



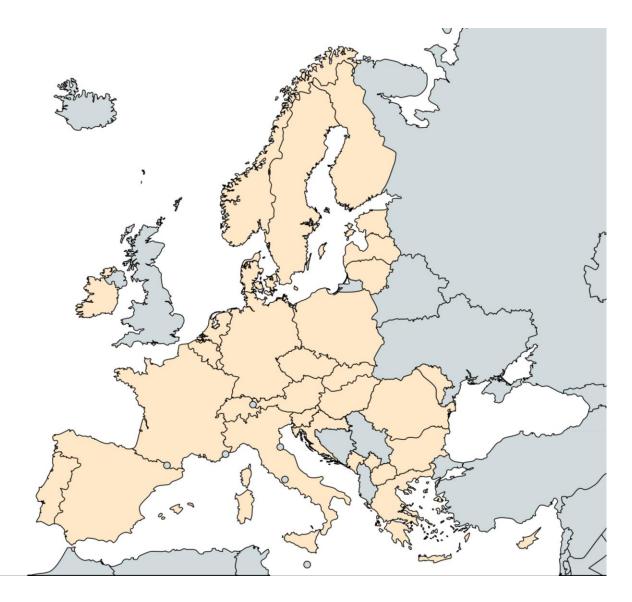
MEMBER STATES (& THIRD COUNTRIES) REPORTS FOR AVIAN INFLUENZA AND NEWCASTLE DISEASE 06/2024 – 06/2025 Based on the response to the questionnaires received

A. Bortolami, I. Zambon and C. Terregino

EURL AI/ND

Istituto Zooprofilattico Sperimentale delle Venezie (IZSVe), Padua, Italy

Responding country

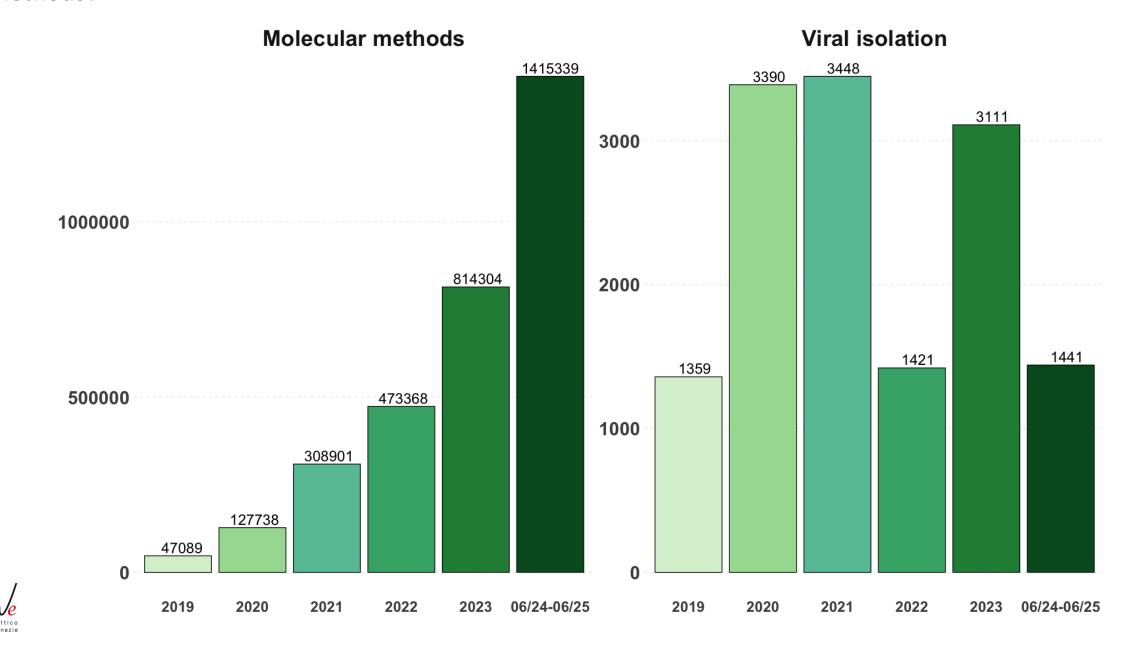


- Austria
- Belgium
- Bulgaria
- Croatia
- Cyprus
- Czech Republic
- Denmark
- Estonia
- Finland
- France
- Germany
- Greece
- Hungary
- Kosovo
- Latvia
- Lithuania

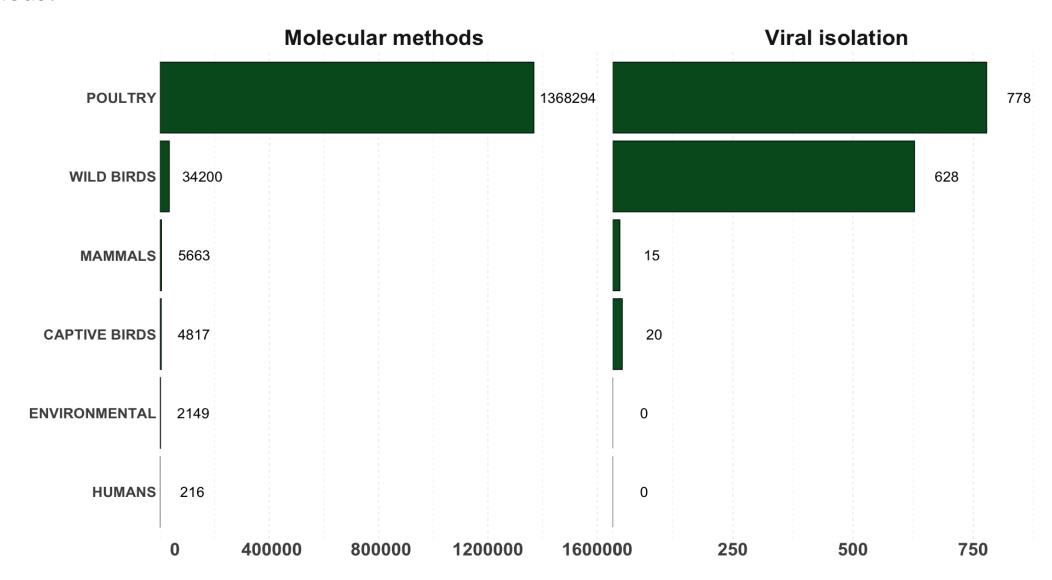
- Luxembourg
- Macedonia
- Moldova
- Montenegro
- Norway
- Poland
- Portugal
- Republic of Ireland
- Romania
- Slovakia
- Slovenia
- Spain
- Sweden
- Switzerland
- The Netherlands



