

The background features abstract, overlapping green geometric shapes in various shades of green, creating a modern and dynamic design. The shapes are primarily triangular and polygonal, with some areas appearing more translucent than others.

PRRS Update

Luc Dufresne DMV

PRRS

- ▶ Historically
- ▶ Type 1: European Strains
- ▶ Less pathogenic
- ▶ Type 2: North American and Asian strain
- ▶ More pathogenic
- ▶ More impact on reproductive performance
- ▶ Impact on the performance of growing pigs
- ▶ Limited to the nursery phase
- ▶ Limited to the acute phase of illness in maternity wards
- ▶ Unfortunately everything has changed

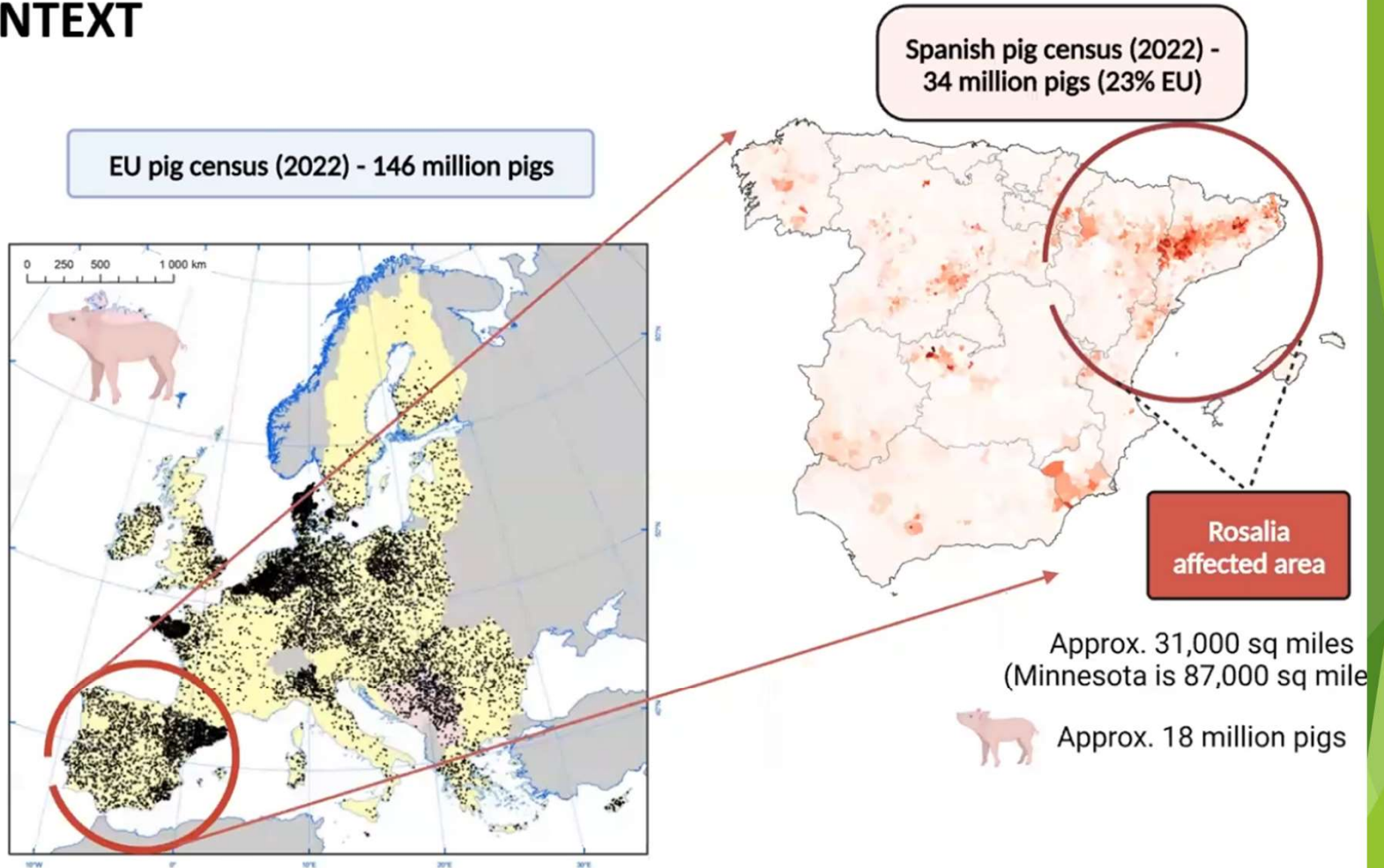
Souche SRRP Rosalia in Spain

Highly pathogenic type 1 strain

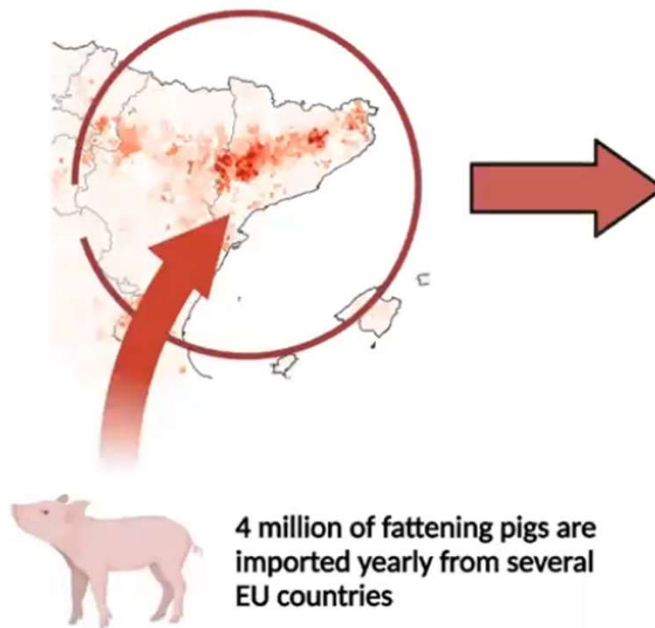
Presentation by Dr. Eric Matau, University of Barcelona






THE CONTEXT

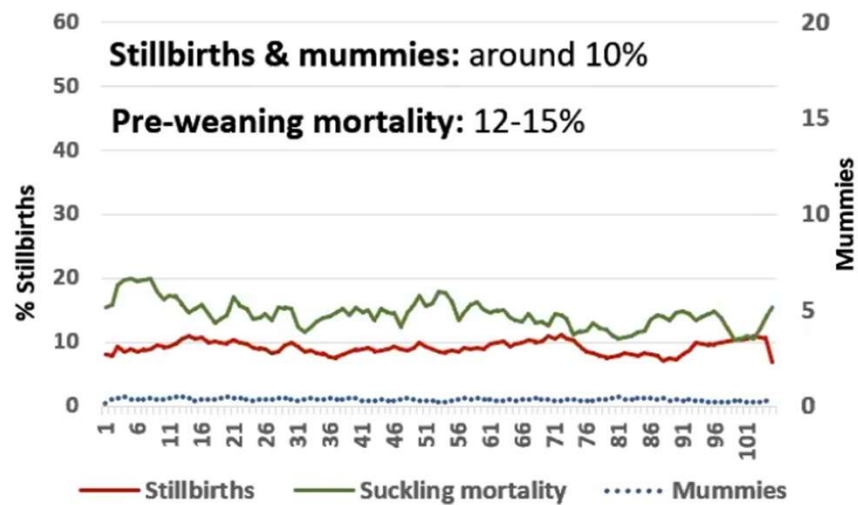
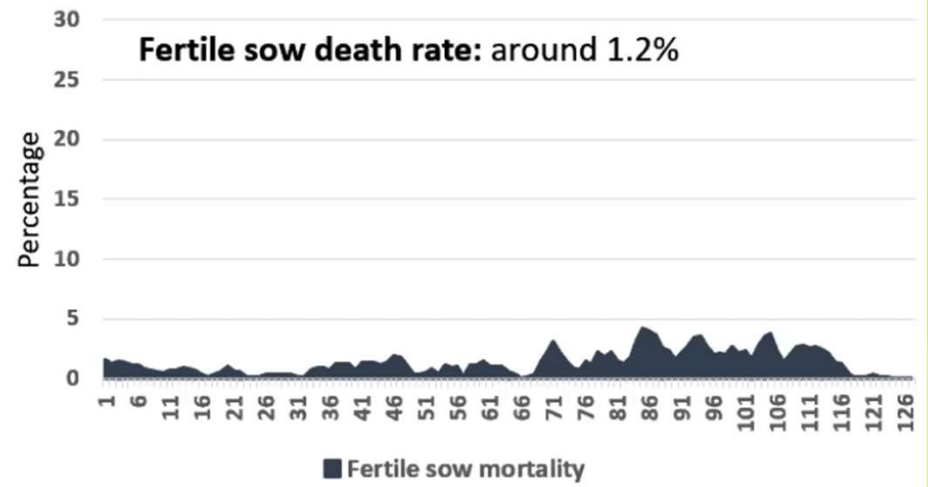
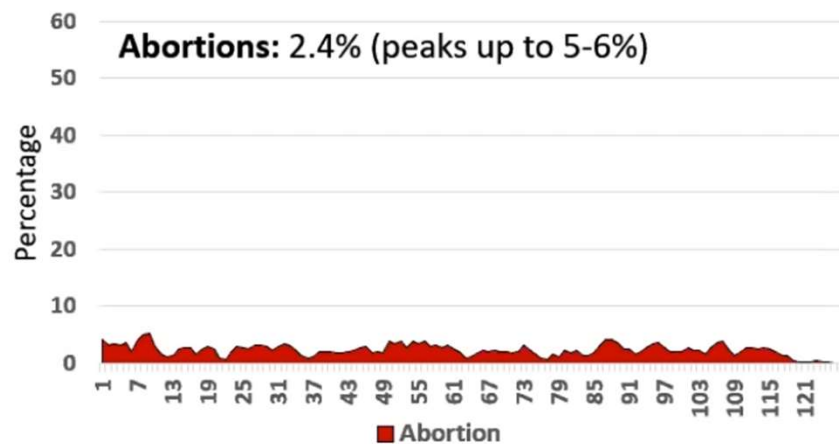


