



VETERINARY MEDICINE



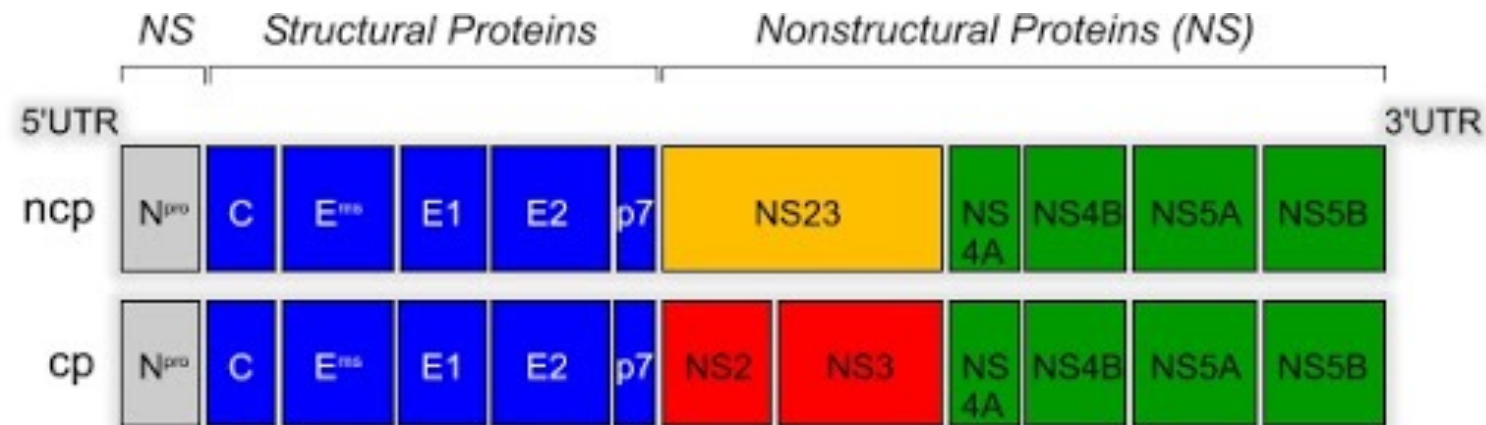
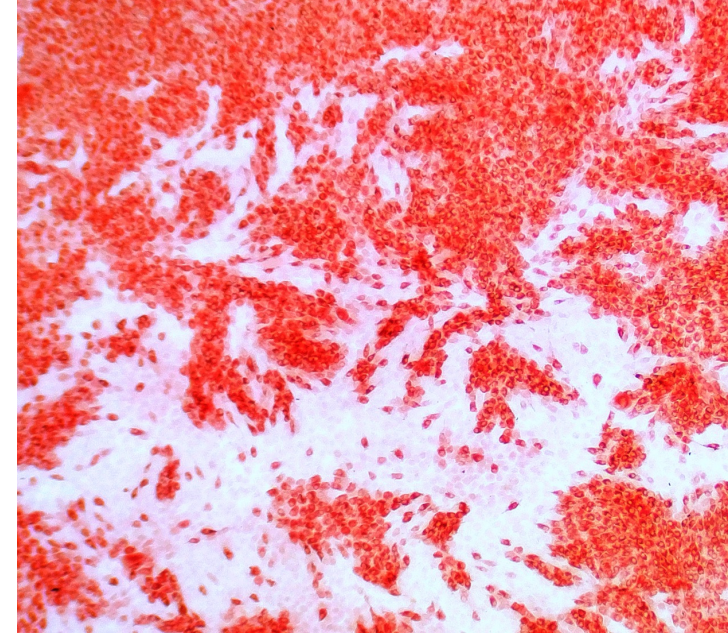
BVDV in Heterologous Species

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Assistant Professor, Food Animal Medicine and Surgery

Bovine Viral Diarrhea Virus

Viral Basics

- Genus *Pestivirus*, family *Flaviviridae*
- Positive-sense, single-stranded RNA genome
- Single open reading frame (ORF)- large polyprotein
 - Four structural: C, E^{ns}, E1, E2
 - Eight non-structural: N^{pro}, NS2, NS3, NS4A, NS4B, NS5A, NS5B
- Order Artiodactyla
 - Acute/Transient Infections
 - Congenital



Worldwide Antigen prevalence 2010-2021

(Su, Wang et al. 2022)

TABLE 2 Antigen prevalence of Bovine viral diarrhea virus of cattle in the word.

	No. studies	No. tested	No. positive	% (95% CI*)
Area*				
Asia	43	27,333	2,957	16.75% (11.27–23.04)
Europe	2	1,138	96	23.27% (0.00–89.41)
North America	1	7,544	24	0.32% (0.20–0.46)
South America	5	10,196	411	10.55% (2.63–22.82)
Sampling years				
Before 2017	30	26,608	1,625	17.18% (11.08–24.27)
After 2017	21	9,513	1,156	17.91% (10.33–26.99)

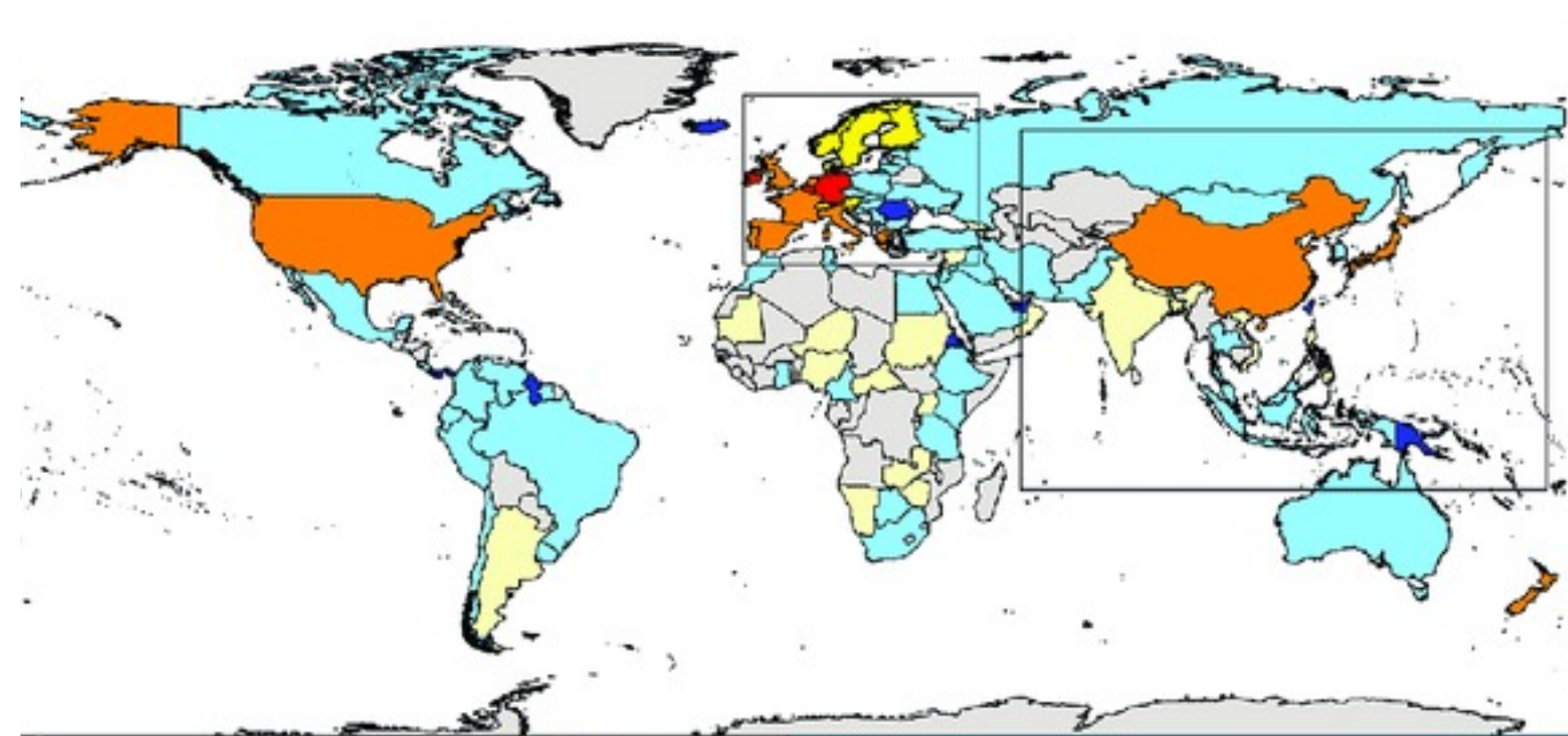
Worldwide Antibody prevalence 2010-2021

(Su, Wang et al. 2022)

TABLE 4 Antibody prevalence of Bovine viral diarrhea virus of cattle in the world.