1.1. How many samples were processed by molecular (RT-PCR or rRT-PCR) or virus isolation methods?



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1.2 Do you or any regional laboratory use commercial PCR kits for diagnostics? If yes,

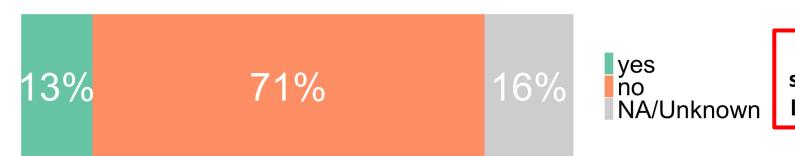
please provide specification of the products and if performance verification has been carried out.



Few NRLs adopting commercial kits, verification of performances obtained by participation to national or EURL PT schemes

1.4. Do you know if a rapid antigen test is used in your country (at field or laboratory level)?

If known, please provide details.



Use by OVS paired with samples submission (1 MS), use at laboratory level (1 MS), used in the field (1 MS)



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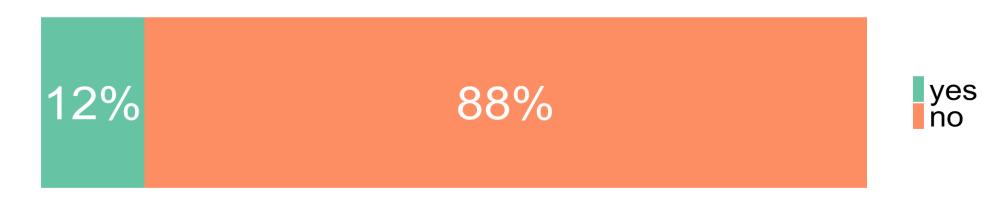
1.5. Do you perform rapid (real time RT-PCR-based) molecular pathotyping for H5 and/or H7 viruses? Please provide details.



All positive answers refers to H5 pathotyping by SOP VIR 1005, modified Naguib et al., 2017 or James et al., 2022

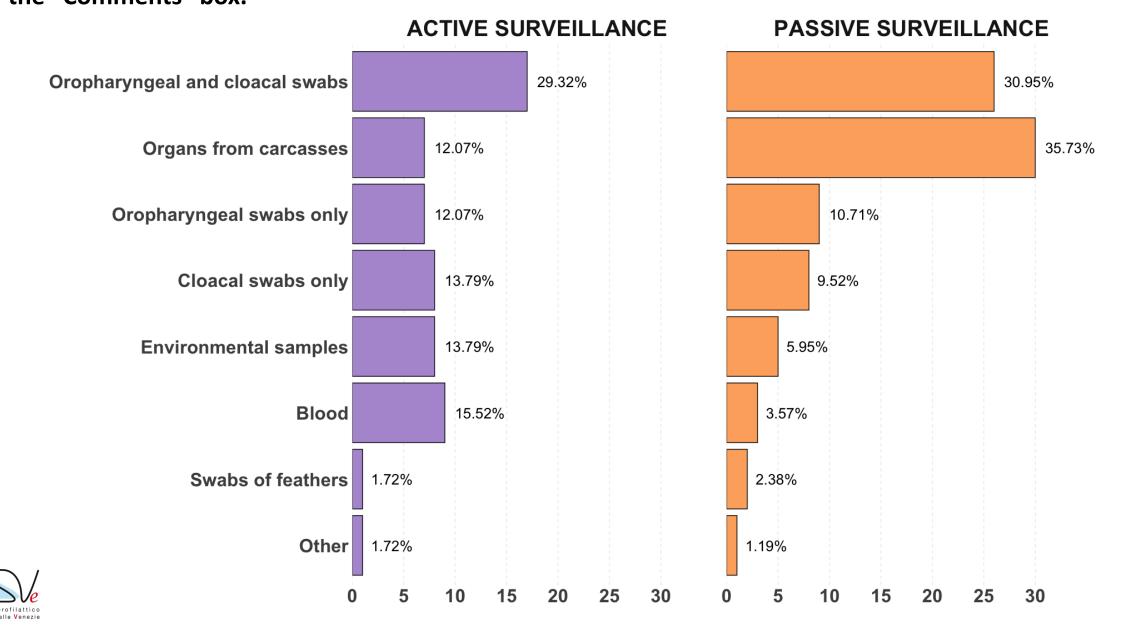


1.6. Have any changes occurred since last year in your access to Next Generation Sequencing (NGS) technology for influenza type A characterization (e.g. new instrument acquisition)? Yes/No. If your answer is yes, please provide details.



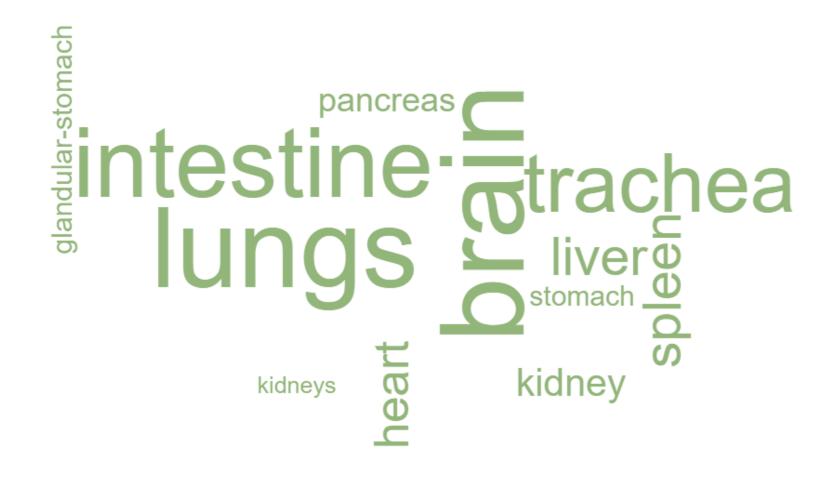
- All positive answers refer to testing or implementing Oxford Nanopore Technologies plc. instruments
- Most labs already implemented NGS in previous years, with no major changes





AI

1.7. Which type of samples from wild birds are submitted to the lab for testing from surveillance activities? Feel free to provide details on the tests performed on such samples in the "Comments" box.