THE CONTEXT

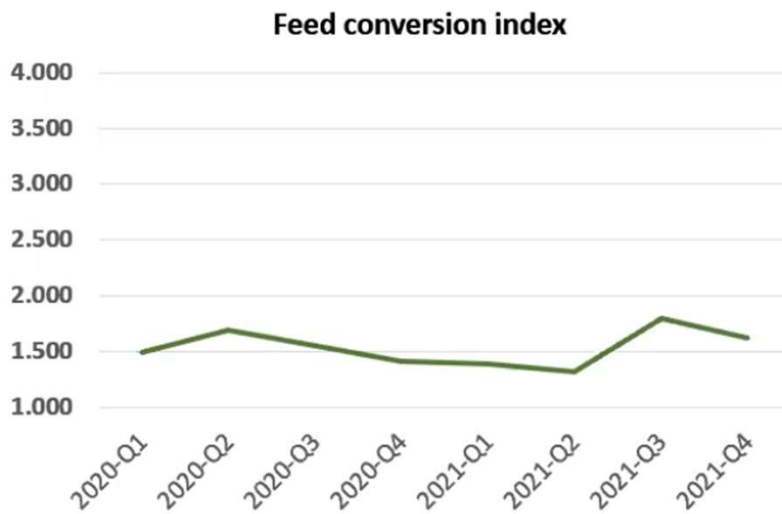


Breeders	Nurseries	Fattening
		
1,000 farms	490 farms	8,000 farms
Average size 1,400 sows		
PRRSV PREVALENCE $\geq 70\%$ PRRSV-1		
Vaccination $\geq 70\%$	Vaccination $\leq 20\%$	Vaccination $\leq 5\%$

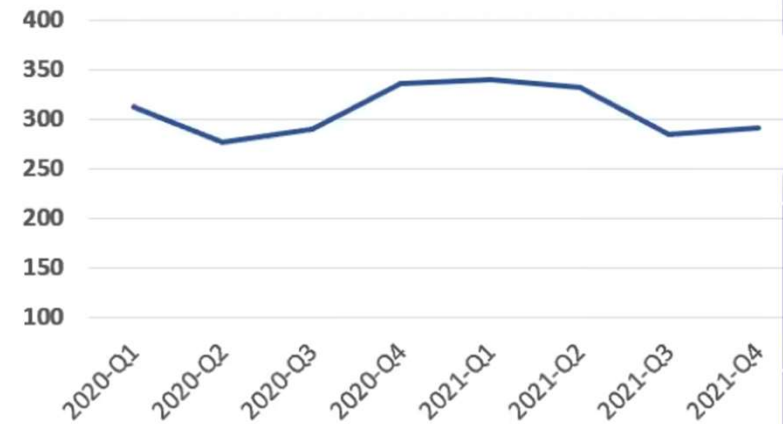
A) Average PRRSV-1 endemic farm in Spain (Breeders)



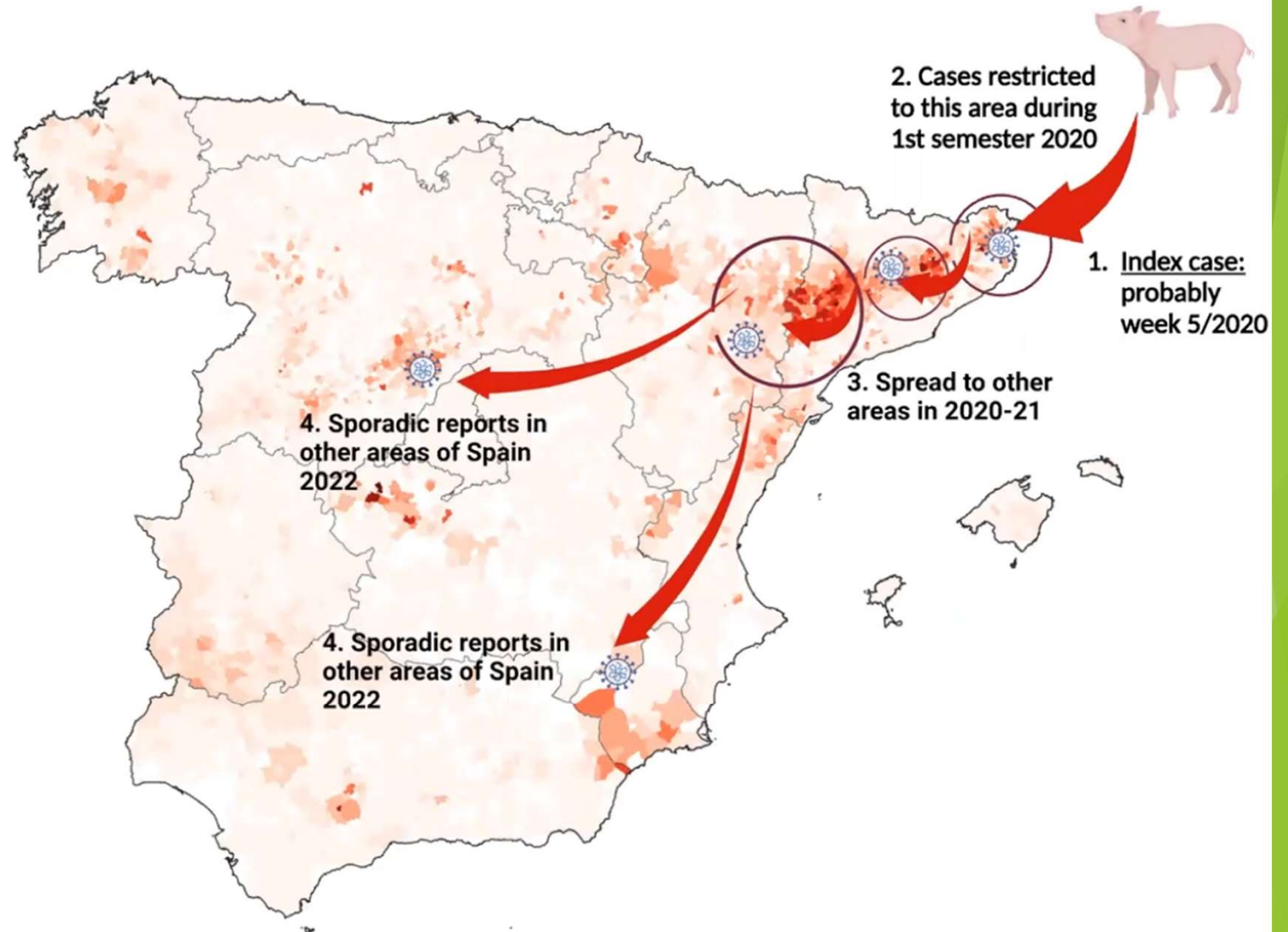
A) Average PRRSV-1 endemic farm in Spain (nurseries)



