

Pandemic prevention Pandemic preparedness what's in the name

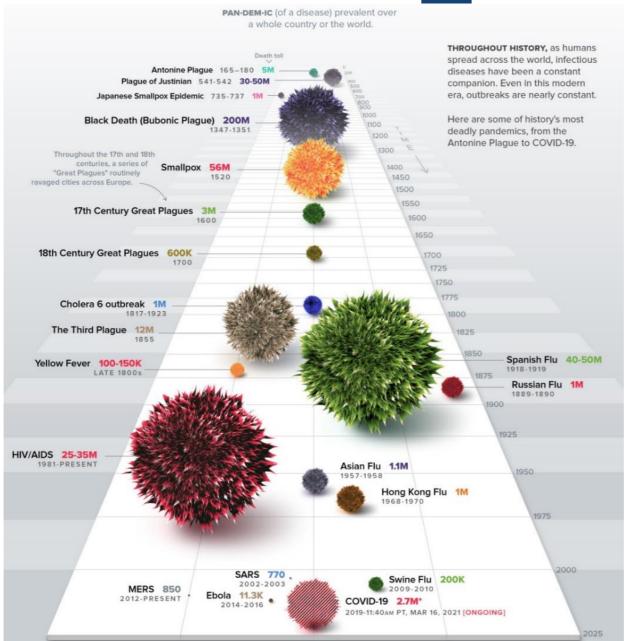
Joke van der Giessen

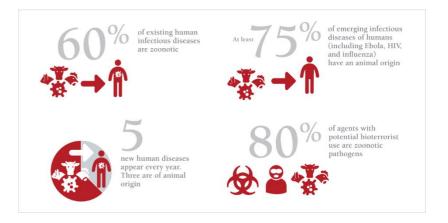
Centrer for Zoonosen & Omgevingsmicrobiologie

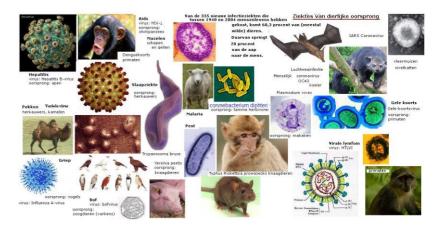
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History of pandemics and role of zoonoses











Zoonoses are priority

Drivers of emergence

Microbiological factors:

Genetic evolution

Human-made factors:

Globalisation

Population growth & urbanisation

Deforestation

Increased animal-human contacts

Socio-economic factors

Climate change

Changing food habits

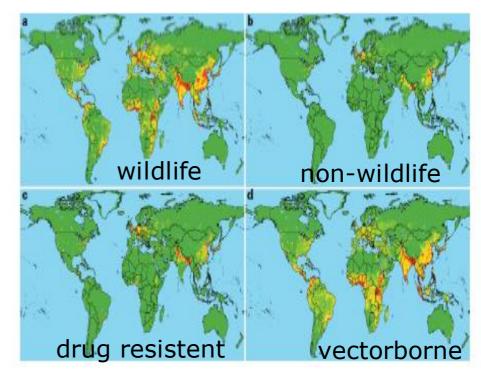


Figure 3 | Global distribution of relative risk of an EID event. Maps are derived for EID events caused by a, zoonotic pathogens from wildlife, b, zoonotic pathogens from non-wildlife, c, drug-resistant pathogens and d, vector-borne pathogens. The relative risk is calculated from regression coefficients and variable values in Table 1 (omitting the variable measuring reporting effort), categorized by standard deviations from the mean and mapped on a linear scale from green (lower values) to red (higher values).

Jones et al., 2008)