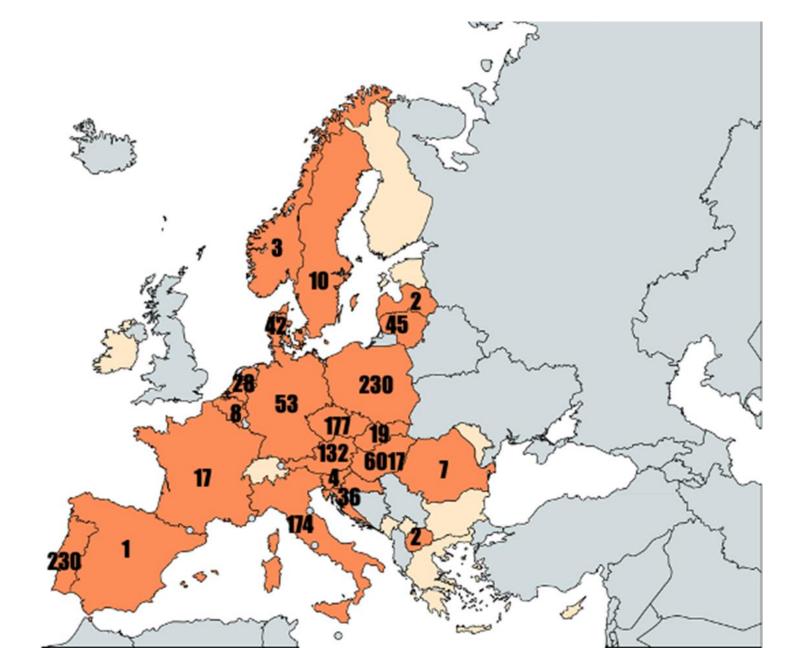




Avian influenza subtype detected in 2024/25 – H5 HPAI domestic

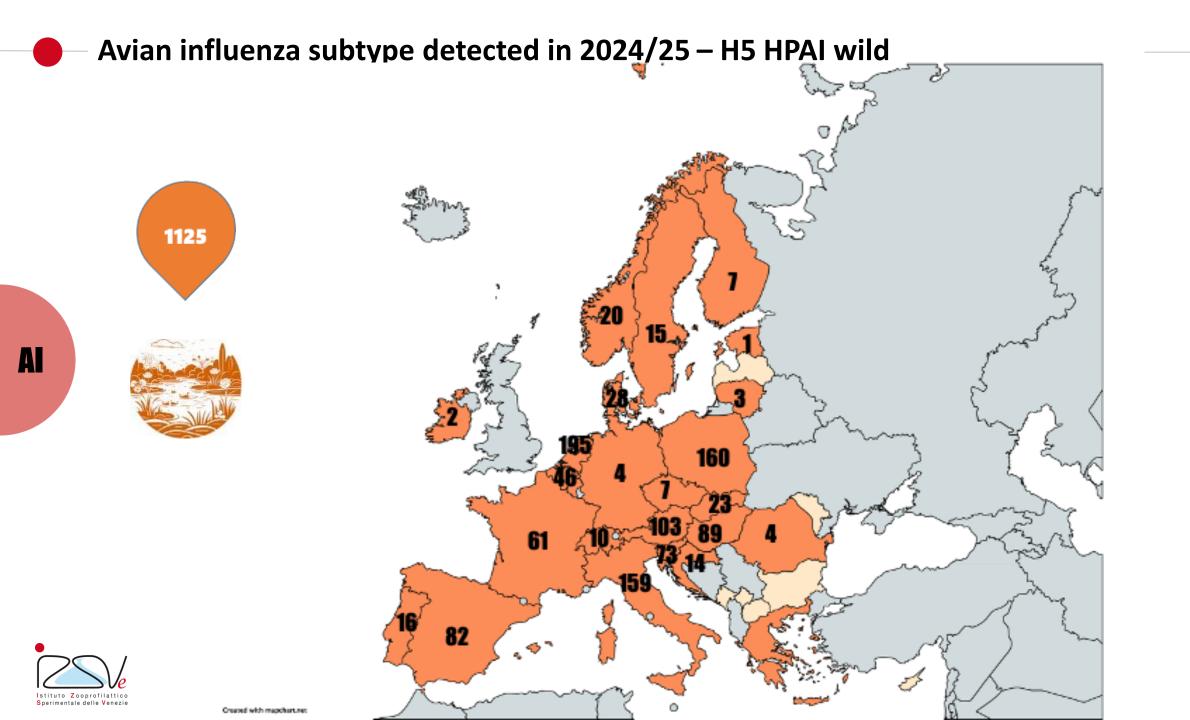




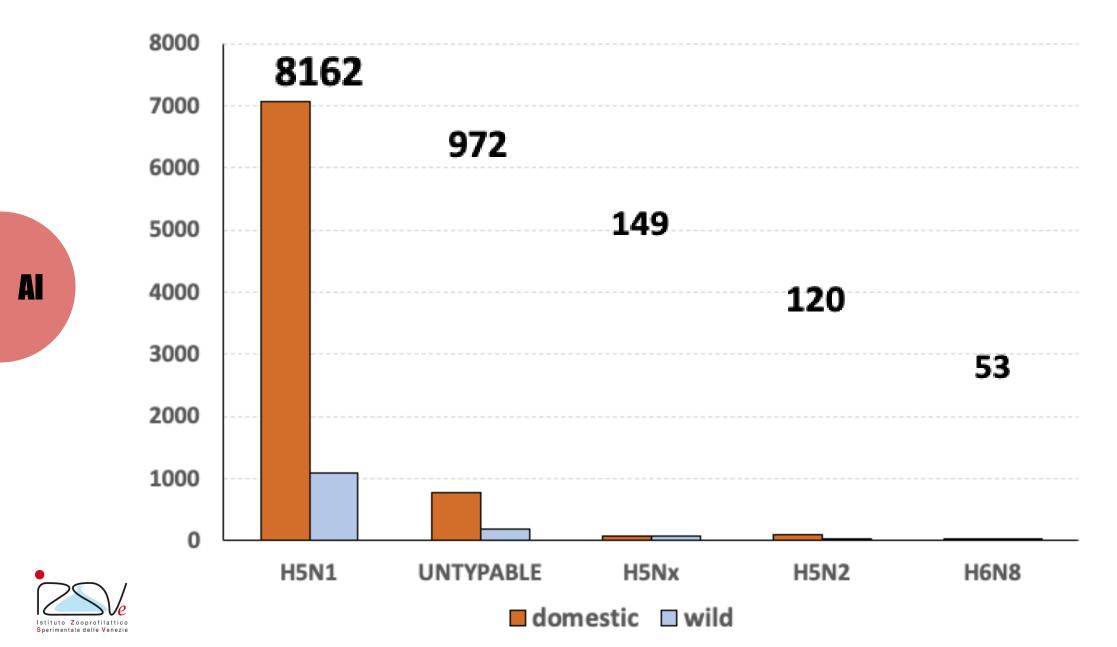


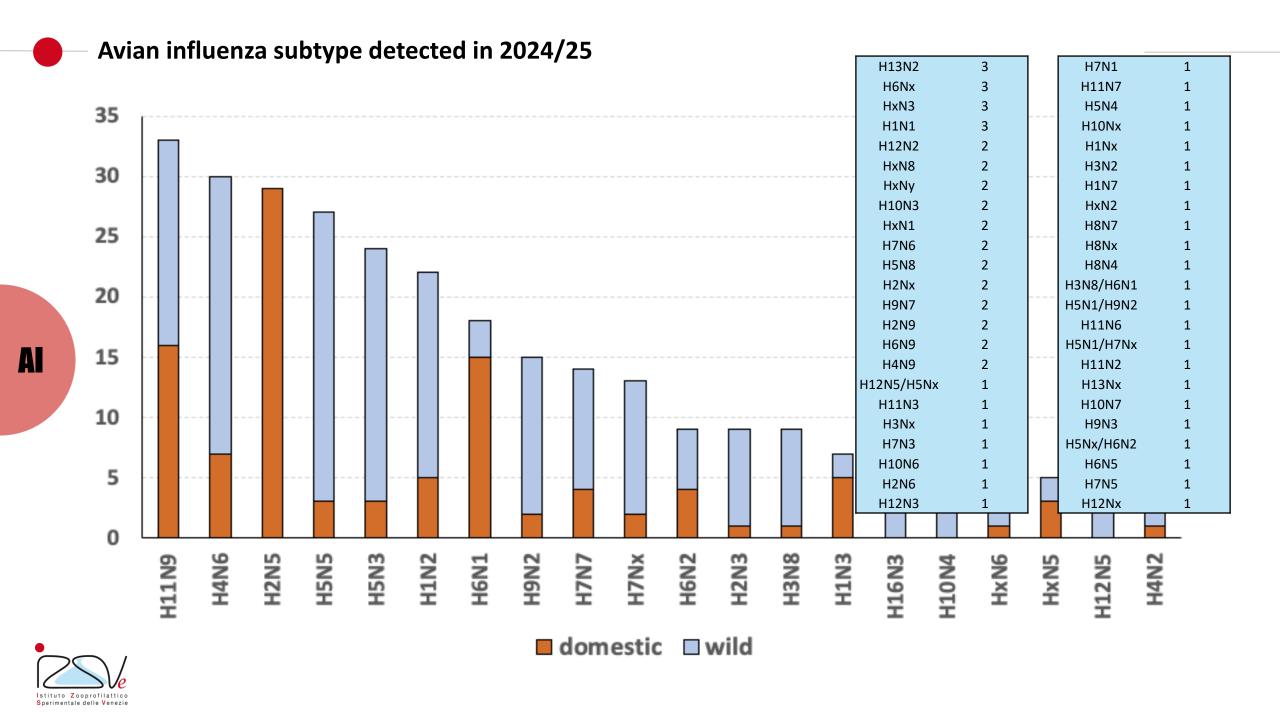


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Avian influenza subtype detected in 2024/25 – TOP 5



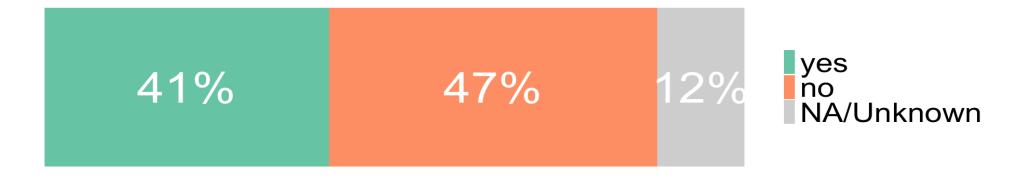


56% 44% both passive

- Mainly passive surveillance (annual program, country-wide)
- Active surveillance only in high-risk areas, periods, or specific bird populations







- Managed by public health authorities, mainly targeting exposed workers and close contacts
- No human cases detected in the reporting period