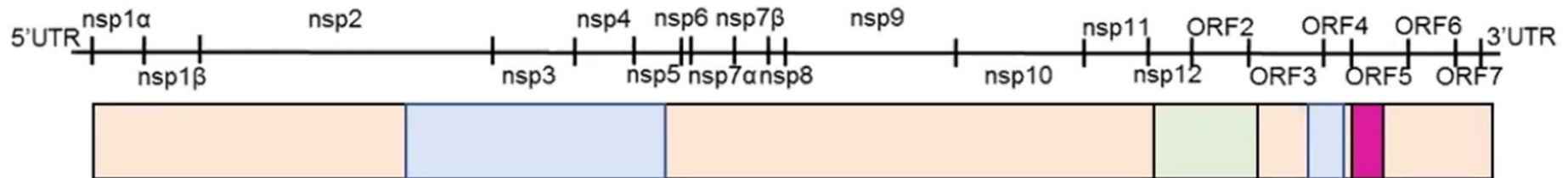
Average daily weight gain (g) between weaning and 10 weeks of age



CHRONOLOGY

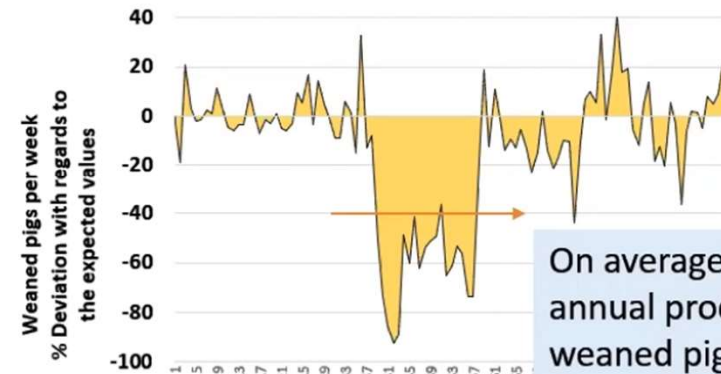
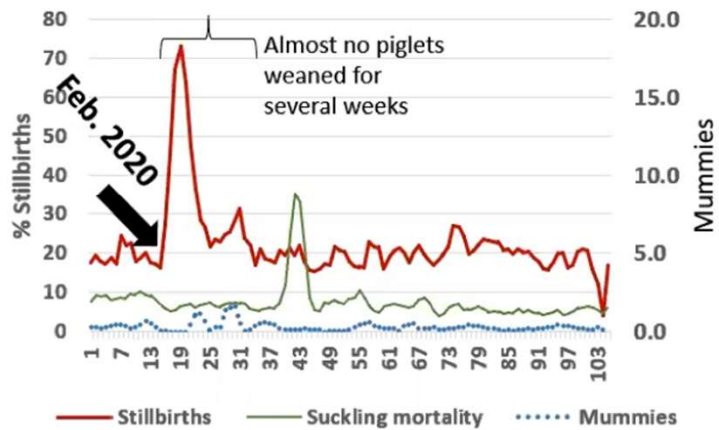
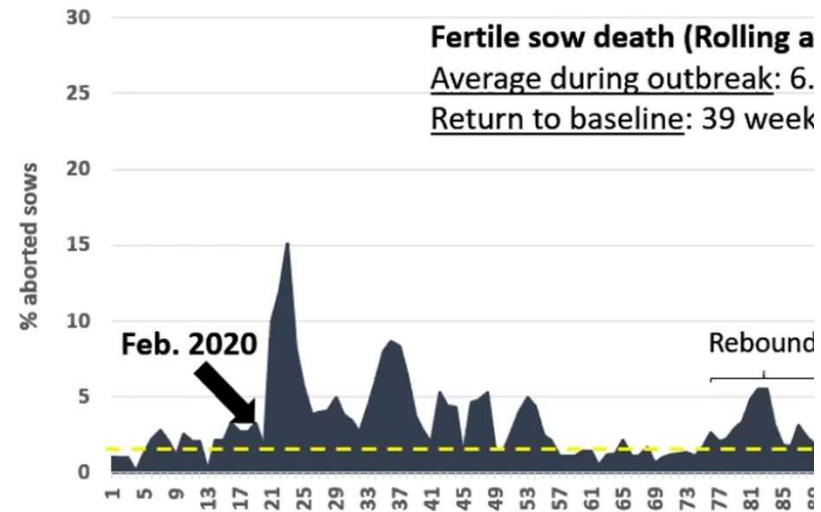
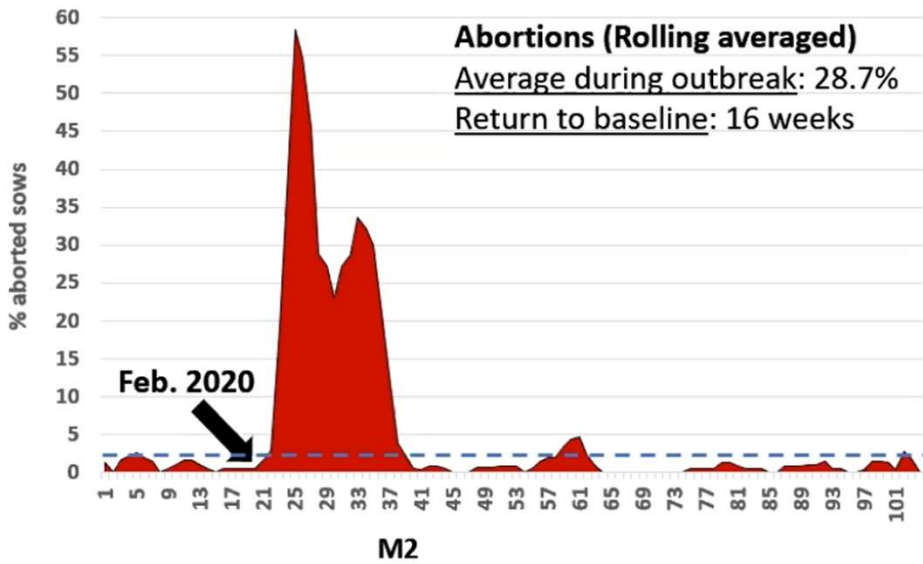


