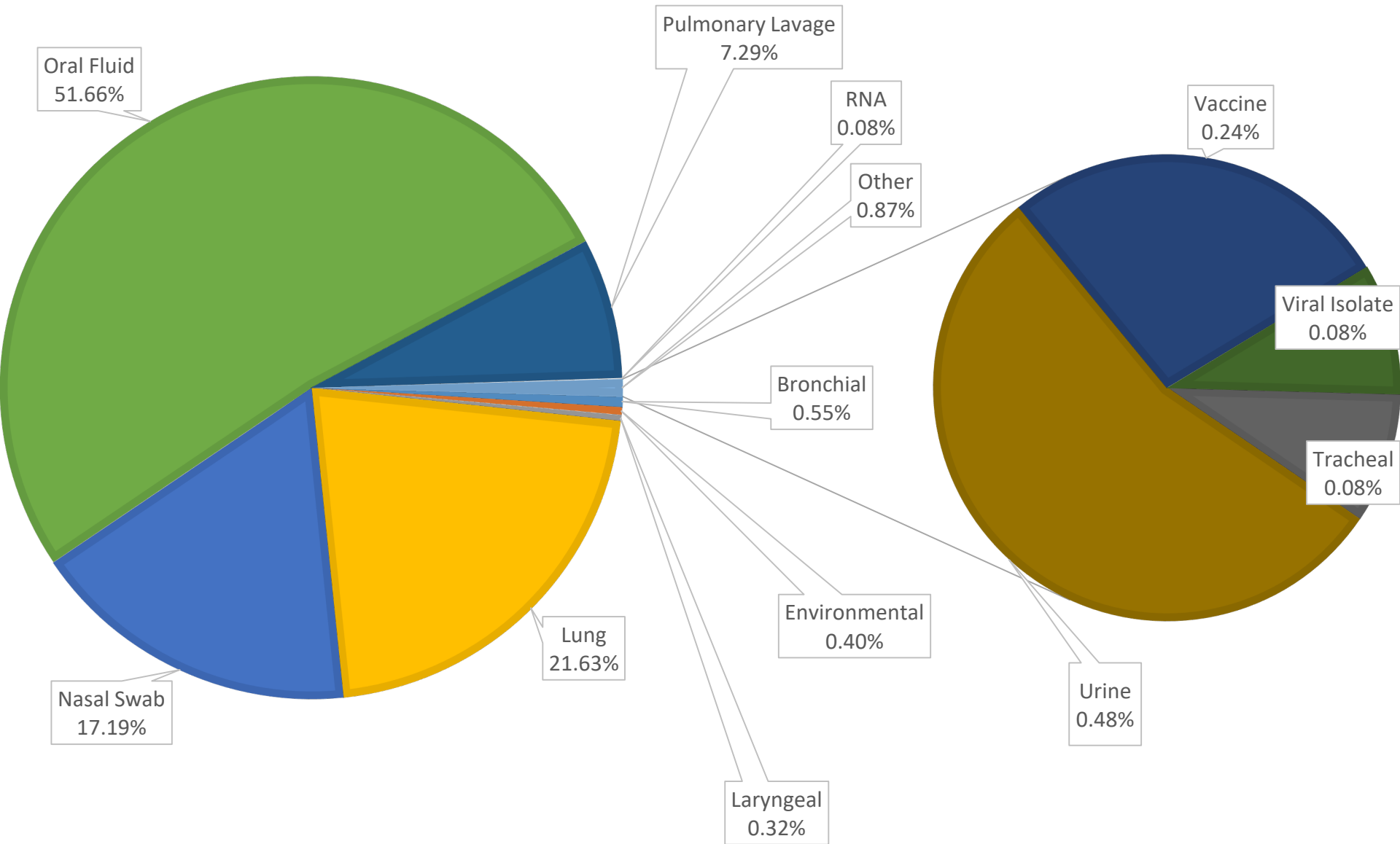


# Influenza A Virus Diversity in US Swine

University of Minnesota Veterinary Diagnostic  
Laboratory Report

Jerry Torrison, Mary Thurn, Marie Culhane

# Types of Specimens Received



# Specimens for influenza surveillance



RESEARCH ARTICLE

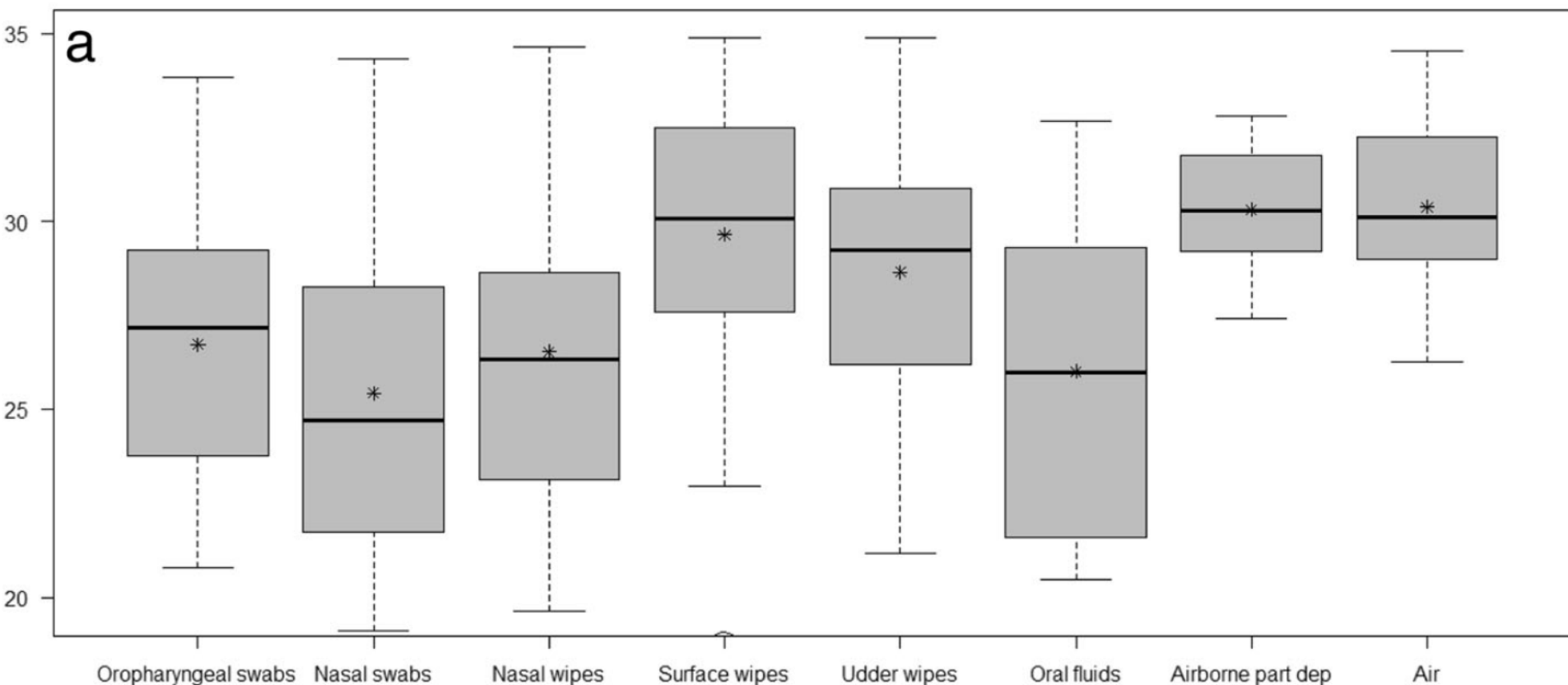
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# Comparison of individual, group and environmental sampling strategies to conduct influenza surveillance in pigs

Jorge Garrido-Mantilla<sup>1</sup>, Julio Alvarez<sup>1,2,3</sup>, Marie Culhane<sup>1</sup>, Jayaveeramuthu Nirmala<sup>1</sup>, Jean Paul Cano<sup>4</sup> and Montserrat Torremorell<sup>1\*</sup>