Content

Development of the zoonoses disease structure in the Netherlands

Pandemic Prevention and Preparedness plans for Zoonoses

Strengthen Zoonoses from signalling to response

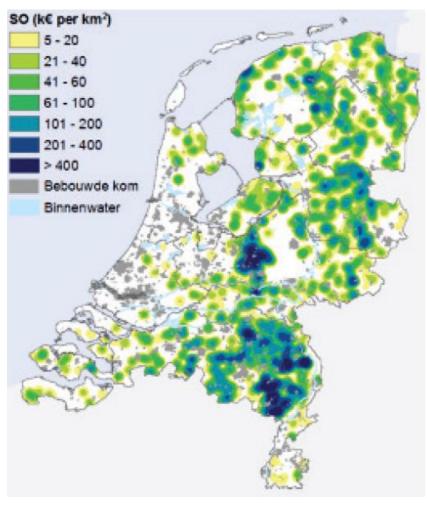
Nine themes and some ongoing results

Example OH surveillance swine influenza

Conclusion



Netherlands: densely populated, intensive livestock and wildlife



- 33,883 km²
- (CBS 2023):
 - 17,8 million habitants
 - 88 million poultry
 - 6,2 million pigs
 - 3,8 million cattle
 - 0,8 million sheep
 - 0,6 million goats

• Wildlife:

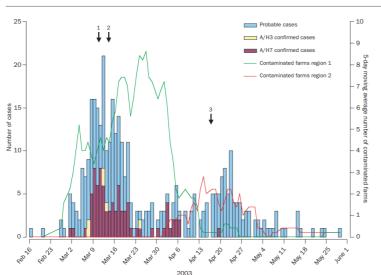
- Migratory and water birds
- Wild mammals: fox, wolf, raccoon dogs, roe deer, rodents, bats, wild boar

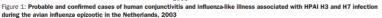




2003: Avian Influenza H7N7 Netherlands







Green and red lines show 5 day moving average of newly diagnosed A/H7 contaminated farms in two regions. Arrows show start of active case finding (1), start of prophylactic treatment with oseltamivir (2) and death of case with HPAI infection (3).





Emerging zoonoses

2004: Advice Health council and WHO Emerging zoonoses



2007-2010: Emerging zoonoses (EmZoo) national programme

- 1. Surveillance animal and human populations: advice on gaps
- 2. Prioritise emerging zoonoses: what's next
- 3. Blueprint for effective One Health collaborative infrastructure: advice





Multicriteria decision analyses: Higher ranked: more impact

1. Probability of introduction

2. Transmission animal reservoirs

3. Economic damage (animal reservoir)

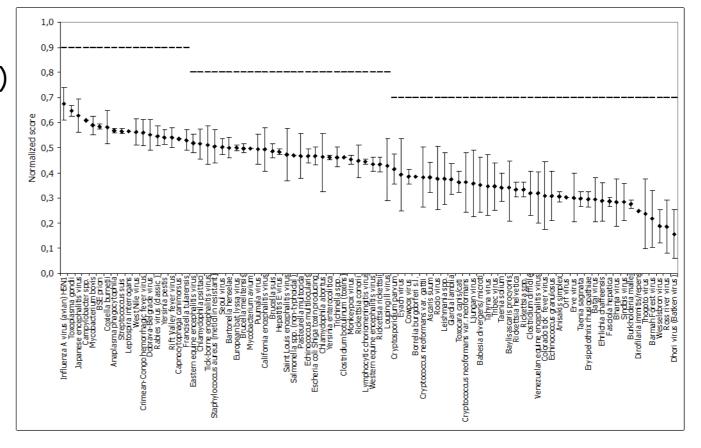
4. Animal-human transmission

5. Transmission between human

6. Morbidity in humans

7. Mortality in humans

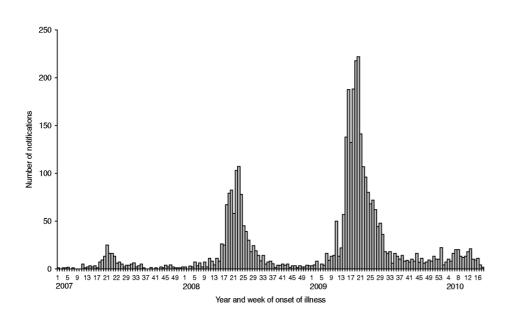
Via livestock 44% 46% Via vectors: Via wildlife: 85% 29% Via companion animals:

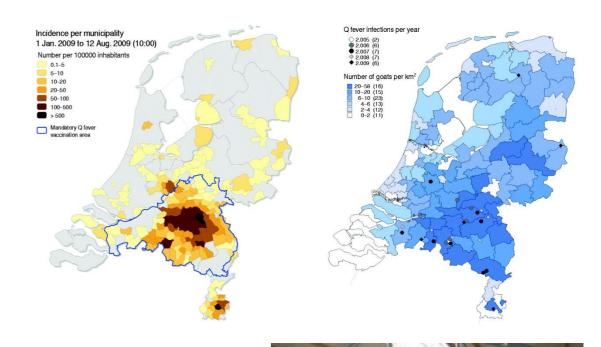


(EmZoo lijst Havelaar et al., Plos One, 2011)