Le virus Rosalia est le résultat de la recombinaison de 4 virus différents

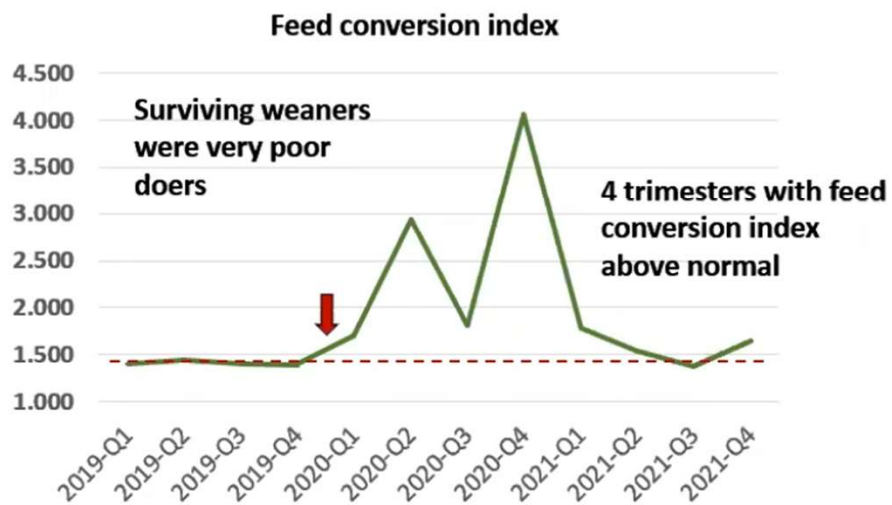
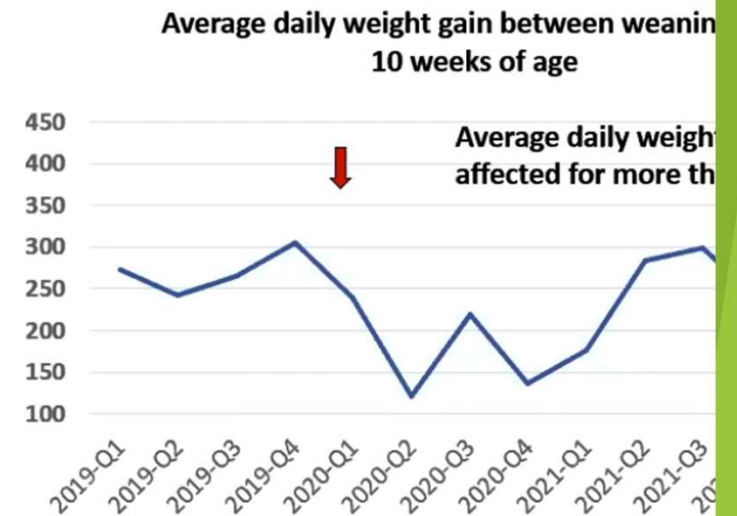
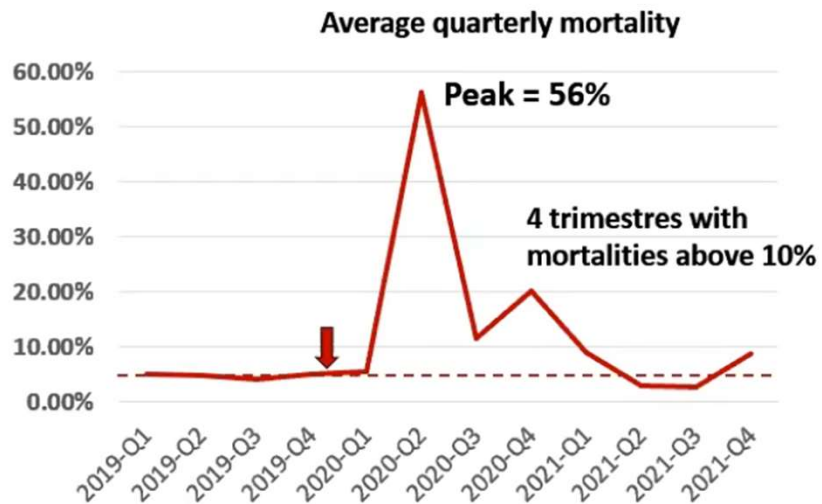


- Derived from PRRSV-1 MF346695 (Italian strain PR40/2014 reported by Canelli et al., 2018)
- Derived from PRRSV-1 local Spanish isolates clustering with KC862570
- Derived from PRRSV-1 isolates clustering with KY434184
- Segment of PRRSV-1 of unknown origin

B) Early outbreaks (Winter-spring 2020). Exmaple: PRRSV-1 stable farm, vaccinated sows



On average, -18% of the annual production of weaned pigs lost during the outbreak



PRRS Lineage 1 variant C

RFLP144

Swine Disease Reporting System Iowa state university

PRRSV RFLP classification - Source ISU, UMN, KSU, and OhioVDL

Classification

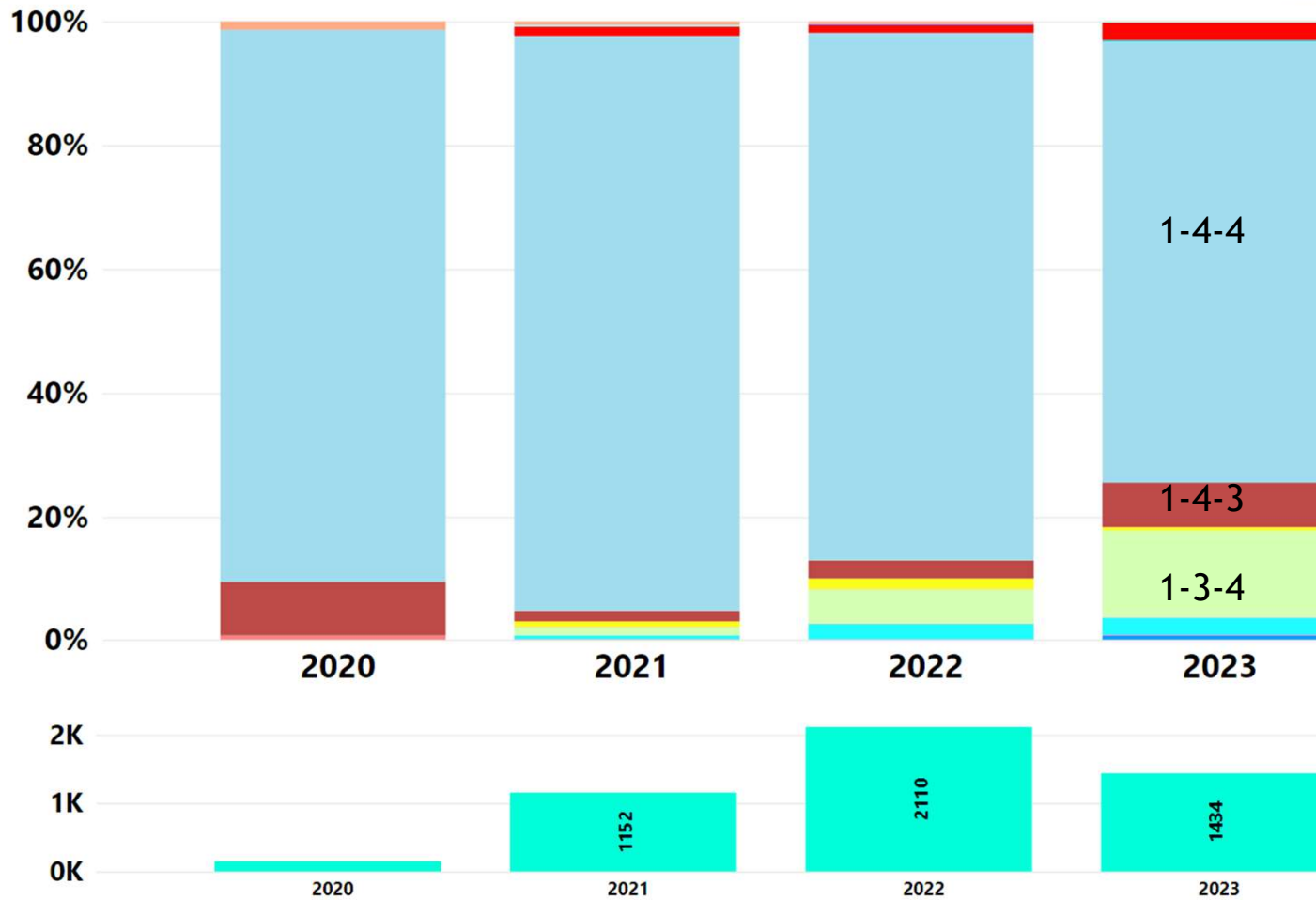
- ☐ Select all
- ☐ L1A
- ☐ L1B
- ☐ L1C
- ☒ L1CVariant
- ☐ L1Dalpha
- ☐ L1Dbeta
- ☐ L1E
- ☐ L1F
- ☐ L1G
- ☐ L1H
- ☐ L5
- ☐ L6
- ☐ L7
- ☐ L8
- ☐ L9
- ☐ PRRSV-1
- ☐ Undetermined

