Influenza A virus in swine in Brazil

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- Brazil



3 - 4 April 2024

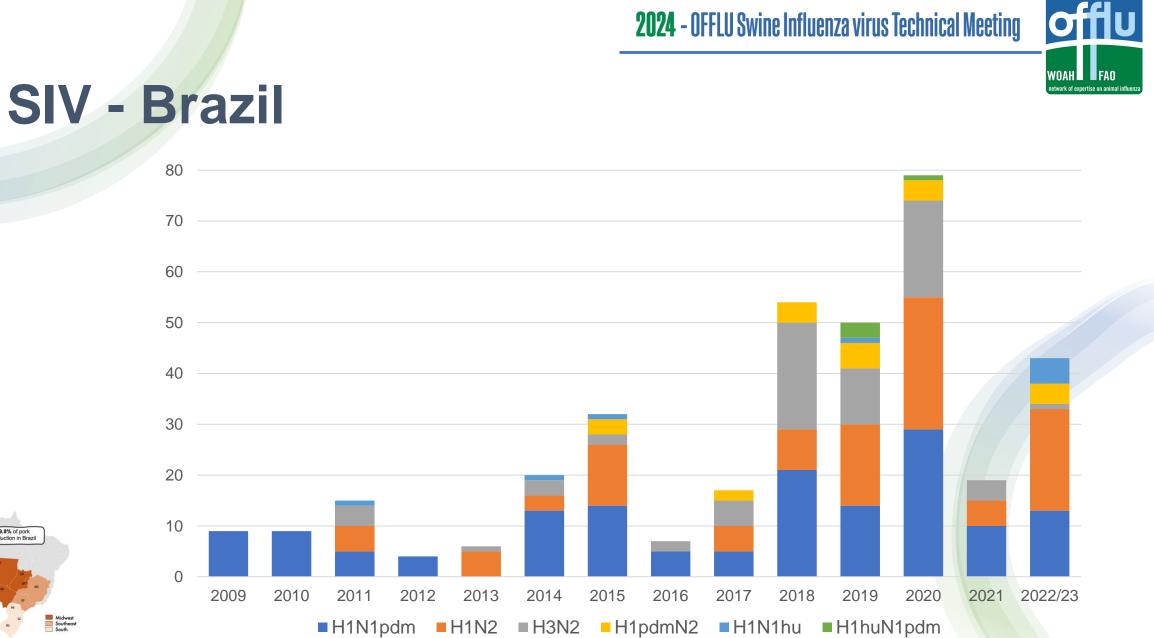
offlu **2024** - OFFLU Swine Influenza virus Technical Meeting twork of expertise on animal influe 99.8% of pork production in Brazil MT GO DF South America MG MS SP PR **Midwest** SC Southeast RS South Powered by Bing © Microsoft, OpenStreetMap

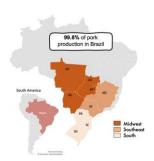
SIV - Brazil



Brazilian Agricultural Research Corporation

- Passive surveillance since late 2009
- Nine states in South, Southeast and Midwest Brazil.





Tochetto et al., 2024. unpublished data

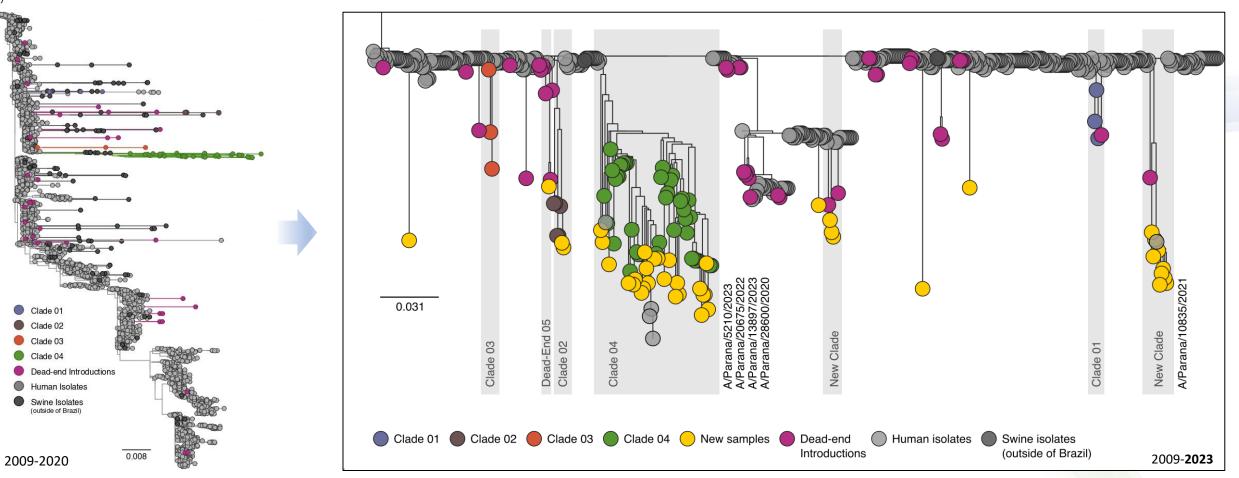


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H1 1A 1.3.3.2 (pdm09)

