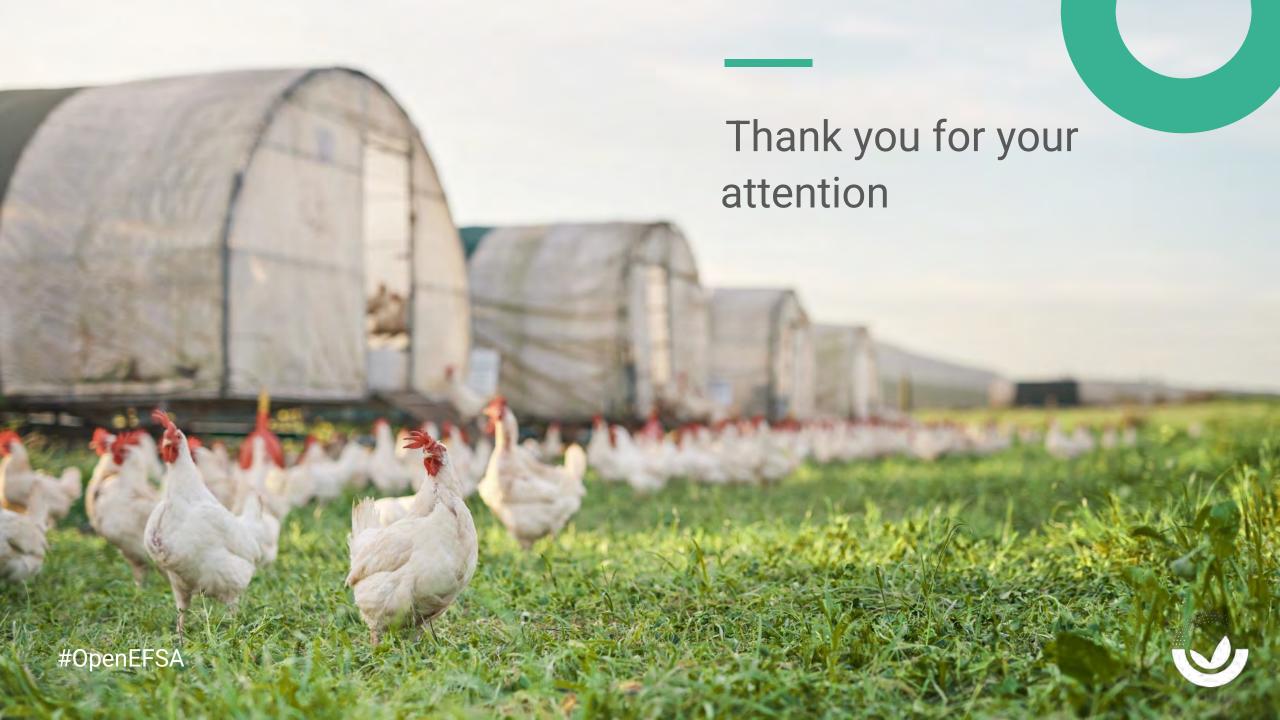
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