Vaccine-like >99%

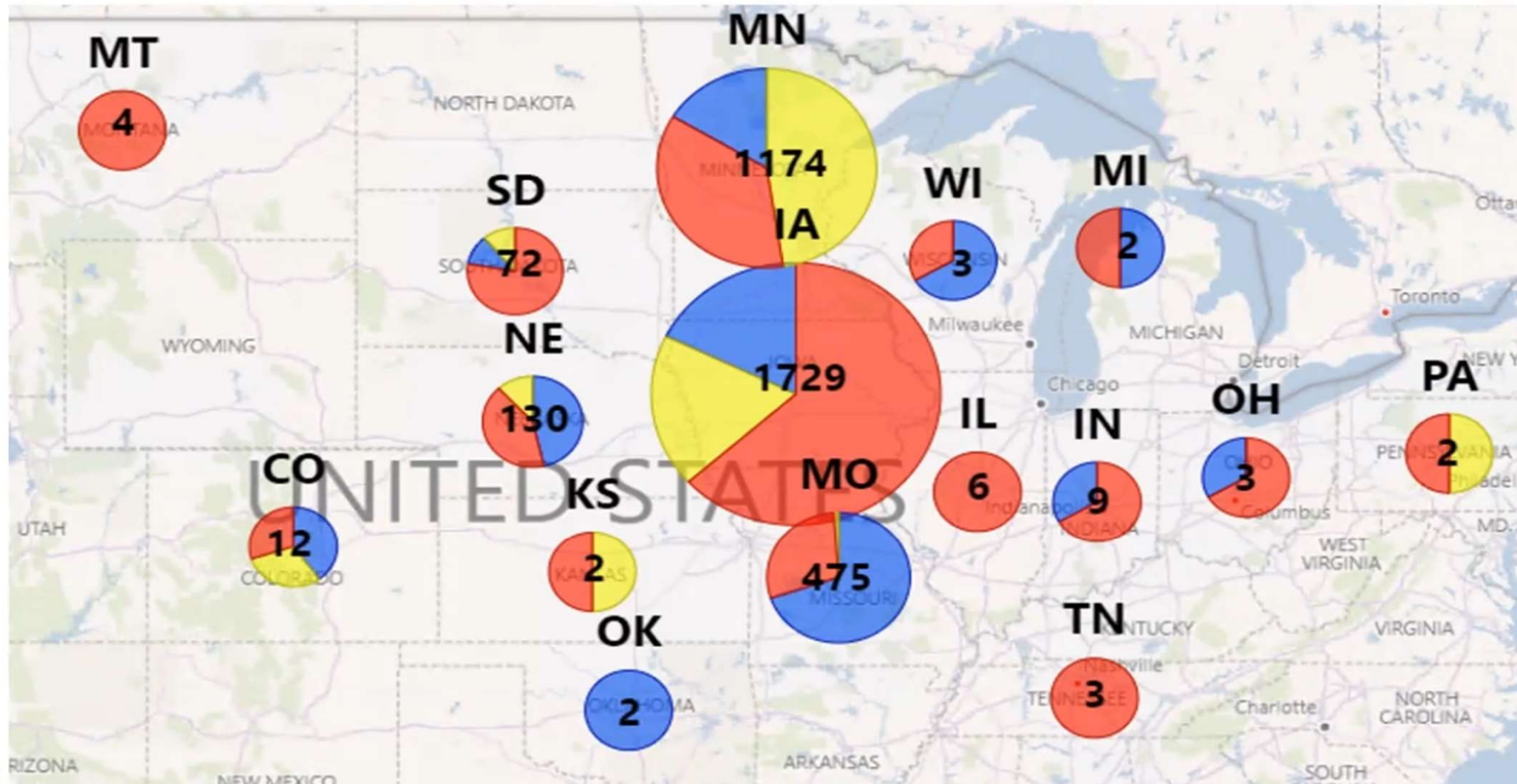
- ☐ Select all
- ☐ Ingelvac MLV like
- ☐ Prevacent like
- ☐ Wild type

Phase

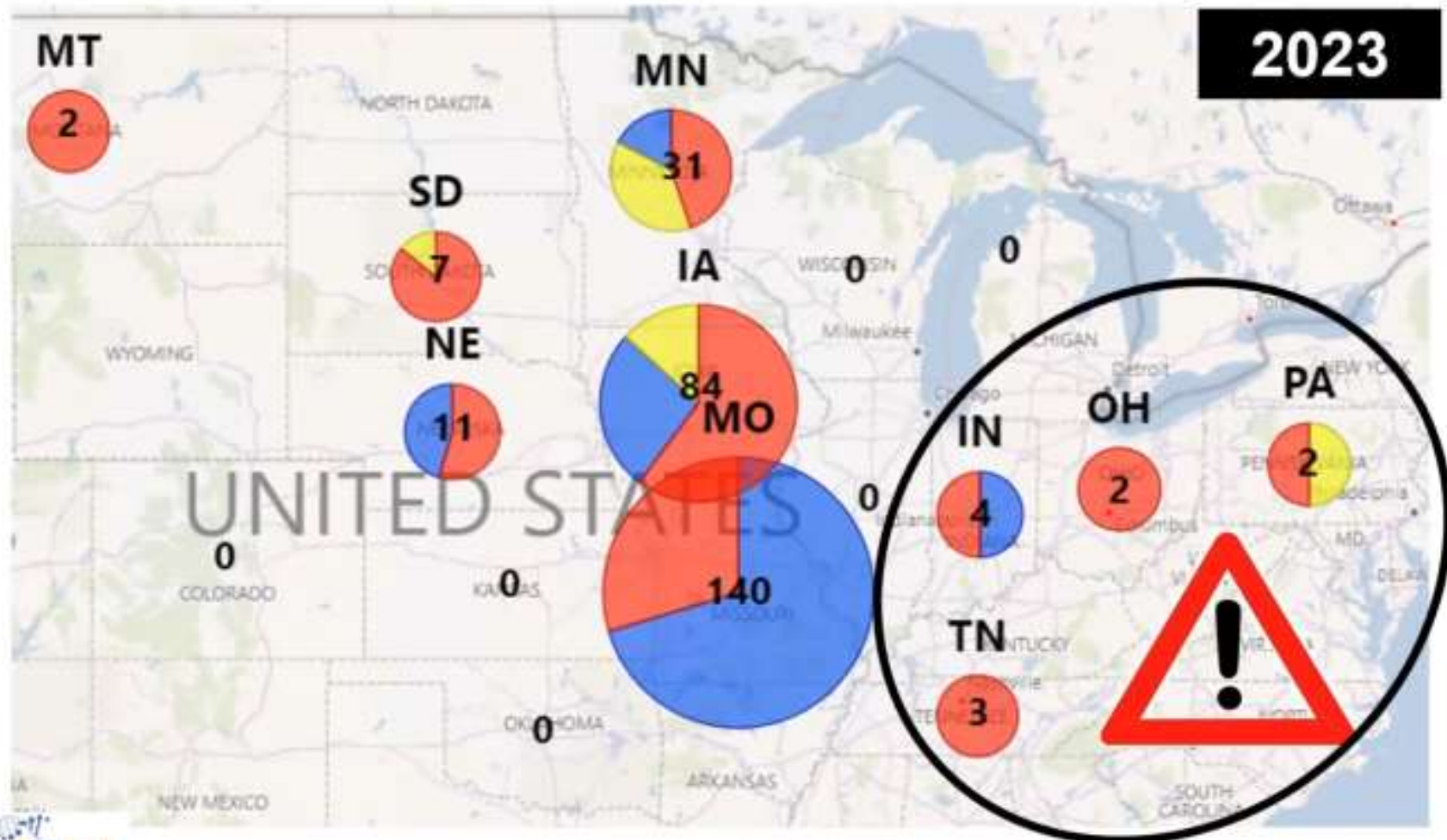
- ☐ Select all
- ☐ Adult/Sow
- ☐ Unknown
- ☐ Wean to Market