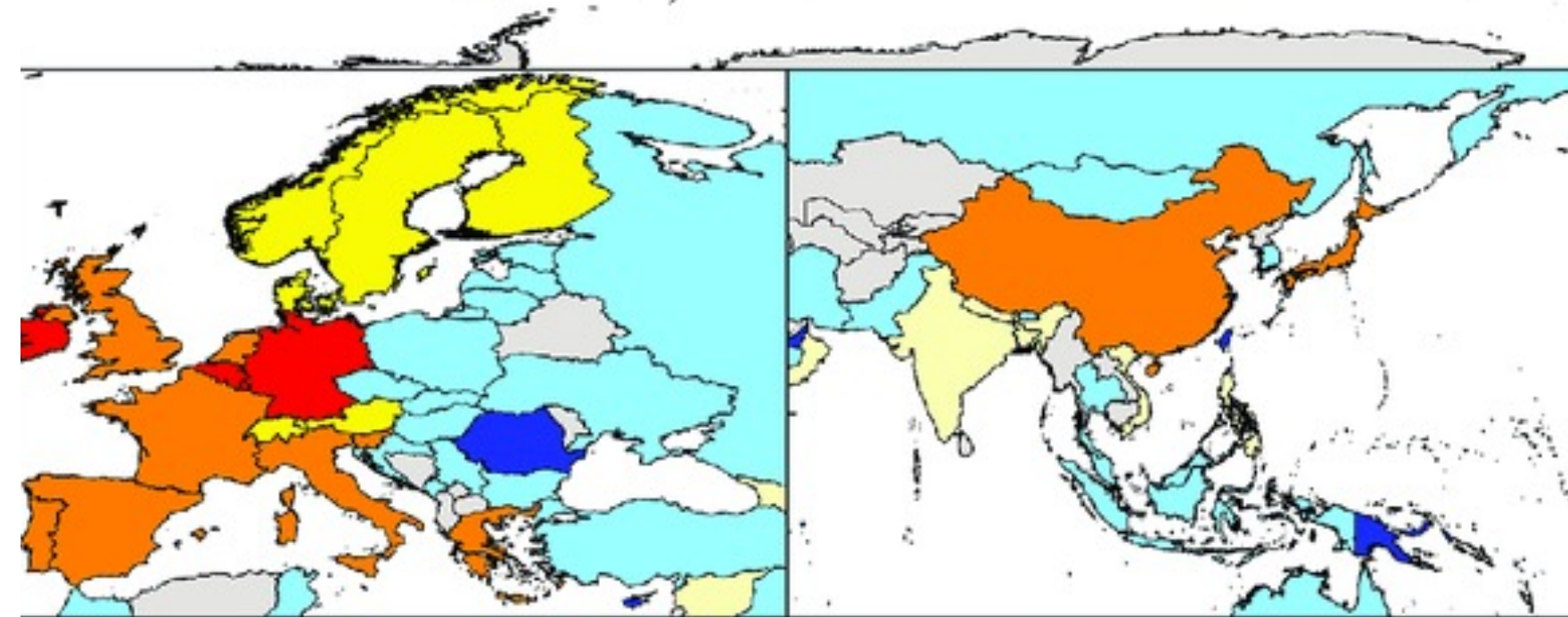
	No. studies	No. tested	No. positive	% (95% CI*)
Area				
Asia	71	53,457	23,597	42.03% (35.99–48.18)
Europe	1	180	82	45.56% (38.32–52.88)
North America	1	385	184	47.79% (42.81–52.79)
South America	1	390	298	76.41% (72.06–80.50)
Africa	3	937	424	46.13% (26.82–66.06)
Sampling years				
Before 2017	55	33,177	15,345	43.63% (37.25–50.13)
After 2017	19	11,619	6,319	41.11% (27.05–55.95)

(Su, Wang et al. 2022)



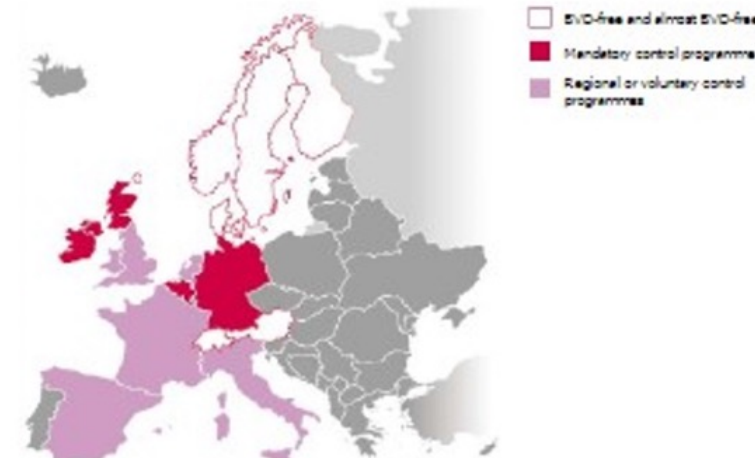
BVDV mitigation programmes

- National compulsory control and/or eradication programme
- Regional compulsory or voluntary control and/or eradication programme
- Compulsory or voluntary control and/or eradication programme changed to surveillance programme
- Herd testing without information about mitigation programmes
- Herd testing without mitigation programmes
- No control and/or eradication programmes; no information about herd testing
- No data



Implications of a resourceful virus

- The economic impact can be evaluated for BVDV by assessing differences in performance in the presence or absence of a PI animal
 - The average direct losses per:
 - naïve dairy cow were \$199.50
 - beef cow \$174.60
 - animal in the feedlot \$93.52
- Cost of testing
- Exposure of heterologous hosts
 - sheep, goats, deer, camels, alpaca, pigs, and a wide array of ungulate wildlife
 - Seroprevalence varied from 0-45% among herding districts for Reindeer in Norway
 - Feral swine seroprevalence ~6%
 - Individual state range 0-6.7%
 - CA, OK, TX highest seroprevalence
- Fetal bovine serum (500 ml bottles)
 - Super value - \$446
 - Value FBS - \$676
 - Premium - \$800 US, \$1656 New Zealand, \$1608 Australia
 - Specialty - FBS call for pricing

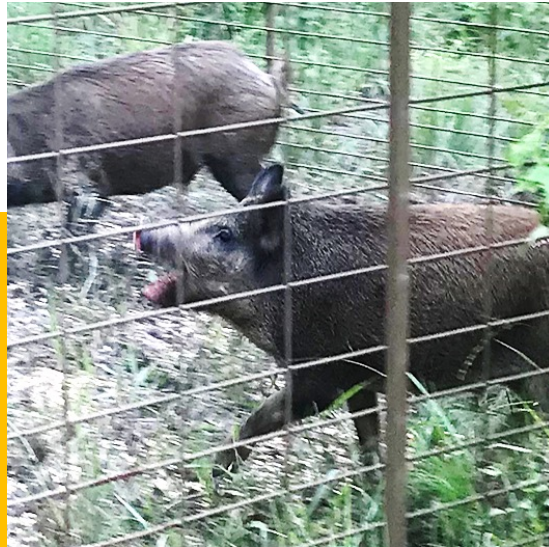


BVDV Heterologous hosts



DOMESTIC SWINE

Genetic Changes PI vs
CI



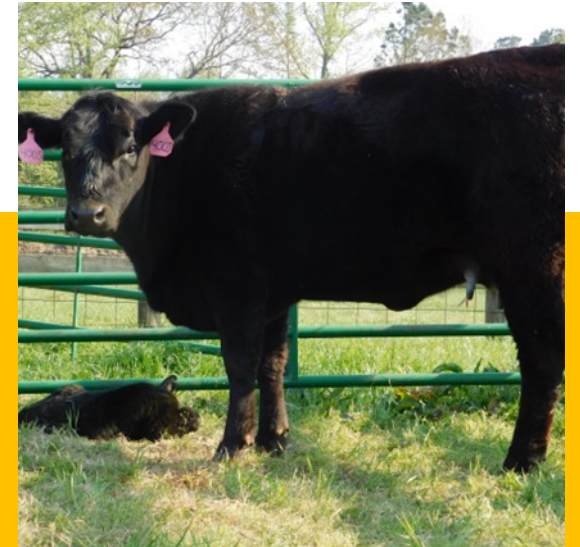
FERAL SWINE/WILD PIGS

Seroprevalence in US



WHITE-TAILED DEER

Commercial Test
Comparisons



CATTLE

Swine vs Bovine BVDV
PI Genetic

BVDV Genetic evaluation: congenitally infected piglets



- Hypothesis:
 - Viral change → clearance of congenital infection
- Specific objectives:
 - Quantify/Characterize nucleotide changes
 - Compare over time, between CI and PI

Chronic Infection Considerations

- CI vs PI
 - Eventual clearance of the virus
 - Serum neutralizing antibodies
- Timing of clearance?
 - Terpstra et al 1997
 - One month of age
 - 6-8 months of age
- Mechanism of clearance?
- Inciting cause of viral clearance?



TIMELINE- Genetic Evaluation Pigs



**DAY 26/27
GESTATION**

GESTATION

FARROWING

**VIREMIA/
SEROCONVERT**

**GENOME
SEQUENCING**

IN and IV
Inoculation
BVDV-1b NCP
AU-526

Gilts
Monthly Trans-abd
US and virus
neutralization

Piglets
Ear notch- ACE
Virus isolation,
whole genome
sequencing, virus
neutralization

Piglets
Every 2 weeks:
Virus isolation and
virus neutralization

PI vs CI piglets V
Inoculum
Next-gen and
Sanger

Results

- 4 Litters- 2019
 - 26 live-born
 - 9 antigen positive at birth
 - Antigen Capture ELISA
 - PCR
 - 4D- chronically infected
 - Seroconverted ~5.5-6 months of age





Results: Repeat

- 3 Litters- 2020
 - 34 live-born piglets
 - 7 antigen positive at birth
 - Antigen Capture ELISA
 - 6 virus positive at birth
 - Virus isolation
- 8H Chronic Infection
 - Seroconverted- ~6-8 weeks

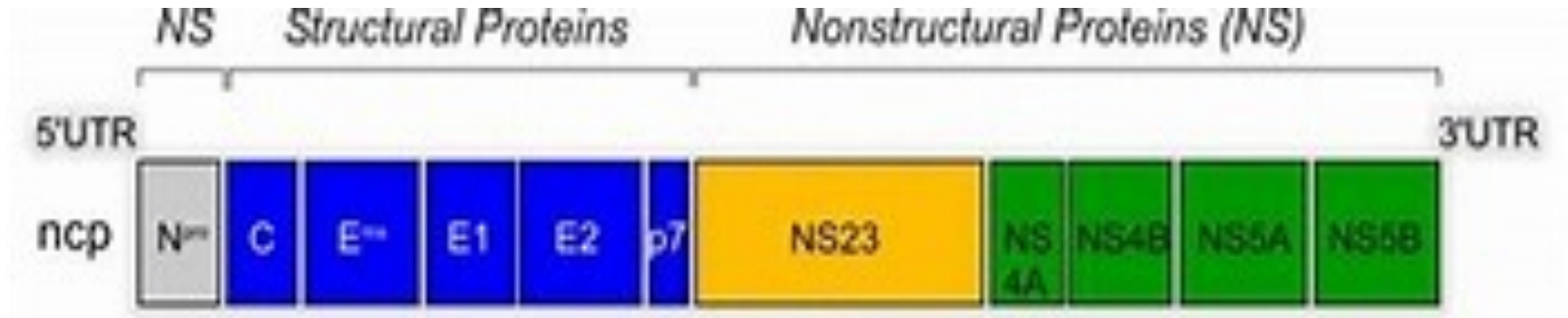
Results- BVDV sequencing

WHOLE-GENOME/NEXT-GEN

- 98.8% Identical to AU526 460F6 inoculum

SANGER

- 15 congenitally infected piglets



Nucleotide difference between Inoculum and Piglets

	Total	Transition	Transversion	Npro	C	Erns	E1	E2
460F6:4A	3	1	2	0	0	0	0	3
460F6:4C	3	2	1	1	0	0	0	2
460F6:4D	6	5	1	1	0	1	0	4
460F6:4F	3	2	1	0	0	0	0	3
460F6:4G	75	52	23	NA	10	6	18	41
460F6:4H	6	5	1	1	0	1	1	3
Median	4.5	3.5	1	1	0	0.5	0	3
Mean	14.4	10.1	4.3	0.7	1.4	1.2	2.7	8.4
	Total	Transition	Transversion	Npro	C	Erns	E1	E2
460F6:8A	2	1	1	0	0	0	0	2
460F6:8B	2	1	1	0	0	0	0	2
460F6:8E	2	1	1	0	0	0	0	2
460F6:8H	2	2	0	0	0	0	0	2
460F6:8L	3	1	2	0	0	0	0	3
Median	2	1	1	0	0	0	0	2
Mean	2.2	1.2	1	0	0	0	0	2.2