2007-2009: Q-fever outbreak Netherlands





ROEST HIJ, TILBURG JJHC, VAN DER HOEK W, et al. The Q fever epidemic in The Netherlands: history, onset, response and reflection. Epidemiology and Infection. 2011;139(1):1-12.

doi:10.1017/S0950268810002268







Recommendations to improve signaling and control of emerging zoonoses: the One Health approach

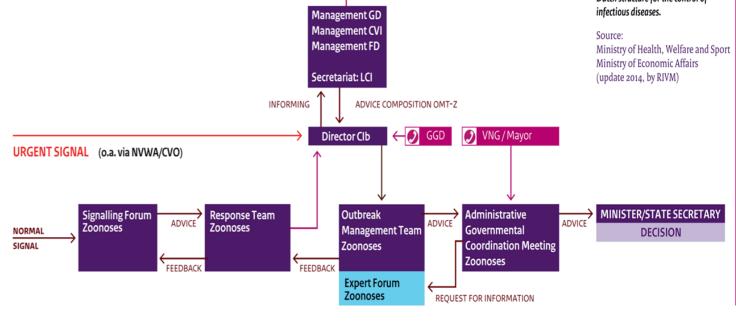


> Implementation medical-veterinary integrated risk analyses structure of zoonoses in 2011

Zoonotic disease structure in the Netherlands from signalling to decision-making







- National Institute of Public Health and the Environment (RIVM), Bilthoven
- Public Health Service, of Area Health Authority (GGD)

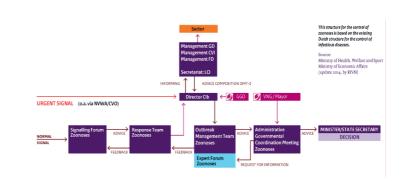
- Netherlands Food and Consumer Product Safety Authority-NVWA
- Wageningen Bioveterinary Research, Lelystad
- Faculty of Veterinary Medicine, Utrecht University, Utrecht
- Dutch Wildlife Health Centre (DWHC), Utrecht
- GD Animal Health (GD), Deventer
- Center Monitoring Vectors-NVWA



OMT-Z during Sars-CoV-2 in minks

- April 2020: 1st notice Sars-CoV-2 in sick/dead minks GD, Deventer
- Direct to CVO ministery LNV
- Confirmation WBVR in Lelystad
- Notification and measurements
- Screening minks and early warning system GD
- New farms infected > Request CVO/ministries OMT-Z risk public health in June en July 2020. Advice OMT-Z
- BAO > Ministries > august 2020 production stop before 2021 en control infected farms
- January 2021: mink production ended





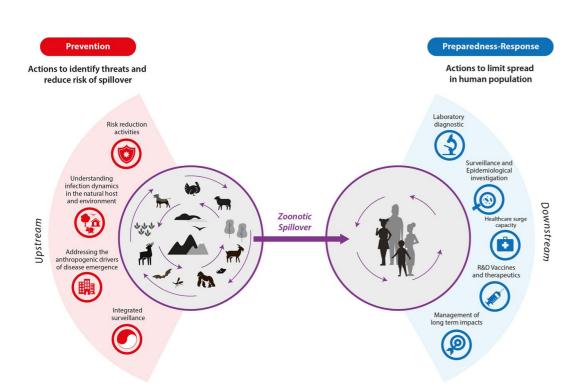




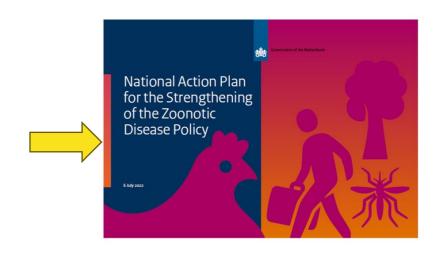
2021 Zoonoses insight and 2022 National Action plan