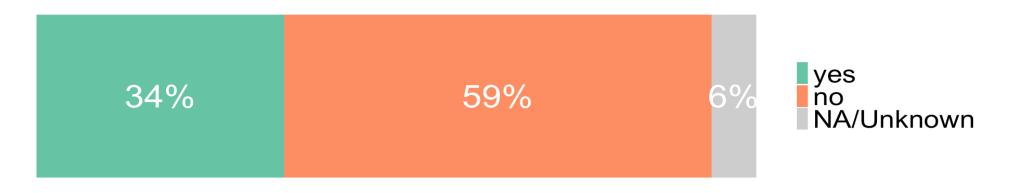
2.3. Were serological and/or virological investigations performed in domestic mammals with history of exposure to HPAI-positive birds in confirmed outbreaks? Yes/No. Please provide details



Jnknown



2.7. Did you perform or plan any surveillance (including retrospective) activities in dairy cattle and/or ruminants? Yes/No. If yes, please provide details

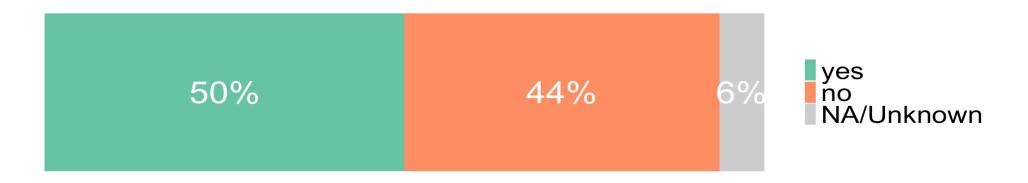


- Most activities involved ELISA testing (on serum or bulk milk samples), sometimes complemented by RT-qPCR.
- A few countries are drafting or planning surveillance schemes for ruminants in contact with infected poultry holdings.
- No positive results have been reported so far.





2.4. Were domestic or wild mammals tested for HPAI in case of active circulation of HPAIVs? Yes/No. If yes, please provide details



 Mainly wild mammals tested, via passive surveillance and the EUfunded One Health project; domestic mammals tested only in outbreaks

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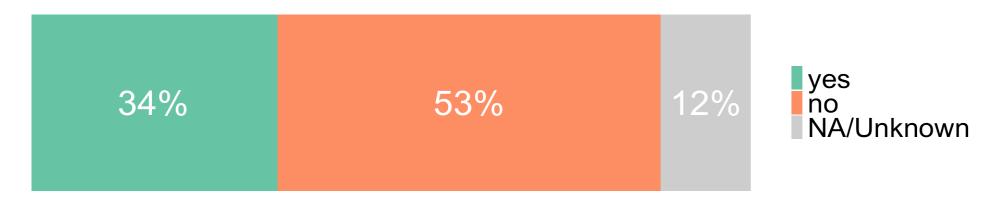
SCIENTIFIC OPINION

Preparedness, prevention and control related to zoonotic avian influenza





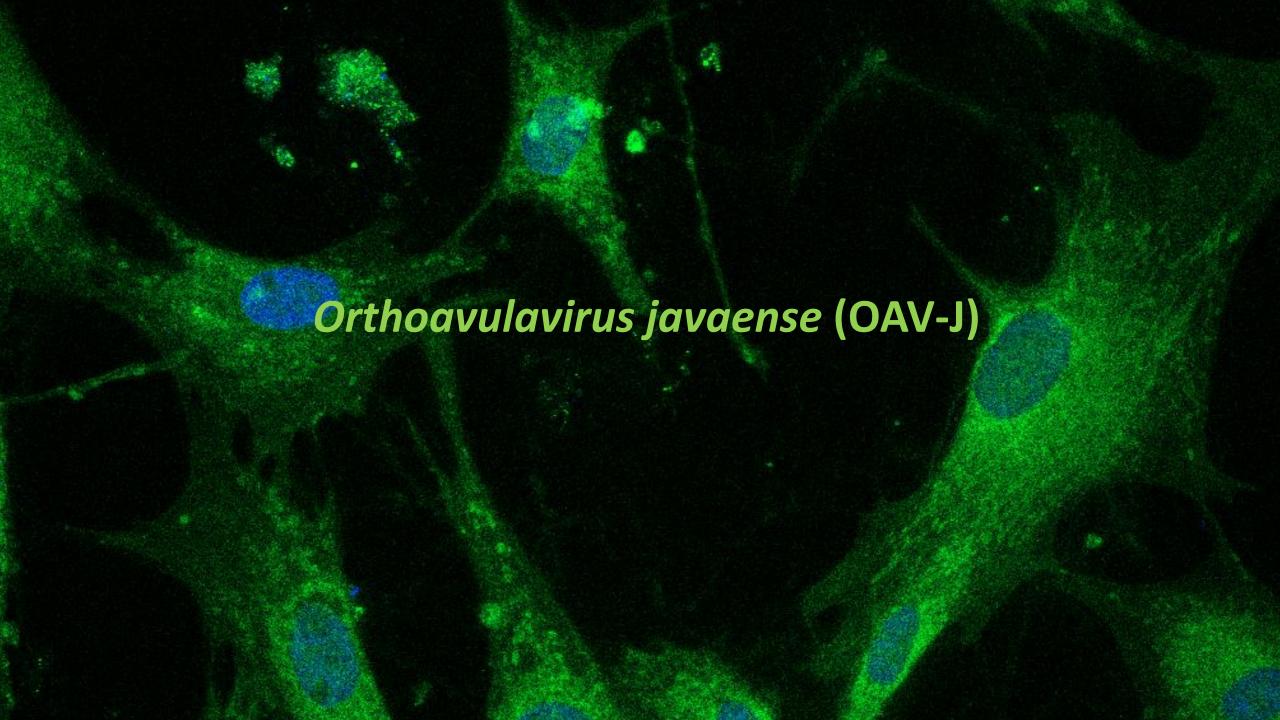
2.6. Did you perform active surveillance before the movement of poultry in case of active circulation of HPAIVs? If yes, please describe strategy of sampling applied