Amino Acid difference between Inoculum and Piglets

	Total	Npro	C	Erns	E1	E2
460F6:4A	13	0	1	2	0	10
460F6:4C	16	0	2	1	0	13
460F6:4D	15	1	1	3	0	10
460F6:4F	11	0	1	1	0	9
460F6:4G	46	10	2	3	1	30
460F6:4H	14	2	1	2	1	8
Median	14.5	0.5	1	2	0	10
Mean	18.5	1.9	1.3	2.0	0.3	12.9
	Total	Npro	C	Erns	E1	E2
460F6:8A	12	0	1	1	0	10
460F6:8B	11	0	1	1	0	9
460F6:8E	11	0	1	1	0	9
460F6:8H	11	0	1	1	0	9
460F6:8L	11	0	1	1	0	9
Median	11	0	1	1	0	9
Mean	11.2	0.0	1.0	1.0	0.0	9.2

Conserved Amino Acids in Piglets

Viral Protein AA Position	E2				
	43	60	103	291	365
460-f6	X	E	L	E	M
460F6:4A	T	E	L	K	L
460F6:4C	T	X	X	K	L
460F6:4D	I	A	L	K	L
460F6:4F	T	E	L	K	L
460F6:4G	T	E	L	K	L
460F6:4H	X	K	L	X	M
460F6:8A	I	K	I	X	M
460F6:8B	I	K	I	X	M
460F6:8E	I	K	I	X	M
460F6:8H	I	K	I	E	M
460F6:8L	I	K	I	X	M

Phylogenetic analysis

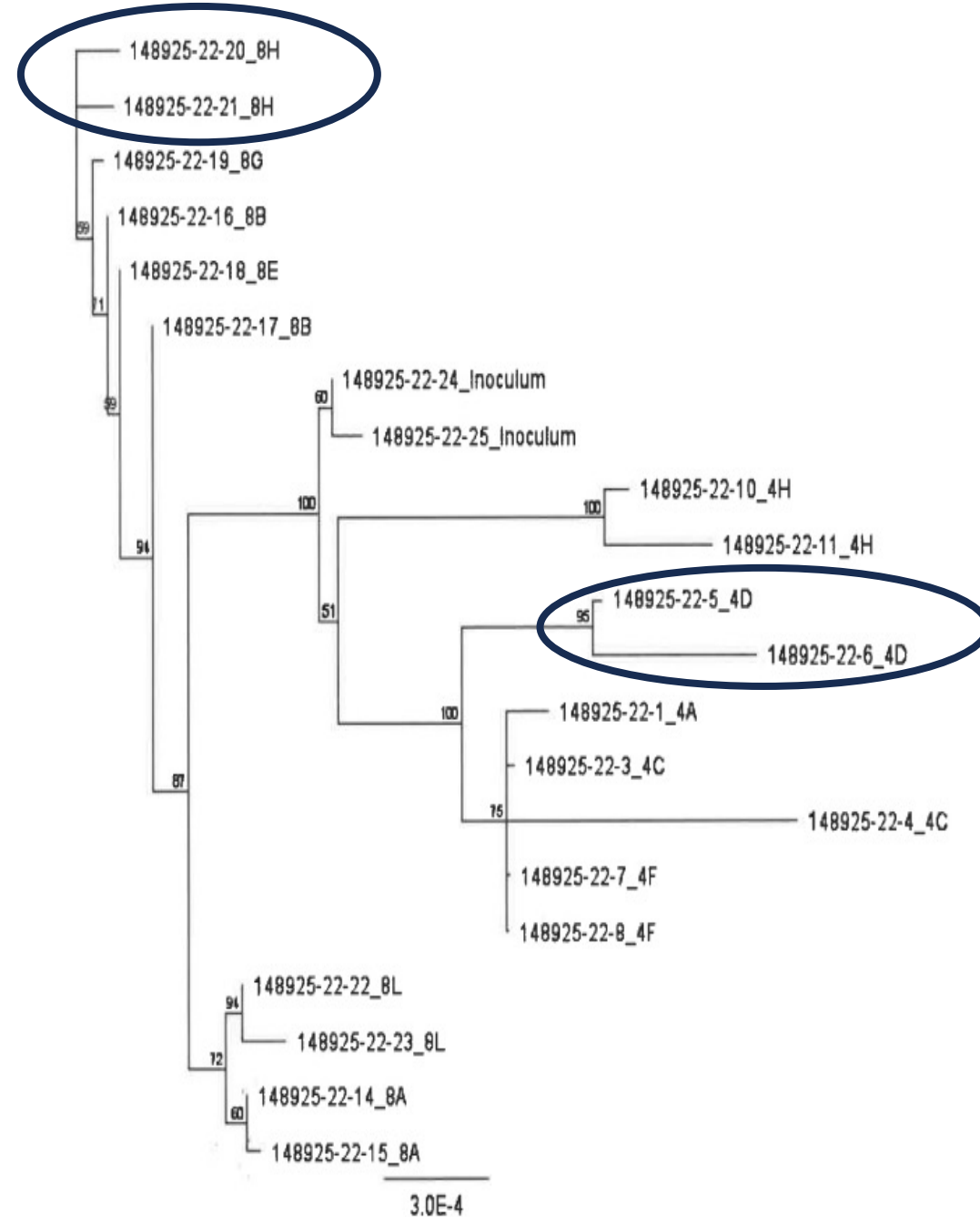


Figure 27: Phylogenetic tree of the whole genome consensus alignments, excludes 148922-22-2, 9, 12, 13 due to having less than 95% coverage.

Seroprevalence of Bovine Viral Diarrhea Virus in Wild Pigs (*Sus scrofa*) in 17 States in the USA

Shari M. Kennedy,^{1,5,6} Thomas Passler,¹ Stephen S. Ditchkoff,² Vienna R. Brown,³ Gage W. Raithel,⁴ Manuel F. Chamorro,¹ Paul H. Walz,⁴ Constantinos S. Kyriakis,⁴ and Shollie M. Falkenberg⁴



USDA Wildlife Services

- Current US Feral Swine/Wild Pig BVDV knowledge:
 - 0% Great Smokey Mountains National Park
 - doi: 10.7589/0090-3558-30.1.103.
- Seroprevalence rate for states with higher feral swine/wild pig or cattle population states

Feral Swine Seroprevalence

- Objective:
 - Determine BVDV seroprevalence in wild pigs in US
 - Determine if age, sex, or location associated with positive antibody titer