Report Zoonoses insight

VWS en LNV juli 2022







Prevention of zoonotic spillover: From relying on response to reducing the risk at source

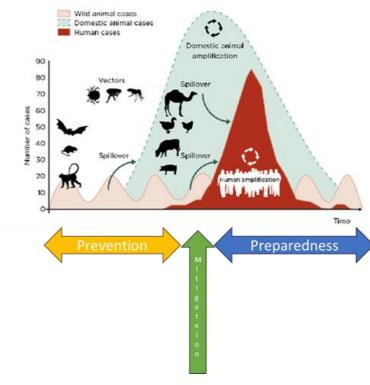
Authored by the members of the One Health High-Level Expert Panel (OHHLEP) . Wanda Markotter . Thomas C. Mettenletter . Wiku B. Adisasmiko, Salama Almuhairi, Casey Barton Behravesh, Pépa Bikrogui, Salome A. Bukachi, Natalia Cases, Natalia Cedies Becerra, Deninique F. Charron, Abhishek Chaudhary, Janice R. Clacci Zanella, Andrew A. Cunningham, Osman Dar, Nitish Debnath, Baptiste Dungu, Elmoubasher Farag, George F. Gao, David T. S. Hayman, Margaret Khaitsa, Marion P. G. Koopmans, Catherine Machalaba, John S. Mackenzie, Serge Morand, Vyacheslav Smolensky, Lei Zhou [Vew less]



Pandemic Prevention & Preparedness 2022-2026

Human Pandemic Preparedness

- Strengthen diagnostic landscape
- Strengthen surveillance landscape
- Strengthen supra-regional collaboration
- Pandemic Prevention
- Strengthen Zoonoses from signalling to response (Zoover)







Strengthen Zoonoses from signalling to response Zoover programme 2022-2026: 9 themes

1H-signaling 1H surveillance

1H datasharing/ICT

Prioritising Zoonoses
Innovation/research
International
Respons

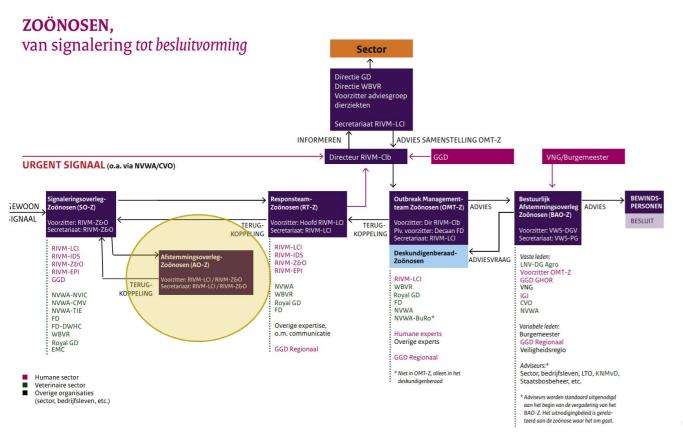
Environment Zoonoses literacy



Zoover: improve 1Health signalling

2022: Evaluation 2011-2021

2023: Recommendations 2024



PERSPECTIVE

Signalling and responding to zoonotic threats using a One Health approach: a decade of the Zoonoses Structure in the Netherlands, 2011 to 2021

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Article submitted on 30 Dec 2021 / accepted on 27 May 2022 / published on 04 Aug 2022

Zoonoses structure including the AO-Z (Avian influenza in cattle USA; psittacosis; alveolar echinococcosis; swine influenza)

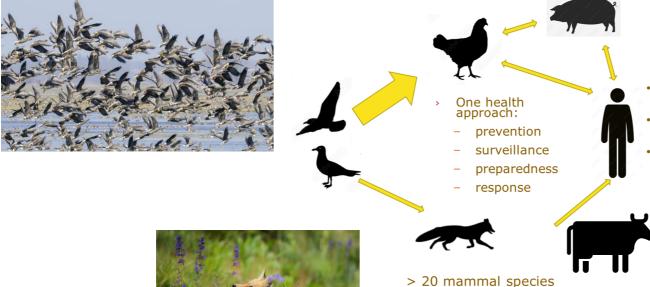
Development of Horizon scanning to focus on drivers of emergence >25 years