## Breeding herds




RESEARCH ARTICLE

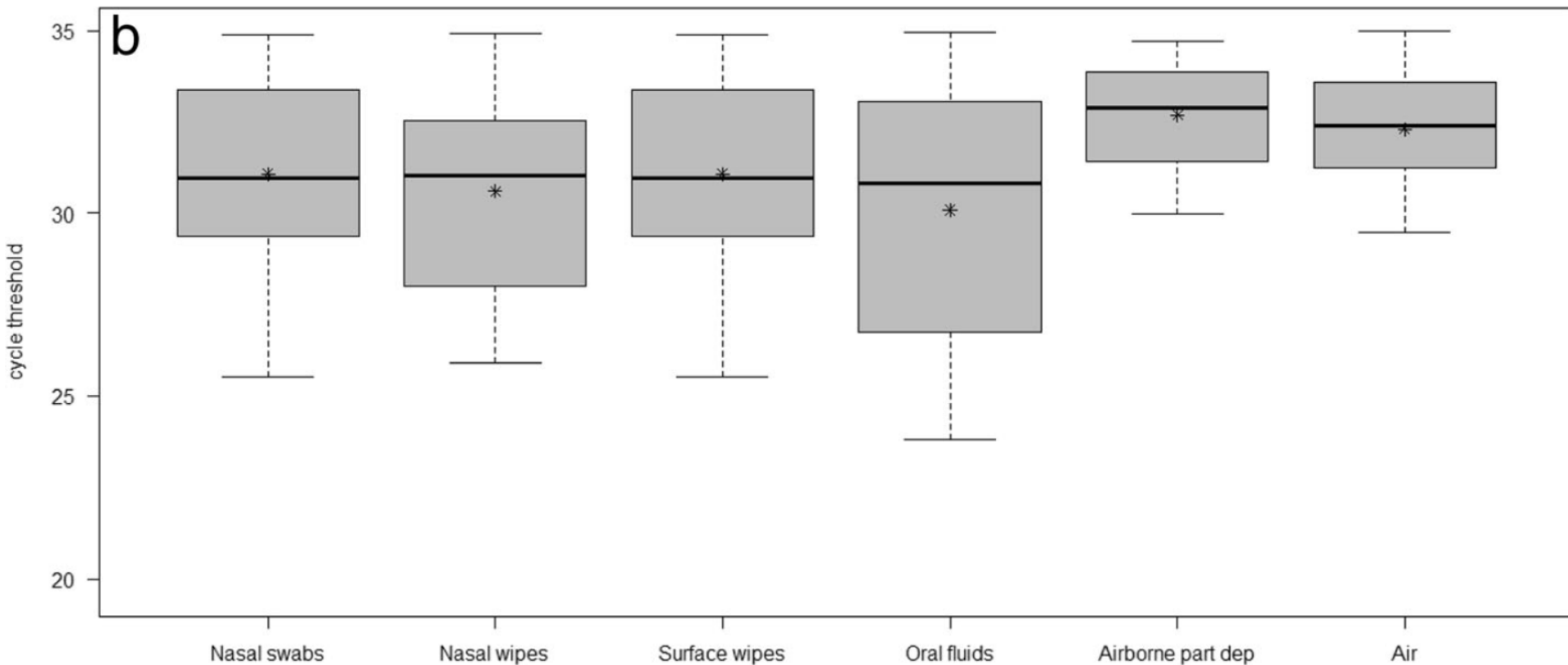
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## Wean to finish




RESEARCH ARTICLE

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**Table 5** Odds ratios (OR) of finding a positive result by sample type using a multivariate analysis considering two random effects (farm and litter or pen) and using the pooled nasal swab as baseline

