

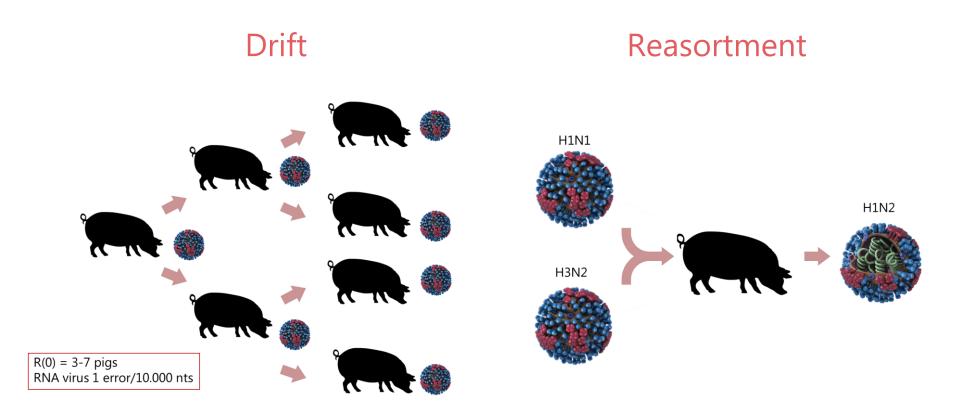
Ten years of swine influenza surveillance: consequences of viral evolution