(A)





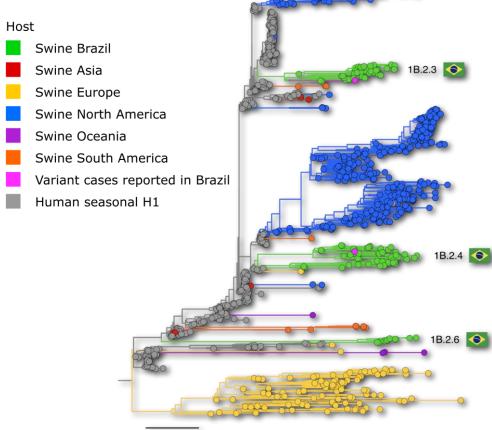
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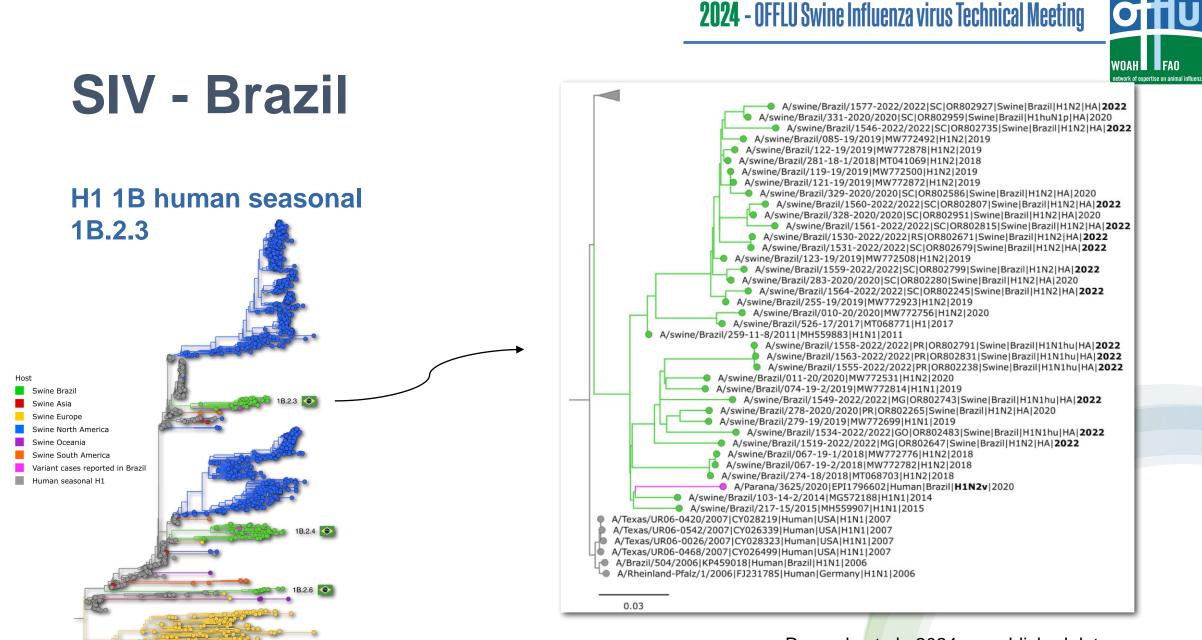
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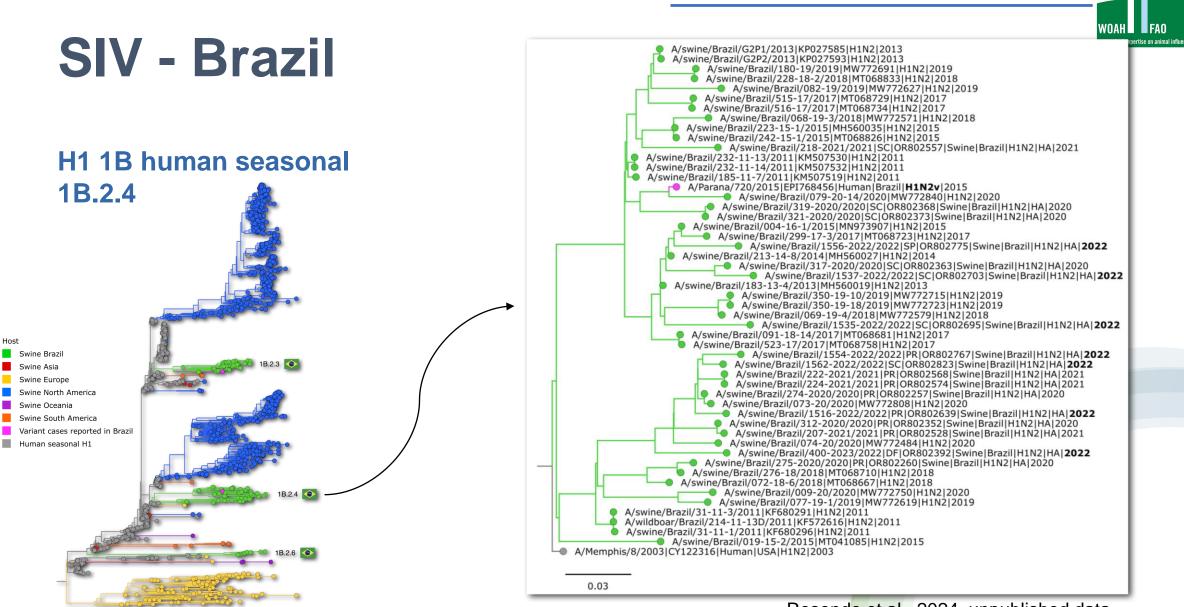
H1 1B human seasonal