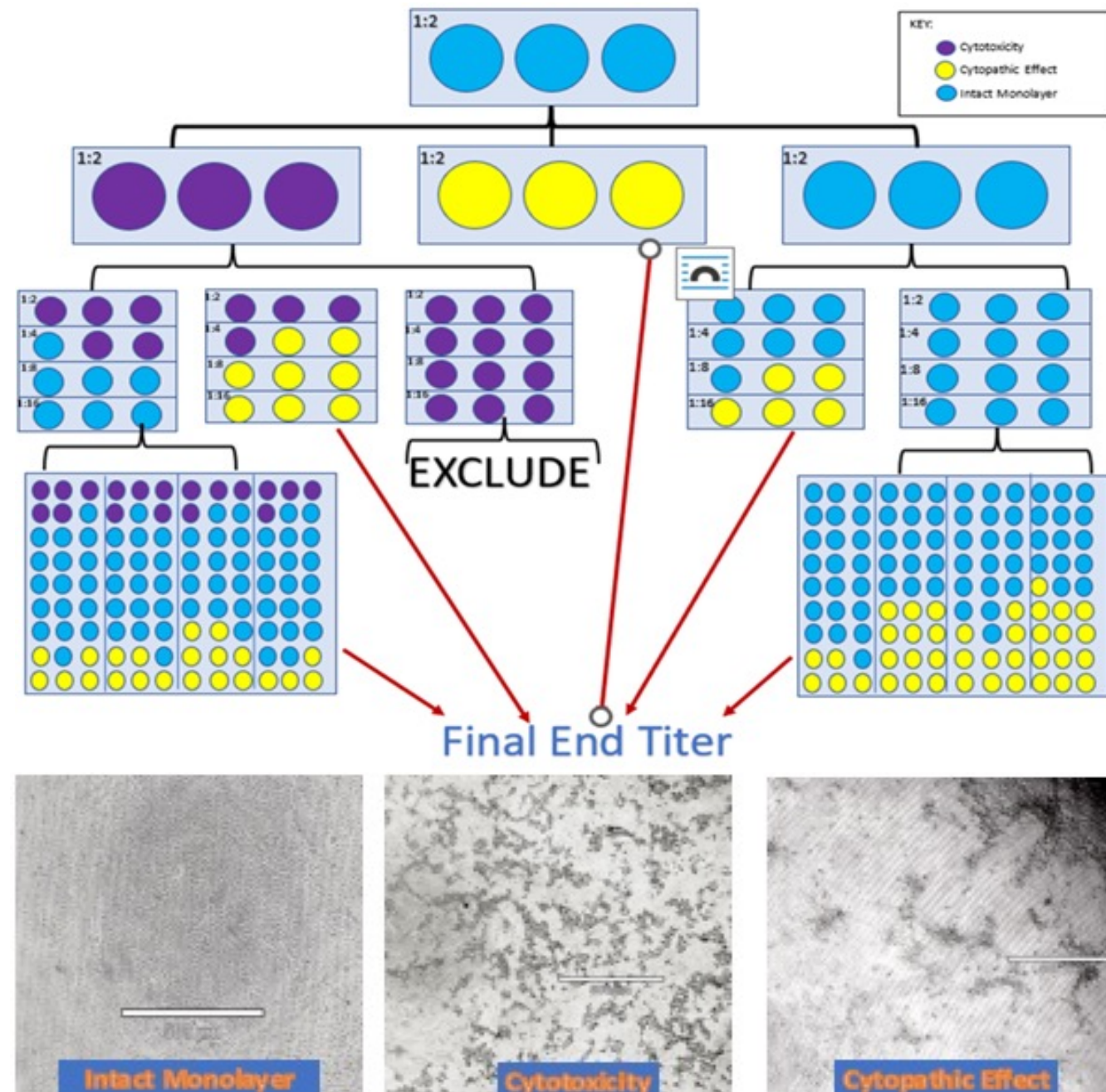


USDA Wildlife Services

Methods

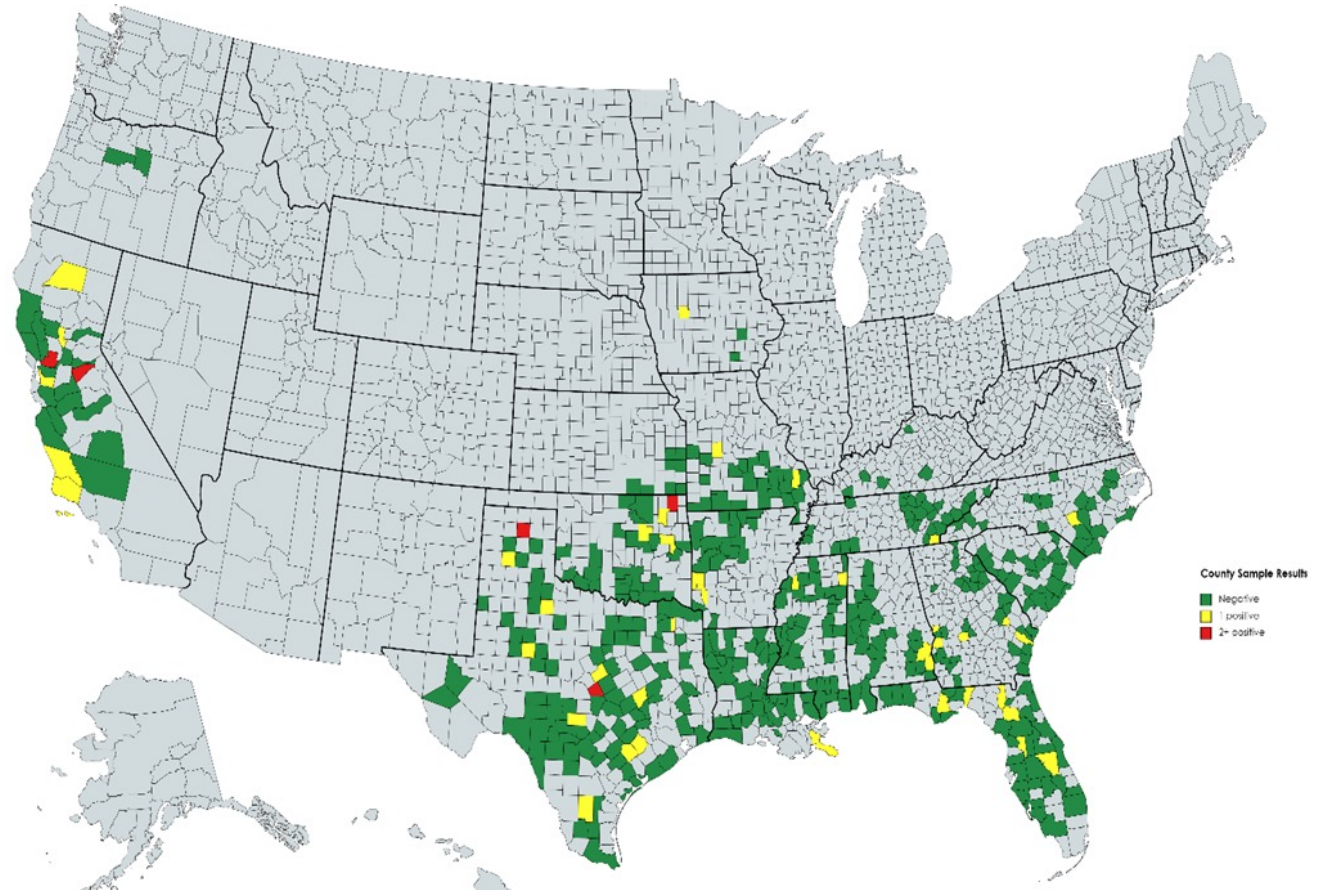
Approach:

- 945 serum convenience samples from 17 states collected by USDA-ARS NFSDM program and Auburn University College of Forestry, Wildlife and Environment
- Virus neutralization- 1:2 serial dilutions until final titer



Feral swine BVDV seroprevalence: Results

- 67 samples excluded from cytotoxicity
- 878 samples analyzed
- Total BVDV seroprevalence= **5.8%**
- BVDV-1b seroprevalence= **4.4%**
- BVDV-2a seroprevalence= **3.6%**
- Individual states 0-16.7%
- No difference in mean antibody titer BVDV-1b/2a by sex or age
- **CA, OK, TX**= greatest seroprevalence
 - Also, largest cattle and most established wild pig population states



Comparative BVDV COMMERCIAL TESTS

Commercially available and
validated for Cattle



RealPCR BVDV RNA Mix

IDEXX SNAP BVDV Antigen Test ★

IDEXX BVDV PI X2 Test

IDEXX BVDV Total Ab Test

IDEXX BVDV Ag/Serum Plus Test

IDEXX BVDV p80 Ab Test

IDEXX BVDV Ag Point-of-Care Test ★

Hypothesis:

- Heterologous hosts would have an elevated false-positive ACE results

★ Could easily be
used by producers
and veterinarians

<https://www.idexx.com/en/livestock/livestock-tests/ruminant-tests/idexx-bvdv-pi-x2-test/>

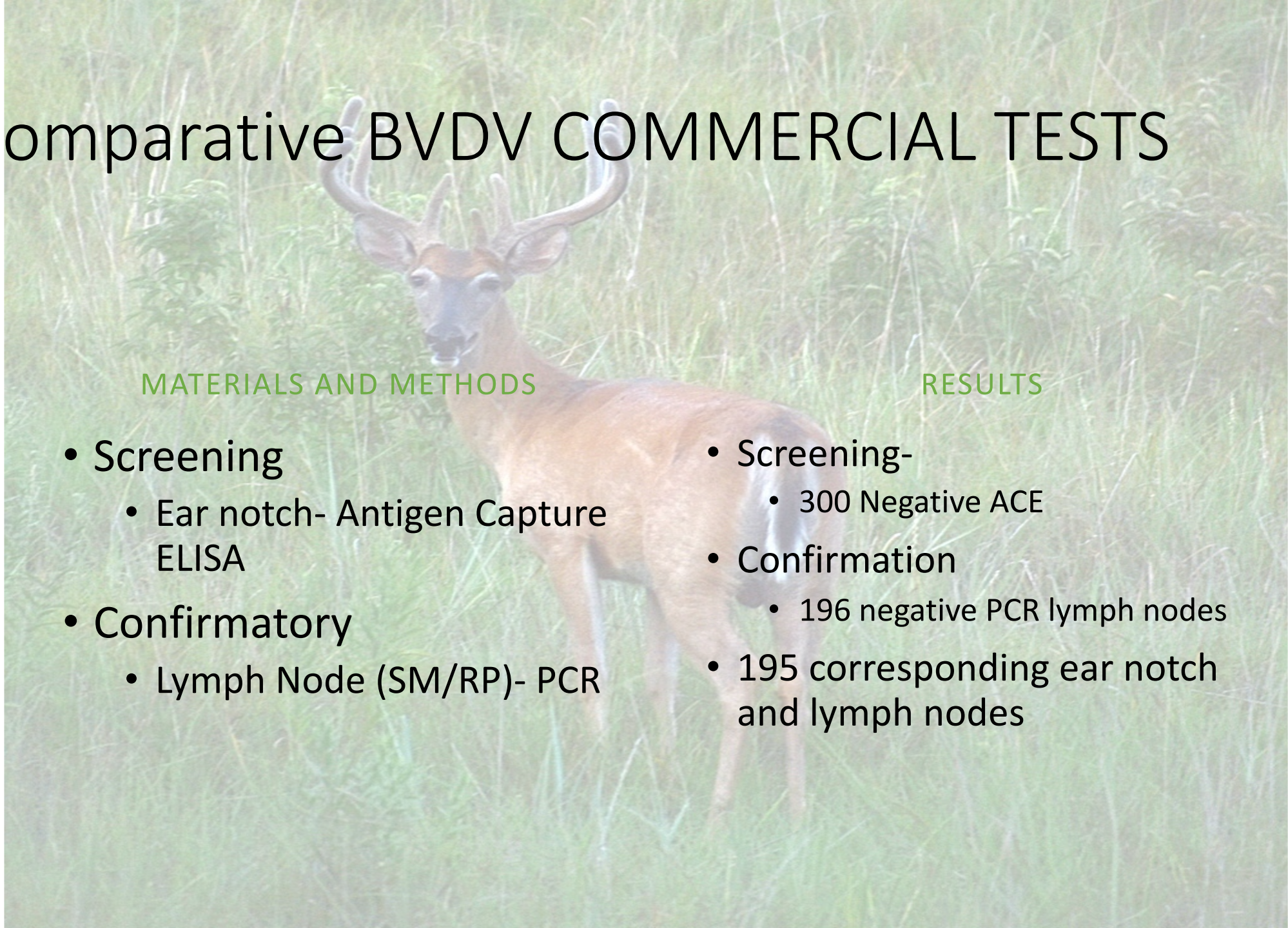
Appropriate testing?- DR. PASSLER

Year	acELISA*	Immunohistochemistry	RT-nPCR
2006/07 440 deer	22 Positive		
2007/08 577 deer	7 Positive, 13 Suspect	5 Positive, 4 Suspect	2 Positive
2008/09 590 deer	4 Suspect, 3 Positive	Pending	0 Positive