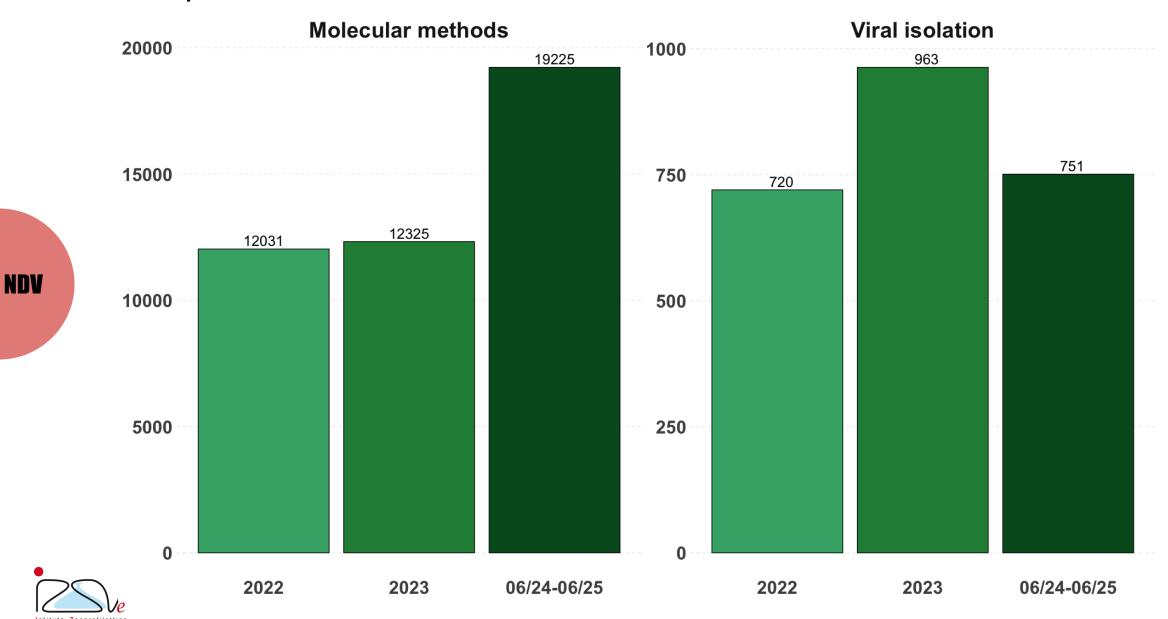
• tracheal/cloacal swabs or serum samples may be collected before movement.



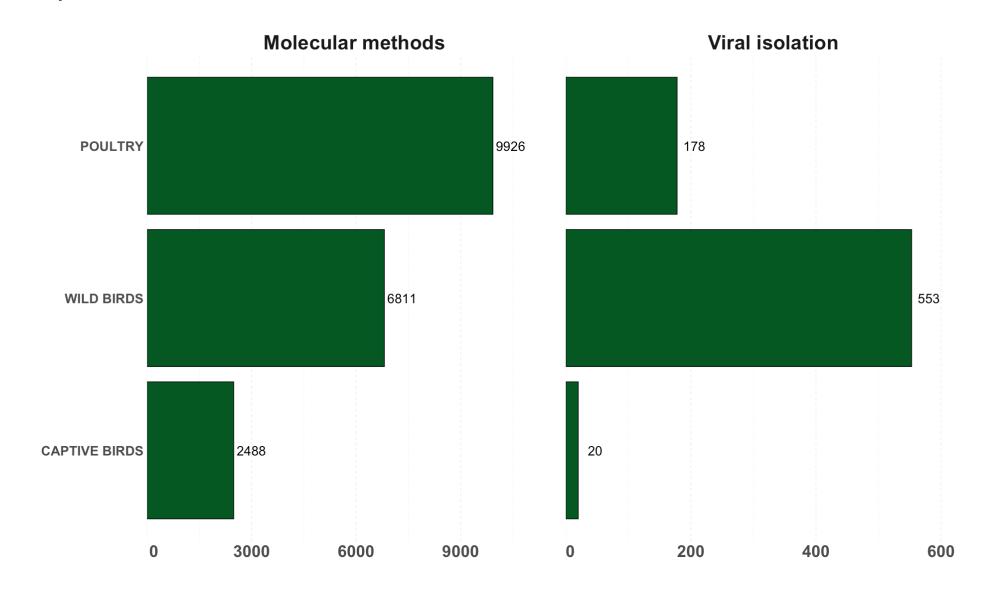




1.1 How many samples from poultry or wild birds were processed by molecular (RT-PCR or rRT-PCR) or virus isolation methods?

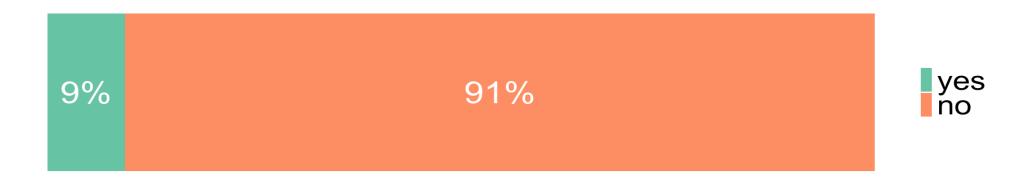


1.1 How many samples from poultry or wild birds were processed by molecular (RT-PCR or rRT-PCR) or virus isolation methods?





1.2 Do you or any regional laboratory use commercial PCR kits for diagnostics? If yes, please provide specification of the products and if and performance verification is carried out.



 Two NRLs reported internal or external use of commercial kits (e.g VetMAX™ NDV).