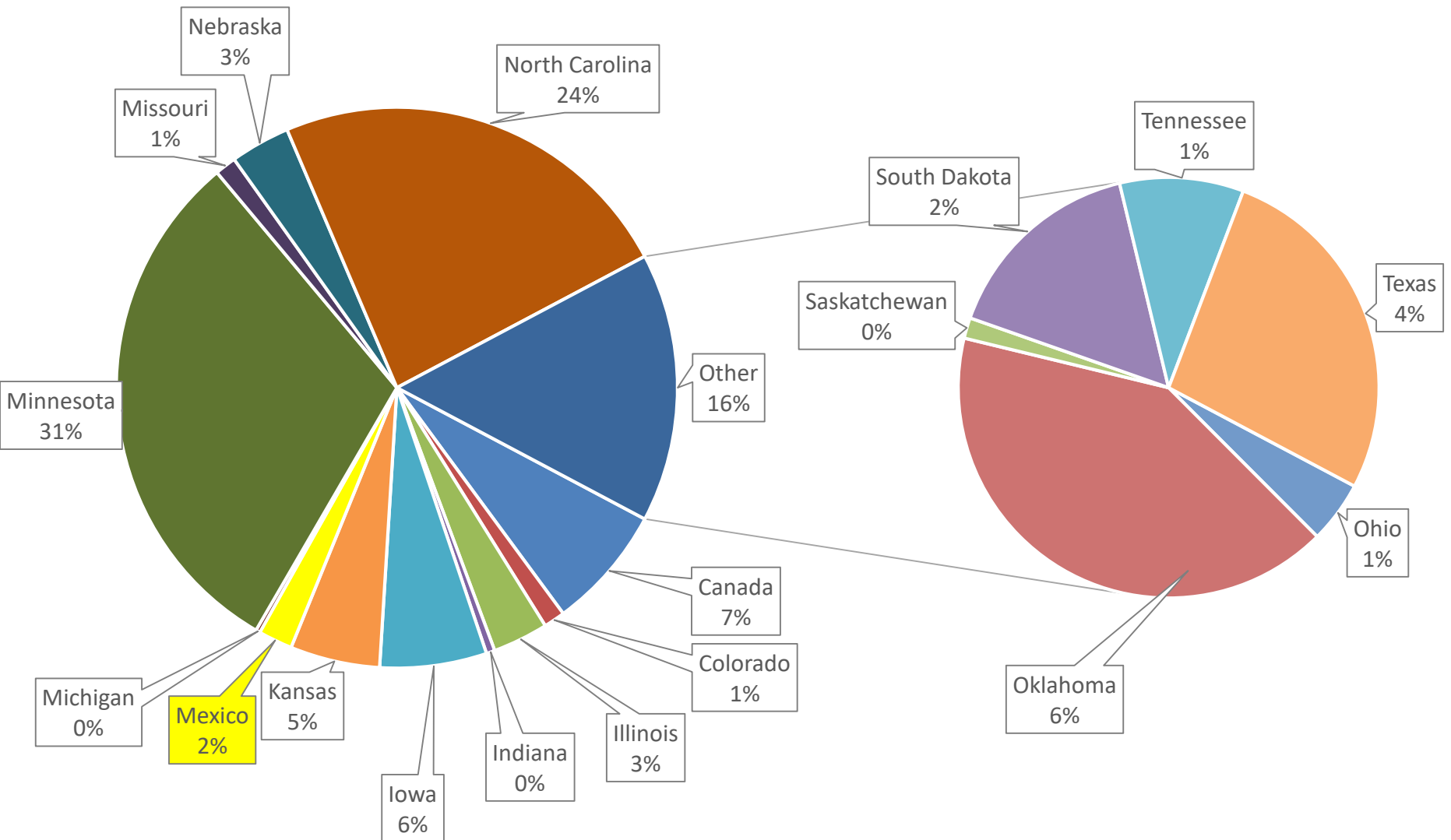
Sample type	N	Positive samples / Total samples (%)	$\beta^a$	SE $\beta^b$	OR (95% CI)	P value
Nasal swab	90	49/90 (54.4)	0.371	0.702	Reference	–
Nasal wipe	90	51/90 (56.7)	0.199	0.447	1.22 (0.51–2.95)	0.66
Oropharyngeal swab	40	32/ 40 (80)	2.088	0.714	8.07 (2.13–35.7)	0.003
Oral fluid	55	41/55 (74.5)	2.284	0.635	9.81 (3.02–37.3)	< 0.001
Surface wipe	90	62/ 90 (68.9)	1.354	0.478	3.87 (1.55–10.2)	0.004
Udder wipe	40	34/40 (85)	2.803	0.79	16.5 (3.88–89.0)	< 0.001