*S/P ratios : Suspect 0.2 – 0.39, Positive: > 0.39

Species	Sample	Test	Results: no. (%)	Confirmation?	Reference
1. Rocky Mountain Bighorn Sheep 2. Mountain goats 3. Mule Deer	Serum	BVDV ACE ^a	Antigen positive 1: 0/10 (0) 2: 0/30 (0) 3: 6/84 (7.1); 0/19 (0)	No	Wolff et al 2016
Algerian Dromedary Camel	Serum	BVDV antibody* ELISA and BVDV antigen [†] ELISA	Antibody positive • 10/111 (9.0) Antigen positive • 46/111 (41.4)	No	Saidi et al 2018
White-tailed deer	Ear notch/ Lymph node	BVDV ACE ^a	Antigen positive • Ear notch 1/311 (0.32) • Lymph node 1/434 (0.23)	Yes • PCR, VI, Sequence analysis- positive samples	Pogranichniy et al 2008
White-tailed deer	Ear Notch	BVDV ACE ^b	Antigen suspect • 3/367 (0.82)	Yes • Immunohistochemistry, VI, and PCR- suspect samples Screening subset • 89/367 PCR negative	Ilha et al 2012
Austrian Red Deer	Ear Notch	BVDV ACE ^a	Antigen positive • 0/764 (0)	Yes • PCR- all samples	Glawischnig et al 2010
Bactrian camels	Serum	BVDV ACE ^a	Antigen positive • 17/56 (30.4)	Yes • PCR- positive samples	Gao et al 2013
Sheep	Serum	BVDV ^{c,d}	Antibody positive • 27/42 (64.3) inoculated sheep	Yes • Agarose gel immunodiffusion (AGID) • All samples	Evans et al 2017

Comparative BVDV COMMERCIAL TESTS



MATERIALS AND METHODS

- Screening
 - Ear notch- Antigen Capture ELISA
- Confirmatory
 - Lymph Node (SM/RP)- PCR

RESULTS

- Screening-
 - 300 Negative ACE
- Confirmation
 - 196 negative PCR lymph nodes
- 195 corresponding ear notch and lymph nodes

Comparative test conclusion

- Reliably detect negative or non-viremic animals.
- High correlation (sensitive) between IDEXX BVDV PI X2 Antigen-capture ELISA with PCR
- High negative predictive value
- Unknown specificity- likely high false positives in previous studies?

	Disease positive	Disease negative	
Result positive	TP	FP	PPV (TP/TP+FP)
Result negative	FN	TN	NPV (TN/FN+TN)
	Sensitivity (TP/TP+FN)	Specificity (TN/FP+TN)	



Swine vs Bovine BVDV isolate PI infection

- Evaluate antigenic changes during immune recognition of pregnant cattle and creation of PI
 - BVDV propagated in heterologous hosts
 - BVDV propagated in cattle hosts



Swine Isolate

460f6

Pre-colostrum viremia PI calf
Acute Infection Heifer
Inoculum

VS



Bovine Isolate

AU-526

Pre-colostrum viremia PI calf
Acute Infection Heifer
Inoculum

Research timeline- swine vs cattle

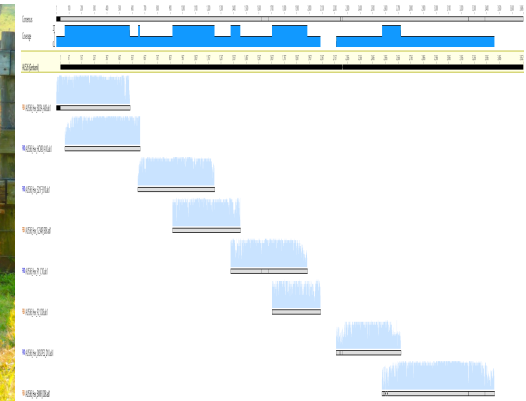
**DAY 75
GESTATION**

**ACUTE
INFECTION**

GESTATION

PARTURITION

SEQUENCING



Nucleotide difference between Inoculum and PI Calves

	Total	Transition	Transversion	Npro	C	Erns	E1	E2
01L	41	21	10	2	5	11	4	19
02L	9	3	0	0	0	1	4	4
03L	27	22	5	NA	3	9	4	11
04L	7	4	3	1	1	1	2	2
05L	35	29	6	5	3	10	5	12
06L	32	26	6	0	4	12	4	12
07L	38	31	7	3	4	11	5	15
08L	6	3	3	1	1	1	1	2
09L	34	25	9	4	3	9	5	13
10L	4	2	2	1	1	0	1	1
Median	29.5	21.5	5.5	1	3	9	4	11.5
Mean	23.9	17.0	5.1	1.8	2.5	6.7	3.5	9.3

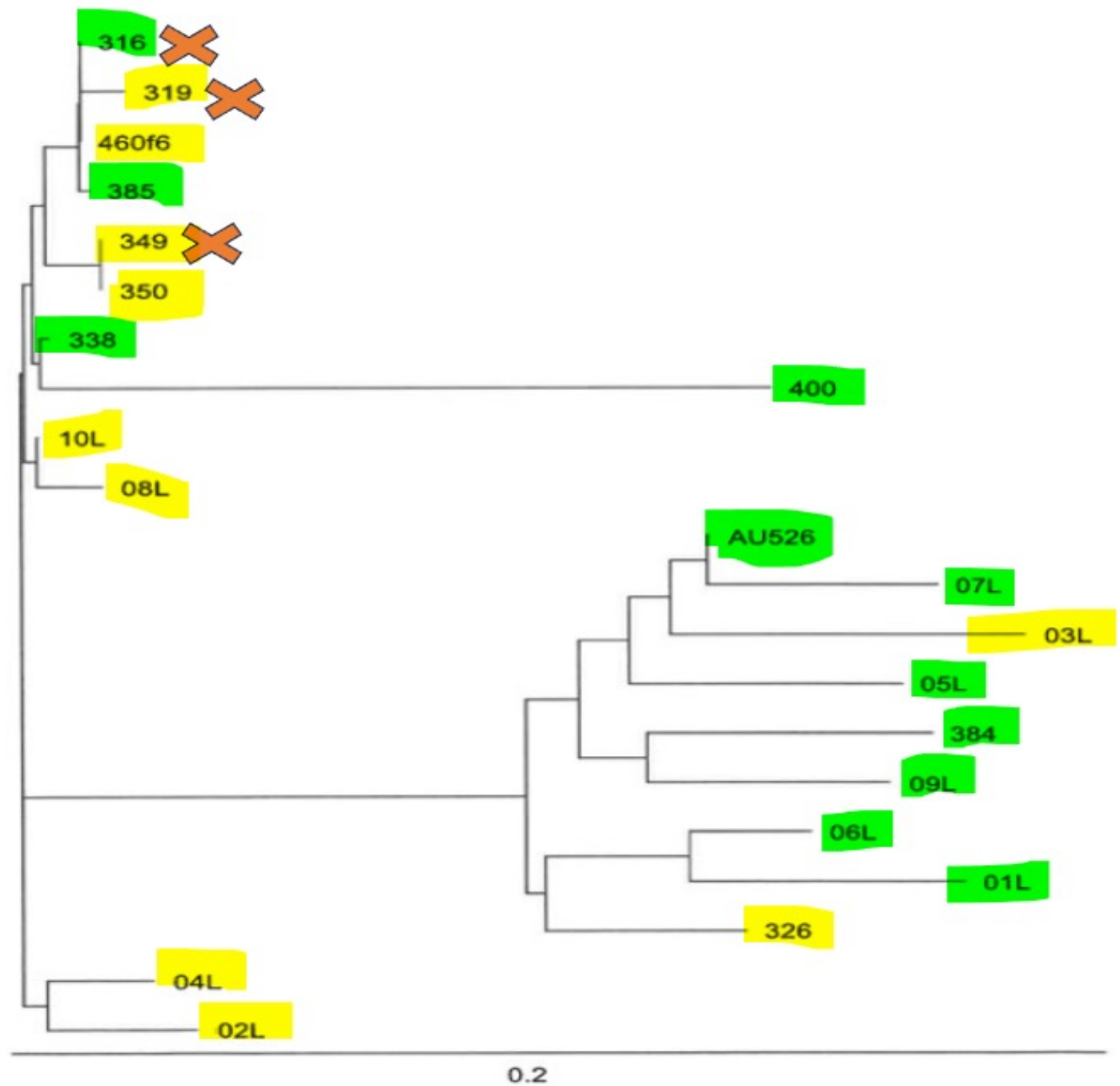
Amino Acid difference between Inoculum and PI Calves

	Total	Npro	C	Erns	E1	E2
01L	19	1	2	7	2	7
02L	8	0	0	1	4	3
03L	11	NA	1	1	2	7
04L	6	0	1	1	2	2
05L	11	0	1	1	2	7
06L	16	0	1	8	2	5
07L	11	0	2	1	2	6
08L	3	0	1	0	1	1
09L	15	1	1	3	3	7
10L	3	0	1	0	1	1
Median	11	0	1	1	2	5.5
Mean	10.4	0.2	1.1	2.2	2.1	4.7