**L1C variant emerged in MN 2020 spread to IA
end of 2020 > South & West 2021/2022 > East in**



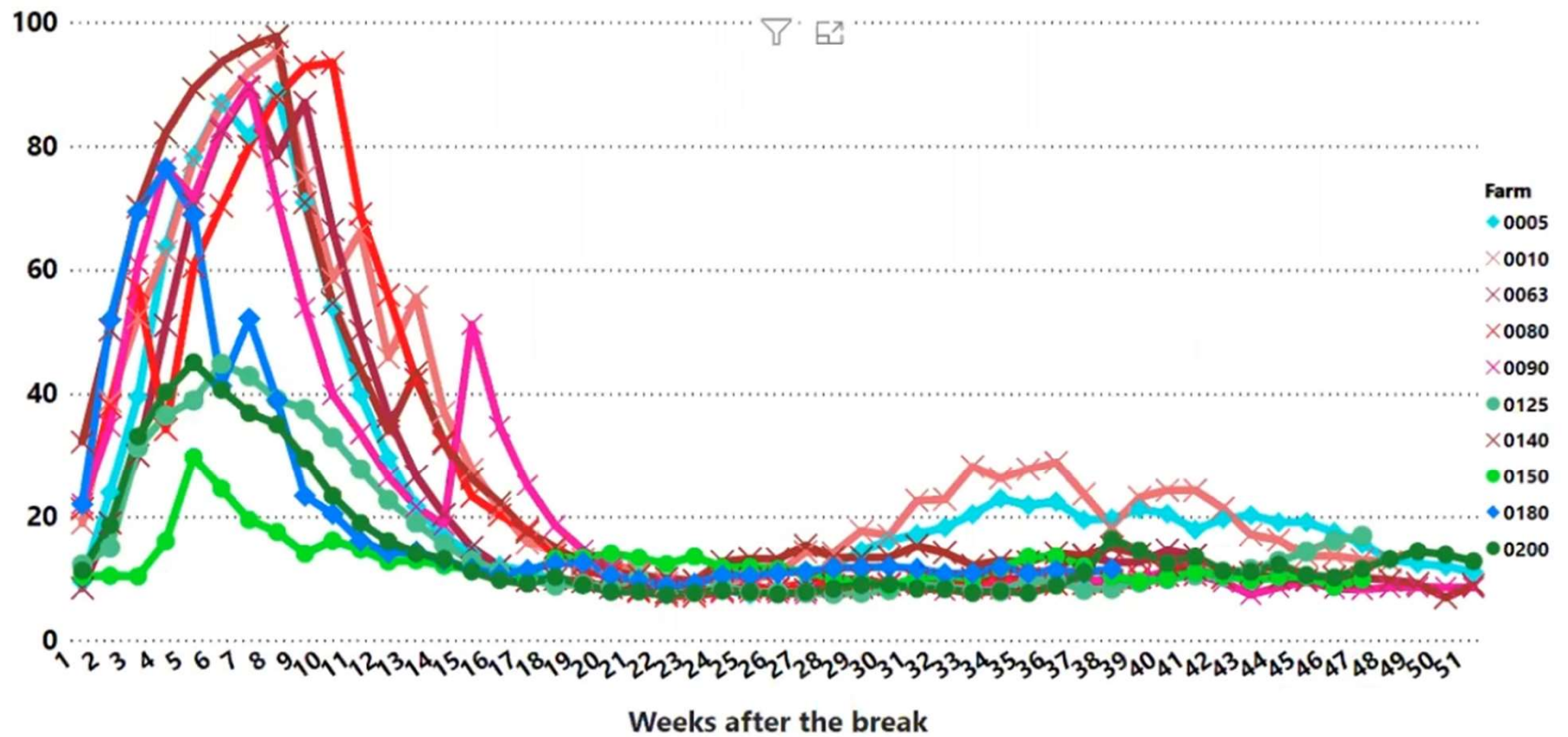
L1C variant is moving east threatening 1.5 + mi



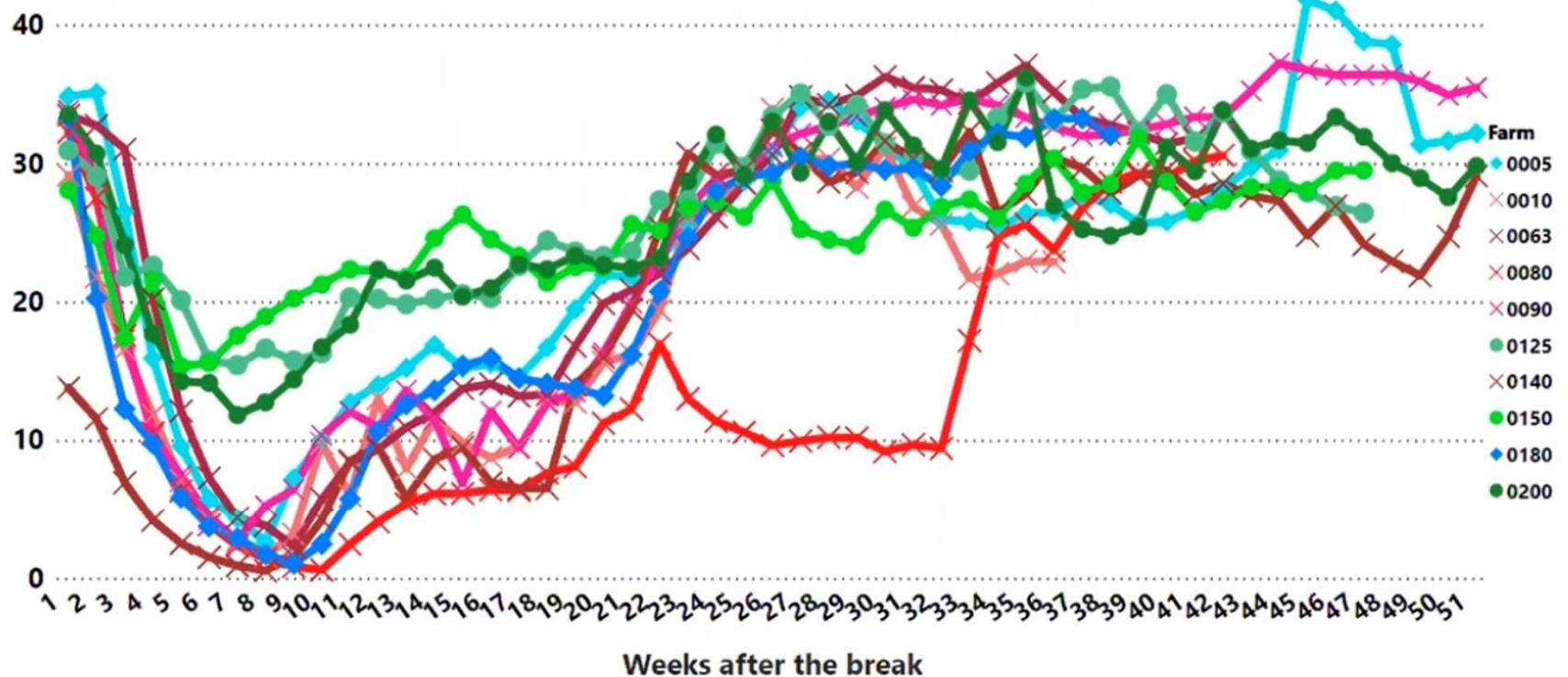
PRRS RFLP 144 L1C variant in Maternity

- ▶ Acute Phase Severe Clinical Signs
- ▶ Sow mortality
- ▶ Abortion
- ▶ Pre-weaning mortality
- ▶ Prolonged chronic phase
- ▶ High mortality and poor zootechnical performance
- ▶ 20-40% post-weaning mortality
- ▶ Do not respond consistently following herd closure
- ▶ Several herds still positive after more than 60 weeks of closure
- ▶ Considers depopulation when infected with its new strains