<sup>a</sup> $\beta$ : regression coefficient

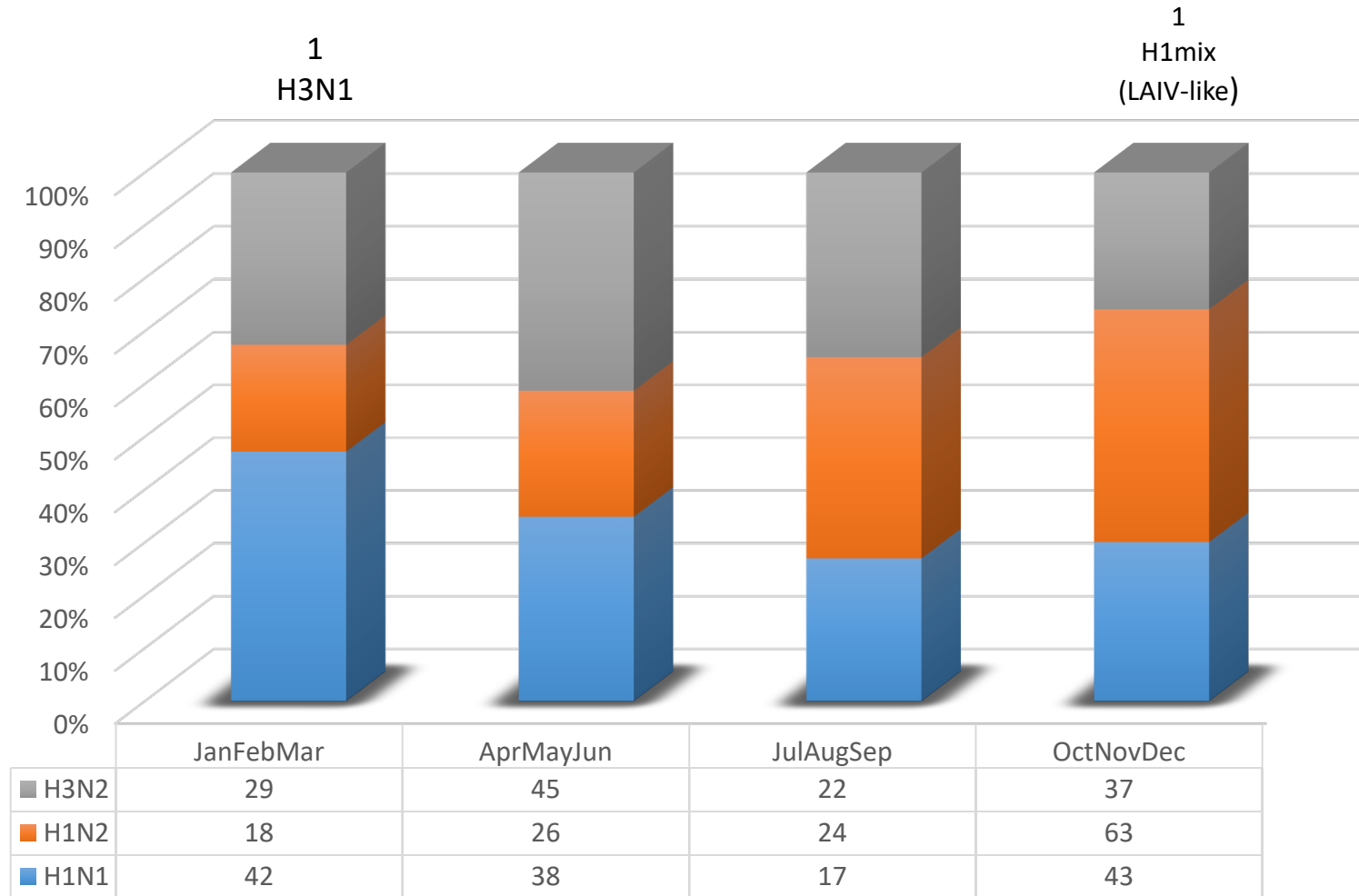
<sup>b</sup>SE  $\beta$ : Standard error of the regression coefficient