Resende et al., 2024. unpublished data



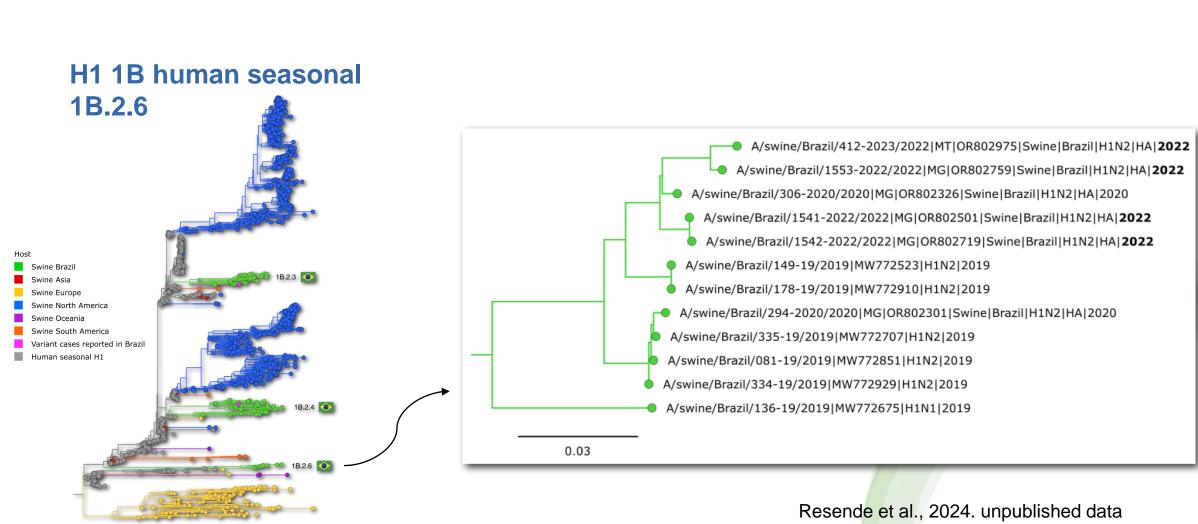
Resende et al., 2024. unpublished data



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Resende et al., 2024. unpublished data