Pia Ryt-Hansen IVH: Enzootic and zoonotic virus research CPH pig 27-01-2022

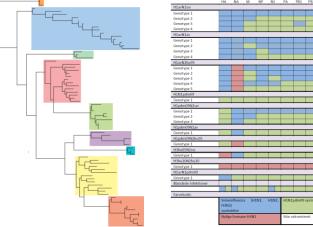
UNIVERSITY OF COPENHAGEN



Swine influenza A virus evolution



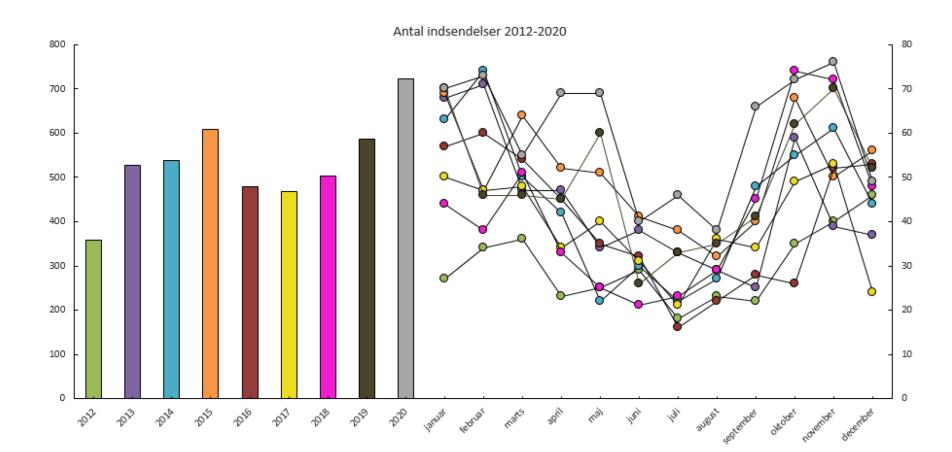
Current swIAV surveillance program



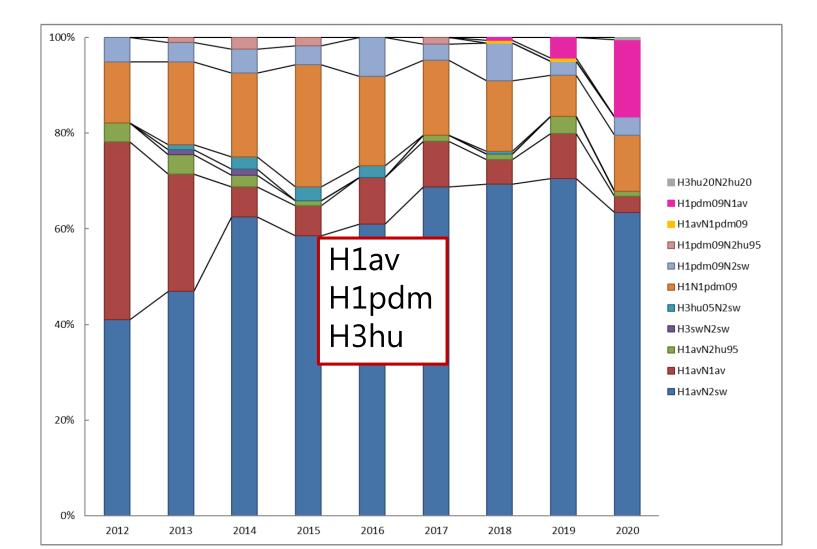


Results of the swIAV surveillance program - submissions

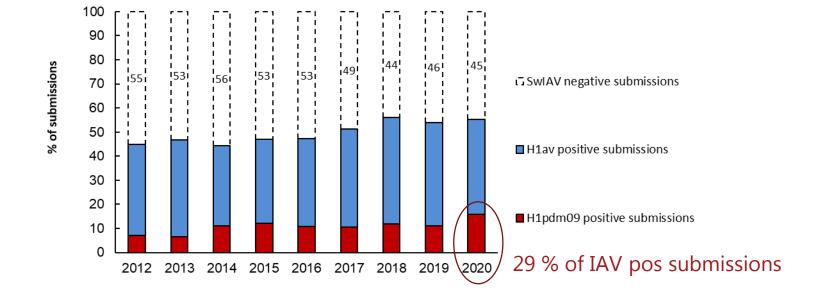
2020: 2512 samples from 517 herds (18 % of Danish herds)



Results of the swIAV surveillance program - subtypes



Results of the swIAV surveillance program – H1pdmNx



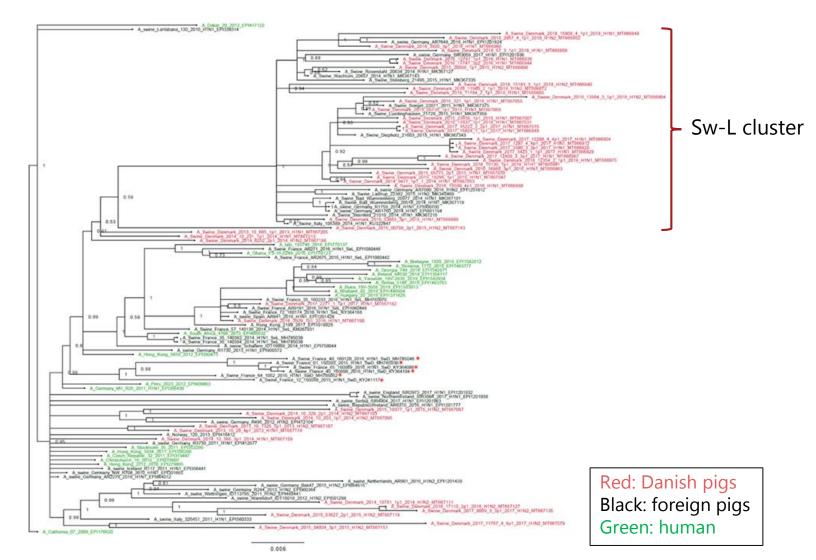
Novel reassortants:

H1pdmN2sw (few submissions/year) H1pdmN2hu (few submissions/year) H1pdmN1av (from 0.6 to 15 % of all subtyped submissions in 3 years!)

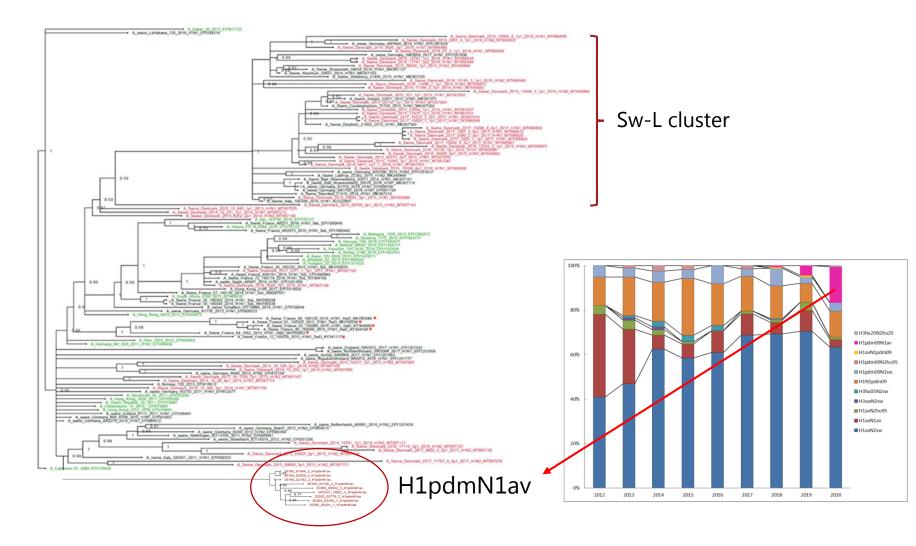
Results of the swIAV surveillance program – H1pdmNx