Nasal swabs, nasal wipes and oropharyngeal swabs were tested in pools of three

# Locations of Submitted Specimens

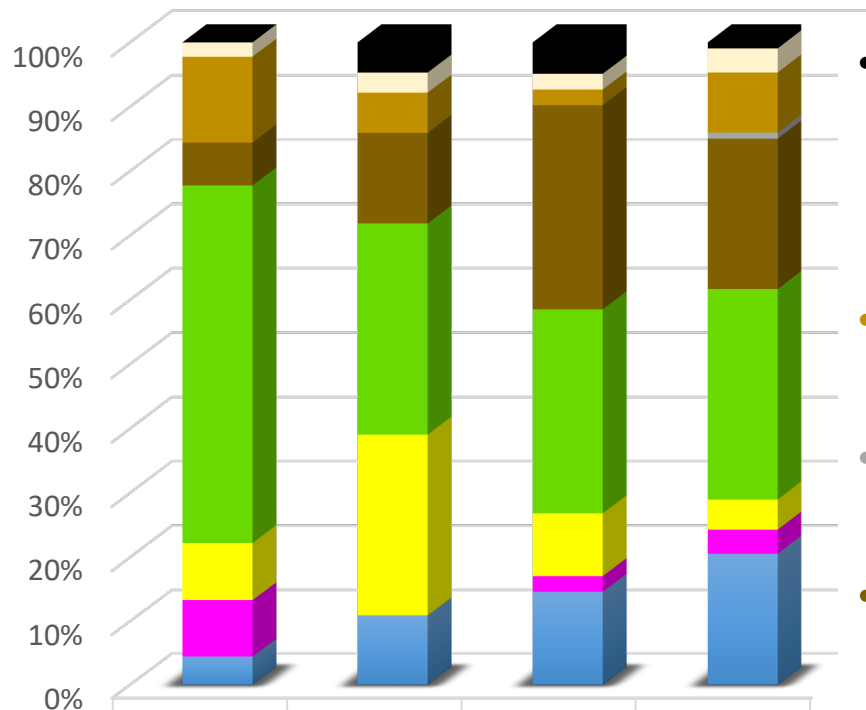


# 2018 Subtypes by Quarter





# H1 Clades 2018



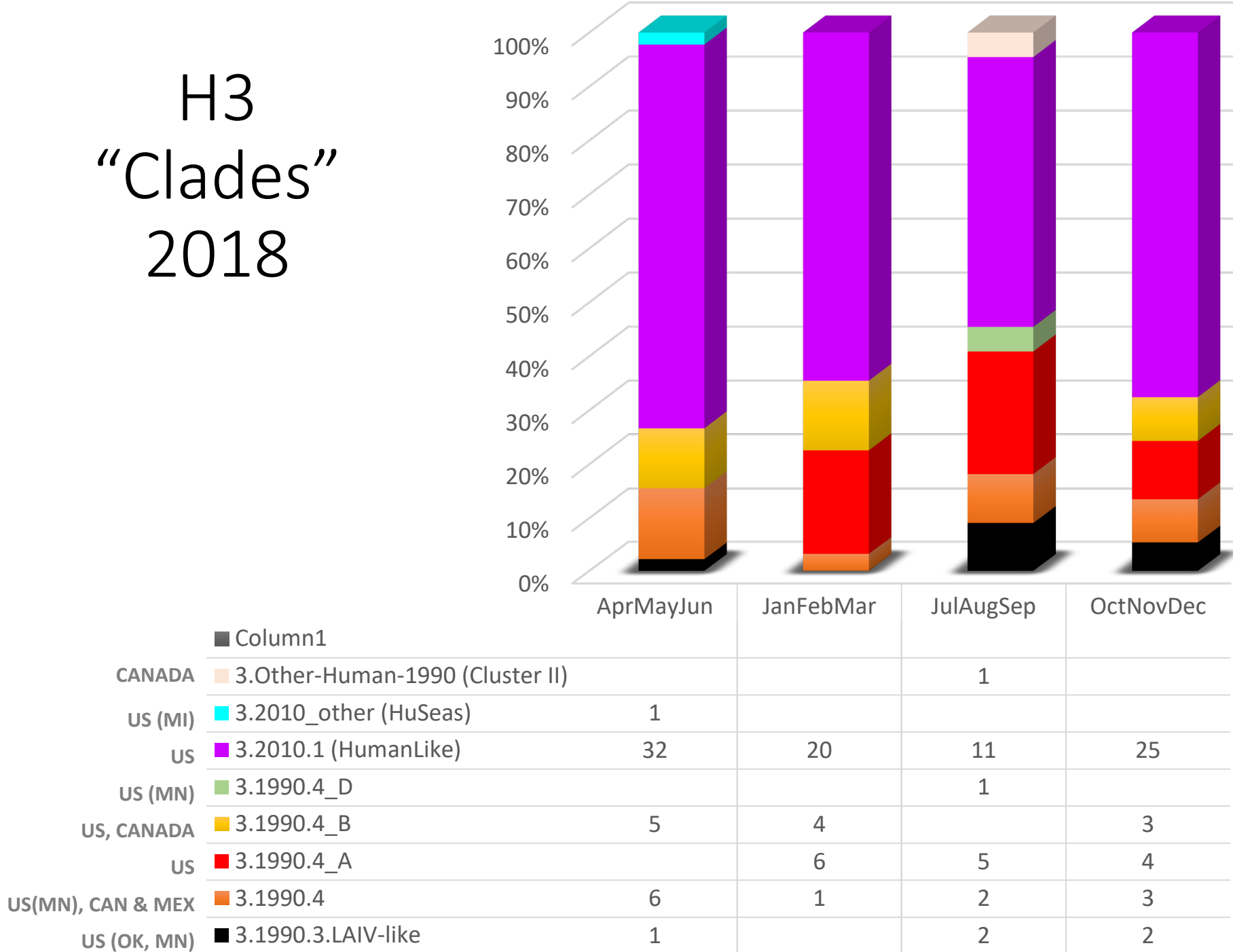
	JanFeb Mar	AprMay Jun	JulAugS ep	OctNov Dec
■ ND gamma2-betalike	0	3	2	1
■ 1B.2.2.2, delta 1b	1	2	1	4
■ 1B.2.2.1, delta 1a	6	4	1	10
■ 1B.2.2, delta 1	0	0	0	1
■ 1B.2.1, delta 2	3	9	13	25
■ 1A.3.3.3, gamma	25	21	13	35
■ 1A.3.3.2, H1N1pdm09	4	18	4	5
■ 1A.2, beta	4	0	1	4
■ 1A.1.1, alpha	2	7	6	22