Conserved Amino Acids in PI calves

Viral Protein AA	C	Erns	E1				E2							
Position	6	36	49	169	178	209	8	36	57	213	254	281	301	373
460-f6	N	K	T	D	A	R	Y	R	T	S	R	H	K	L
01L	D	R	I	H	T	K	Y	G	T	F	R	Y	T	S
02L	N	K	T	D	A	G	Y	R	T	S	Q	H	K	L
03L	D	K	T	H	T	K	Y	G	I	F	R	Y	S	S
04L	D	K	T	D	A	G	Y	R	T	S	Q	H	K	L
05L	D	K	T	H	T	K	F	G	I	F	R	Y	R	S
06L	D	R	I	H	T	R	Y	G	T	F	R	Y	T	S
07L	D	K	T	H	T	K	F	G	I	F	R	Y	S	S
08L	D	K	T	D	A	R	Y	R	T	S	R	H	K	L
09L	D	K	T	H	T	G	F	G	T	F	L	H	R	S
10L	D	K	T	D	A	R	Y	R	T	S	R	H	K	L

Swine vs Cattle Genetics Results

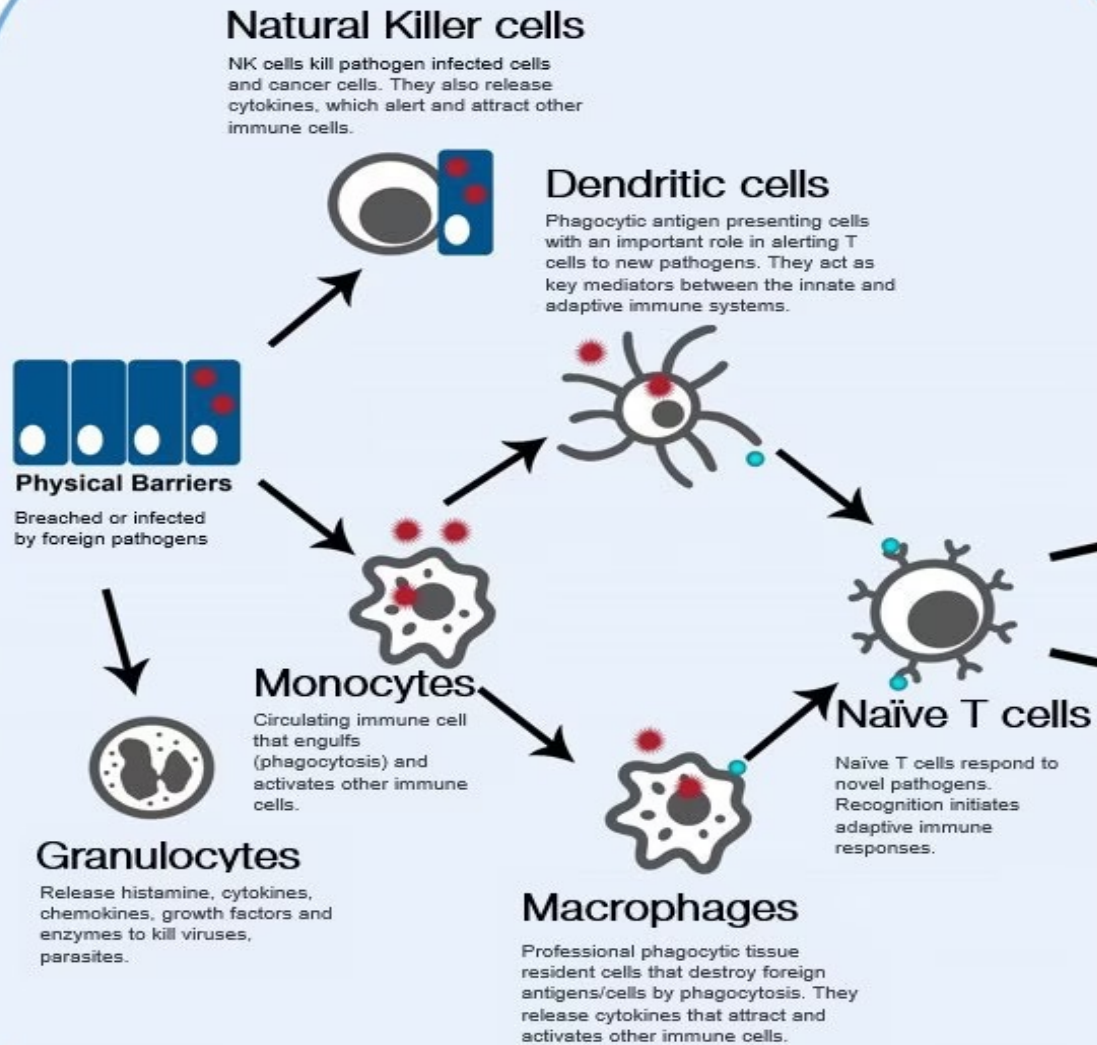


Cell-mediated immunity during pregnancy recognition with a PI fetus

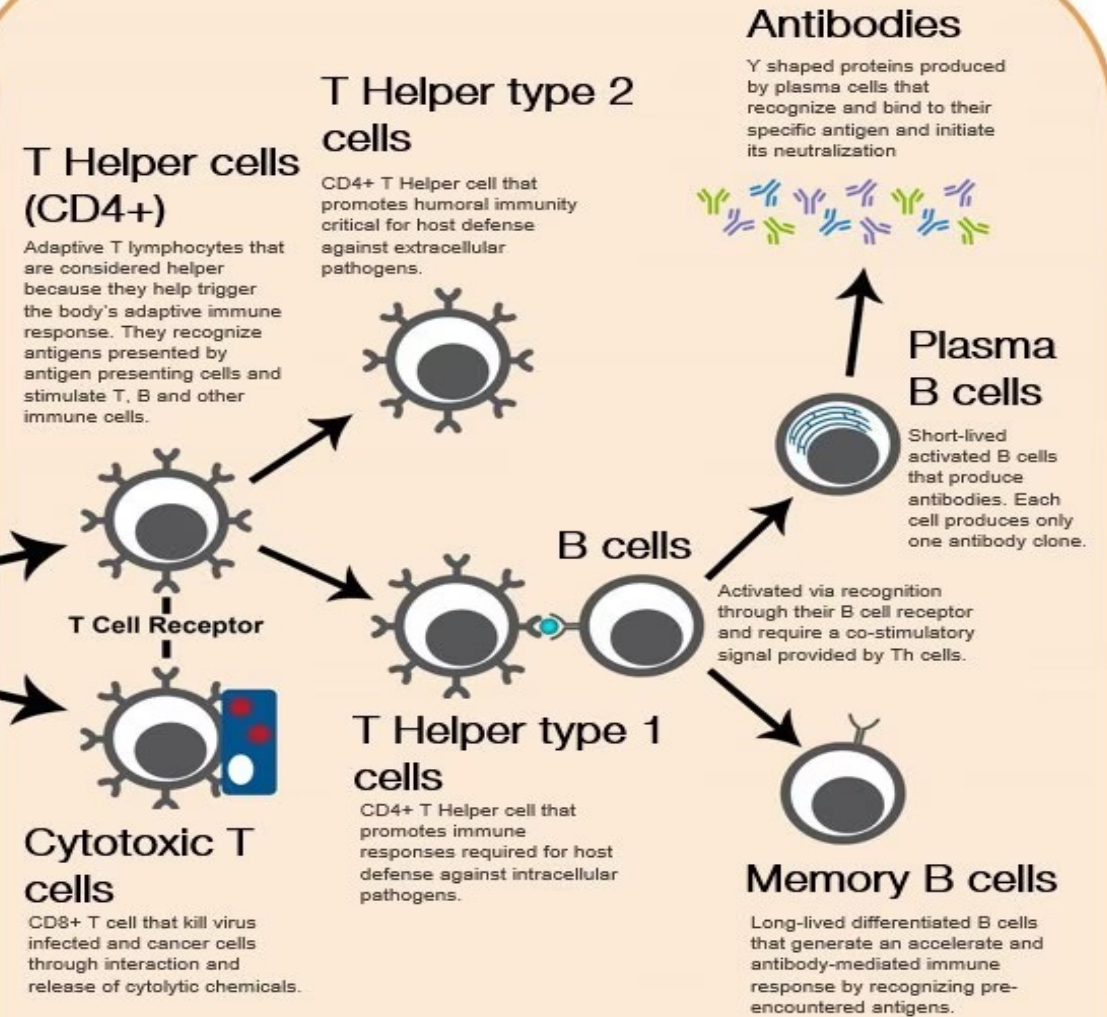


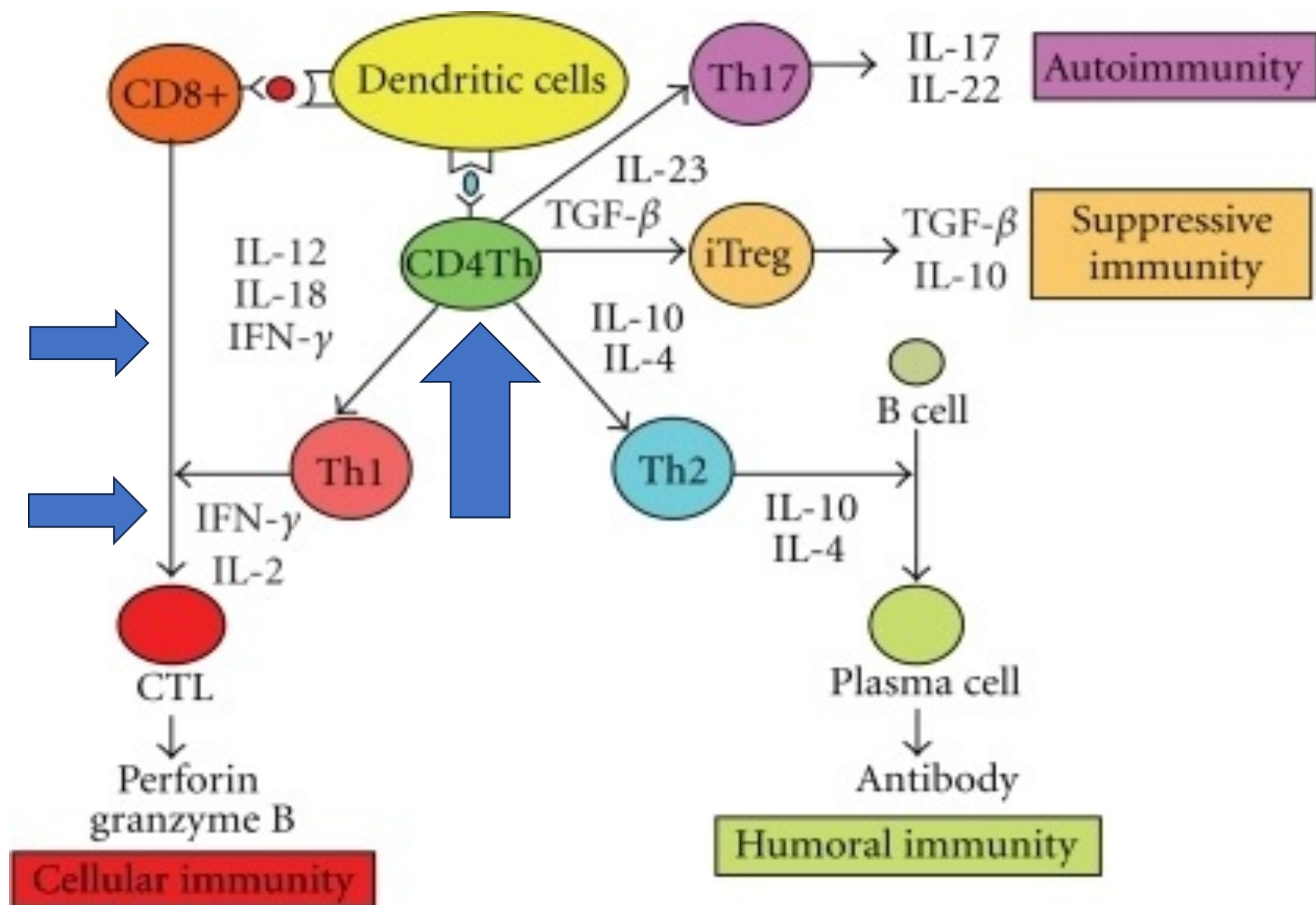
- Does pregnancy with a PI fetus alter the cell mediated immunity and the ability for the adaptive immune system to respond to both specific and non-specific challenges?