Avian influenza (H5N1)





affected



HPAI virus detections in wild bird categories 2023 - 15 March 2024

Colony-breeding seabird species (34) **Mixed* (41) **Others (27) **Raptors (53) **Waterfowl (296)

**Waterfowl (296)

**Face of the Version of t

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

sporadic human infections

- public health risk in the EU is low
- risk is low to moderate for occupationally or otherwise exposed people

March 2024: Avian influenza in dairy cattle and goats USA Signalling forum: risk profile to ministeries

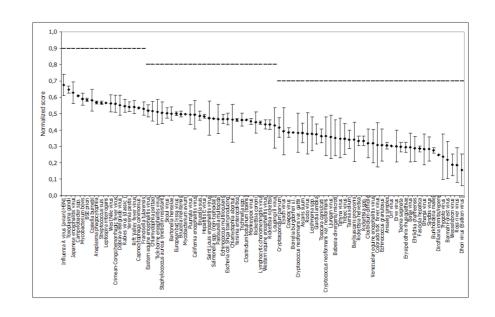


Zoover: update prioritizing list zoonotic pathogens

Multicriteria decision analyses (MCDA):

- 1. Probability of introduction
- 2. Transmission animal reservoirs
- 3. Economic damage (animal reservoir)
- 4. Animal-human transmission
- 5. Transmission between human
- 6. Morbidity in humans
- 7. Mortality in humans

Implement other 1H criteria: ??



Scoring and weighting criteria for new list of 81 zoonotic pathogens



Zoover: 1Health surveillance systems/datasharing

Development One Health surveillance surveillance systems:

Psittacosis

Porcine influenza

HEV

West Nile

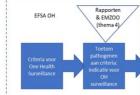
Foodborne pathogens

Sewage

Legionella

Criteria for new OH surveillance systems:







Disease X: syndrome surveillance and lab prepredness

Identify list of pathogens (EMZOO)

Framework building

Identify criteria for indication OH surveillance

Expert opinion SURVEY I

Expert opinion SURVEY II

Score pathogens against the criteria

MCDA modelling

Rank pathogens for indication OH surveillance

MCDA



Example One Health surveillance porcine influenza

Aim: implement surveillance of influenza A viruses in pigs with clinical samples collected by veterinarians 2022-2023 and comaper with human influenza A viruses.

Design: sentinel veterinary network collecting 200 pooled saliva and nose swabs tested by PCR. PCR +: cell culture, WGS, antiviral susceptibility, antigenic characterization

Set up a joint online database/platform for the sequences and a small set of contextual data.

Add a set of public reference sequences from humans and pigs as background for the molecular analysis.











Results OH surveillance porcine influenza

90 pig farms included national distributed north-middle-south part 57.9% of the saliva samples was PCR positive 28.5% of the nose swabs were PCR positive

Influenza subtype	Subtype
H1N1	74
H1N2	51
H3N2	1
Onbekend (geen HA sequentie verkregen)	3
Totaal	129

H1viruses different clades:

Clade 1A.3.3.2 (Classical Swine lineage)

Clade 1C.2.1 en 1C.2.2 (Eurasian Avian lineage)

Clade 1B.1.2.1 (Humane Seasonal lineage)

H3 virus close related circulating in Germany in pigs in 2015-2016











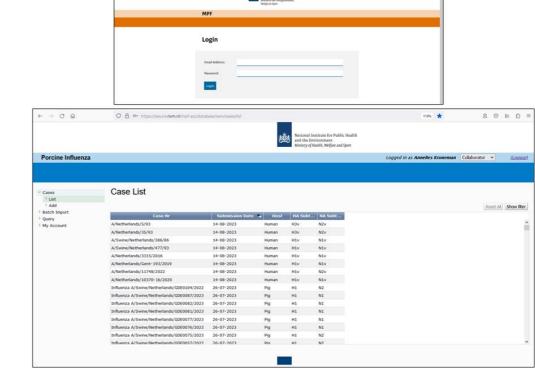
Online database/platform

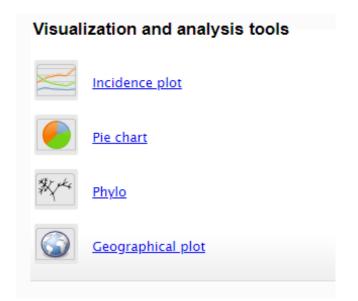
Host
Sampling date
Region (North/Middle/South)