Co-circulation of multiple influenza A reassortants in swine harboring genes from seasonal human and swine influenza viruses

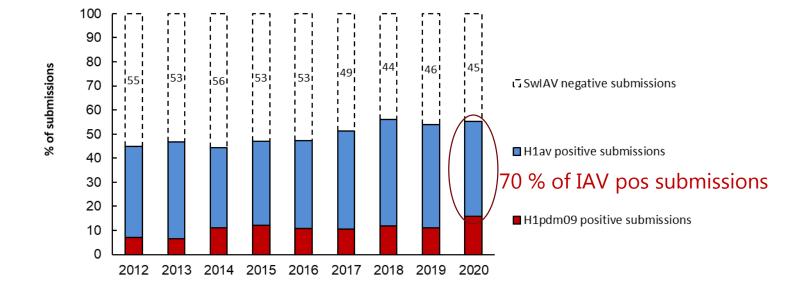
Pia Ryt-Hansen^{1,2}*, Jesper Schak Krog³, Solvej Østergaard Breum³, Charlotte Kristiane Hjulsager³, Anders Gorm Pedersen⁴, Ramona Trebbien³, Lars Erik Larsen^{1,2}



Results of the swIAV surveillance program – H1pdmN1av



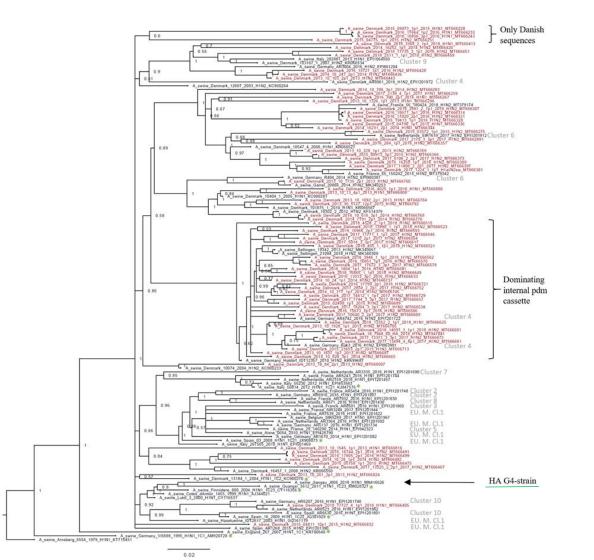
Results of the swIAV surveillance program – H1avNx



Reassortants:

H1avN2sw (most prevalent in Denmark ~60 %) H1avN2hu (few submissions/year) H1avN1pdm (one submission)

Results of the swIAV surveillance program – H1avNx

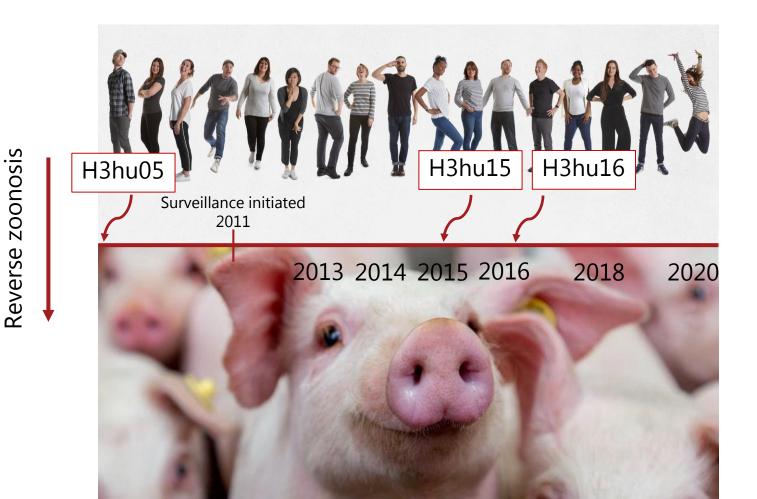


Co-circulation of multiple influenza A reassortants in swine harboring genes from seasonal human and swine influenza viruses

Pia Ryt-Hansen^{1,2}*, Jesper Schak Krog³, Solvej Østergaard Breum³, Charlotte Kristiane Hjulsager³, Anders Gorm Pedersen⁴, Ramona Trebbien³, Lars Erik Larsen^{1,2}