Innate immune response



Adaptive immune response





CMI Assay Heifers

- Materials and Methods

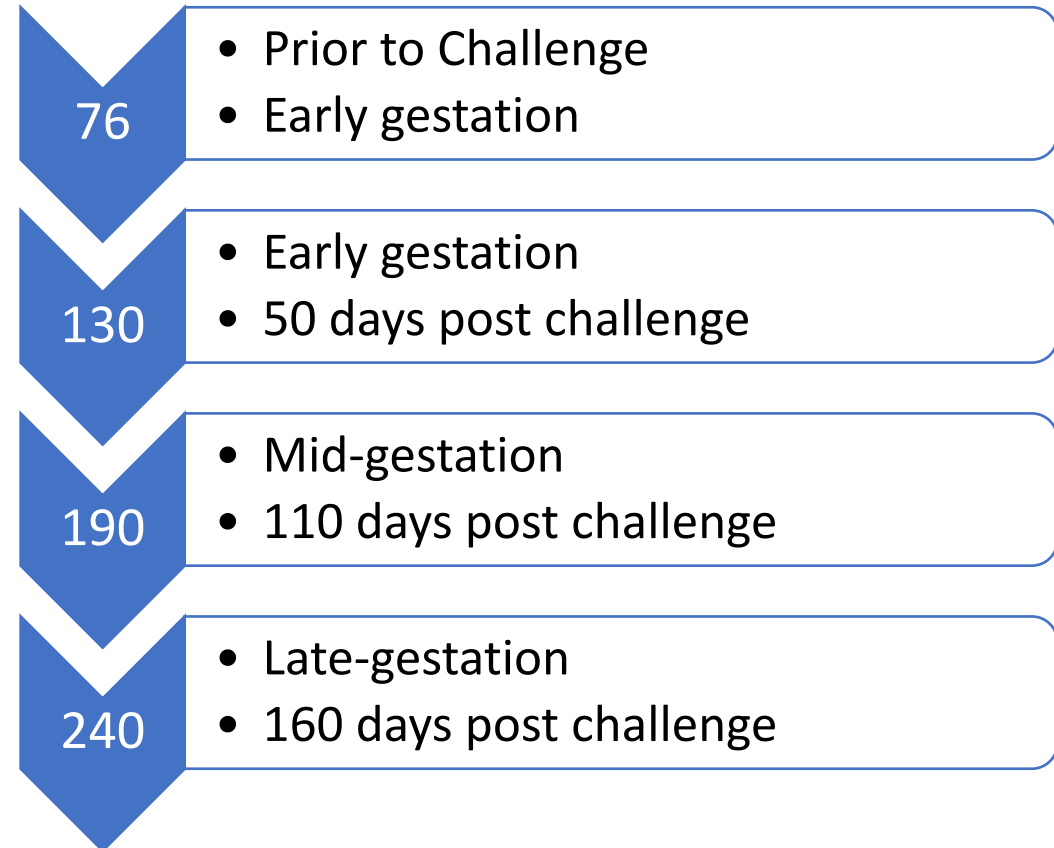
Measuring CMI responses using the PrimeFlow RNA assay: A new method of evaluating BVDV vaccination response in cattle

Shollie M. Falkenberg^{a,*}, Rohana P. Dassanayake^a, John D. Neill^a, Paul H. Walz^b, Eduardo Casas^a, Julia F. Ridpath^a, James Roth^c

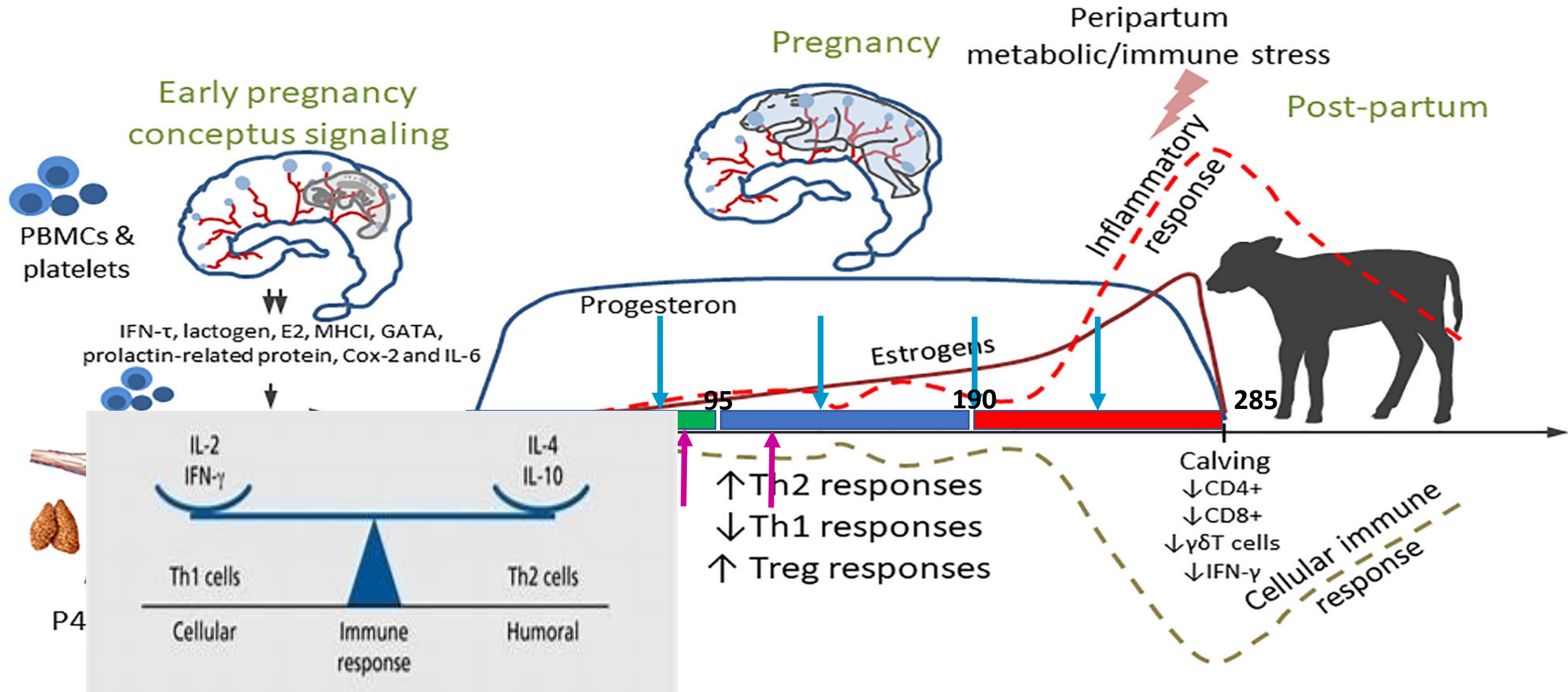
^a Ruminant Disease and Immunology Research Unit, National Animal Disease Center, USDA, Agricultural Research Service, Ames, IA, 50010, United States

^b Department of Pathobiology, College of Veterinary Medicine, 129 Sugg Laboratory, Auburn University, AL, 36849, United States

^c Department of Veterinary Microbiology and Preventive Medicine, College of Veterinary Medicine, Iowa State University, IA, United States