Typing tool: HA and NA subtype, internal segment lineage

Project sequences: 126 porcine strains

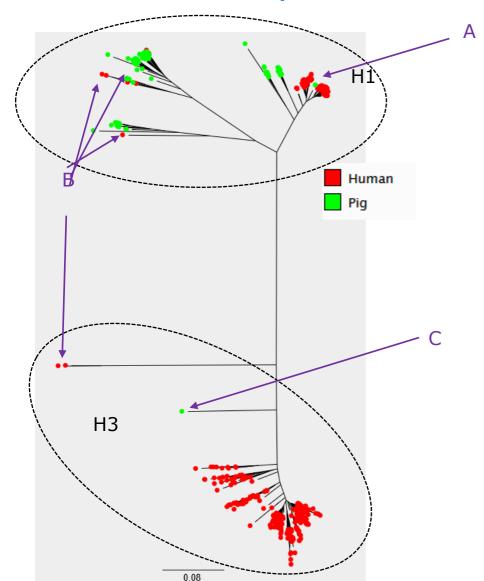
Reference sequences (public):
Porcine vaccine strains
Recent Dutch human strains H1N1 and H3N2
10 Dutch strains of known infections of humans
with porcine strains (1993-2023)
Porcine influenza strains from other countries







Swine HA sequences compared to human reference sequences



The diversity within porcine sequences is larger than within recent human H1 strains

- A: Infection of a pig with a human influenza A(H1N1) pdm09 virus
- B: Strains of humans infected with porcine influenza
- C: The one porcine H3N2 strain found during the project.



Antigenic and antiviral tests

- Not all porcine strains could be antigenically characterized with the available sera
- No resistance was found against Oseltamivir or Zanamivir











Next steps OH surveillance porcine influenza

- Structural One Health surveillance porcine influenza

 Combining online platform with Strong1H (EU4H) project on Avian Influenza in (wild) mammals and outdoor pigs

- In depth phylogenetic analyses, reassortment analysis

Strengthening OH Zoonoses Strong1 H

One Health surveillance in animals and the environment

Direct grant EU4H single application

The Netherlands

- Update of the sera needed for antigenic characterization











Response to zoonotic threats

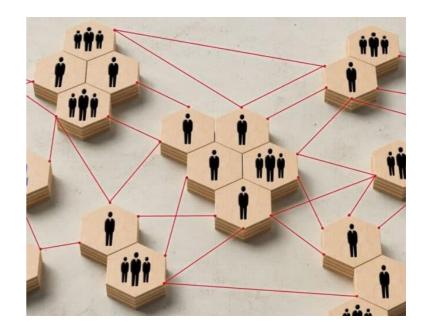
- Timely response is important, especially for zoonotic diseases with pandemic potential: One Health approach
- Lessons learned after a simulation exercise of HPAI outbreak.
- Needs between domains :
 - Strong networks
 - Regular contacts in a quite phase
 - Timely sharing of essential data
 - Cooordinated communication strategy





Zoover: Strengthening response 2023-2026

- Stimulate partnerships and network activities between the different domains at regional and national level
- > Initiate training and simulation exercises at regional and national level to strengthening the response to zoonotic threats
- Improve the link between national and regional level, stimulate knowledge sharing and cooperation
- Renew national response guidelines,
 Vademecum zoonosis and cooperation agreements