Preweaning Mortality



Pigs Weaned/Mated Female/Year

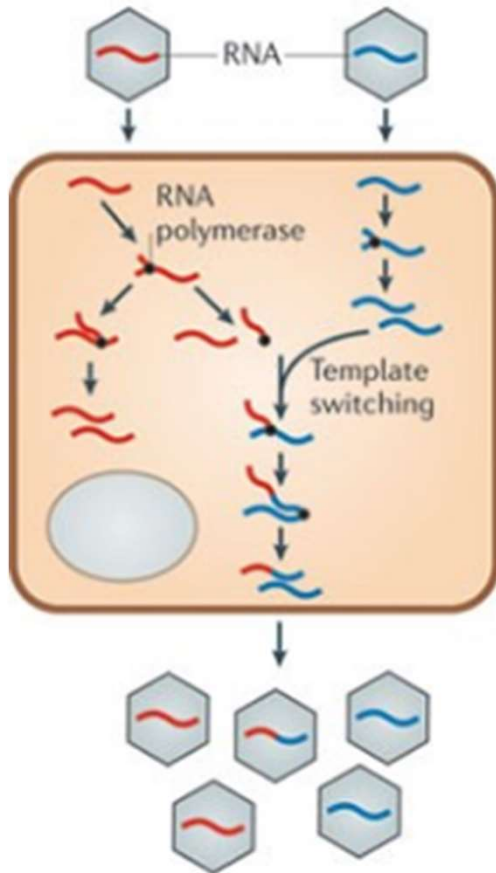


PRRS RFLP144 L1C variant : Growing Pig

- ▶ Very high viral replication and shedding
- ▶ More Enhanced Aerosol Transmission
- ▶ Post-weaning piglet performance is affected for a long time
- ▶ Very high transmission rate of growing pigs in the U.S. Midwest
- ▶ High pig density
- ▶ Few biosecurity measures
- ▶ Severe clinical sign and very high mortality in pigs from negative herds
- ▶ 20-50% mortality
- ▶ Little positive impact of PRRS vaccination
- ▶ Evidence of viral recombination



a Unsegmented RNA viruses



Viral recombination

- ▶ A process that allows the genesis of a new variety of virus by mixing the genetic program of two viruses of the same or unrelated family.
- ▶ The result of this recombination is referred to as a reassorted virus
- ▶ Mostly discussed at the level of influenza

Viral recombination

- ▶ Viral recombination can only occur if two different viruses simultaneously infect the same cell
- ▶ Viral recombination gives more or less viable, more or less virulent results
- ▶ In the vast majority of cases, the result is unsustainable
- ▶ However
 - ▶ the more genetically close the different virus variants are
 - ▶ The greater the number of infected animals
 - ▶ And the longer the viral replication phase
- ▶ The greater the likelihood of viable recombination occurring

PRRS Viral Recombination

- ▶ The characteristics of the SRRP virus combined with some modern production practices used in the United States and Spain
- ▶ Multi-site production
- ▶ Movement of pigs between regions (countries at the level of Spain)
- ▶ Increases the number of pigs simultaneously infected with different strains of PRRS
- ▶ Whole-genome sequencing of the PRRS virus shows that this phenomenon is much more frequent than previously estimated
- ▶ Several evidences of recombination
 - ▶ between wild strains
 - ▶ Between wild-type and vaccine strains
 - ▶ Between vaccine strains
- ▶ Those farming practices are also present in Ontario
- ▶ This should be consider in Control/elimination strategies

PRRS OUTBREAK MANAGEMENT PROGRAM (POMP)

- POMP Database Overview
 - 235 farms total for TTS, TTBP, TL covering 2013-2021
 - 67 completed farms with survey information (2018-2021)
 - 29 farms currently enrolled awaiting stabilization
- Some preliminary data analysis

Time to low prevalence (weeks)	Cohort 2011	Cohort 2021
10 th percentile	15	26
25 th percentile	22	32
median	27	36
75 th percentile	33	50
90 th percentile	46	73

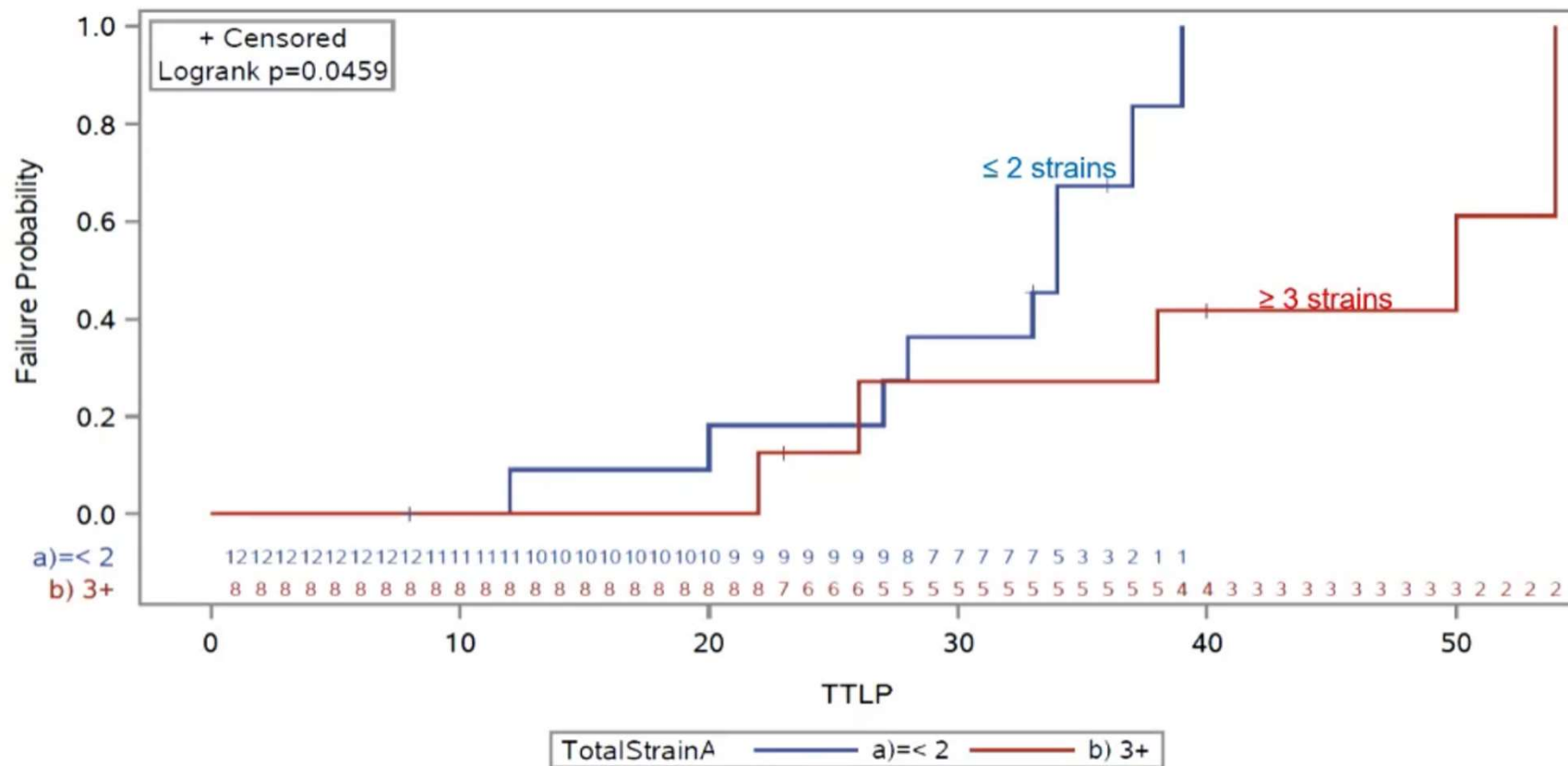
Time to baseline productivity (weeks)	Cohort 2011	Cohort 2021
10 th percentile	0	7
25 th percentile	8	16
median	16	22
75 th percentile	18	26
90 th percentile	28	31

Compared to the 2009-2012 cohort, the 2020-21 POMP herds had:

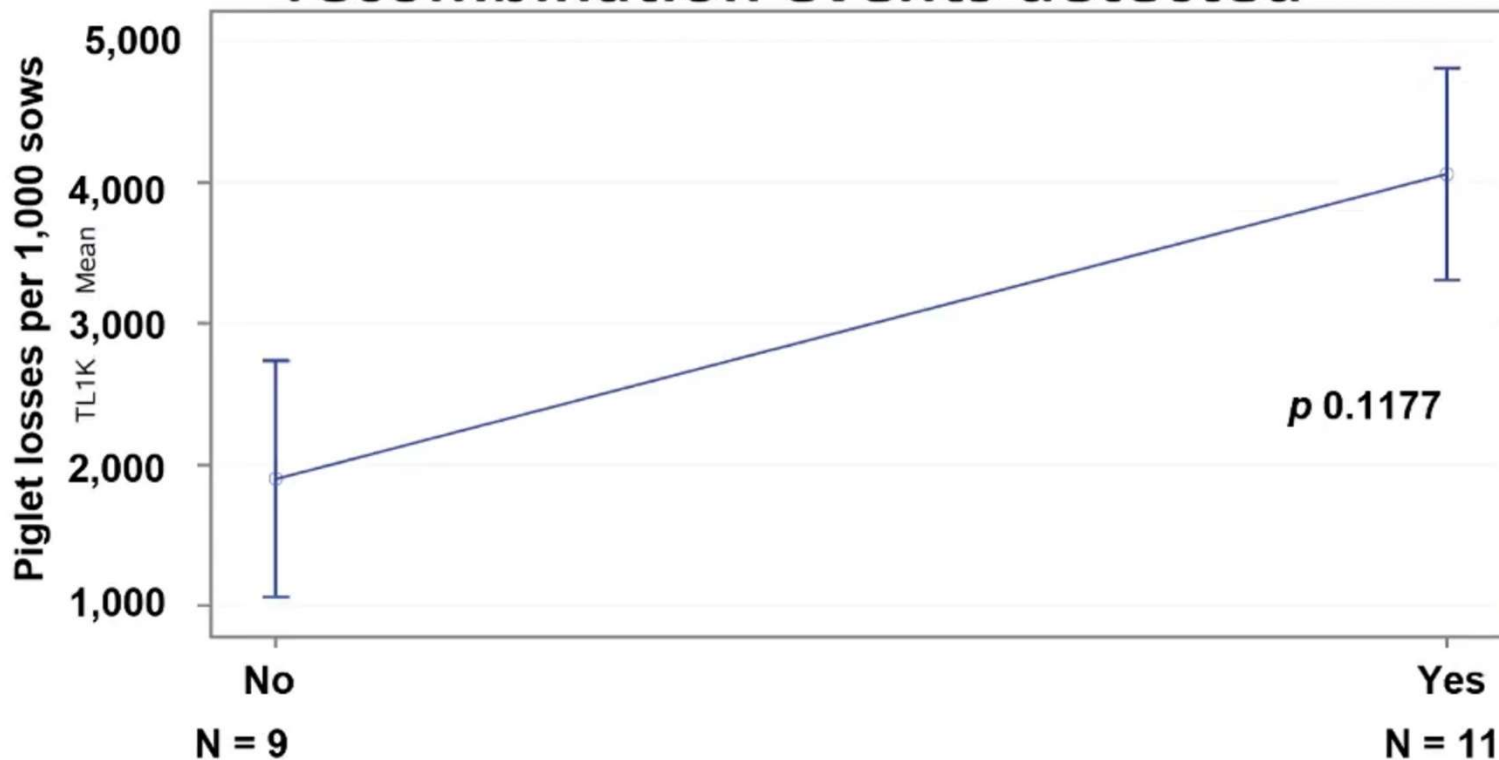
- Longer TTLP (+9 weeks average)
- Longer TTBP (+6 weeks average)
- More severe losses +1,355 pigs not weaned/1k sows

Total loss per thousand sows	Cohort 2011	Cohort 2021
10 th percentile	43	1,910
25 th percentile	1,174	2,363
median	2,789	4,144
75 th percentile	4,755	5,557
90 th percentile	6,087	7,195

Farms with ≥ 3 strains reached low prevalence 12 weeks after farms with ≤ 2 PRRSV strains



Farms with recombination events detected had 1,827 higher piglet losses / 1,000 sows than farms with no recombination events detected



<https://www.fieldepi.org/POMP>

Preliminary Analysis

Factors associated with shorter TTS & Production impact:

- Prior immunity (still) matters
 - Negative herds take longer to recover and have higher losses
- Herd closure: increases the success rate to achieve Stability
- Reporting 'seeking elimination':
 - 80% achieved stability compared to 40%
 - Longer TTBP, higher TL
- Batch farrowing system was associated with quicker TTLP, TTBP (n=4 herds)
- Timing bio-management practices: sooner = better
- PRRSV genotype ("different" or "new" = worse)
- Multiple strains or recombination events = Longer TTS and higher TL

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New Diagnostic Technique

Collection des bouts de langues (mort-nés et mortalité préseuvrage)

CONSIDERATIONS:

- Risk-based sample – an additional tool
- Great to assess PRRSV circulation in the gestation herd (stillborn)
- Practical, cheap, time-efficient

SUPPLIES:

- ☐ Scissors and forceps.
- ☐ Disposable plastic bag.
- ☐ Conical tube.
- ☐ Freezer (-20°C).



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www.field^{epi}.org

COLLECTION INSTRUCTIONS:



STEP 1.

From dead piglets, collect ~1 inch of tongue tips with the help of scissors and forceps.

MORE VIDEOS



STEP 2.

Place 20+ tongue tips in a disposable bag.



STEP 3.

(optional)

Freeze the bag, followed by thawing (freeze-thaw) immediately prior to submission.



STEP 4.

Squeeze the bag of thawed tongue tips and place the fluid in a conical tube.



STEP 5.

Ship the fluid to the VDL for PRRSV RNA detection by RT-qPCR. Keep samples refrigerated during shipment.

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3:02 / 20:29

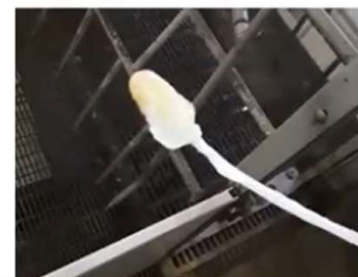
Tongue tips

- ▶ As sensitive if not better than processing fluid
- ▶ Better quality sample than oral fluids
- ▶ Can be collected and frozen for an extended period of time
- ▶ Minimum equipment needed
 - ▶ Scissor
 - ▶ Forceps or pliers
 - ▶ Freezer bags



New Diagnostic Tools in Process

- Developed to help with PRRS Elimination projects
 - Better understand gestation herd
 - Throat samples



	Serum	Tonsil Scraping	Throat sampling
PCR positivity	5 of 30 (10%)	X of 30 (70%)	30 of 30 (100%)
Need for straining	Yes	Yes	No
Collection time	2 min	3 min	30 sec





Conclusion

- ▶ PRRS takes advantage of modern production systems to become
- ▶ More pathogenic
- ▶ More contagious
- ▶ At the clinical level, these new viruses
- ▶ Create More Severe Losses
- ▶ Easier to transmit
- ▶ Harder to get rid of
- ▶ More efforts must be put in place to limit the transmission but also the creation of new strains



Conclusion

- ▶ Can we keep new variant from emerging?
 - ▶ Reduce the rate of recombination
 - ▶ Reduce at risk behavior
 - ▶ Comingled multiple unstable sources
 - ▶ Same barn
 - ▶ Same region
 - ▶ Reduce transmission to contain evolution
 - ▶ Movement of highly infectious animal
 - ▶ Kept biosecurity up to speed on the main risk of infection and transmission
 - ▶ Emphasis on Biocontainment
 - ▶ Animal movement



PED

- ▶ Acute diarrhea in all age groups
- ▶ 100% mortality in lactating piglets
- ▶ Very strong maternal immunity three weeks after exposure
- ▶ Following the devastation of 2013 and 2014, PED became endemic in the U.S. pig herd
- ▶ Sporadic hatching on sow farms
- ▶ Especially in areas dense with pigs
- ▶ Circulation in the grow-finish population is often undetected
- ▶ Lack of washing and disinfection on market transport is a significant problem
- ▶ The virus can remain infectious for an extended period of time
- ▶ In manure (more than 60 weeks)
- ▶ In feed ingredients (several weeks in some ingredients)
- ▶ Several systems use viral inoculation to reduce the risk of recirculation

PED risks of infection in naïve area

- ▶ Movement of highly infectious animal
- ▶ Manure handling of infected site
- ▶ The port of entry in negative region
 - ▶ Cull market
 - ▶ If site positive possibility of moving highly infectious animal
 - ▶ If site is never depopulating high risk of maintaining infection in clean zone
 - ▶ Packing plant
 - ▶ If the site receives positive animal
 - ▶ Cross contamination of finisher late in phase
 - ▶ Movement of highly infectious animal
- ▶ It is critical to monitor both Packing plant and cull market to detect and act on early infection
 - ▶ Cheap (pool and 1 daily test)

Questions?