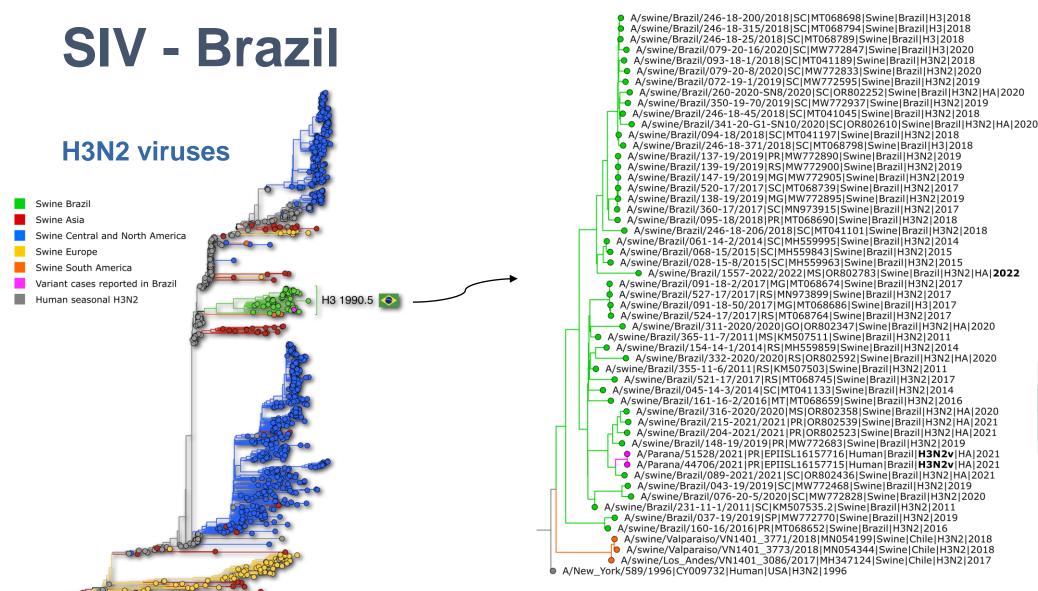
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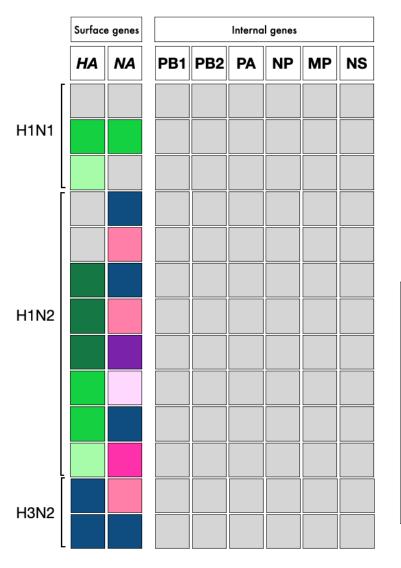
Resende et al., 2024. unpublished data



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Genotypes

H1N1, H1N2 and H3N2



Origin and genetic clade	
	2009 pdm H1N1
	Human seasonal H1N1 (early 2000s) / H1 1B.2.3
	Human seasonal H1N2 (early 2000s) / H1 1B.2.4
	Human seasonal H1N1 (late 1980s) / H1 1B.2.6
	Human seasonal H3N2 (late 1990s) / H3 1990.5
	Human seasonal H3N2 (2014-2015) / N2-#1
	Human seasonal H3N2 (middle 2010s) / N2-#2
	Human seasonal H3N2 (late 1990s) / N2-#5
	Human seasonal H3N2 (late 1990s) / N2-#6

Each genetic clade (N2-#) represents a distinct genetic clade derived from different introductions of human seasonal IAVs into swine in Brazil.

Tochetto et al., 2024. unpublished data

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Antigenic analysis



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Antigenic and genetic diversity of HI and H3 influenza A viruses in swine in Brazil

Sara Lopes, ¹⁰ Tavis K.Anderson, ¹⁰ Rejane Schaefer, ¹⁰ Caroline Tochetto, ¹⁰ Danielle Gava, ¹⁰ Mauricio E. Cantao, ¹⁰ Janice R. Ciacci-Zanella, ¹⁰ Amy L.Vincent Baker, Nicola S. Lewis **doi:** https://doi.org/10.1101/2023.12.01.569635

- Antigenic characterization of IAVs circulation in Brazilian swine between 2010-2018 was performed.
- Limited cross-reactivity between circulating swine lineages, and significant antigenic variation within lineage.
- No human vaccine strains were antigenically similar enough to provide significant protection against potential zoonotic infections by Brazilian H1 or H3 swine IAV.



Final remarks and ongoing projects

- The genetic diversity of swIAV in Brazil is increasing in the last years;
- Since 2015, nine variant IAVs were detected in humans in Southern Brazil;
- Continue to monitor influenza in pigs through sequencing and antigenic analyses;
- New research partnerships;
- Scientific divulgation (Publications).



Acknowledgements



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Federal University of Santa Maria



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Agricultural Research Service



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Thank you

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