1.3 Do you perform serological testing for ND? If yes, please provide details on the methods and kits used.

75% 25% yes no

- Mainly to check vaccination titres.
- Methods: HI test and ELISA





1.4 Do you perform rapid (real time RT-PCR-based) molecular pathotyping for APMV-1? Yes/No. If yes, please provide details



Protocols: Fuller et. al. 2009; Sabre et al. 2017; SOP VIR 1006



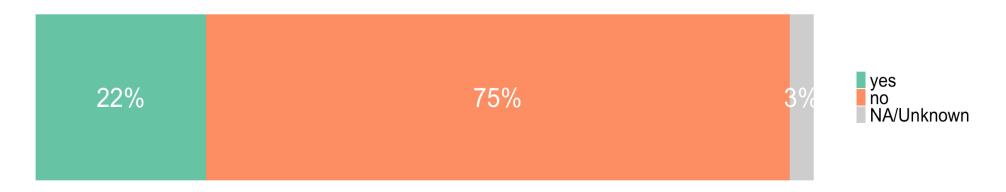
1.5 If an APMV-1 was detected, was sequencing of the F gene performed? Yes/No. If yes, please specify whether a partial or complete sequence of the F gene was performed and deposited



- Most laboratories perform partial sequencing of the F gene, mainly targeting the cleavage site.
- Some laboratories also perform full F gene sequencing or whole genome sequencing (WGS).



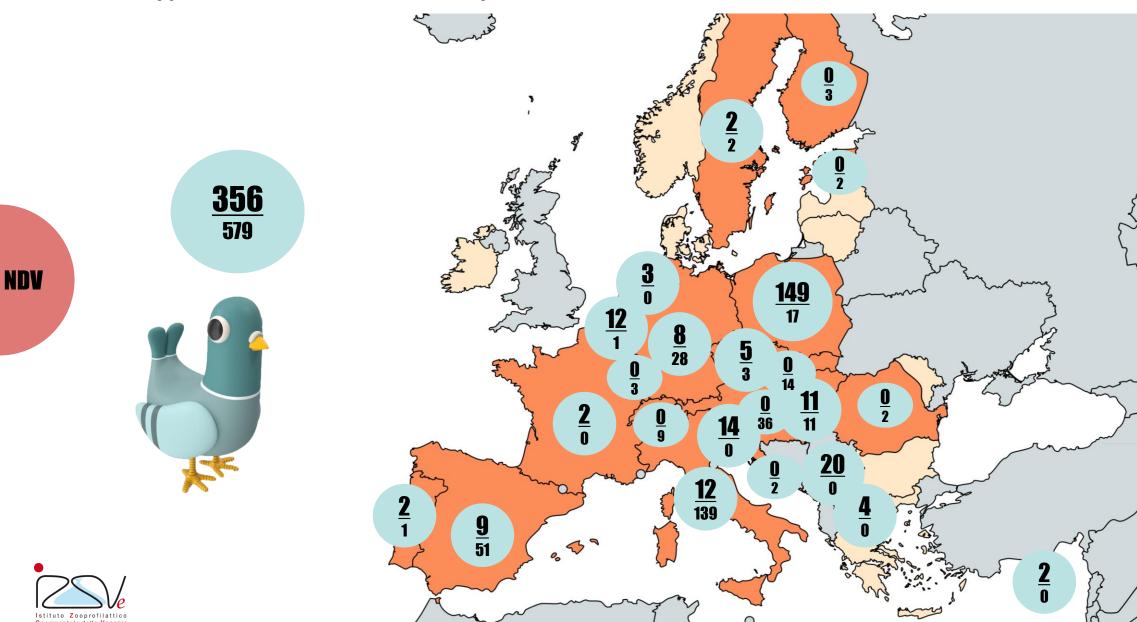
1.6 Did you perform sequencing also of other genes (other than F gene)? Yes/No. If yes, please provide details



- Some laboratories perform whole genome sequencing (WGS) or sequencing of other genes, mostly for selected isolates or as part of epidemiological investigations.
- Systematic WGS is not routine, and sequencing of additional genes is done only when viral RNA concentration allows it.



1.7 Indicate the number of APMV-1 isolated/detected, their serotype/genotype and the type of bird from which they were identified



OAV-J viruses detected in poultry (excluding pigeons) in 2024/25

Country	Species	ICPI	CS	Genotype	Patotype	Number of outbreaks
Belgium	chicken	nd	GRQGRL	n.a.	Avirulent	4
Czech Republic	chicken	nd	RRQKRF	VII.1.1	Virulent	5
Germany	poultry	nd	GRQGRL	2.11	Avirulent	2
Italy	chicken	nd	GRQGRL	I 1.1	Avirulent	2
Italy	chicken	nd	GRQGRL	I 1.2.2	Avirulent	3
Italy	chicken	nd	GRQGRL	1.2	Avirulent	1
Poland	chicken	na	RRQKRF	VII 1.1	Virulent	149
Republic Of North Macedonia	chicken	no	RQKRF	VII.2	Virulent	4
Slovenia	chicken	1,98	RRQKRF	VII.1.1	Virulent	2
Slovenia	chicken	nd	GKQGRL	I.1.1	Avirulent	1
Slovenia	chicken	nd	GRQGRL	II	Avirulent	5
Slovenia	turkey	nd	GRQGRL	II	Avirulent	1
Spain	mallard	no	GKQGRL	na	Avirulent	2
Spain	egyptian goose	no	GKQGRL	na	Avirulent	1
Sweden	chicken	no	RRQRRF	XIII.1.1	Virulent	1



