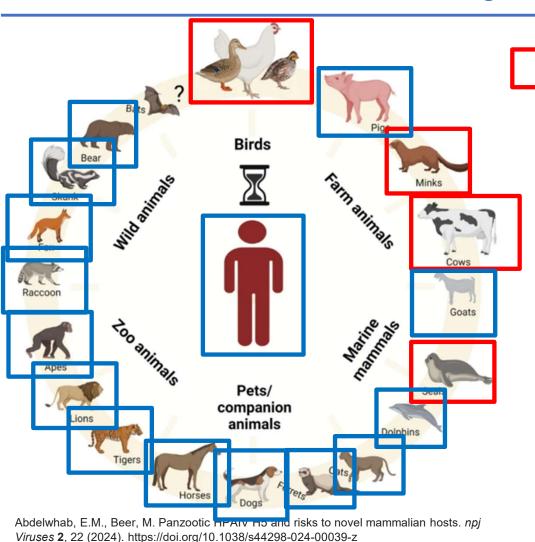


HPAIV H5N1 2.3.4.4b - broad host range - spill-over - zoonotic risks



Circulation



"dead-end"-host

Research questions:

Species-Specific Susceptibility and Pathogenesis

Immune Response (adaptive/innate) and Inflammation

Zoonotic Risks and Transmission Models

Testing of Vaccines and Antivirals

Viral Evolution and Host Adaptation

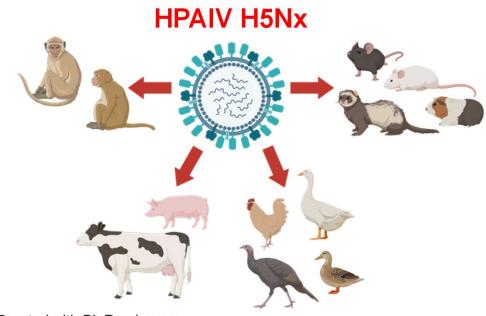
Comparative Models

Chronic and Long-Term Effects

Ethical and Practical Considerations



HPAIV H5N1 2.3.4.4b animal models – selection of key examples



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Pathogenesis in target animals:

- Chicken/ducks (lethal infection)
- Ferrets (high susceptibility)
- Pigs (low susceptibility)
- Cows (replication in the mammary gland)

Zoonotic potential:

- MxA mouse model
- Ferrets (cave: encephalitis!)
- Non-human primates

Transmission models (direct/aerosol):

- Ferrets
- Guinea pigs
- (Pigs)

Vaccines/antivirals/cross immunity:

- Mouse models
- Ferrets
- Non-human primates
- (Pigs)



H5N1 clade 2.3.4.4b: Key results - research needs/gaps (1)



- ➤ <u>General:</u> **Genotype differences** in virulence and transmission (more than 200 genotypes/reassortants of clade 2.3.4.4b!)
- Ferrets: Moderate to fatal disease, no or inefficient aerosol transmission
 Protective cross immunity shown after H1N1pdm infection
- ➤ Human MxA mice: 2.3.4.4b viruses show no escape from MxA response
- > Pigs: Low susceptibility, low level replication (dose/route dependent), no transmission
- Calves/cows: Intranasal infection inefficient, no nasal spread; highly efficient replication in the mammary gland, high level milk shedding
- ➤ Non-human primates: Mild to severe disease (dose-dependent, route of infection)
- Vaccine testing: Successful in mice, ferrets, non-human primates

H5N1 clade 2.3.4.4b: Key results - research needs/gaps (2)



- > Harmonization e.g. of inoculation titers, inoculation routes or group size
- > Pathogenesis data in different in vivo models
- Capacity to test the large number of genotypes / reassortants
- ➤ Need for **alternative in vitro methods** (e.g. organoid cultures) to allow improved preselection (following 3R principles)
- Correlates to determine zoonotic/pre-pandemic potential (improved risk analysis/prediction; potential of possible ressortants, GoF)
- ➤ More data on **genotype-phenotype** association
- Analysis of neurotropism (birds, carnivores) and exceptions (humans, cows)
- > Susceptibility of other ruminant species (goats, sheep, deer)
- Factors associated with low susceptibility in pigs
- > Barriers to efficient respiratory replication and transmission (e.g. in cattle)
- > Availability of field data and linking to trial data (e.g. missing field data from cows)

Thank you for your attention!