- gamma2-beta-like 1A.2.3-like
  - USA – MN, OK, NE
- **delta 1b 1B.2.2.2**
  - USA
- **delta1a 1B.2.2.1**
  - USA
- **delta1, 1B.2.2**
  - USA (TX only)
- **delta 2 1B.2.1**
  - USA
- **Gamma 1A.3.3.3**
  - USA
- **npdm 1A.3.3.2**
  - Canada, USA, Mexico
- **beta 1A.2**
  - USA (KS, OK, TX)
- **alpha 1A.1.1**
  - CANADA, USA

# H1 – NA Pairings

	1998 N2	2002 N2	2002 N2	classical swine N1	pandemic N1
1A.1.1, alpha	8%	92%	0%	0%	0%
1A.2, beta	0%	0%	0%	100%	0%
1A.2-3, gamma 2-beta-like	0%	0%	0%	100%	0%
<b>1A.3.3.2, H1N1pdm09</b>	0%	0%	0%	0%	100%
1A.3.3.3, gamma	0%	4%	0%	96%	0%
1B.2.1, delta 2	94%	6%	0%	0%	0%
1B.2.2, delta 1	0%	0%	100%	0%	0%
<b>1B.2.2.1, delta 1a</b>	22%	78%	0%	0%	0%
<b>1B.2.2.2, delta 1b</b>	0%	100%	0%	0%	0%

# H3 “Clades” 2018



# H3 – NA Pairings

	1998 N2	2002 N2	2002 N2	2002 N2subset
<b>3.1990.4</b> <b>(Cluster IV)</b>	0%	0%	0%	<b>100%</b>
<b>3.1990.4_A</b> <b>(Cluster IVA)</b>	0%	<b>100%</b>	0%	0%
<b>3.1990.4_B</b> <b>(Cluster IVB)</b>	0%	<b>100%</b>	0%	0%
<b>HumanLike</b>	7%	<b>90%</b>	3%	0%



*OFFLU meeting*  
*27 February 2019*  
*Paris, France*

**It will likely continue to be difficult to control influenza due to high IAV-S genetic diversity in pig populations.**