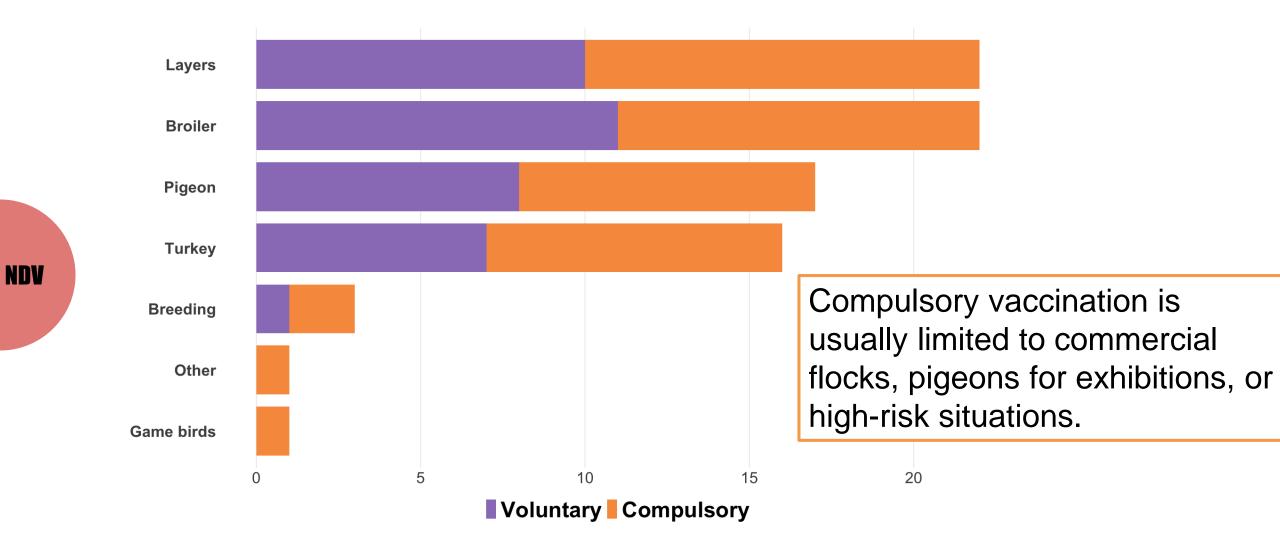
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Italy	chicken	nd	GRQGRL	I 1.1	Avirulent	2
Italy	chicken	nd	GRQGRL	I 1.2.2	Avirulent	3
Italy	chicken	nd	GRQGRL	1.2	Avirulent	1
Poland	chicken	na	RRQKRF	VII 1.1	Virulent	149
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Slovenia	chicken	nd	GKQGRL	I.1.1	Avirulent	1
Slovenia	chicken	nd	GRQGRL	II	Avirulent	5
Slovenia	turkey	nd	GRQGRL	II	Avirulent	1
Spain	mallard	no	GKQGRL	na	Avirulent	2
Spain	egyptian goose	no	GKQGRL	na	Avirulent	1
Sweden	chicken	no	RRQRRF	XIII.1.1	Virulent	1



umbiformes	wild wild wild domestic wild wild domestic wild wild wild wild wild wild wild wild	icpi no	KRRKR/FIG RRQKRF RRQKR*FIG KRQKR*FIG KRQKRF KRQKRF	genotype VI.1.1 (L4b**) XXI.1.1 (L4d**) XXI.1.1 (L4d**) PPMV-1 VI.2.1.1.2.2 VI.2.1.1.2.2 Not yet sequenced XXI.1.1 XXI.1.1 VI.2.1.1.2.2 VI.2.1.1.2.2 VI.2.1.1.2.1 na na VI.2.1.1.2.1 PPMV-1 XXI.1.1 PPMV-1 XXI.1.1 VI	patotype PPMV-1 virulent PPMV-1 virulent PPMV-1 virulent velogenic PPMV-1 PPMV-1 PPMV-1 PPMV-1 PPMV-1 PPMV-1 PPMV-1 virulent PPMV-1 virulent PPMV-1 virulent velogenic velogenic velogenic velogenic velogenic velogenic velogenic
umbiformes	wild wild domestic wild wild domestic wild wild wild wild wild wild wild wild	no no na no no na no	RRQKR/FIG KRRKR/FIG RRQKRF RRQKR*FIG KRQKR*FIG KRQKR*FIG KRQKR*FIG RRQKR*FIG RRQKRF RRQKRF RRQKRF RRQKRF SGGRRRKR/FIG GRRQKRFIG GRRQKRFIG GRRQKRFIG KRQKRF RRQKRF RRQKRF RRQKRF RRQKRF RRQKRF RRQKRF RRQKRF RRQKRF RRQKRF	XXI.1.1 (L4d**) XXI.1.1 (L4d**) PPMV-1 VI.2.1.1.2.2 VI.2.1.1.2.2 Not yet sequenced XXI.1.1 XXI.1.1 VI.2.1.1.2.2 VI.2.1.1.2.2 VI.2.1.1.2.1 na na VI.2.1.1.2.1 PPMV-1 XXI.1.1 YI.XII.1 VI.XII.1 PPMV-1 XXI.1.1 VI.XII.1 PPMV-1 XXI	PPMV-1 virulent PPMV-1 virulent velogenic PPMV-1 PPMV-1 velogenic PPMV-1 PPMV-1 PPMV-1 PPMV-1 virulent PPMV-1 virulent Mesogenic/Velogenic na na Velogenic velogenic velogenic velogenic Velogenic Velogenic
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umbiformes umbiformes	wild			na	Velogenic
umbiformes			RRQKRF	VI 2.1.1.2.2	velogenic PPMV-1
	WIIU	no	RRQKRF	VI 2.1.1.2.2 VI 2.1.1.2.2	velogenic PPMV-1
annonnes	wild	no	RRQKRF		_
umbiformes				na PRAM/1	na
		na	RRQKRF	PPMV1	velogenic
		nd	KRQKRF	na	virulent
		nd	RERRQKR*FVG	na	na
		nd	RGRRQKR*FVG	na	na
			na	pigeon paramyxovirus-1	na
					velogenic
			KRQKRF	XXI.1.1	velogenic
		na			
	domestic	na			Velogenic
	domestic	na			Velogenic
umbiformes	domestic	no	RRQKR*FIG	VI.2.1.1.2.2	PPMV-1
umbiformes	domestic	no	KRQKR*FIG	XXI.1.1	PPMV-1
umbiformes	domestic	na	na	pigeon paramyxovirus-2	na
umbiformes	domestic			VI.2.1.1.2.2	Velogenic
umbiformes	domestic	no	RRQKRF	VI.2.1.1.2.2	Velogenic
			RRQKRF	VI.2.1.1.2.2	velogenic
	domestic	no		VI	Velogenic
				VI.1.2.1.2	Velogenic
					Velogenic
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2.2 If vaccination is performed in your country, please provide releavant details





Thank you for your contribution!