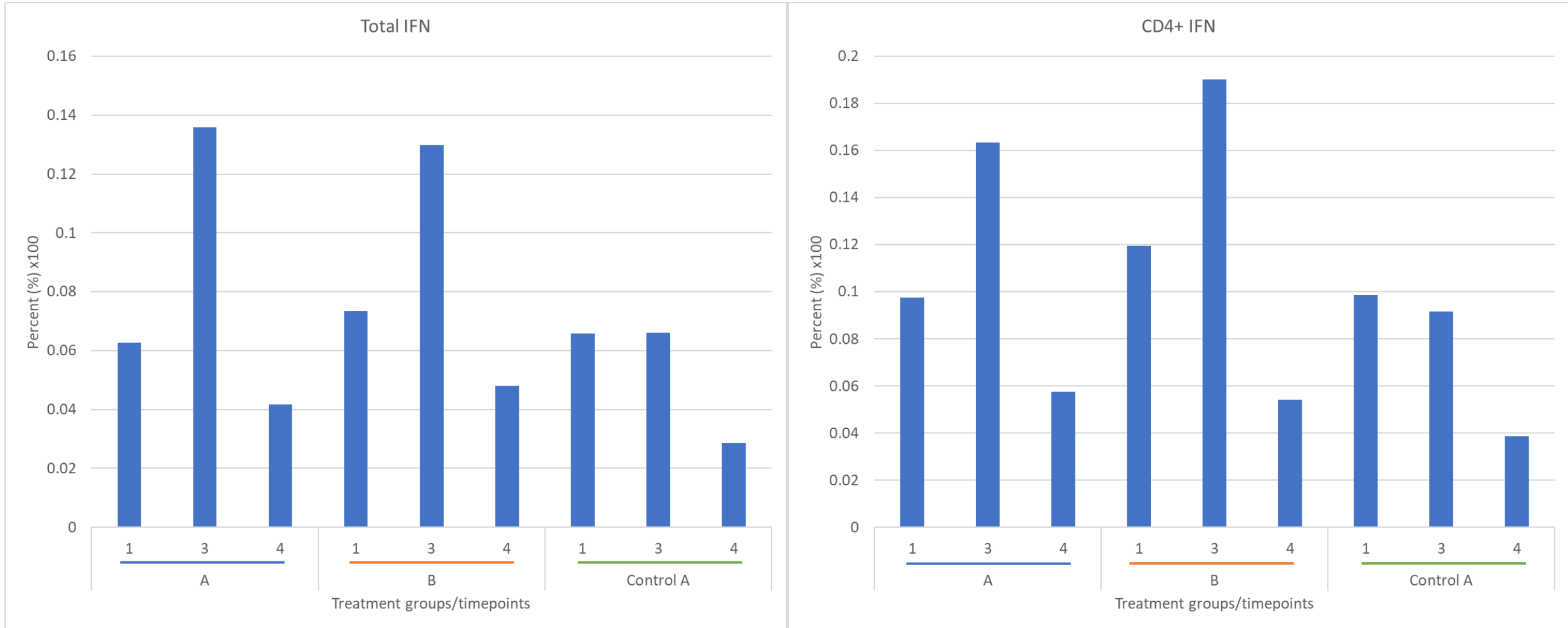


Immune response during pregnancy



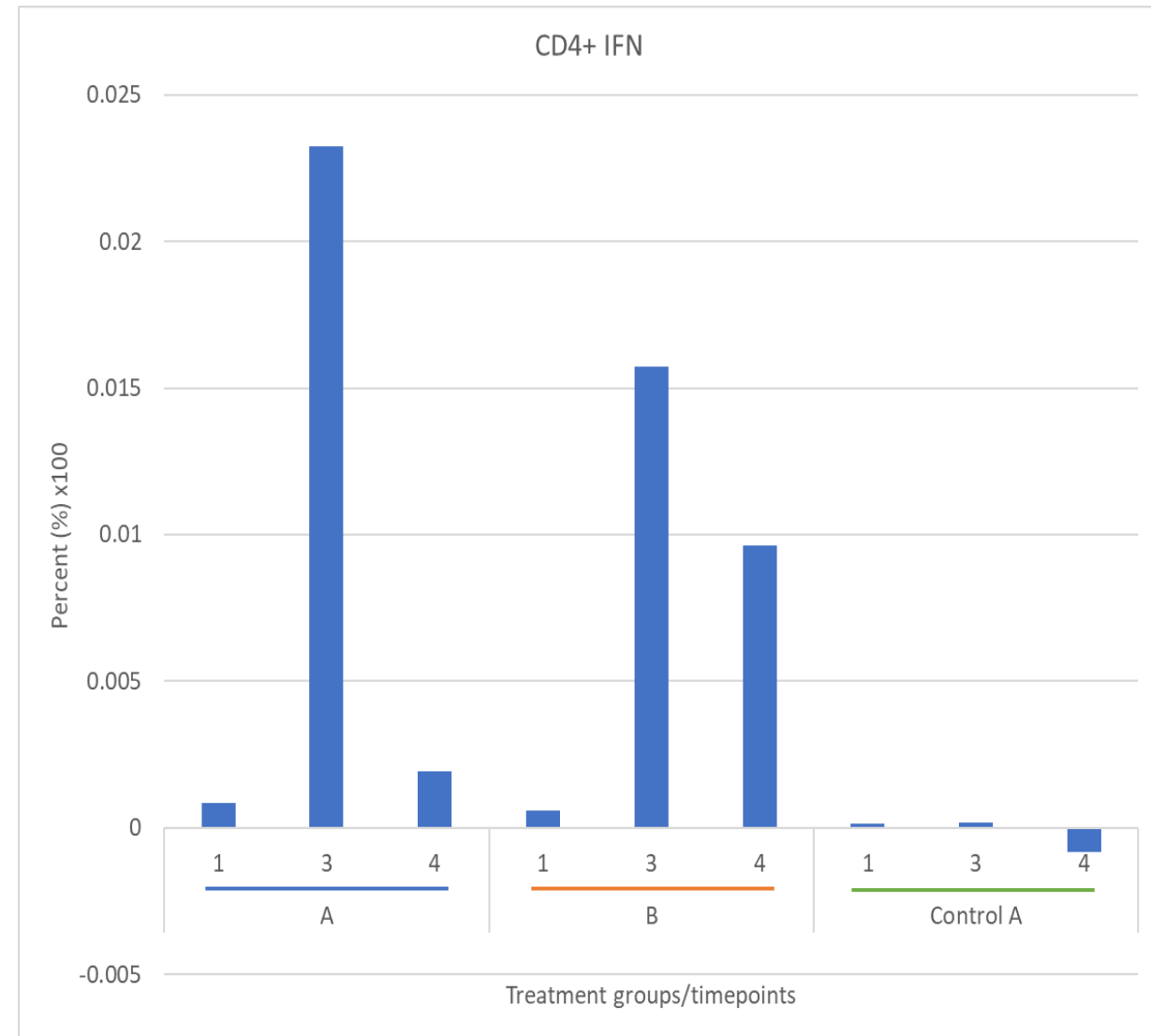
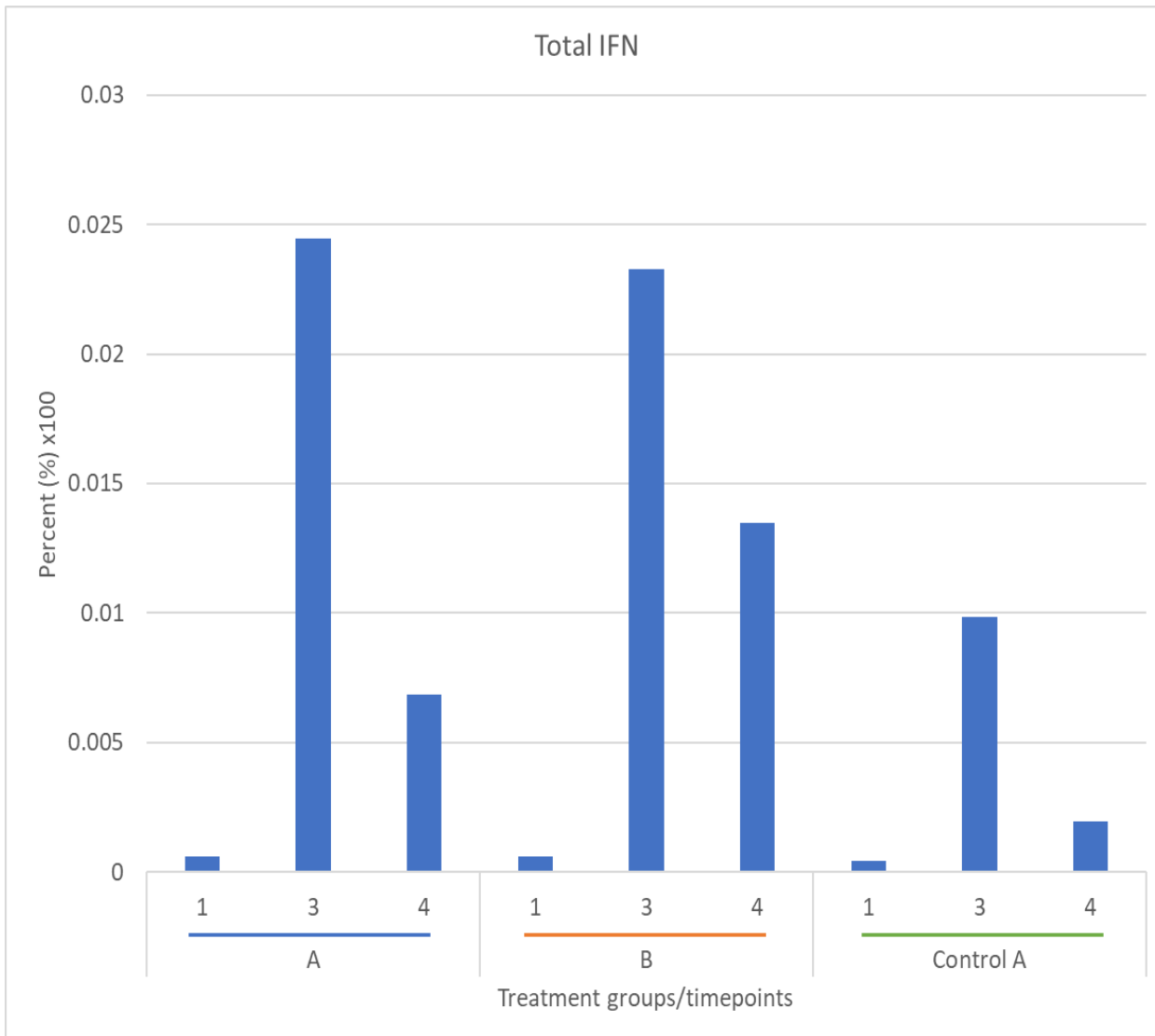
Heifers with PI and non-PI fetuses

Mitogen stim- Total IFN and CD4s



Heifers with PI and non-PI fetuses

BVDV stim – Total and CD4





Thank you

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