> Red: Danish pigs Black: Foreign pigs

Results of the swIAV surveillance program – H3huN2



Consequences of viral evolution for humans

High risk swIAV strains:

H1pdmNx Sw-L cluster

H1pdmN1av

H3huN2sw

H1avNx with MxA resistens

No cross-reaction with human vaccine sera

Enkeltstående tilfælde af ny influenza virus-variant hos en borger I januar blev der fundet et tilfælde af influenza hos en borger med en variant, der formentlig stammer fra svin. Information NS M KUNSTEN AT HOLDE SIG VARM

Forskere frygter, at influenzavirus i svin kan udløse ny pandemi hos mennesker

Tyske forskere har fundet influenzastammer i den europæiske svineproduktion, der kan føre til pandemi hos mennesker. Flere af virusstammerne er fundet i danske svin. Eksperter betegner risikoen for en ny svineinfluenzapandemi som alvorlig

lo fiere dyr, der er samlet, desto fiere virusser kan sprede sig, desto fiere mutationer sker der i virussern, desto stærre er risikoen for, at to virusser, der er unteret i hver sin retning. Inder sammen il et rekombinant virus, som ingen kender i forvien. Der er risikoen ved moderne, industriel husdyrproduktion, siger professor Hans Jern Kolmos. G siger Negaard



H3hu20N2hu20 H1pdm09N1av H1avN1pdm09 H1pdm09N2hu95 H1pdm09N2sw H1pdm09N2sw H1pdm09N2sw H1pdm09N2sw H1avN2sw H1avN2sw H1avN2hu95 H1avN1av H1avN2sw

Kommentarer (7)

Consequences of viral evolution for swine

H1pdm

Respiporc FLUpan

- Human-like strain from 2009
- Limited cross protection to the Sw-L cluster
- Limited cross protection to the H1pdmN1av cluster?

H1av

Respiporc FLU3

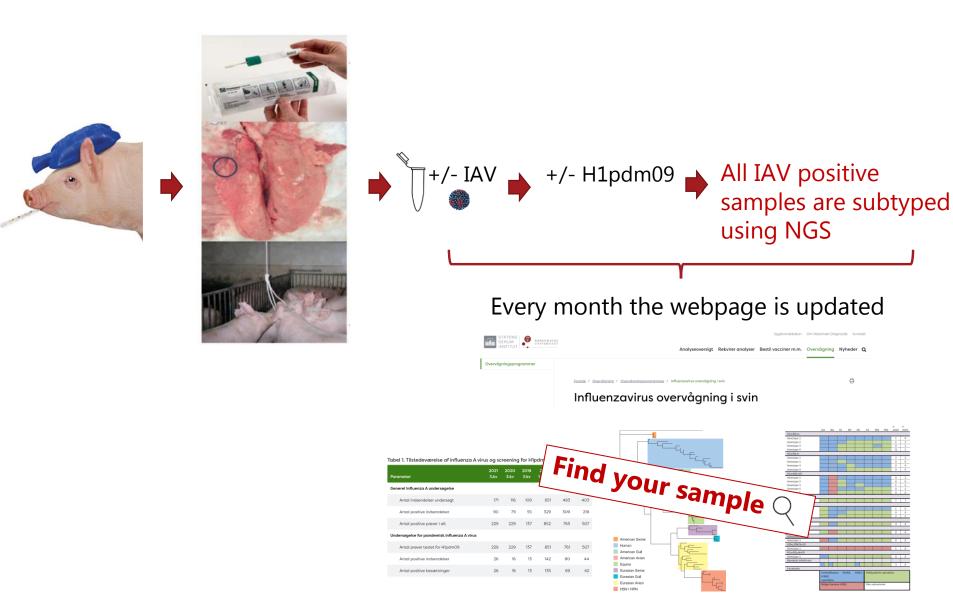
- Strains form 2000-2003
- Limited cross protection among different drifted H1av strains

11/11/11

H3hu

- Limited herd immunity
- No vaccine available

The future of the swIAV surveillance program



Acknowledgements

Lars Erik Larsen Charlotte Kristiane Hjulsager Jesper Schak Krog Ramona Trebbien Solvej Østergaard Breum Jakob Nybo Nissen Nicole Bakkegård Goecke Sophie Joanna George Anders Gorm Pedersen Nina Dam Grønnegaard Hue Thi Thanh Tran Sari Mia Dose



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STATENS SERUM INSTITUT



Thank you for your attention