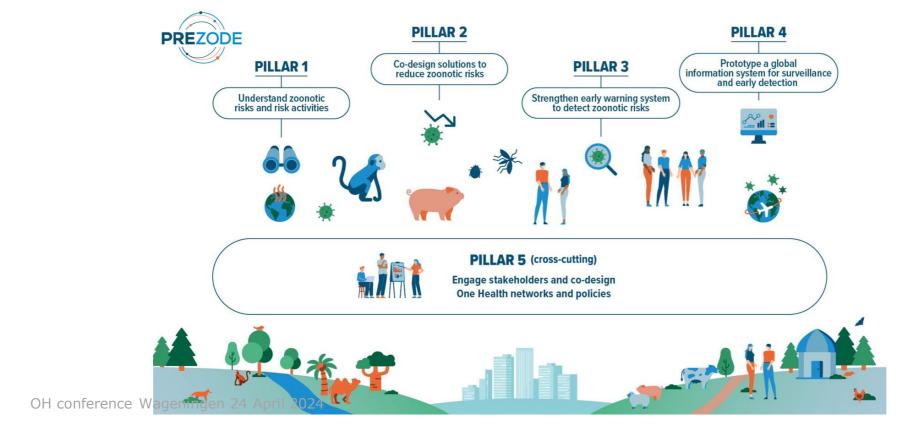


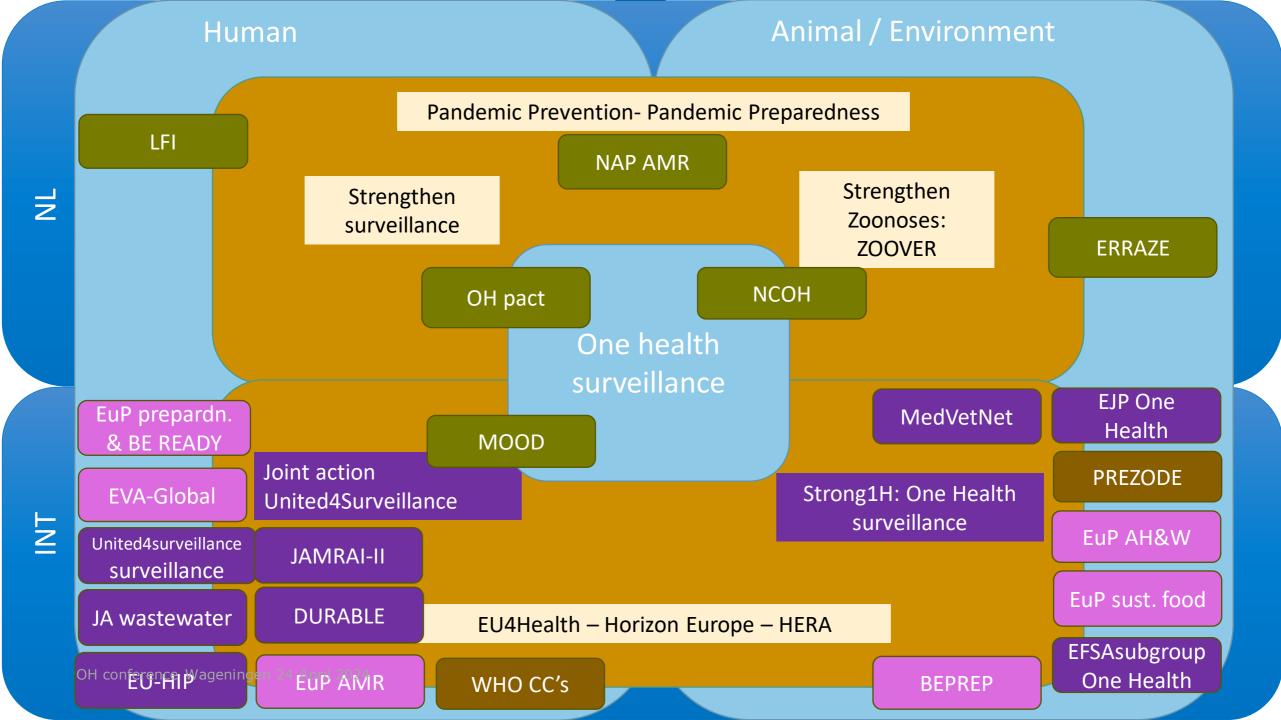
Zoover: international



Global initiatives to prevent Emerging Zoonoses

Prezode: preventing zoonotic disease emergence (RIVM/WBVR)







- Zoonoses diseases structure in the Netherlands 2011 formalised
- Zoover: to improve zoonoses from signalling to response
 - Develop One Health surveillance systems to share and jointly assess risk of zoonoses
 - Data sharing between domains complicated, One Health legal expertise needed
- Pandemic prevention and preparedness both relevant
- International collaborations important, challenge to combine all collaborations

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- PhD students: Frits Vlaanderen, Laura Derks, Sara Schneeman, Petro Mnyagatwa
- And many others



One Health surveillance swine influenza:



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